

# **Inducing immunity to liver stage malaria through endogenous tissue resident memory cells**

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## ABBREVIATIONS

### Special characters

%	Percent
#	Number
°C	Degree Celsius
α	alpha
β	beta
γ	gamma
μg	Microgram
μl	Microlitre

### A

A.	<i>Anopheles</i>
Ab	Antibody
AMA1	Apical membrane antigen-1
αClec9A-Pb-1	Rat anti-Clec9A mAb genetically coupled to the Pb-1 epitope
APC	Antigen presenting cell

### B

B6	C57Bl/6 mice
BSA	Bovine serum albumin fraction V

### C

CSP	Circumsporozoite protein
CD	Cluster of differentiation
cDC	Conventional dendritic cell
CDN	Cyclic dinucleotides
cGAS	Cyclic GMP–AMP (cGAMP) synthase
ChAd	Chimpanzee adenovirus
ChAd63	Chimpanzee adenovirus serotype 63
CpG	Cytosine-phosphate-Guanine
CTL	Cytotoxic T Lymphocytes
CTV	Cell-Trace Violet™
cVac	Chemoprophylaxis vaccination
CXCL	CXC chemokine ligand
CXCR	CXC chemokine receptor

### D

D	Day
DC	Dendritic cells
dLN	Draining Lymph node
DMSO	Dimethyl Sulphoxide
DNA	Deoxyribonucleic Acid
dsDNA	Double-stranded DNA
dsRNA	Double-stranded RNA



<b>E</b>	
ECM	Experimental cerebral malaria
e.g.	<i>exempli gratia</i>
<b>F</b>	
FCS	Fetal Calf Serum
FNRG	FAH <sup>-/-</sup> NOD Rag1 <sup>-/-</sup> IL2R $\gamma$ <sup>NULL</sup>
<b>G</b>	
GAP	Genetically-attenuated parasites
GFP	Green fluorescent protein
<b>H</b>	
HBSS	Hanks' buffered salt solution
HBV	<i>Hepatitis B virus</i>
HPLC	<i>High-performance liquid chromatography</i>
HPV	<i>Human papilloma virus</i>
Hr	Hour(s)
HSV	Herpes Simplex Virus
<b>I</b>	
i.e.	<i>Id est</i>
IFN	Interferon
Ig	Immunoglobulin
IL	Interleukin
i.m.	Intramuscular
iRBC	Infected red blood cells
IRF	Interferon-Regulatory Factor
iRT	Indexed retention time
ISG	Interferon-Stimulated-Gene
i.v.	Intravenously
<b>K</b>	
KC	Kupfer cells
KLRG1	Killer cell lectin-like receptor subfamily G member 1
KO	Knock out
<b>L</b>	
LC-MS/MS	Liquid chromatography coupled to tandem mass spectrometry
LN	Lymph node
LPS	Lipopolysaccharide
LSA1	Liver-stage antigen 1
LSAP2	Liver stage-associated protein 2
<b>M</b>	

MAVS	Mitochondrial antiviral-signalling
MDA5	Melanoma Differentiation-Associated gene 5
MAb	Monoclonal antibody
ME	Multi-epitope string
MHC	Major Histocompatibility Complex
MSP1	Merozoite surface protein-1
MVA	Modified vaccinia virus Ankara
MyD88	Myeloid Differentiation factor 88

**N**

NCY	NCYDFNNI
NF- $\kappa$ B	Nuclear Factor kappa B
NK	Natural killer
Np	Nanoparticles
NVY	NVYDFNLL

**O**

ODN	Oligodeoxynucleotides
OVA	Ovalbumin

**P**

<i>P.</i>	<i>Plasmodium</i>
PAMP	Pathogen-Associated Molecular Pattern
PbA	<i>Plasmodium berghei</i> ANKA
PBMC	Peripheral blood mononuclear cells
PBS	Phosphate Buffered Saline
PCR	Polymerase chain reaction
pDC	Plasmacytoid Dendritic Cell
PD-1	Programmed death 1
PE	Pre-erythrocytic
PfCSP	<i>P. falciparum</i> CSP
PfSPZ	<i>P. falciparum</i> sporozoites
PI	Propidium iodine
(p.i.)	Post infection
Poly(I:C)	Polyinosinic-polycytidylic acid
PRR	Pattern-Recognition Receptor

**R**

RAS	Radiation-attenuated sporozoites
rAAV	Recombinant adeno-associated virus
RBC	Red blood cells
RIG-I	Retinoic acid Inducible Gene I
RPL6	Ribosomal protein L6
RT	Room temperature

**T**

T <sub>CM</sub>	Central memory T (cells)
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TCR	T cell receptor
T <sub>EM</sub>	Effector memory T (cells)
Th1	Type 1 T helper
Th2	Type 2 T helper
Tim-3	T-cell immunoglobulin mucin 3
TLR	Toll-like receptor
TNF $\alpha$	Tumour necrosis factor alpha
TRAP	Thrombospondin-related adhesion protein
TRIF	TIR-domain-containing adapter protein inducing interferon IFN- $\beta$
T <sub>RM</sub>	Tissue-resident memory T (cells)

**S**

SLO	Secondary lymphoid organs
ssRNA	Single stranded RNA

**W**

WHO	World Health Organization
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**ABSTRACT**

Tissue-resident memory CD8 T ( $T_{RM}$ ) cells provide effective tissue surveillance and can respond rapidly to infection due to their strategic location. Within the liver,  $T_{RM}$  cells can induce effective protection against liver-stage *Plasmodium* infection. Recently, members from our group identified a highly immunogenic peptide (named Pb-1) within the putative 60S ribosomal protein L6 of *P. berghei* ANKA. Experiments conducted and presented in this thesis aimed to assess the suitability of Pb-1 for the induction of endogenous liver  $T_{RM}$  cells that confer sterilizing protection in B6 mice. To this end, a series of different immunisation strategies targeting the Pb-1 epitope were implemented and specific CD8 T cell responses were assessed. Results revealed that the number of naïve specific CD8 T cell precursors for the Pb-1 epitope was very large. Substantial expansion and formation of specific liver  $T_{RM}$  cells was achieved by two different immunisation strategies: i) Single injection with Clec9A mAb plus adjuvant and ii) Prime-and-trap, both targeting the Pb-1 epitope. While mice vaccinated with Clec9A mAb developed partial protection, almost all mice vaccinated with prime-and-trap targeting Pb-1 were sterily protected against liver-stage challenge.

Inflammation favours the formation  $T_{RM}$  cells and adjuvants can affect their numbers. Accordingly, a second focus of this thesis sought to investigate how to enhance liver  $T_{RM}$  cell formation by using TLR and RIG-I-like receptors agonists as adjuvants. For this, 8 different agonists were assessed for the generation of liver  $T_{RM}$  cells induced by Clec9A targeted immunisation with the Pb-1 epitope. Data from this screen showed that CpG-based adjuvants were most effective at inducing the formation of  $T_{RM}$  cells in the livers of vaccinated mice and that the transfection reagent DOTAP enhanced this effect. Based on this understanding, we then investigated the potential of CpG and its encapsulation in DOTAP to improve  $T_{RM}$  cell generation by other vaccination strategies. Surprisingly, these studies revealed that CpG-based adjuvants did not improve liver  $T_{RM}$  cell

generation by vaccination with radiation-attenuated sporozoites. The basis for this outcome is discussed.

Altogether, these findings provide insights into elements that favour the generation of protective liver T<sub>RM</sub> cells; information that can be used for the design of T<sub>RM</sub> cell-based subunit vaccines against *Plasmodium* infection.



## DECLARATION

The work presented in this thesis was conducted at The Rheinische Friedrich-Wilhelms-Universität Bonn, in the laboratory of Dr. Winfried Barchet and at The University of Melbourne, in the laboratory of Prof. William Heath. This work was supported by the Deutsche Forschungsgemeinschaft (DFG, German Research Council) within the Bonn & Melbourne Research and Graduate School (Immunosciences, IRTG2168). Ana Maria Valencia Hernandez was supported by the Deutsche Forschungsgemeinschaft, The Medical Faculty of the The Rheinische Friedrich-Wilhelms-Universität Bonn, the Melbourne International Research Scholarship and grants from the National Health and Medical Research Council (NHMRC) of the Australian Government.

This is to certify that,

- (i) the thesis comprises only my original work towards the PhD except where indicated in the preface;
- (ii) due acknowledgement has been made in the text to all other material used;
- (iii) the thesis is less than 100,000 word limit in length, exclusive of tables, maps, bibliographies and appendices as approved by the Research Higher Degrees Committee.

Melbourne, October 2020

Ana Maria Valencia Hernandez

PREFACE

My contribution to the experiments within each chapter was as follows:

**Chapter 3:** 90%

**Chapter 4:** 100%

I acknowledge the important contributions of these people:

**Chapter 3:**

Results of Figure 3.5 (**A.**) Dr. Nazanin Ghazanfari (5%) and (**B.**) Dr. Maria Menezes (5%).

**Research declaration**

Some of the content of Chapter 2 and Chapter 3 is also represented elsewhere in the reference below.

**Valencia-Hernandez AM**, Ng WY, Ghazanfari N, Ghilas S, de Menezes MN, Holz LE, et al. A Natural Peptide Antigen within the Plasmodium Ribosomal Protein RPL6 Confers Liver TRM Cell-Mediated Immunity against Malaria in Mice. *Cell Host Microbe* 2020:1–13. doi:10.1016/j.chom.2020.04.010.

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## LIST OF PUBLICATIONS

1. Ghilas S., **Valencia-Hernandez, A. M.**, Enders, M. H., Heath, W. R & Fernandez-Ruiz, D. (2020). Resident memory T cells and their role within the liver. *International Journal of Molecular Sciences*, 1-15. <https://doi.org/10.3390/ijms21228565>
2. **Valencia-Hernandez, A. M.**, Ng, W. Y., Ghazanfari, N., Ghilas, S., de Menezes, M. N., Holz, L. E., Huang, C., English, K., Naung, M., Tan, P. S., Tullett, K. M., Steiner, T. M., Enders, M. H., Beattie, L., Chua, Y. C., Jones, C. M., Cozijnsen, A., Mollard, V., Cai, Y., ... Fernandez-Ruiz, D. (2020). A Natural Peptide Antigen within the Plasmodium Ribosomal Protein RPL6 Confers Liver TRM Cell-Mediated Immunity against Malaria in Mice. *Cell Host & Microbe*, 1–13. <https://doi.org/10.1016/j.chom.2020.04.010>
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## CHAPTER 1

Literature review

## 1.1 Introduction:

Malaria is a disease caused by protozoan parasites of the *Plasmodium* spp. Six plasmodial species infect humans, and *Plasmodium falciparum* and *P. vivax* are responsible for the largest number of malaria cases (Cowman et al., 2016; Phillips et al., 2017; White et al., 2014). Over the past years, the international community have undertaken great efforts to prevent, control and eliminate malaria. In 2016, the World Health Organization (WHO) reported that 57 countries reduced their malaria cases by 75% compared to 2000. For the African region, the location with the highest burden worldwide, mortality fell 66% among all age groups and by 71% among children under five; the group with the highest mortality (WHO, 2016). Insecticide-treated bed nets, antimalarial medication and indoor residual insecticide spraying, were the three scaled-up interventions that effectively reduced malaria burden (WHO, 2016).

Additionally, in 2016, the WHO and other international partners established a target for 2030 to achieve at least 90% reduction in global malaria mortality and cases compared to 2015. Despite these encouraging results, the WHO revealed in 2019 that there was no significant reduction on the global malaria cases from 2015-2017. On the contrary, high endemic areas reported a growing number of cases for the same period of time, which translated to an estimated 217 and 219 million cases in 2016 and 2017, respectively (WHO, 2018). Thus, there is an urgent need for strategies that contribute to the reduction of malaria cases. Besides the current strategies to accomplish the objectives towards a malaria free-world, the international community have focused their funding into five research areas: drugs, basic research, vaccines, vector control and diagnostics, in that order of priority (WHO, 2018).

The literature review of this thesis is focused on vaccine development, especially those strategies that target the liver-stage of the infection. In the first part of this chapter, general information about the life cycle of the parasite,

pathogenesis and how humans acquired immunity in endemic areas will be revised.

## 1.2 Biology of the malaria disease:

*Plasmodium* parasites have a complex life cycle characterized by alternation within two different hosts; female *Anopheles* mosquitoes and vertebrate hosts. Thus, parasites undergo a series of substantial cellular differentiations that allows them to survive in different organs, temperatures and environments within those hosts.

Malaria transmission and epidemiology relies greatly on the vector fitness. Longevity, density and resilience to environmental challenges determine the efficiency of the transmission by *Anopheles* spp. (Smith and McKenzie, 2004). *A. gambiae* is a robust and efficient vector distributed in sub-Saharan Africa, where *P. falciparum* predominates (Gething et al., 2011; Snow et al., 2005). In this high endemic region, infectious bites can reach 1000 per year. Morbidity and mortality are prevalent in children > 5 years, and clinical immunity to malaria can be developed during adulthood (Dondorp et al., 2008; Von Seidlein et al., 2012).

During the blood meal of an infected mosquito, sporozoites – the motile form of the parasite – are injected into the skin of the mammalian host. After they reach the bloodstream and infect hepatocytes, parasites are released to the blood stream once again where they replicate asexually. Sexual forms, which are able to infect mosquitoes later develop from blood stage parasites and the cycle is complete. The life cycle of the parasite is depicted in Figure 1.1.

### 1.2.1 Malaria liver stage

Experimental murine malaria infection starts with a blood meal of a malaria infected female *Anopheles* mosquito during which approximately 50-100 highly



motile sporozoites are inoculated into the skin of the murine host (Kebaier et al., 2009). Within a couple of hours, approximately half of the parasites glide out of the bite site and invade dermal blood or lymphatic vessels (Fig. 1.1) (Amino et al., 2006; Kebaier et al., 2009; Sidjanski and Vanderberg, 1997). The sporozoites that remain at the bite site beyond 3 hours have a decreased motility and are eventually destroyed by host factors (Amino et al., 2006; Kebaier et al., 2009). Trap-like protein (TLP), highly expressed by sporozoites (Howick et al., 2019; Moreira et al., 2008), has been found to be important during migration from the skin to the blood vessels, as mutant sporozoites fail to traverse epithelial cells *in vitro* and have lower infectivity *in vivo* (Moreira et al., 2008). Once in the blood stream, sporozoites can rapidly access the liver to establish infection (Yamauchi Lucy M. et al., 2007).

Before infecting hepatocytes, sporozoites cross the liver sinusoidal barrier by traversing through fenestrated endothelial cells and Kupffer cells (KC) (Tavares et al., 2013). Cell traversal activity involves breaching the host cell plasma membrane, gliding through the cytosol forming transient vacuoles and exiting the host cell again (Mota et al., 2001; Risco-Castillo et al., 2015). It has been recently demonstrated that GAPDH, a protein highly expressed across different stages of the *Plasmodium* life cycle, including sporozoites (Howick et al., 2019; Lindner et al., 2013), interacts with the KC receptor CD68 and promotes cell traversal (Cha et al., 2016). Cell-passage alters the plasma membrane integrity and can lead to cell death, preventing clearance by the KC, which might be exploited as a mechanism of immune evasion (Tavares et al., 2013). In fact, antibodies blocking *Plasmodium* GAPDH reduces hepatocyte invasion by sporozoites (Cha et al., 2016). Some other proteins involved in cell traversal include the sporozoite microneme protein essential for traversal (SPECT), SPECT2 and the cell traversal protein for ookinetes and sporozoites (CelTOS). SPECT is expressed by liver-infective sporozoites and it is required for the cell-passage through the liver sinusoidal cell layer (Ishino et al., 2004). SPECT2 is important for the egress of sporozoites from the transient vacuoles, which is thought to be associated with the presence of a pore forming complex domain

within the protein (Risco-Castillo et al., 2015). CelTOS is produced by liver-infective sporozoites, important for cell-passage activity *in vitro* and hepatocyte invasion *in vivo* (Kariu et al., 2006). Sporozoites can also traverse multiple hepatocytes before invading the final hepatocyte where they further develop.

During the last few decades, several molecular mechanisms involved in the host-parasite interactions during hepatocyte invasion have been described. Here, some of them will be discussed. Sporozoites are densely coated with several copies of the circumsporozoite protein (CSP) which is highly expressed by sporozoites and liver-stage parasites and is required for binding to highly sulphated forms of the heparan sulphate proteoglycans (HSPG) on the surface of hepatocytes (Herrera et al., 2015; Pinzon-Ortiz et al., 2001). After sporozoite binding to HSPG, CSP is cleaved by the parasite kinase CDPK-6, a step thought to be involved in a switch from migration to invasion mode of the sporozoite (Coppi et al., 2011, 2007). The thrombospondin-related anonymous protein (TRAP) is a transmembrane protein expressed by liver-infective sporozoites. It also binds HSPG and it is important for gliding motility *in vitro* and hepatocyte invasion *in vivo* (Kappe et al., 1999; Sultan et al., 1997). TRAP requires specific cleavage by rhomboid proteases to ensure the active invasion of hepatocytes by sporozoites (Baker et al., 2006; Ejigiri et al., 2012). The apical membrane antigen-1 (AMA-1) is known as a merozoite protein important for erythrocyte invasion (Anders et al., 1998; Thomas et al., 1984; Triglia et al., 2000). Additionally, AMA-1 is also expressed by sporozoites and while it seems to be dispensable for *P. berghei* liver infection (Giovannini et al., 2011), blocking antibodies can reduce hepatocyte invasion by *P. falciparum* sporozoites (Silvie et al., 2004; Yang et al., 2017).

A major component of the hepatocyte infection is the establishment of the parasitophorous vacuole membrane (PVM) of hepatocyte origin that sustain the further development of the parasite inside the host cell (Meis et al., 1983). The PVM not only provides a barrier between the developing parasite and the host cell cytoplasm but also mediates acquisition of nutrients. Some of the parasitic

proteins involved in the establishment of the PVM include P52 and P36 (Ishino et al., 2005; Labaied et al., 2007; van Schaijk et al., 2008). From the host side, three proteins expressed by hepatocytes have been found to play a role in cell invasion and the formation of the PVM. Transgenic mice and human cell lines lacking the scavenger receptor BI and the tetraspanin CD81, as well as the use of blocking antibodies can reduce sporozoite cell infection and affect the establishment of the PVM (Rodrigues et al., 2008; Silvie et al., 2007, 2003). Furthermore, antibodies binding the hepatocytic transmembrane receptor Ephrin 2 (EphA2) reduce the formation of the PVM and the subsequent development within the hepatocyte (Kaushansky et al., 2015). Evidence suggests that the *Plasmodium* protein P36 interacts with EphA2 (Kaushansky et al., 2015).

Once sporozoites have established hepatocyte invasion and reside within the PV, the PVM is extensively modified and parasites began to dedifferentiate into liver trophozoites (Jayabalasingham et al., 2010). Some of the proteins involved in the PVM modification include UIS3 and UIS4. They are highly transcribed by salivary sporozoites and translated during *Plasmodium* liver-stage development (Howick et al., 2019; Kaiser et al., 2004). Indeed, deletion of either UIS3 or UIS4 leads to early liver stage arrest but has no effect in the initial sporozoite invasion (Jobe et al., 2007; Mueller et al., 2007, 2005). Another protein present on the PVM is Hep17 in rodent malaria, also known as exported protein 1 (EXP1) in *P. falciparum* (Charoenvit et al., 1995; Doolan et al., 1996). This protein is not only expressed by schizonts and trophozoites during blood stage development but also by merozoites and liver stage parasites (Charoenvit et al., 1995; Howick et al., 2019). Defective parasites fail to develop intracellularly and it has been shown that EXP1 is important for nutrient intake (Mesén-Ramírez et al., 2019).

As liver trophozoites mature, they differentiate into schizonts containing merozoites (Fig. 1.1). PVM-associated proteins expressed during late liver stage development include liver specific protein 1 (LISP1) and LISP2 (Ishino et al., 2009; Orito et al., 2013). LISP1 is dispensable for blood stage and mosquito

development and it is required for PVM rupture of mature schizonts and the release of merozoites (Ishino et al., 2009). LISP2 can be exported to the host cytoplasm through the PVM and is important for merozoite development, as mutants have a reduce capacity to invade erythrocytes (Orito et al., 2013). It has been estimated in mice that up to 30,000 merozoites are released from every schizont into the bloodstream (Baer et al., 2007). The liver stage is usually asymptomatic, lasting approximately 2 days in mice and about 7 to 10 days in humans (Sturm et al., 2006). During *P. vivax* and *P. ovale* infections, parasites can remain in the liver for months or years in a dormant state (called the hypnozoite) that can cause relapses (Krotoski et al., 1982; Markus, 2011; Mikolajczak et al., 2015; Phillips et al., 2017; Soulard et al., 2015).

It is important to consider that during *Plasmodium* development, parasites display different sets of proteins at different stages of infection (Howick et al., 2019). Some of the proteins are stage-specific while others are expressed at multiple stages of the parasite development (Franke-Fayard et al., 2004; Koning-Ward et al., 1999). As a consequence of differences in temporal protein expression, antigens derived from individual proteins will also be subject to temporally regulated presentation. In this sense, the kinetics of presentation of each antigen is a dynamic process. Some antigens are only expressed by sporozoites, and as a consequence these may be presented shortly after hepatocyte invasion, but their presentation may wane as the antigen source is depleted. Other antigens, for example those derived from proteins only expressed by liver-stage parasites, might not be presented immediately upon sporozoite entry into hepatocytes as some time will be needed for protein synthesis. Presentation of this type of protein may thus be limited to the mid to late stages of liver-stage development. Blood-stage proteins will be expressed by merozoites as they develop at the end of the liver stage and might therefore represent antigens presented late in the liver stage. Finally, other proteins constitutively expressed in multiple stages might be presented both early and later during the liver stage infection. Thus, T cell control of liver-stage infection may be shaped not only by antigen expression levels, but by the kinetics of

antigen presentation for proteins expressed at different stages of the pre-erythrocytic development.

### *1.2.2 Malaria blood stage*

During the blood stage, parasite replication inside red blood cells (RBC) is extreme, cell death is abundant, and symptoms begin to occur. Merozoite invasion of RBC is a fast three-step process – involving pre-invasion, internalization and echinocytosis (Weiss et al., 2015) – that happens in approximately 2 minutes. The pre-invasion step is induced by low affinity protein interactions and the second step is mediated by several specific ligand-receptor interactions of molecules expressed on the surface of merozoites and host erythrocytes (Paul et al., 2015; Weiss et al., 2015). The echinocytosis occurs after internalization, when newly infected RBC (iRBC) shrink and adopt a shape with spikes.

Within erythrocytes, parasites remain in a parasitophorous vacuole and export a large number of effector proteins into the host cell cytoplasm (Charpian and Przyborski, 2008; Maier et al., 2009; Tilley et al., 2008). These molecules modulate parasite nutrient intake and cytoadherence of the host cell. Cell adhesion allows *P. falciparum*-infected erythrocytes to sequester in major organs, preventing circulation to the spleen, increasing the adhesion to other erythrocytes, and potentially leading to blood flow obstruction linked to severe clinical outcomes (Boddey and Cowman, 2013; Spillman et al., 2015). Malaria blood stage is a cyclic process of four differentiation steps (Fig. 1.1). During the first 5 h, the invading merozoite form a ring and start exporting proteins. From 10 to 40 h, parasites develop in trophozoites characterized by the enlargement of the food vacuole and maximized hemoglobin consumption. Next, parasites undergo nuclear division forming the schizont (40–48 h). Lastly, the host cell membrane collapses, realizing between 16 and 32 *P. falciparum* merozoites per infected erythrocyte and the cycle starts again (Boddey and Cowman, 2013; Spillman et al., 2015). In humans, after a week of asexual replication, when the

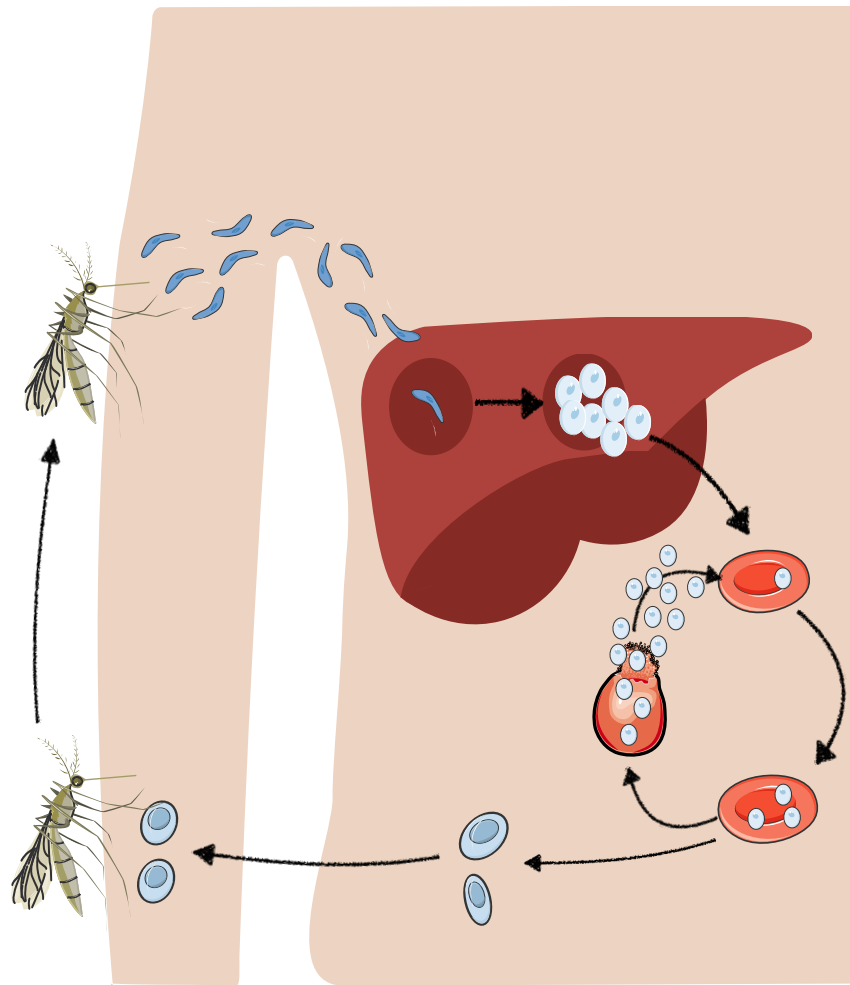


Figure 1.1. Plasmodium life cycle within the mammalian host. During the blood meal of a female infected mosquito, sporozoites are injected into the skin of the mammalian host. Within a few hours sporozoites reach the bloodstream and migrate to the liver. Before infecting hepatocytes, sporozoites cross the liver sinusoidal barrier by traversing through endothelial cells and KC (Tavares et al., 2013). Only a few sporozoites establish infection within hepatocytes, where they mature into schizonts producing merozoites that are then released into the bloodstream. The liver stage is usually asymptomatic, lasting approximately 2 days in mice and about 7 to 10 days in humans (Cowman et al., 2016). During blood stage infection, merozoites replicate asexually inside erythrocytes in a cyclic manner realising more merozoites each time with the capacity to infect other RBC. In humans, after a week of asexual replication, when the number of parasites in the blood is high cell death is abundant, and symptoms begin to occur. Eventually, some merozoites undergo a special differentiation step to become gametocytes with the capacity to infect mosquitoes. Within the mosquito gut, haploid gametes fuse to form zygotes (Sinden, 1983), which develop into oocysts, that produce sporozoites. Then they migrate to the salivary glands of the mosquito and get injected into the mammalian host during a blood meal to complete the cycle.

number of parasites in the blood stream reaches around 100 million, it is possible to detect these parasites by microscopy of blood smears and the symptoms begin to occur (Simpson et al., 2002).

Some merozoites undergo a special differentiation step to become gametocytes, a longer-lived sexual form with the capacity to infect mosquitoes (Fig. 1.1). Within the mosquito gut, haploid male and female gametes fuse to generate zygotes (Sinden, 1983), that develop into oocysts, which in turn divide asexually (Vlachou et al., 2006). Oocysts then release sporozoites that migrate to the salivary glands to start the transmission cycle again (Josling and Llinás, 2015).

### 1.3 Pathogenesis:

Uncomplicated malaria causes fever, mild anemia and jaundice, which is more common in adults, while an enlarged spleen and liver is more common in infants (Bartoloni and Zammarchi, 2012). Episodes of high fever have been associated with the systemic release of pro-inflammatory cytokines including tumour necrosis factor alpha (TNF $\alpha$ ) (Karunaweera et al., 1992; Wijesekera et al., 1996). Hemozoin, a detoxification product derived from the lysis of the haemoglobin, is accumulated in the parasite food vacuole inside infected erythrocytes (Coronado et al., 2014). One study revealed that after erythrocytes rupture releasing several merozoites, the hemozoin is up taken by myeloid cells, inducing the production of TNF- $\alpha$  in a toll-like receptor 9 (TLR9)-dependent manner (Coban et al., 2005).

#### 1.3.1 Severe malaria

Complicated malaria cases and mortality depend on age and transmission intensity. Most deaths in sub-Saharan Africa happen in children under 5 years of age where transmission is high. Severe malaria may develop in non-treated or

under-treated malaria and is mainly caused by extensive sequestration of infected erythrocytes in capillaries of vital organs. Four different syndromes are the main manifestation of severe malaria, cerebral malaria, severe anemia, respiratory distress and acidosis (MacKintosh et al., 2004; Wassmer et al., 2015).

Cerebral malaria can cause coma and although the precise pathogenic mechanisms remain to be understood, several studies suggest that the sequestration of iRBC is associated with limited blood flow through the brain capillaries and coma (MacPherson et al., 1985; Pongponratn et al., 2003; Silamut et al., 1999). Additionally, autopsy reports have demonstrated substantial accumulation of platelets in microvasculature of brains from children that died from cerebral malaria (Dorovini-Zis et al., 2011; Grau et al., 2003; Hochman et al., 2013). Blood-brain barrier disruption and brain swelling have also been implicated in the pathogenesis of human cerebral malaria (Dorovini-Zis et al., 2011; Greiner et al., 2015; Seydel et al., 2015). Based on murine studies, a simplified model has been suggested to explain the mechanisms underlying cerebral malaria (Ghazanfari et al., 2018). In this model, sequestration of iRBC and accumulation of platelets in the brain vasculature occurs first. This process is amplified by the intravascular accumulation of monocytes, neutrophils, T cells and natural killer (NK) cells in brains of mice with experimental cerebral malaria (ECM) (Belnoue et al., 2008, 2002; Hansen et al., 2007). This accumulation increases blood pressure and the risk of blood-brain barrier breakdown, leading to edema and swelling of the brain (Ghazanfari et al., 2018).

Severe anemia is mainly developed by infants and is caused by overlapping processes including excessive splenic removal of uninfected RBC (Buffet et al., 2011), lysis of iRBC by schizogony disruption (Perkins et al., 2011; Price et al., 2001), and ineffective erythropoiesis (Dörmer et al., 1983; Phillips et al., 1986). Respiratory distress is particularly present in pregnant woman infected with *P. falciparum* or *P. vivax* (Nosten et al., 1999; Taylor et al., 2012). It has been suggested that pulmonary edema may be mediated by inflammation driven by endothelial and cytoadherence damage (Charoenpan et al., 1990; Corbett et al.,



1989; Pongponratn et al., 2003). Metabolic acidosis develops from the excessive production of organic acids, such as lactic acid, that result from anaerobic glycolysis in tissues and organs with low supply of blood and oxygen (Maitland and Newton, 2005). Patients with acidic breathing have poor prognosis, as at this stage acidosis is suggestive of major organ failure (Hanson et al., 2010; Marsh et al., 1995).

#### 1.4 Naturally acquired immunity:

Despite decades of efforts to control malaria, it still is a major public health concern, which can damage economic development in many disadvantaged communities. Aspects that limit the progress in the field are: (i) the development of parasite resistance to almost all antimalarial drugs, (ii) the mosquito resistance to vector insecticides and the impracticality of those interventions for application in certain regions lacking economic support, (iii) the social instability in war and conflict areas where the health system is fragile, (iv) and failure to develop an efficient vaccine.

The complex life cycle of the parasite and the variety of clinical manifestations of the disease in the host, represent a major challenge to understand how people in endemic regions become protected against severe malaria, and hence, how to develop effective vaccine strategies. It has been suggested that two major factors contribute to the development of naturally acquired immunity, the age of the host and the parasite transmission intensity (Doolan et al., 2009).

Virtually all naïve individuals of all ages develop clinical malaria at relatively low levels of parasitemia (numbers of circulating parasites) during the first infection (Doolan et al., 2009; Langhorne et al., 2008). In the case of *P. falciparum*, naturally acquired immunity to clinical malaria develops with age in countries with high parasite transmission. Remarkably, children under 6

months of age are protected from severe malaria through inhibitory factors found in the breast milk (Kassim et al., 2000), and the passive transfer of maternal immunoglobulin (Ig) G before birth (Hviid and Staalsoe, 2004). Morbidity and mortality increases after 6 months of age and cerebral malaria peaks in infants ranging from 2 to 4 years of age and is usually associated with high levels of parasitemia (Okiro et al., 2009; Reyburn et al., 2005). Beyond 5 years of age, the risk of severe malaria begins to diminish although parasitemia levels may still be high. During adolescence, clinical disease frequency decreases, deaths caused by malaria are rare and the average number of circulating parasites decreases as well (Carneiro et al., 2010; Griffin et al., 2015; Roca-Feltrre et al., 2010). Adults become asymptomatic carriers with low parasitemia levels (Doolan et al., 2009; Marsh and Kinyanjui, 2006). In adults, naturally acquired immunity protects against clinical malaria by maintaining low parasitemia levels, helping to prevent the onset of severe disease (Baird et al., 2003; Owusu-Agyei et al., 2001). Sterilizing immunity, i.e. no blood stage parasites, never develops and clinical immunity is not long lasting (Doolan et al., 2009; Keegan and Dushoff, 2013). Importantly, studies on communities migrating from low to hyper-endemic areas revealed that adults developed naturally acquired immunity more rapidly than children (Baird, 1995; Baird et al., 1998), suggesting that immunity depends on intrinsic factors related to age, like immune system maturity (Kurtis et al., 2001).

The intensity of parasite transmission also influences the development of acquired immunity. In areas of low to intermediate transmission, the acquisition of clinical immunity is delayed (Carneiro et al., 2010; Marsh and Kinyanjui, 2006). Children living in these areas remain at a constant risk of severe malaria even after 5 years of age (Carneiro et al., 2010; Mbogo et al., 1995, 1993; Rodriguez-Barraquer et al., 2018; Snow et al., 1997). Some adults may develop severe malaria in rare cases when transmission intensity transitions from high to extremely high (Doolan et al., 2009; Marsh and Kinyanjui, 2006).

The exact immunological mechanisms involved in the development of naturally acquired immunity remain to be elucidated. Many studies have reported

a correlation between IgG antibodies (Ab) against blood-stage parasites and the reduced risk of severe malaria (Dodoo et al., 2001; Magistrado et al., 2007; Ofori et al., 2002; Yone et al., 2005). Ab against sporozoites, the liver-stage parasitic form, seem to have a limited or negligible role in naturally acquired immunity (Hoffman et al., 1987; Schofield et al., 1987). In fact, immune adults from highly endemic areas were able to control parasitemia and avoid symptoms after direct injection of blood-stage parasites (Bruce-chwatt, 1963; Langhorne et al., 2008).

### 1.5 Malaria vaccines:

As mentioned previously, exceptional progress has been made to control malaria and a substantial reduction in mortality has been achieved in endemic countries. The scale up of interventions such as the distribution of antimalarial drugs, insecticide-treated bed nets, and indoor insecticide spraying, are the major contributors to this progress (WHO, 2016). However, emerging resistance to both insecticides and antimalarial medications have stalled the mortality reduction and threaten the sustainability of these interventions (WHO, 2018). Thus, the development of a highly efficacious vaccine is urgent to complement the current malaria elimination efforts.

Despite decades of intense research, a highly effective malaria vaccine for application in endemic areas remains elusive (Coelho et al., 2017; Draper et al., 2018). Currently, a wide range of vaccine approaches are in pre-clinical development and clinical trials. These strategies target different stages of the *Plasmodium*-life cycle, including pre-erythrocytic (PE) forms, blood-stage parasites and gametocytes.

#### 1.5.1 Blood-stage vaccines:

The rationale behind this type of vaccines is that they aim at triggering a similar immune response induced by naturally acquired immunity. These strategies face several challenges related to immune evasion mechanisms, such

as antigenic variation of both free merozoites as well as in the surface proteins of infected erythrocytes (Ferreira et al., 2004).

While subunit approaches targeting merozoite derived antigens have failed to confer protection (Miura, 2016), whole parasite strategies have conferred immunity in murine and primate models (Good et al., 2013; Raja et al., 2016). In these studies, blood-stage parasites were chemically attenuated by incubation with compounds that irreversibly alkylates the DNA of the parasite (Good et al., 2013). Immunised mice developed long-lasting protection against other strains and *Plasmodium* and it was mediated by CD4 T cells (Good et al., 2013; Raja et al., 2017, 2016). Promising results in *Aotus* monkeys using chemically attenuated blood-stage *P. falciparum* (De et al., 2016) prompted the implementation of a pilot experiment in malaria-naïve volunteers. Preliminary results revealed that the vaccine is safe, well tolerated, and that it induces the production of homologous and heterologous antibodies as detected in the sera of participants (Stanisic et al., 2018).

In addition to the whole parasite approach, recent studies on the novel antigen *P. falciparum* RH5 have demonstrated its capacity to induce neutralizing Ab that block merozoite invasion *in vitro* (Douglas et al., 2015, 2011; Illingworth et al., 2019). The production of neutralizing Ab was also observed *in vivo* in studies with *Aotus* monkeys (Hjerrild et al., 2016). Now this protein is being targeted in phase I and II clinical trials. The expression of the RH5 protein has proven to be particularly difficult and therefore a different approach has been taken to successfully target it. For this, a replication-deficient chimpanzee adenovirus (ChAd) and an attenuated modified vaccinia virus Ankara (MVA) have been engineered to encode the RH5 *P. falciparum* protein sequence (Payne et al., 2017). Preliminary results in malaria naïve volunteers confirmed the induction of RH5 Ab, B cells and T cell responses, and some degree of growth inhibition *in vitro* mediated by Ab found in the sera of vaccinated volunteers (Payne et al., 2017).

### 1.5.2 Transmission-blocking vaccines:

Vaccines that aim at inducing immunity against parasite stages that infect mosquitoes, e.g. gametocytes, are known as transmission-blocking vaccines. Less progress has been made in this field but there are some antigens that have been subjected to investigation in the recent years. In animal models, epitopes of the Pfs25 surface protein expressed in mosquito stage of *P. falciparum* have demonstrated capacity to induce Ab with transmission blocking activity (Miura et al., 2007). In preliminary human trials, soluble Pfs25 has proved to be poorly immunogenic and in formulation with the adjuvant Montanide<sup>1</sup> ISA 51, induced unexpected adverse events (Wu et al., 2008). During a second effort, Pfs25 was conjugated to the carrier protein ExoProtein A from *Pseudomonas aeruginosa* and formulated together with the adjuvant Alhydrogel<sup>®</sup> (Radtko et al., 2017; Talaat et al., 2016). While this vaccine was well tolerated, functional activity of Ab in serum was only achieved after four vaccine doses and Ab titres waned rapidly after each dose (Talaat et al., 2016).

### 1.5.3 Pre-erythrocytic (PE) vaccines:

PE vaccine strategies are directed to induce immune responses against sporozoites prior to hepatocyte invasion and liver-stage parasites before being released to the blood stream. These vaccine strategies have priority as they i) avoid the occurrence of disease symptoms, ii) stop the development of clinical malaria and iii) block transmission. Prevention of egress to the blood-stage is called sterilizing immunity.

Immunization against PE stages has been investigated since the 1960s in murine models (Nussenzweig et al., 1967) and 1975 when Clyde and collaborators achieved protection in human volunteers that were previously immunized by bites from irradiated mosquitoes (Clyde 1990). At that time, vaccination was only possible via mosquito bites, which represented a major obstacle for the design and implementation of safer approaches. During the

1980s, efforts switched to a more suitable and safer approach based on the production of recombinant proteins derived from sporozoites, for use as subunit vaccines.

To date, the most advanced sub-unit vaccine candidate for *P. falciparum* sporozoites is the RTS,S/AS01. In fact, a pilot implementation program for this vaccine is currently being undertaken in Ghana, Kenya and Malawi (Adepoju, 2019). This vaccine targets the CSP of *P. falciparum* (PfCSP) and is a virus-like particle with several copies of the C-terminal domain of PfCSP fused to hepatitis B virus surface antigen and a potent liposomal adjuvant called AS01 (RTSS Clinical Trials Partnership, 2015). The RTS,S/AS01 vaccine demonstrated partial clinical immunity during a phase III clinical study performed in seven countries of sub-Saharan Africa (RTSS Clinical Trials Partnership, 2015). For a period of two years, children (5-17 months age) and young infants (6-12 weeks age) were vaccinated with 3 doses of RTS,S/AS01 or control vaccine. In children, vaccine efficacy against clinical and severe malaria was 36% and 32%, respectively; while in young infants it only reached 25% and 17%. Importantly, results also indicated that the protection wanes rapidly over time (Olotu et al., 2013). In fact, vaccine efficacy after 7 years, in a small cohort of children enrolled in the phase II, was only 4.4% (Olotu et al., 2016). Antibodies against the CSP protein have been implicated as the potential mechanism mediating the immunity induced by the RTS,S/AS01 vaccine (Oyen et al., 2017; White et al., 2015).

Other PE vaccines are based on administration of whole attenuated sporozoites and have demonstrated high efficacy in mice and humans. These strategies include immunisation with radiation-attenuated sporozoites, genetically attenuated parasites and infection with living sporozoites in conjunction with chemoprophylaxis.

#### 1.5.4 Radiation-attenuated sporozoites (RAS) based vaccines:

Early studies in mice and humans demonstrated the efficacy of RAS to confer protection through mosquito bites (Clyde, 1990; Nussenzweig et al., 1967). Since then, substantial efforts have been made to develop a safe product that can be used in clinical trials. The PfSPZ vaccine (*P. falciparum* sporozoite) are sporozoites manually isolated from the salivary glands of infected and irradiated mosquitoes, aseptically and cryopreserved (Hoffman et al., 2010).

In 2013, Seder and collaborators demonstrated that whereas four doses protected 33% of the volunteers, five doses of  $1.35 \times 10^5$  PfSPZ protected 100% of the subjects. They also showed that high doses of RAS enhanced protection since 55% of individuals vaccinated with four doses of  $2.7 \times 10^7$  PfSPZ remained protected after 21 weeks of vaccination (Seder et al. 2013). These results suggest that there is a critical threshold of RAS numbers required to achieve effective immunity. The researchers also showed that immunity was correlated with Ab titres and specific T cell responses (Seder et al. 2013). Moreover, the PfSPZ vaccine must be administered intravenously as other routes have limited immunogenicity (Ishizuka et al., 2016). While these studies demonstrated substantial protection in malaria-naïve volunteers, RAS vaccination demonstrated limited efficacy in malaria-exposed subjects from Mali and Equatorial Guinea (Olotu et al., 2018; Sissoko et al., 2017). Two major factors may influence the immunity mediated by RAS vaccination in malaria-exposed subjects. Firstly, malaria-naïve volunteers were protected against homologous challenge, while subjects from Mali and Equatorial Guinea were exposed to naturally transmitted heterologous parasites (Epstein et al., 2017). Second, some studies support the idea that pre-existing blood-stage infections are able to induce immune suppression, thus affecting the establishment of appropriate immune responses against sporozoites (Rénia and Goh, 2016).

### 1.5.5 Genetically-attenuated parasites (GAP) based vaccines:

Contrary to RAS, genetically-attenuated parasites arrest at specific stages of the pre-erythrocytic development. It has been demonstrated in murine models that late arrest of liver-stage parasites potentiates protection by enhancing the breadth of the immune response (Butler et al., 2011). Pilot experiments in humans have demonstrated that administration of a triple knock out (KO) parasite (known as PfGAP3KO) through mosquito bites was safe and able to induce the production of inhibitory Ab to sporozoites found in the sera of vaccinated volunteers (Kublin et al., 2017). The PfGAP3KO parasite lacks the *P. falciparum* *p52*, *p36* and *sap1* genes which induces early pre-erythrocytic arrest (Kublin et al., 2017; Mikolajczak et al., 2014). Engineering human parasites that arrest late during liver-stage development has been difficult because these parasites often become unviable in the mosquito. Moreover, there are safety concerns related to breakthrough infections with GAP that could lead to blood stage infections. However, substantial efforts are currently being made to provide such vaccine (Vaughan et al., 2018)

### 1.5.6 Chemoprophylaxis vaccination (cVac):

These strategies involve infections with susceptible live wild type sporozoites under host drug treatment, which eliminates blood stage parasites. This approach is considered one of the most potent strategies since it allows full liver development – increasing the spectrum of antigens that the host is exposed to. In fact, murine studies have demonstrated the superior capacity of cVac to induce protection, since fewer and lower doses were required to induce sterilizing immunity compared to RAS immunisation (Belnoue et al., 2004). Clinical trials confirmed the superior capacity of cVac as only three doses of  $5.12 \times 10^4$  sporozoites were sufficient to confer 100% protection (Mordmüller et al., 2017). This is 10 to 100 times less than with the PfSPZ immunisations (Seder et al. 2013). Assessment the cVac approach in endemic areas is underway.



## 1.6 Malaria liver-stage immunity:

While whole parasite-based vaccines are efficient at inducing protection in humans, these strategies have major limitations that question their suitability for routine use in endemic areas. Some of those challenges are associated with technical and logistical issues of large scale sporozoite production and distribution in endemic areas. There are also safety issues related to breakthrough infections with GAP or the use of a live fully virulent pathogen that demands strict attachment to drug administration regimes, as is the case with cVac.

Animal studies using pre-erythrocytic vaccines have provided essential knowledge about the immune mechanisms that govern the induction of sterilizing immunity. Deep understanding of these mechanisms provide important insights for the design and development of vaccine strategies that are more suitable for implementation in endemic areas.

The next section of the chapter will be focused on how animal studies have revealed memory CD8 T cell responses as the major mediator of sterilizing immunity. Additionally, innate mechanisms involved in the recognition of liver-stage parasites will be briefly discussed.

### *1.6.1 Innate response to liver-stage infection:*

For many years, the liver-stage of malaria was considered immunologically silent because it is asymptomatic, and the liver is considered as an immunoprivileged organ (Prudêncio et al., 2006). Recent murine studies have proved this is not the case, showing that liver-stage parasites are recognised after infection (Epiphany et al., 2008; Leiriao et al., 2005; Liehl et al., 2014; Van De Sand et al., 2005). While parasites inside hepatocytes are poorly recognized by macrophages and dendritic cells (DC), hepatocytes recognize them through cytosolic pattern recognition receptors (PRR) and are able to initiate an antipathogen type I interferon (IFN) response (Liehl et al., 2014; Miller et al.,

2014). Liehl and collaborators showed that the innate sensing of *Plasmodium* nucleic acids activates the innate immune response and restricts parasite replication in the liver. They first found that the replication of parasites inside mouse livers was accompanied by a significant induction of interferon stimulated genes (ISGs) compared to non-infected mice. Additionally, they showed that *Plasmodium* RNA triggered a type I IFN response that is dependent of melanoma differentiation-associated protein 5 (MDA5) and independent of retinoic acid-inducible gene I (RIG-I), TLR3 and TLR4. In the same study, they also found that live sporozoites induced a higher type I IFN response compared to heat inactivated or live irradiated parasites (Liehl et al., 2014). The activation of type I IFN response leads to a wide range of events, including not only expression of ISG, but also production of chemokines that mediate recruitment of macrophages, neutrophils and lymphocytes to the site of infection. It also induces generation of IFN- $\gamma$  that promotes infiltration of NK cells to the liver (Liehl et al., 2014; Miller et al., 2014).

NK cells are important effector cells of the innate immune system and defend the host during the early phase of pathogenic infections (Korbel et al., 2004). Studies with a murine malaria model have shown that after *P. yoelli* sporozoite infection, the numbers of NK cells decreases in the spleen and increases in the liver (Roland et al., 2006). Functional *in vitro* assays have revealed their cytotoxic capacity to inhibit parasite development within hepatocytes but not against blood-stage parasites (Roland et al., 2006). NKT cells express effector mechanisms similar to NK cells and a semi-invariant T cell receptor (TCR). In steady state, 20 to 30% of murine liver lymphocytes are NKT cells (Bendelac et al., 1997), and when activated, they exert potent antitumor responses (Nakagawa et al., 1998). Infection with high numbers of genetically-attenuated *P. yoelli* sporozoites lead to the recruitment of NKT cells to the liver (Miller et al., 2014). When they were stimulated *in vivo* by treating mice with  $\alpha$ -galactosylceramide ( $\alpha$ -GalCer), an NKT cell-specific ligand, they mediated reduction of liver-stage parasite development for a short period of time (Gonzalez-Aseguinolaza et al., 2000). Moreover, it has been also suggested that

activation of NKT cells promote the accumulation of parasite-specific CD8 T cells (Gonzalez-Aseguinolaza et al., 2000; Kortzen et al., 2005). Even though NKT cells were initially suggested to reduce liver-stage development, studies in CD1d KO mice, which lack NKT cells, have shown that CD8 T cell immune responses and levels of protection induced by RAS vaccination is independent of NKT cells (Carvalho et al., 2002; Overstreet et al., 2011; Romero et al., 2001).

### *1.6.2 T cell priming:*

During natural infection by mosquito bites, T cell activation is first detected in the skin draining lymph node (LN) at 48 h post infection and not in spleen, liver or liver-draining LN (Chakravarty et al., 2007). The removal of the skin draining LN prior to mosquito bite-infection reduced the number of activated T cells in the liver by over 60%; and in the absence of both LN and spleen, this reduction reached 80% (Chakravarty et al., 2007). Remarkably, sporozoites can be detected by PCR in draining LN as soon as 3 h post intradermal inoculation (Yamauchi Lucy M. et al., 2007). After intradermal inoculation of murine sporozoites, antigen-specific CD8 T cells cluster around lymph-node resident XCR1 cDC in draining LN at early time points (Radtke et al., 2015). Using immunofluorescence microscopy, Radtke et al. revealed that the two cells are in direct physical contact, indicating that CD8 T cell priming may be mediated by LN-resident CD8<sup>+</sup> DC. Interestingly, when sporozoite antigen presentation is restricted to the skin draining LN (by using a mouse strain in which sporozoites cannot establish liver stage infection), T cell priming at this site was sufficient to induce protective immunity against malaria (Obeid et al., 2013). Priming of CD8 T cells has been observed in the spleen and liver after intravenous (i.v.) administration of sporozoites. In these studies, activated CD8<sup>+</sup> T cells accumulate in spleen between days 4 and 5 after i.v. injection with either live sporozoites or RAS (Lau et al., 2011; Sano et al., 2001).

Although, priming of CD8 T cells can occur in the liver (Bertolino et al., 2001), the role of liver-resident cells as antigen presenting cells (APC) of

*Plasmodium* antigens is controversial. Liver DC are located in close proximity to portal veins, relatively far from non-fenestrated endothelial cell layer (Lau and Thomson, 2003). With this distribution it is unlikely liver DC can mediate intrahepatic naïve CD8 T cell activation (Holz et al., 2010). Additionally, hepatic DC exhibit an immature phenotype, suggesting these cells collect antigen from blood circulating in the liver and then migrate to draining LN for further maturation and effector functions, rather than inducing T cell activation *in situ* (Thomson et al., 2002). In hepatocytes, the sporozoite protein CSP can be degraded and presented on major histocompatibility complex (MHC) I to CD8 T cells (Bongfen et al., 2007; Cockburn et al., 2011), however they cannot prime the naïve CD8 T cells. Recently, a population of liver monocyte-derived CD11c<sup>+</sup> APC was found to acquire *Plasmodium* parasites from dying infected hepatocytes (Kurup et al., 2019). Importantly, hepatic CD11c<sup>+</sup> cells harbouring *Plasmodium* parasites were also found in the liver draining LN where they primed CD8 T cells responses to liver-stage malaria antigens (Kurup et al., 2019).

Following vaccination, naïve CD8 T cells can be primed and then form memory T cells in the liver for detection of infected hepatocytes when natural infection occurs. Although exclusive liver homing molecules remained to be identified, it has been suggested that CXCL16, which is highly expressed by liver sinusoidal endothelial cells, may influence the retention of immune cells in this organ, including NK T cells expressing CXCR6 (Geissmann et al., 2005). In a murine vaccination study, wild-type (WT) mice were transferred with WT or *Cxcr6*<sup>-/-</sup> OT-I cells, transgenic CD8 T cells specific for an ovalbumin (OVA) antigen, and then vaccinated with RAS expressing SIINFEKL, the antigen recognised by OT-I cells. Results revealed 75% reduction of OT-I cell numbers in the liver of mice transferred with CXCR6-deficient cells compared to mice receiving WT cells (Tse et al., 2014). Moreover, two studies have demonstrated the importance of antigen presentation by hepatocytes for the immunity mediated by CD8 T effector cells. During the first study, bone marrow chimera experiments revealed that APC are essential for priming of parasite-specific CD8 T cells and are dispensable for CD8 T cell effector function in the liver, which is mediated by hepatocytes (Chakravarty

et al., 2007). For the second study, transporters associated with antigen processing (TAP) knockout mice, which lack antigen presentation by APC were co-transferred with infected hepatocytes and effector CSP-specific T cells. These mice were efficiently protected from infection, suggesting that effector cells are able to exert their function by hepatic antigen presentation independently of APC antigen presentation (Balam et al., 2012).

### 1.6.3 Memory CD8 T cells:

Early studies in 1980s first demonstrated the critical role of CD8 T cell in immunity mediated by RAS vaccination (Schofield et al., 1987; Weiss et al., 1988). Since then, several animal studies have confirmed these findings (Cockburn et al., 2014; Doll et al., 2016; Epstein et al., 2011; Fernandez-Ruiz et al., 2016; Ishizuka et al., 2016; Nganou-Makamdop et al., 2012). Other reports also confirmed that CD8 T cells are crucial for the immunity induced by a GAP vaccine based on a double-knockout *P. berghei* parasite for *UIS3* and *UIS4*, which are highly expressed by sporozoites (Jobe et al., 2007).

The number of infected hepatocytes under a normal course of infection has been estimated to be only 1 in one million cells in mice and one in 100 million cells in humans (Van Braeckel-Budimir and Harty, 2014). Accordingly, the generation of highly effective parasite-specific CD8 T cells required for the surveillance of the whole liver and the elimination of every single infected hepatocyte (prerequisite of sterilizing immunity) is essential. In 2008, Schmidt and collaborators induced sterilizing immunity in BABL/c mice by immunisations with mature DC coated with a *P. berghei* antigen and boosting with a recombinant attenuated *L. monocytogenes* strain expressing the same antigen. They found that sterilizing immunity was correlated with the number of specific CD8 T cells and estimated that the threshold of specific cells is one million in the spleen and 20,000 in the liver to achieve protection against malaria. These numbers correspond to a 100-1000-fold higher dose of specific T cells required for sterilizing immunity against bacterial and viral infections (Schmidt et al., 2008).

In the context of a primary infection, productive activation of naïve T cells induces clonal expansion, providing a large army of specific effector cells (Joshi et al., 2007). After the elimination of infected cells, the effectors contract and memory T cells survive – providing long-term protection against reinfection (Sarkar et al., 2008). In fact, parasite-specific short-lived effector cells peak in spleen shortly after *P. yoelii* infection and decrease with clearance, while memory precursor effector cells remain and develop into true memory T cells over time (Chandele et al., 2011). Memory cells are functionally and phenotypically different and can be grouped into central, effector and tissue resident memory T cells (Fig. 1.2).

Central memory T ( $T_{CM}$ ) cells home to secondary lymphoid organs (SLO), express high levels of CD62L and CCR7, have the ability to circulate to the blood, retaining highly proliferative features, providing a stem-like pool of antigen-specific memory T cells (Sallusto et al., 1999). Little evidence supports the idea that  $T_{CM}$  cells have an important role in immunity against sporozoites (Holz et al., 2016).

Effector memory T ( $T_{EM}$ ) cells express low levels of CD62L and CCR7, important molecules for homing to SLO, but possess receptors that allow them to enter non-lymphoid tissues (Sallusto et al., 1999). These cells patrol the body and are loaded with proteins to exert several effector functions. In C57BL/6 (B6) mice vaccinated with *P. berghei* RAS, the majority of circulating memory cells (96%) are  $T_{EM}$  cells (Berenzon et al., 2003). BALB/c mice, which are relatively easy to protect, generate high frequencies of CSP-specific  $T_{EM}$  cells, whereas B6 mice, which are more difficult to protect, produce a lower proportion of specific  $T_{EM}$  cells and these are not specific for CSP (Schmidt et al., 2010).

Tissue-resident memory T cells ( $T_{RM}$  cells) are found in several organs and tissues including intestines (Masopust et al., 2010), skin (Gebhardt et al., 2009), brain (Wakim et al., 2010), lungs (Anderson et al., 2012; Teijaro et al., 2011), female reproductive tract (Shin and Iwasaki, 2012), salivary glands

(Hofmann and Pircher, 2011), thymus (Hofmann et al., 2013), liver (Fernandez-Ruiz et al., 2016; Tse et al., 2013), kidneys, pancreas and heart (Casey et al., 2012). T<sub>RM</sub> cells remain in the tissues, maintaining an effective local surveillance at sites of previous infection, especially at barrier sites in the body (Steinbach et al., 2018). These cells have been effectively studied in parabiosis experiments where the circulatory systems of two mice are joined, circulatory cells balanced between the mice while a proportion of T cells do not recirculate and thus are tissue resident (Jiang et al., 2012; Klonowski et al., 2004). In the case of secondary infections, the strategic location of the T<sub>RM</sub> cells allows them to mediate effector responses more rapidly than their counterpart, the circulating memory cells (Mueller and Mackay, 2016). T<sub>RM</sub> cells have a unique molecular profile that distinguishes them from the circulatory memory cells. They express low levels of Killer cell lectin-like receptor subfamily G member 1 (KLRG1) (Gebhardt et al., 2009; Mackay et al., 2012a; Steinert et al., 2015). Additionally, high levels of CD103 has been observed on many cells with some exceptions (Carbone, 2015), and CD69, which is also a marker of recent activation (Gebhardt et al., 2009). The differential expression of these markers promotes the residency of these cells in different tissues.

T<sub>RM</sub> cells exert effective killing of infected cells through many effector mechanisms, including the expression of granzyme B (Casey et al., 2012; Mackay et al., 2012b; Schenkel et al., 2014) and the production of proinflammatory cytokines such as IFN- $\gamma$  (Glennie et al., 2015; Schenkel et al., 2013), promoting the recruitment of circulating T cells and B cells (Schenkel et al., 2014). The production of IFN- $\gamma$  can also alert neighbouring cells and promote the upregulation of innate functions that prevent viral infections (Gil et al., 2001; Karst et al., 2003; Kimura et al., 1994; Shrestha et al., 2006). Upon re-activation with cognate stimulus, CD8 T<sub>RM</sub> also express TNF $\alpha$  and promote DC maturation, leading to granzyme B upregulation in NK cells (Schenkel et al., 2014). The presence of T<sub>RM</sub> cells has been associated with tumour and infection control (Masopust and Soerens, 2019). The generation and maintenance of T<sub>RM</sub> cells is

the target of intensive research efforts for the development of more potent vaccines.

Liver T<sub>RM</sub> cells have been recently described and differently from T<sub>RM</sub> cells from other tissues, these cells are found in the blood patrolling the liver sinusoids and are much faster than skin T<sub>RM</sub> cells (Fernandez-Ruiz et al., 2016; Mackay et al., 2016; Steinert et al., 2015). Intravital microscopy and sequencing data of after RAS vaccination, have revealed that the elimination of *Plasmodium*-infected hepatocytes is associated with the effector function of CD8 T cells that display matching features with T<sub>RM</sub> cells (Cockburn et al., 2013; Tse et al., 2013). In fact, Fernandez-Ruiz and collaborators confirmed by parabiosis experiments in B6 mice that *P. berghei* ANKA (PbA) RAS induced the formation of liver T<sub>RM</sub> cells (Fernandez-Ruiz et al., 2016). They first investigated the transcriptional profile of different memory T cell subpopulations present in the liver and the spleen after vaccination with PbA RAS. A specific subpopulation only present in the liver was found to express CXCR6, CD69, CXCR3 and low levels of KLRG1 and CD62L, common markers for T<sub>RM</sub> cells (Fernandez-Ruiz et al., 2016). Subsequently, parabiosis experiments confirmed the tissue resident status of this subpopulation and intravital images showed that they patrol the liver sinusoids. Importantly, formation of large numbers of liver T<sub>RM</sub> cells can be promoted through different vaccination strategies (Fernandez-Ruiz et al., 2016). Depletion using anti-CXCR3 antibodies abrogated the sterile protection mediated by vaccination, providing evidence that liver T<sub>RM</sub> cells are essential for protection against liver-stage parasites (Fernandez-Ruiz et al., 2016).



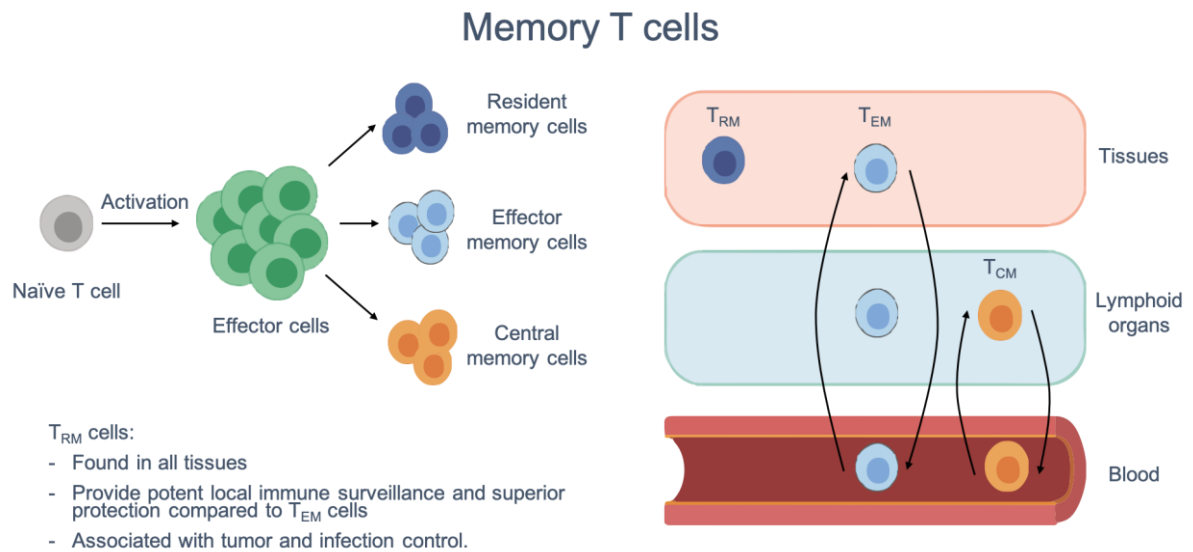


Figure 1.2. Subtypes of memory T cells. In the context of a primary infection, productive activation of naïve T cells induces clonal expansion, providing a large army of specific effector cells. After the elimination of infected cells, the effectors contract and memory T cells survive providing long-term protection against reinfection. Memory cells are functionally and phenotypically different and can be grouped into central, effector and tissue resident memory T cells.  $T_{CM}$  cells home to SLO, express high levels of CD62L and CCR7, have the ability to circulate to the blood, retaining highly proliferative features, providing a stem-like pool of antigen-specific memory T cells (Sallusto et al., 1999).  $T_{EM}$  cells express low levels of CD62L and CCR7, important molecules for homing to SLO, but possess receptors that allow them to enter non-lymphoid tissues (Sallusto et al., 1999). These cells patrol the body loaded with proteins to exert several effector functions.  $T_{RM}$  cells remain in the tissues, maintaining an effective local surveillance at sites of previous infection, especially at barrier sites in the body (Steinbach et al., 2018). They can be virtually found in all tissues and induce a superior protection compared to effector memory cells. The presence of  $T_{RM}$  cells has been associated with tumour and infection control.

In healthy adults,  $\gamma\delta$  T cells constitute 1 to 5 % of circulating lymphocytes, whereas higher frequencies can be found in epithelial-rich tissues, such as the reproductive track, skin and intestine (Hayday, 2000; Holtmeier and Kabelitz, 2005). These cells display features that are common for innate and adaptive immune cells and the exact biological functions remain to be elucidated (Carding and Egan, 2002). Emerging studies have suggested the importance of  $\gamma\delta$  T cells in the induction of protective responses mediated by RAS vaccination in humans and murine models (Ishizuka et al., 2016; Zaidi et al., 2017).

All these data demonstrate that sterile protection against malaria mediated by vaccination largely depends on the formation of parasite-specific memory CD8 T cells that kill infected hepatocytes displaying cognate antigen. Of the several memory CD8 T cell subpopulations, liver T<sub>RM</sub> cells appear to be essential for sterile protection induced by RAS vaccination against sporozoite challenge (Fernandez-Ruiz et al., 2016). These cells are strategically located in liver sinusoids to provide potent surveillance and are armed with effector mechanisms to eliminate infected cells. Because liver T<sub>RM</sub> cell formation can be induced through vaccination, the design of strategies aiming at the generation of effective T<sub>RM</sub> cells is a priority.

### 1.7 Liver-stage subunit vaccines:

Liver-stage subunit vaccines are designed to generate CD8 T cell immune responses against specific antigens derived from liver-stage parasites. These vaccines overcome the manufacturing challenges of whole-parasite approaches but face their own difficulties, such as limitations in the number of known immunogenic and protective CD8 T cell epitopes derived from *Plasmodium* parasites (Draper et al., 2018; Schmidt et al., 2010). Noting the shortage of known epitopes, this section of the chapter will cover the existing development of liver-stage subunit vaccines.

Viral vectored vaccines are to-date the most clinically tested liver-stage subunit vaccines, since they are known to be particularly effective at inducing T cell responses. They use replication-deficient viral particles, in particular recombinant chimpanzee adenovirus serotype 63 (ChAd63) for priming and modified vaccinia Ankara (MVA) for boosting (Ewer et al., 2015). These viral particles are modified to encode for the TRAP linked to a multi-epitope string (ME-TRAP). The ME is a string of T and B cell epitopes from several expressed during pre-erythrocytic stage (Ewer et al., 2015; Gilbert et al., 1997), and TRAP is a protein that plays an important role in hepatocyte cell invasion (Akhouri et al., 2008). Vaccine efficacy ranges from 20% to 25% in malaria naïve volunteers (Ewer et al., 2013), and was found to reduce the risk of infection in adults in Kenya by 67% during 8 weeks of monitoring (Ogwang et al., 2015). However, no efficacy was observed in Senegalese adults (Mensah et al., 2016). Vaccination of children in two cohorts from Gambia and Burkina Faso revealed high immunogenicity (Bliss et al., 2017), but no efficacy was observed in the highly endemic malaria transmission cohort from Burkina Faso (Tiono et al., 2018). Interestingly, other studies in malaria naïve subjects revealed that co-administration of ChAd63-MVA encoding for other well-known epitopes, such as apical membrane antigen-1 (AMA1) and merozoite surface protein-1 (MSP1) together with the ME-TRAP vaccine, failed to improve protection (Hodgson et al., 2015; Sheehy et al., 2012).

Another clinically tested strategy uses DNA plasmids for priming and adenovirus for boosting (Sedegah et al., 2011; Tamminga et al., 2011; Wang et al., 2001; Webster et al., 2005). Immunisations with DNA plasmids and adenovirus vector encoding for CPS and AMA1 resulted in 27% of sterile protection in malaria naïve volunteers; this immune response was associated to effector CD8 T cells (Chuang et al., 2013).

#### 1.7.1 Prime-and-trap vaccine:

“Prime-and-trap” is a three-component vaccine strategy used to induce parasite-specific liver  $T_{RM}$  cell formation through the priming of naïve immune cells and

the trapping of those cells in the liver (Fig. 1.3) (Fernandez-Ruiz et al., 2016). The first component is based on the use of an  $\alpha$ Clec9A monoclonal Ab (mAb) coupled to a relevant epitope to achieve epitope delivery to conventional DC-1 (cDC1) (Caminschi et al., 2008; Lahoud et al., 2011); which is the main antigen presenting cell stimulating T cells in malaria (Fernandez-Ruiz et al., 2017; Lau et al., 2014). The Clec9A molecule, also known as DNGR-1, improves cross-presentation of dead-cell-associated antigens (Ahrens et al., 2012; Iborra et al., 2012; Sancho et al., 2008). Its expression is restricted to DCs and is highly expressed by CD8<sup>+</sup> DCs and CD103<sup>+</sup> DCs in mice and in humans by the BDCA-3<sup>+</sup> DCs (Caminschi et al., 2008; Desch et al., 2011; Huysamen et al., 2008; Sancho et al., 2008; Schreibelt et al., 2012). Moreover, the in vivo delivery of exogenous antigens by targeting Clec9A with mAb has been exploited successfully in the context of tumour therapy and vaccine development (Caminschi et al., 2012a; Palucka et al., 2010; Tacke et al., 2007). Importantly, cytotoxic T lymphocyte (CTL) responses in mice injected with  $\alpha$ Clec9A coupled to specific antigens is mediated by co-injection with adjuvants such as cytosine-phosphate-Guanine (CpG), polyinosinic-polycytidylic acid (p I:C) and anti-CD40 (Caminschi et al., 2012b; Lahoud et al., 2011; Sancho et al., 2008). In the absence of adjuvant, this form of immunisation leads to MHC-II presentation, follicular helper T cell activation and efficient antibody production, but no CD8 T cell response (Caminschi et al., 2012b; Kato et al., 2015; Lahoud et al., 2011; Li et al., 2015).

The second component of the prime and trap vaccine is a recombinant adeno-associated virus (rAAV) engineered to selectively infect hepatocytes, inducing the presentation of antigen on hepatocytes and trapping of the T cells to the liver. This vector is also modified to express the relevant epitope under the control of a hepatocyte-specific promoter (Tay et al., 2014). This enables the presentation of antigen on hepatocytes to facilitate the trapping of the T cells to the liver for formation of T<sub>RM</sub> cells. The third and last component, is a CpG adjuvant used for the licensing of DC involved in CD8<sup>+</sup> T cell priming (Lahoud et al., 2011) and to promote a proinflammatory liver-microenvironment

## Prime-and-trap

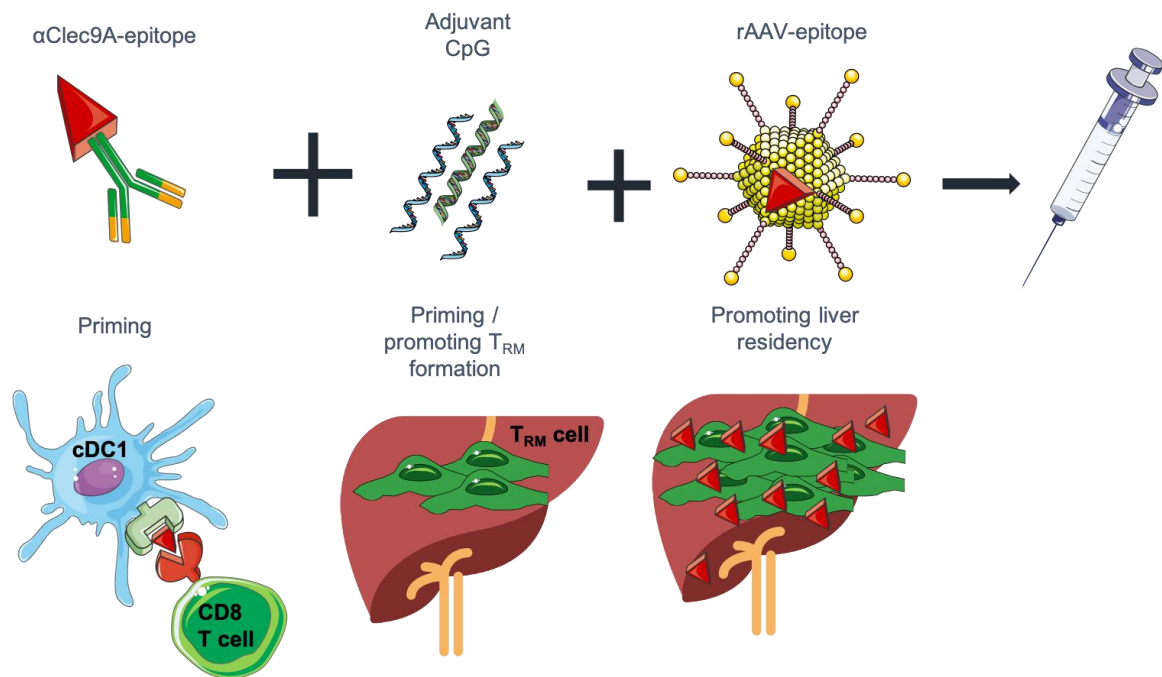


Figure 1.3. Prime-and-trap vaccine. Prime-and-trap is a three-component vaccine strategy for the induction of liver  $T_{RM}$  cell formation. The first component is an  $\alpha$ Clec9A mAb coupled to a relevant epitope for delivery to cDC1. The adjuvant CpG is used for the licensing of DC and for the induction of a proinflammatory liver-microenvironment that promotes the accumulation and formation of  $T_{RM}$  cells. The trap component is based on the use of a recombinant adeno-associated virus (rAAV) that expresses the antigen in the liver enhancing the differentiation of  $T_{RM}$  cells in the liver. Three components are delivered i.v. to improve targeting the antigen to the liver.

(Huang et al., 2013), which aids T<sub>RM</sub> cell formation (Holz et al., 2018; Mackay et al., 2012a). Importantly, all three components are delivered i.v., which has been associated with improved protection (Seder et al., 2013). Fernandez-Ruiz and collaborators demonstrated that relatively high numbers of liver T<sub>RM</sub> cells can be generated by prime-and-trap vaccination and that these cells can effectively confer sterilizing immunity (Fernandez-Ruiz et al., 2016).

### 1.7.2 Prime-target vaccine:

The prime-target approach (Gola et al., 2018) is similar to “prime-and-trap” and uses a modified adenovirus for the priming step and nanoparticles (Np) or a modified viral vector (MVA) for cell targeting to the liver. In detail, mice were primed intramuscularly (i.m.) with a modified adenovirus expressing OVA and two weeks later Np containing OVA were administered i.v. (Gola et al., 2018). They observed increased numbers of specific-CD8 T cells expressing IFN- $\gamma$  in the livers of vaccinated mice, as well as the formation of cell clusters displaying T<sub>RM</sub> cell markers, and colocalization of IFN- $\gamma$ <sup>+</sup> granzyme B<sup>+</sup> CD8 T cells in proximity to sporozoites. Vaccinated mice were sterilely protected from challenge with sporozoites expressing OVA and protection was abrogated by the use of anti-CD8-depleting Ab (Gola et al., 2018). In an effort to translate this approach using clinically relevant epitopes, they also investigated protection induced by the *P. falciparum* antigens TRAP, liver-stage antigen 1 (LSA1) and liver stage-associated protein 2 (LSAP2). For this, the ChAd63 vector was engineered to express the respective antigens, mice were primed i.m. and two weeks later mice received either same ChAd63 i.v. or MVA i.m. for targeting the CD8 T cells to the liver. Three weeks later mice were challenged with *P. berghei* sporozoites expressing the relevant *P. falciparum* antigens (Gola et al., 2018). Again, mice were sterilely protected, these encouraging results prompted a phase I clinical trial to assess the safety and immunogenicity of this approach by vaccinating volunteers i.v. with ChAd63.ME-TRAP (Gola et al., 2018). Preliminary results of this phase I clinical trial showed that intravenous administration of ChAd63.ME-TRAP was safe and well tolerated. Moreover, they also showed that IFN- $\gamma$

production measured by enzyme-linked immunospot (ELISPOT) picked at day 14 after vaccination and was still detectable on day 42 in volunteers that received the highest dose of the viral vector ( $5 \times 10^{10}$  vp) (Gola et al., 2018).

### *1.7.3 CD8 epitopes derived from PbA:*

As previously mentioned, one of the major limitations for the development of liver-stage subunit vaccines is the scarcity of known protective T cell epitopes derived from *Plasmodium* parasites. The paucity of known epitopes could be partially explained by the difficulty in isolating infected hepatocytes and then identify epitopes via elution and mass spectrometry. As mentioned before, under a normal course of liver-stage infection, the percentage of infected hepatocytes is low (Van Braeckel-Budimir and Harty, 2014), and therefore the number of MHC I molecules that can be obtained from these cells is also low. Recently, efforts have been made to generate a source of sufficient epitopes, e.g. via use of humanized liver-chimeric mouse models and liver-stage cell culture methods (Longley et al., 2015).

To date, only two MHC-I restricted epitopes from PbA have been described to confer some protection against liver stage infection in B6 mice, a mouse strain that has been considered difficult to protect (Doolan and Hoffman, 2000; Schmidt et al., 2010). The first characterized epitope was the PbTRAP<sub>130-138</sub>. Hafalla and collaborators found that B6 mice vaccinated with a modified adenovirus expressing PbTRAP<sub>130-138</sub> showed 95% reduction in liver parasite load, but this vaccine did not induce sterile protection (Hafalla et al., 2013).

The second epitope, from PBANKA\_1031000<sub>631-639</sub>, is termed Kb-17 and is contained within a protein of unknown function; the MIF4G-like protein (Pichugin et al., 2018; Speake et al., 2016). Pichugin and colleagues showed that B6 mice primed intramuscularly with Kb-17-DNA plasmid and boosted intravenously with a recombinant adenovirus expressing Kb-17 had 7-fold less

liver parasite burden compared to control mice, but no sterile protection was observed (Pichugin et al., 2018).

PbT-I cells are transgenic MHC I-restricted cells specific for PbA, which can confer sterile protection against sporozoite challenge in B6 mice (Lau et al., 2014). These cells have the ability to respond to blood-stage PbA, *P. yoelii* XNL and *P. chabaudi*, and to PbA RAS parasites. Initial attempts to identify the authentic epitope recognized by the PbT-I cells were unsuccessful (Lau et al., 2014). From initial screenings, an agonist mimic epitope for activation of PbT-I cells was identified but was not present in the *Plasmodium berghei* genome (Fernandez-Ruiz et al., 2016). This epitope was then used as the target antigen for the prime-and-trap vaccine developed by Fernandez-Ruiz and collaborators (Fernandez-Ruiz et al., 2016). They adoptively transferred B6 mice with PbT-I cells and vaccinated them with  $\alpha$ Clec9A mAb conjugated to the mimic epitope, together with CpG and the modified adenovirus expressing this epitope. Mice immunised with this prime-and-trap vaccine formed substantial numbers of liver PbT-I  $T_{RM}$  cells and developed high levels of sterile protection against PbA (Fernandez-Ruiz et al., 2016). The authentic cognate epitope for the PbT-I cells has now been identified – and Chapter 3 of this thesis will focus on the characterization of the memory response induced by prime-and-trap immunisation using this epitope.

### 1.8 Adjuvants:

As a general principle, the majority of vaccines administered to humans already contain or are formulated together with an adjuvant. Vaccine adjuvants are substances that enhance the immunogenicity of a vaccine when added or mixed to it (Apostólico et al., 2016; Di Pasquale et al., 2015). They are used for several purposes including (i) to reduce the number of doses and amount of antigen required to induce protective immunity, (ii) increase the rate of



seroconversion and induce protective responses more rapidly (Coffman et al., 2010).

In many cases, adjuvants were widely used based on empirical data due to the lack of understanding of the mechanisms of action. However, over the last years, several mechanisms of action have been described. Adjuvants enhance immunity by (i) modulating the kinetics of the vaccine release, increasing the antigen lifetime, (ii) stimulating the release of cytokines by immune or non-immune cells, (iii) increasing antigen uptake and maturation of APC, and (iv) inducing local inflammation for cellular recruitment (Awate et al., 2013; Coffman et al., 2010; Di Pasquale et al., 2015). Based on the mechanism of action, adjuvants can be classified into delivery systems and immune potentiators.

#### *1.8.1 Delivery systems*

These types of adjuvants are employed as antigen carriers and they can induce local inflammation for the recruitment of immune cells to the site of injection. These adjuvants include aluminium salts, microparticles and lipid particles. Alum is used with vaccines against hepatitis A, B, diphtheria/tetanus/pertussis (DTP), human papillomavirus (HPV) and *Haemophilus influenzae* type B (FDA, 2018). For several years it was thought that alum works by inducing the formation of a depot of antigen reducing the rate of antigen release (Gupta et al., 1995). On the contrary, recent studies revealed that independently of the depot formation, alum can activate an innate immune response by stimulating the NLRP3/NALP3 inflammasome complex (Eisenbarth et al., 2008; Kool et al., 2008). Moreover, adjuvants that contain aluminium preferentially induce humoral over cellular immune responses (Lindblad, 2004).

Microparticles including virus-like particles are generated by structural viral proteins and are non-infective and non-replicative (Huret et al., 2013). They can deliver antigen directly to immune cells such as DC (Bosio et al., 2004; Da Silva

et al., 2007), activate B cells (Zhang et al., 2009), induce cross-presentation to CD8 T cells (Dalba et al., 2007), and thus promote a broad range of humoral and cellular immune responses (Buonaguro et al., 2002; Smith et al., 2013). Some vaccines against hepatitis B (GlaxoSmithKline, 2012) and HPV (GlaxoSmithKline, 2015) are based in virus-like particles. Poly (lactic acid) (PLA) and poly (lactic-co-glycolic acid) (PLGA) are biodegradable polymers that can form micro and nanoparticles to encapsulate antigen and adjuvants in same particle, increasing the delivery efficiency (Danhier et al., 2012). PLGA particles are taken up by endocytosis and can reach the cytosol soon after administration (Vasir and Labhassetwar, 2007). They have also been associated with increased uptake by APC (Beaudette et al., 2009), and antigen cross-presentation (Shen et al., 2006). PLGA particles have been used in clinical trials to deliver antigens from *Bacillus anthracis* (Manish et al., 2013) and hepatitis B virus (HBV) (Thomas et al., 2011).

MF59 is a water-in-oil emulsion and the mechanism of action remains unknown. It induces powerful cellular and humoral responses (Stephenson et al., 2005) by stimulating macrophages, monocytes and DC to secrete chemokines which in turn induce leukocyte recruitment (Seubert et al., 2008). MF59 is currently used as adjuvant for the flu vaccine (Wilkins et al., 2017) and it was approved for the H1N1 pandemic vaccine (O'Hagan et al., 2013). AS03 is an oil-in-water emulsion that contains  $\alpha$ -tocopherol and induces the production of NF- $\kappa$ B, recruitment of immune cells and the induction of high Ab titres (Garçon et al., 2012; Morel et al., 2011). AS03 has shown promising results in clinical trials of vaccines against influenza in immunised children (Díez-Domingo et al., 2015).

### 1.8.2 Immune potentiators

Immune potentiators activate innate immune responses by stimulating pattern-recognition receptors (PRR) that are widely expressed on immune and non-immune cells. Some of those PRR include TLR, nucleotide-binding oligomerization domain-like receptors (NLR) and RIG-I like receptors (RLRs). Binding of PRR with their respective agonists triggers the activation and

maturation of immune cells which in turn modulate the production of cytokines and chemokines that ultimately can lead to induction of adaptive immune responses (Akira et al., 2006; Kumar et al., 2011). For the following paragraphs, the function of several PRR ligands will be discussed in the context of vaccine adjuvants. Based on their location, PRR can be classified as plasma membrane, endosomal or cytosolic sensors (Fig. 1.4).

TLR4 is expressed on the plasma membrane of human macrophages and dendritic cells (Vaure and Liu, 2014). A broad range of pathogen-associated molecular patterns (PAMP) are able to trigger TLR4 mediated responses. However, the most common ligand is lipopolysaccharide (LPS) which enhances type 1 T helper (Th1) immune responses (Steinhagen et al., 2011). Stimulation of TLR4 signals via the adaptor protein myeloid differentiation primary response protein 88 (MYD88) and the TIR-domain-containing adapter protein inducing interferon IFN- $\beta$  (TRIF), which in turn lead to the production of cytokines (Vaure and Liu, 2014). LPS is quite toxic, even at low doses, therefore safe TLR4 ligands has been recently designed. Monophosphoryl lipid A (MPL) is a non-toxic LPS derivate that induces the production of high amounts of TNF $\alpha$ , IL-12 and IFN- $\gamma$  (Mata-Haro et al., 2007). AS04 is an approved adjuvant consisting of MPL and alum used in vaccines against stage IV melanoma (Melacine®) (Cluff, 2013), HPV (Giannini et al., 2006) and HBV (Tong et al., 2005), and is part of clinical trials for leishmaniasis, malaria and Herpes antigens (NIH, 2020).

### *1.8.3 Endosomal sensors:*

Endosomal PRR include TLR3, TLR7/8 and TLR9 (Fig. 1.4), which are selectively expressed by a subset of innate immune cells (Junt and Barchet, 2015). Poly I:C is a well-known TLR3 ligand that leads to NF- $\kappa$ B activation and IL-12 production but can also bind MDA5 inducing IFN $\alpha$  release (Kato et al., 2006; Kumar et al., 2006). Stimulation with p I:C activates DC for the upregulation of MHC II, production of IL-12 and type I IFN (Choi et al., 2012; Davey et al., 2010; Longhi et al., 2009). While IL-12 activates NK cells for the production of

IFN- $\gamma$ , which in turn activates T and B cells (Lieberman et al., 1991), type I IFN responses have been associated with Th1 responses and antigen cross-presentation (Schulz et al., 2005). Altogether, p I:C modulates APC maturation, antigen presentation and ultimately T and B cell responses. Moreover, p I:C has been tested as an adjuvant for vaccines against several pathogens including HIV (Trumpfheller et al., 2008), dengue (Henriques et al., 2013) and cancer (Ammi et al., 2015). These studies demonstrated that p I:C enhances the generation of a robust immune response mediated by CD4 (Trumpfheller et al., 2008) and CD8 T cells (Henriques et al., 2013; Trumpfheller et al., 2008) in mice vaccinated with antigen DC-targeted antibodies, increasing protection (Henriques et al., 2013; Trumpfheller et al., 2008).

TLR7 detects single stranded (ss) GU-rich RNA (Lund et al., 2004) and is differentially expressed by human pDC and B cells, and by murine pDC, B cells, monocytes, cDC2 and neutrophils (Junt and Barchet, 2015). Ligand binding of TLR7 induces signalling through MyD88, activating the NF- $\kappa$ B pathway (Kawai and Akira, 2007). Imiquimod is a synthetically produced molecule that mimics ssRNA and can trigger TLR7 (Harrison et al., 1994; Shukla et al., 2010) and it has been associated with Th1, CD8 T cell and CTL responses (Stanley, 2002). Imiquimod has been approved for topical use in the treatment of skin conditions (Berman and Caperton, 2011) and as an adjuvant against melanoma cancer (Adams et al., 2008; Shackleton et al., 2004; Swetter et al., 2015).

CpG oligodeoxynucleotides (ODN) are synthetic DNA molecules containing unmethylated CG motifs able to trigger responses via TLR9. The pattern of TLR9 expression differs between humans and mice. For instance, in mice, TLR9 is expressed by B cells and various DC subtypes, including cDC and pDC (Edwards et al., 2003; Krieg et al., 1995), whereas in humans, TLR9 is mainly expressed by pDC and B cells (Hornung et al., 2002; Jarrossay et al., 2001; Kadowaki et al., 2001; Rothenfusser et al., 2001). Given these variations in the expression profile of TLR9 and the distinctive role played by diverse DC subtypes, different immunological outcomes could be expected in humans and

mice. Moreover, early studies have shown that CpG ODN sequences are species-specific, meaning that mice and humans have different optimal ligands (Bauer et al., 1999, 2001; Hartmann et al., 1999; Krieg et al., 1995). Therefore, it is important to carefully distinguish between types of CpG ODN, their respective roles and different immunological outcomes when used as vaccine adjuvants.

There are four types of CpG ODN, A-, B-, C- and P-class. The following data is based on studies investigating the role of different CpG ODN in B cells and pDC, which are known to express TLR9 in both, humans and mice (Edwards et al., 2003; Hornung et al., 2002; Kadowaki et al., 2001; Krieg et al., 1995). A-class CpG ODN induce strong production of IFN $\alpha$  by pDC, leading to the activation of cellular antiviral state as shown by studies performed in human PBMC and murine splenocytes (Guiducci et al., 2006; Hemmi et al., 2003). B-class ODN triggers pDC differentiation and the production of TNF $\alpha$ , which in turn leads to T cell activation. Additionally, B-class ODN induces B cells to proliferate and secrete IgM (Hartmann and Krieg, 2000; Hemmi et al., 2003; Rothenfusser et al., 2001; Verthelyi et al., 2001). C-class combines features from both A and B-class, as it stimulates B cells to secrete IL-6 and pDC to produce IFN $\alpha$  (Hartmann et al., 2003; Marshall et al., 2003). P-class ODN is the last described CpG ODN and has been shown to produce more IFN $\alpha$ , IL-6 and the CXCR3 ligand CXCL10 compared to B- or C-class ODN (Samulowitz et al., 2010). Stimulation with CpG ODN ultimately leads to the production of proinflammatory cytokines such as IL-6, IL-12, IL-18 and TNF $\alpha$ , inducing the maturation of DC and the upregulation of costimulatory molecules such as CD40, CD80 and CD86 (Bayik et al., 2016; Klinman, 2004).

From all the endosomal sensor ligands, CpG ODN have been extensively tested in several clinical trials. In fact, CpG ODN have been demonstrated to induce strong humoral responses in vaccine studies against HBV (Davis et al., 1998) and cellular T cell responses against *Leishmania* (Raman et al., 2012; Verthelyi et al., 2002), influenza (Beignon et al., 2002), anthrax (Minang et al., 2014) and tumours (Muraoka et al., 2010; Speiser et al., 2005; Stern et al., 2002). Moreover,

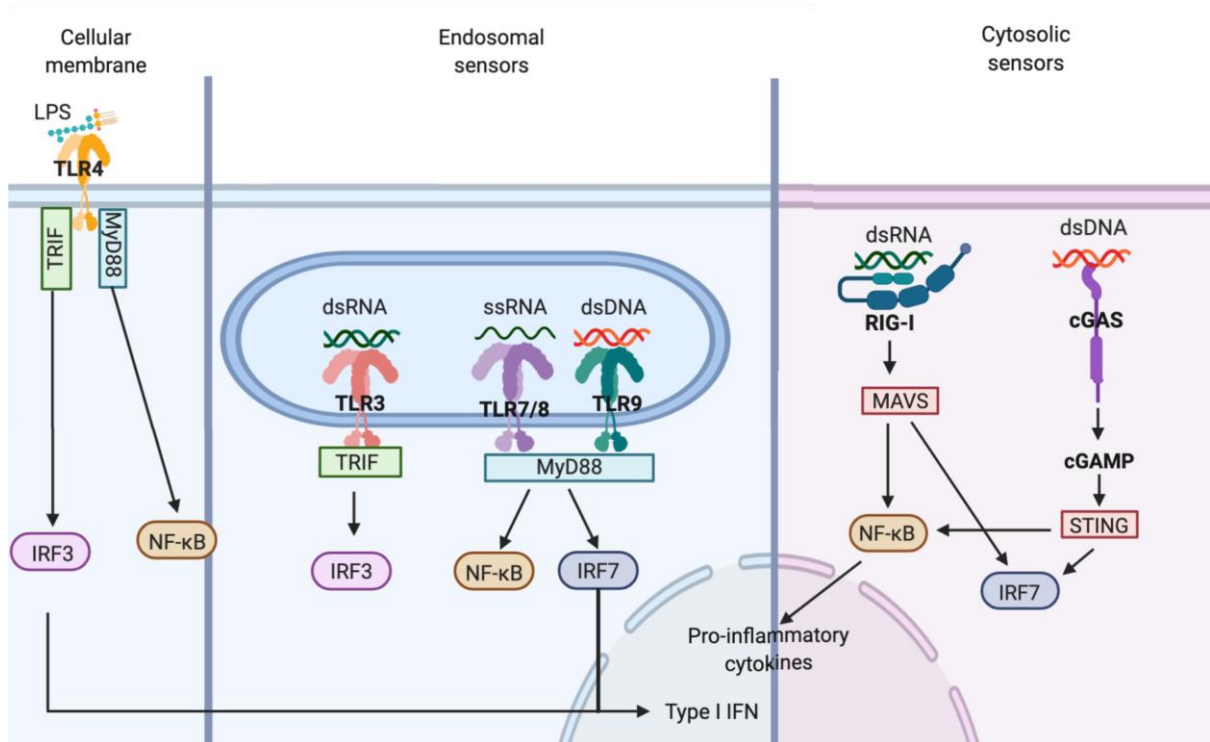


Figure 1.4 Immune potentiators: TLR4 is expressed on the plasma membrane and can get activated by LPS. Endosomal sensors include: TLR3 that recognizes dsRNA structures like p I:C, TLR7/8 binds to ssRNA and TLR9 which recognizes dsDNA with CpG motifs. Some of the best described cytosolic receptors include i) RIG-I-like receptors that get activated with dsRNA and signal through the adaptor protein MAVS; and ii) cGAS that upon recognition of dsDNA produces cGAMP, which is an activator of STING. TLR4 and TLR3 stimulates TRIF which in turn activates the transcription factor IRF3. TLR7/8, TLR9 and TLR4 signal through MyD88 protein leading to the activation of the transcription factors IRF7 and NF-κB, which can also get stimulated by MAVS and STING. Ultimately, activation of IRF3 and IRF7 induces the expression of type I IFN stimulated genes and NF-κB of pro-inflammatory cytokines. (Figure made in BioRender®).

CpG ODN also has the ability to promote the formation of intrahepatic aggregates of myeloid cells that facilitate the expansion of CD8<sup>+</sup> T cells in the liver (Huang et al., 2013).

#### *1.8.4 Cytosolic sensors:*

In contrast to the TLRs, cytosolic receptors are ubiquitously expressed in immune and non-immune cells. Cytosolic sensors include RIG-I-like receptors (RLR), cyclic GMP–AMP (cGAMP) synthase (cGAS), stimulator of interferon genes (STING), and absent in melanoma 2 (AIM2).

RLR are a family of DExD/H box RNA helicases that upon activation signal through a cascade pathway leading to an antiviral response mediated by type I IFN (Loo and Gale, 2011). RIG-I and MDA5 are members of this family. RIG-I recognizes short double stranded (ds) RNA bearing 5' triphosphorylated or 5' diphosphorylated ends (Hornung et al. 2006; Goubau et al. 2014); whereas self mRNA is excluded by the presence of a methyl group at the first base pair (Schuberth-Wagner et al. 2015). MDA5 binds p I:C, which mimics the structure of a high molecular weight dsRNA (Kato et al., 2008). After RIG-I and MDA5 engagement, the adaptor protein mitochondrial antiviral-signalling (MAVS) is triggered. This event leads to activation of the transcription factors interferon-regulatory factor 3 (IRF3), IRF7 and NF- $\kappa$ B, which in turn induce the production of IFN $\beta$ , IFN $\alpha$  and interferon stimulated genes (Loo and Gale, 2011). Studies have demonstrated that use of a RIG-I ligand as adjuvant for an influenza vaccine enhanced Ab affinity, conferred protection against challenge, and reduces the amount of antigen required for protection in mice (Kulkarni et al., 2014). Additionally, a recent report revealed that activation RIG-I agonists promotes cross-presentation enhancing CTL responses in mice vaccinated against an adenovirus (Hochheiser et al. 2016).

cGAS binds dsDNA and upon activation it produces cGAMP (Ablasser et al., 2013; Diner et al., 2013; Gao et al., 2013), which functions as a second

messenger that binds the adaptor STING (Burdette et al., 2011; Burdette and Vance, 2013; Danilchanka and Mekalanos, 2013). Cyclic dinucleotides are PAMP derived from bacteria (Tamayo et al., 2007) and like cGAMP, can stimulate STING, which in turn activates the transcription factors NF- $\kappa$ B and IRF3 to induce type I IFN and cytokines (Tanaka and Chen, 2012; Wu et al., 2013). *In vivo* studies showed that mice treated with cyclic dinucleotides controlled bacterial infections with *Klebsiella pneumoniae* and *Staphylococcus aureus* by activating innate immune cells (Karaolis et al., 2007b, 2007a). Immunisation with the model antigen OVA in conjunction with cyclic dinucleotides as adjuvant enhanced T cell activation and Ab production (Li et al., 2013). Of clinical relevance, it has been demonstrated that cyclic dinucleotides used as adjuvant in murine vaccine studies promotes humoral immunity against extracellular bacteria such as *S. aureus* and *Streptococcus pneumoniae* (Ebensen et al., 2007; Hu et al., 2009; Ogunniyi et al., 2008; Yan et al., 2009). Additionally, cyclic dinucleotides has been used as adjuvant for the generation of protective T cell responses against *Mycobacterium tuberculosis* (Van Dis et al., 2018), influenza virus (Madhun et al., 2011) and breast cancer metastasis (Chandra et al., 2014). Importantly, *in vitro* studies with human PBMC demonstrated that cGAMP derivatives enhance antigen specific expansion and functional maturation of effector CD8 T cells (Gutjahr et al., 2019).

Finally, AIM2 binds dsDNA, triggering the activation of the inflammasome machinery resulting in activation of caspase-1, the release of IL-1 and IL-18 and programmed cell death (Bürckstümmer et al., 2009; Fernandes-Alnemri et al., 2009; Hornung et al., 2009; Roberts et al., 2009). Data from studies in *Aim2*-deficient mice suggest the inflammasome activation mediated by AIM2 is essential for host defence against the cytosolic bacteria *Francisella tularensis* and *Listeria monocytogenes* (Fernandes-Alnemri et al., 2010; Sauer et al., 2010) and DNA viruses such as vaccinia virus and mouse cytomegalovirus (Rathinam et al., 2010). Although this research implicates AIM2 as an important mediator of defence against infection, the role of AIM2 ligands as vaccine adjuvants have not been addressed.



### 1.8.5 Adjuvants for malaria vaccines:

While adjuvants from the delivery system category have been tested in several clinical trials of vaccines against malaria, only a few studies have investigated the effect of immune potentiators. Early studies in malaria naïve volunteers revealed that RTS,S vaccine in an oil-in-water emulsion enhanced the production of IgG compared to RTS,S containing alum. Importantly, when the vaccine was in emulsion plus the immune stimulant MPL (TLR4 ligand), vaccine efficacy was further enhanced (Stoute et al., 1997). In other studies including emulsion adjuvants, strong humoral and T cell responses were triggered, but they failed to enhance protection against sporozoite challenge or the trials had to be discontinued due to reactogenicity (Malkin et al., 2008; McCarthy et al., 2011; Pierce et al., 2010; Wu et al., 2008).

In a phase I clinical trial, epitopes of the CSP were co-expressed with HBV core structure particles that self-assemble into virus-like particles. Immunised volunteers developed high Ab titres and elicited CS-specific T cells that produce IFN- $\gamma$  (Nardin et al., 2004). Pre-clinical studies in mice also revealed that immunisation with PLGA nanoparticles containing VMP001 *P. vivax* antigen induced Ab with enhanced avidity and affinity, and stronger T cell responses compared to the mice vaccinated with soluble antigen (Moon et al., 2012).

In the context of immune potentiators, studies in non-human primates revealed that p I:C used as vaccine adjuvant for CSP immunisation, enhanced the production of long-lasting high affinity IgG, and reduced the liver parasite load in mice (Tewari et al., 2010). Recently, in a macaque study, CpG or R848 (a TLR7/8 ligand) encapsulated in PLGA based particles were co-administered together with the poorly immunogenic Pfs25 antigen. Results demonstrated that both TLR-ligands enhanced the production of long-lasting Ab that also exhibit higher affinities, and higher numbers of B and CD4 T cells (Thompson et al., 2018). Additionally, in a phase I clinical trial against blood-stage *P. falciparum* malaria, CpG ODN was used as an adjuvant together with AMA1 antigen,

inducing significantly higher titres of IgG compared to participants that received vaccine alone (Mullen et al., 2008). The naturally occurring glycolipid,  $\alpha$ Gal-Cer, has been also used as an adjuvant for a whole parasite-based vaccine (Gonzalez-Aseguinolaza et al., 2002).  $\alpha$ Gal-Cer has the capacity to activate NKT cells which are present at high frequencies in the liver. It was shown that co-administration of this compound together with RAS enhances the number of IFN- $\gamma$  producing splenocytes and vaccine efficacy in mice (Gonzalez-Aseguinolaza et al., 2002).

#### *1.8.6 Adjuvants for $T_{RM}$ cell formation:*

There is minimal research that suggests immune potentiators can act as adjuvants for driving  $T_{RM}$  cell formation. Early studies demonstrated that in the absence of in situ antigen recognition, inflammation driven by external agents such as 1-fluoro-2,4-dinitrobenzene (DNFB) is enough for the generation of long-lasting  $T_{RM}$  cells in the skin (Mackay et al., 2012a). In fact,  $T_{RM}$  cell formation can be driven by inflammatory stimuli in the absence of cognate antigen in other tissues such as the female reproductive tract (Shin and Iwasaki, 2012), gut (Bergsbaken and Bevan, 2015), lung (Caminschi et al., 2019) and liver (Holz et al., 2018).

It has been demonstrated that CpG ODN enhance the formation of liver  $T_{RM}$  cells in mice vaccinated with antigen targeted to DC (Fernandez-Ruiz et al., 2016; Holz et al., 2018). This enhancement might occur because CpG ODN induces a pro-inflammatory microenvironment in the liver that supports the accumulation of CD8 T cells (Huang et al., 2013). To the author's knowledge there are not further studies on adjuvants that induce the formation of liver  $T_{RM}$  cells. CpG has been also used as adjuvant in a skin model, where mice were vaccinated subcutaneously with OVA and CpG ODN was administered epicutaneously as adjuvant. Results revealed that these vaccinated mice had an increased number of skin  $T_{RM}$  cells and enhanced protection against melanoma-

OVA expressing cells compared to mice immunised without CpG adjuvant (Lai et al., 2019).

Cytokines can also be used as immune potentiator adjuvants. In fact, it has been recently reported that mice immunised intranasally with adenoviral replication-deficient vectors expressing the influenza antigen hemagglutinin and nucleoprotein together with IL-1 $\beta$  developed more lung specific T<sub>RM</sub> cells and increased immunity against viral challenge (Lapiente et al., 2018).

In a pre-clinical study in young macaques, a TLR7/8 agonist was assessed as an adjuvant for the generation of HIV-specific T<sub>RM</sub> cells in the vagina. The prime and boost vaccine regime consisted of three components, first, viral vectors encoding for HIV-Gag protein for the generation of specific CD8 T cells, HIV-Env protein for the generation of Ab and PLGA nanoparticles containing a TLR7/8 agonist (Petitdemange et al., 2019). They demonstrated that macaques vaccinated with the three components (including the adjuvant) developed substantial protection against intravaginal challenge with heterologous HIV virus strain compared to macaques that received the Gag protein or the Env protein alone. Animals vaccinated with the three components developed higher numbers of vaginal T<sub>RM</sub> cells and the protection was correlated with the production of Env-specific neutralizing Ab and Ab-dependent cell-mediated virus inhibition (Petitdemange et al., 2019).

### 1.9 Thesis aims:

Despite the substantial progress that has been made in controlling malaria, reduction in morbidity and mortality has stalled in the last 5 years. Therefore, malaria is still a major health public concern causing 400,000 deaths annually (WHO, 2018). One way to improve the control malaria is to develop an effective vaccine. Vaccines designed to stop parasite infection within the liver are ideal because they avoid the occurrence of symptoms associated with blood-stage infection and would also block transmission. Studies have demonstrated that pre-erythrocytic vaccines based on administration of whole attenuated sporozoites have high efficacy in animal models and humans (Butler et al., 2011; Clyde, 1990; Fernandez-Ruiz et al., 2016; Ishizuka et al., 2016; Mordmüller et al., 2017; Nussenzweig et al., 1967; Seder et al., 2013; Tse et al., 2013). However, these vaccines face major limitations that question their suitability for routine use in endemic areas. Some of those challenges are associated with poor efficacy in malaria pre-exposed individuals, logistical limitations for large scale sporozoite production and safety concerns when using live parasites. Liver-stage subunit vaccines overcome many of these problems, in particular, the manufacture limitations and safety issues related to delivering infective sporozoites.

*Plasmodium* liver-stage infection has constrained biology and it is critical that a robust immune response in the liver is triggered to identify the few infected hepatocytes in the entire liver before parasites progress to the blood-stage and induce symptoms (one week in humans and 2 days in mice) (Sturm et al., 2006). Several studies have demonstrated that sterile protection mediated by vaccination largely depends on the formation of effective memory CD8 T cell responses in the liver (Cockburn et al., 2014; Doll et al., 2016; Epstein et al., 2011; Fernandez-Ruiz et al., 2016; Ishizuka et al., 2016; Nganou-Makamdop et al., 2012). Moreover, liver T<sub>RM</sub> cells, a recently described subpopulation of memory T cells, have proved to be essential for protection against liver-stage challenge due to their strategic location and the potent surveillance they provide (Fernandez-Ruiz et al., 2016).

Recently, substantial efforts have been made to develop effective liver-stage subunit vaccines that promote the formation of liver associated CD8 T cells (Ewer et al., 2013; Fernandez-Ruiz et al., 2016; Gola et al., 2018). However, progress in this area is hampered by the limited number of identified protective CD8 T cell epitopes derived from *Plasmodium* parasites (Schmidt et al., 2010). To date, only two MHC-I restricted epitopes from PbA have been described to reduce liver parasite load in B6 mice and both failed to confer sterilising immunity (Hafalla et al., 2013; Pichugin et al., 2018). Accordingly, it is important to (i) identify new immunogenic MHC-I restricted epitopes and (ii) use those peptides in effective vaccine strategies that can generate robust liver CD8 T cell responses that in turn induce sterilizing immunity.

Our team was previously able to generate a transgenic CD8 T cell specific for PbA, which can be expanded and differentiated into liver T<sub>RM</sub> cells to confer effective protection against sporozoite challenge in B6 mice (Fernandez-Ruiz et al., 2016). Recently, the cognate epitope for these transgenic cells has been identified and shown to be derived from the ribosomal protein L6 (RPL6) of *P. berghei* ANKA (Valencia-Hernandez et al., 2020). This epitope is termed Pb-1 in this thesis. Experiments conducted and presented in this thesis aim to investigate the suitability of Pb-1 for induction of protection through different immunisation strategies.

As previously mentioned, liver T<sub>RM</sub> cells are effective and essential for protection against malaria liver-stage infection (Fernandez-Ruiz et al., 2016). Therefore, it is important to explore how to enhance the generation of liver T<sub>RM</sub> cells. Vaccine immunogenicity is usually enhanced by adjuvants, but limited information is currently available about adjuvants or molecules that could enhance the formation of liver T<sub>RM</sub> cells. Studies have shown that inflammation favours the formation of T<sub>RM</sub> cells in general (Bergsbaken and Bevan, 2015; Caminschi et al., 2019; Mackay et al., 2012a; Shin and Iwasaki, 2012), thus molecules that generate local inflammation in the liver may be particularly effective at promoting liver T<sub>RM</sub> formation. In fact, studies have shown that CpG

ODN promote local inflammation in the liver that supports the accumulation of CD8 T cells (Huang et al., 2013) and also enhances the generation of T<sub>RM</sub> cells (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Based on these findings, we hypothesized that nucleic acid sensor agonists that generate liver inflammation can act as adjuvants for the formation of liver T<sub>RM</sub> cell. To test this hypothesis, a series of synthetic nucleic acid agonists were assessed for the formation of T<sub>RM</sub> cells in the liver of mice vaccinated with the novel epitope Pb-1.

More precisely, the studies contained within this dissertation address the following aims:

- (1) To investigate the CD8 T cell memory responses mediated by different immunisation strategies using the Pb-1 epitope and to assess the suitability of these vaccines to evoke sterilizing immunity in mice with a normal endogenous T cell repertoire (Chapter 3).
- (2) To analyse various nucleic acid sensor agonists as novel adjuvants for the induction of liver T<sub>RM</sub> cell formation induced by antigen targeted immunisation (Chapter 4).

## Chapter 2

### Materials and Methods

## 2.1 Materials

### 2.1.1. Experimental models

Table 2.1: List of the experimental organisms and strains used in this study. The table shows the experimental organisms and strains and the origin/source.

Organisms / Strains	Origin / Source
Mouse: C57BL/6 (H2-Db) (B6)	Janvier LABS (Australia and Germany), Charles River (Australia and Germany), Animal Resources Centre, Australia, bred in house at The Peter Doherty Institute for Infection and Immunity.
Mouse: PbT-I/uGFP	(Lau et al., 2014), bred in house at The Peter Doherty Institute for Infection and Immunity.
<i>Anopheles stephensi</i> (STE2/MRA-128)	The Malaria Research and Reference Reagent Resource Center; BEI resources, bred in house at the School of Biosciences, University of Melbourne
<i>Plasmodium berghei</i> ANKA (MRA-871)	The Malaria Research and Reference Reagent Resource Center; BEI resources, bred in house at the School of Biosciences, University of Melbourne.

### 2.1.2. Immunisation reagents

Table 2.2: List of the immunisation reagents used in this study. The table includes names, features, sequences and source of the immunisation reagents used in this study including adjuvants and transfection reagents.

Reagent	Sequence (5'-3')	Source
rAAV-NVF (recombinant Adeno-	rAAV encoding for the following sequence: NVFDFNNL	Patrick Bertolino, David G Bowen; University of Sydney



Associated Virus encoding NVF)		
Rat, anti-mouse Clec9A-NVY mAb (clone 24/04-10B4)	mAb linked to the following sequence: HSLSNVYDFNLLLERD	Irina Caminschi, Mireille H. Lahoud; Monash University
Rat, anti-mouse Clec9A-NVF mAb (clone 24/04-10B4)	mAb linked to the following sequence: STNVFDFNLS	Irina Caminschi, Mireille H. Lahoud; Monash University
RAS γ-irradiated PbA sporozoites		Fresh sporozoites provided by Geoff McFadden and Anton Cozijnsen, University of Melbourne
<b>Adjuvants</b>		
cGAS-ligand (D3250 P*) dsDNA	AGCATAGGCCTATGCTG TGATGTATACATCAC 3' phosphorylated	Integrated DNA technologies, Coralville, IA, USA.
CpG-combo (CpG-2006 linked to CpG-21798) TLR9 ligand	T*C*G*T*C*G*T*T*T*T*G* T*C*G*T*T*T*T*G*T*C*G* T*T*T*CG*T*CG*A*CG*A* T*CG*G*C*G* (*Phosphorothioate backbone)	Integrated DNA Technologies, Coralville, IA, USA.
CpG-1668 (ODN 1668) TLR9 ligand	TCCATGACGTTCCCTGAT GCT	Geneworks, Thebarton, SA, Australia
CpG-2006 (PTO 2006) TLR9 ligand	T*C*G*T*C*G*T*T*T*T*G* T*C*G*T*T*T*T*G*T*C*G* T*T* (*Phosphorothioate backbone)	Integrated DNA technologies, Coralville, IA, USA.
LPS		Sigma-Aldrich, USA

TLR4 ligand		
Poly (I:C) (p I:C) TLR3 ligand		GE Health Care, Chicago, IL, USA
RIG-I-ligand (IVT4) 5' 3p ssRNA	5'- ppp GGGACGCUGACCCAGA AGAUCUACUAGAAUAG UAGAUCUUCUGGGUCA GCGUCCC (p phosphate)	Winfried Barchet, Patrick Müller, University Hospital Bonn, Germany.
TLR7-ligand (R3023), GU rich ssRNA	5'- C*C*rUrCrCrUrCrCrUrCrCr GrGrGrGrGrGrGrGrGrGrG rGrGrG*G*G*G*G*G -3' (*Phosphorothioate backbone)	Integrated DNA technologies, Coralville, IA, USA.
<b>Transfection reagent</b>		
DOTAP Liposomal transfection reagent N-[1-(2,3-Dioleoyloxy)propyl]-N,N,N- trimethylammonium methyl-sulfate		Sigma-Aldrich, USA
Glucose (10% Ultra-pure)		Polyplus transfection SA, Illkirch- Graffenstaden, Francia
In-vivo Jet PEI ®		Polyplus transfection SA, Illkirch- Graffenstaden, Francia

### 2.1.3 Cell biology reagents and materials

#### 2.1.3.1 Antibodies

Table 2.3: List of antibodies. The table shows the antibody name, clone, application and company used in this study.

Antibodies	Company	Application
Rat, anti-mouse CD4 (RM4-5)	BD Biosciences, San Jose, CA, USA	Flow cytometry
Rat, anti-mouse CD8 $\alpha$ (53-6.7)	BD Biosciences, San Jose, CA, USA	Flow cytometry
Rat, anti-mouse CD11a (clone 2D7)	BD Biosciences, San Jose, CA, USA	Flow cytometry
Armenian hamster, anti-mouse TCR $\beta$ (H57-597)	BD Biosciences, San Jose, CA, USA	Flow cytometry
Rat, anti-mouse V $\alpha$ 8.3 TCR (B21.14)	BD Biosciences, San Jose, CA, USA	Flow cytometry
Rat, anti-human/mouse B220 (RA3-6B2)	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry
Rat, anti-mouse CD11b (M1/70)	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry
Rat, anti-human/mouse CD44 (IM7)	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry
Rat, anti-mouse CD62L (MEL-14)	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry
Armenian hamster, anti-mouse CD69 (H1.2F3)	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry
Rat, anti-mouse F4/80 (BM8)	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry

Rat, anti-mouse CXCR6 (SA05D1)	BioLegend, San Diego, CA, USA	Flow cytometry
Mouse, anti-mouse NK1.1 (PK136)	BioLegend, San Diego, CA, USA	Flow cytometry
Live/Dead Aqua Blue	ThermoFisher Scientific, Waltham, MA, USA	Flow cytometry
Rat, anti-mouse Tim-3 (B8.2C12)	BioLegend, San Diego, CA, USA	Flow cytometry
Rat, anti-mouse PD1 (29F.1A12)	BioLegend, San Diego, CA, USA	Flow cytometry
Anti-mouse CD4 (GK 1.5)	Recombinant antibody facility, WEHI, Australia	CD8 T cell enrichment cocktail
Anti-mouse F4/80	Recombinant antibody facility, WEHI, Australia	CD8 T cell enrichment cocktail
Anti-mouse Gr1 (RB6-8C5)	Recombinant antibody facility, WEHI, Australia	CD8 T cell enrichment cocktail
Anti-mouse I-A/E (M5/114)	Recombinant antibody facility, WEHI, Australia	CD8 T cell enrichment cocktail
Anti-mouse Mac-1 (M1/80)	Recombinant antibody facility, WEHI, Australia	CD8 T cell enrichment cocktail
Anti-mouse erythrocytes (Ter119)	Recombinant antibody facility, WEHI, Australia	CD8 T cell enrichment cocktail

### 2.1.3.2 Cell biology chemicals and reagents

Table 2.4 List of cell biology reagents and materials.

Reagents and materials	Company
Anti-PE microbeads	MACS, Miltenyi Biotec, Bergisch Gladbach, Germany
Anti-APC microbeads	MACS, Miltenyi Biotec, Bergisch Gladbach, Germany
BioMag® Goat anti-rat IgG beads	Qiagen, Germany

Bovine serum albumin fraction V (BSA)	Sigma-Aldrich, USA
Cell strainer, 70 µm	MACS, Miltenyi Biotec, Bergisch Gladbach, Germany
Desatinib	Cell Signalling Technology, Danvers, MA, USA
Dimethyl Sulphoxide (DMSO)	Sigma-Aldrich, USA
EDTA	Sigma-Aldrich, USA
FACS buffer	PBS 5% w/v BSA, 5% EDTA 5mM.
Fetal Bovine Serum (FBS)	Gibco – Life Technologies
Glycerol	Sigma-Aldrich, USA
H2-K <sup>b</sup> (for tetramers)	Jie Lin, The University of Melbourne
Hanks' buffered salt solution (HBSS)	Media preparation Unit – The Peter Doherty Institute for Infection and Immunity
Hoechst 33258	ThermoFisher Scientific, USA
LS column MACS	MACS Miltenyi Biotec, Bergisch Gladbach, Germany
Nylon mesh, 70 µm	Madison Filter Pty Ltd., Australia
Percoll <sup>®</sup>	GE Health Care, Chicago, IL, USA
Phosphate buffered saline (PBS)	Media preparation Unit – The Peter Doherty Institute for Infection and Immunity
Polypropylene tubes (5 ml) (in this thesis called FACS tubes)	BD Bioscience, USA
Propidium iodine (PI)	Sigma-Aldrich, USA
RPMI1640	Gibco – Life Technologies, Media preparation Unit – The Peter Doherty Institute for Infection and Immunity

RPMI 1640 – 2.5 % FBS (RPMI 2.5%)	RPMI 1640 supplemented with 2.5 % of heat inactivated FBS
Red blood cell lysis buffer (RBCL)	826 g/L NH <sub>4</sub> Cl, 1g/L KHCO <sub>3</sub> , 0.037g/L EDTA in dH <sub>2</sub> O; pH7.3
Sphero blank calibrations beads (6.0 – 6.4 µm)	BD Bioscience, USA
Streptavidin-PE	ThermoFisher Scientific, Waltham, MA, USA
Trypan Blue	Sigma-Aldrich, USA

#### 2.1.4 Molecular biology reagents and materials

Table 2.5 List of molecular biology reagents and materials.

Reagents and materials	Company
BD Microtainer® blood collection tubes	BD Bioscience, USA
Chloroform	Carl Roth GmbH, Karlsruhe, Germany
DBL™ Heparin Sodium (porcine mucous)	Hospira, Australia
Donkey anti-Rb-HRP	Jackson ImmunoResearch Laboratories, INC.
Ethanol	Carl Roth GmbH, Karlsruhe, Germany
EvaGreen polymerase mix	Bio-Budget, Krefeld, Germany
Isopropanol	Carl Roth GmbH, Karlsruhe, Germany
Mouse IFNa	PBL assay sciences, USA
Oligo (dT) <sub>18</sub> primer (0.5 µg/µL)	ThermoFisher Scientific, Waltham, MA, USA
Rabbit polyclonal anti-IFNa	PBL assay sciences, USA
RevertAid RT Reverse Transcription kit	ThermoFisher Scientific, Waltham, MA, USA
RMMA-1 antibody	PBL assay sciences, USA

TMB substrate reagent set	BD Bioscience, USA
TRIzol®	ThermoFisher Scientific, Waltham, MA, USA
UltraPure™ DEPC-treated water	ThermoFisher Scientific, Waltham, MA, USA
Zirconia beads (2.0 mm)	Carl Roth GmbH, Karlsruhe, Germany

### 2.1.5 Specialised equipment

Table 2.6 List of specialised equipment.

Equipment	Company
FACS Canto	BD Immunocytometry Systems, San Jose, CA, USA
Fortessa or Fortessa X20	BD Immunocytometry Systems, San Jose, CA, USA
Precellys 24 homogenizer	Bertin Technologies
Cytation 3 Imaging reader	BioTek Instruments, Inc, USA
Quant Studio 5	Applied Biosystems, by Thermo Fischer Scientific

### 2.1.6 Software

Table 2.7 List of software packages.

Software	Company
FACSDiva	BD Immunocytometry Systems
Flowjo software	Tree Star, Ashland, OR, USA
GraphPad Prism 8	GraphPad Software, San Diego, CA, USA

## 2.2 Methods

### 2.2.1 Mice, mosquitoes and parasites:

All animal experiments were approved by the local government authorities LANUV, NRW, Germany (Animal ethics number Az. 84-02.04.2016.A403) and The University of Melbourne Animal Ethics Committee (Animal ethics number 1714302, 1814522). All protocols were performed in strict accordance with the recommendations of the national and institutional guidelines for the care and use of animals for scientific purposes.

Mice used in this study were bred and maintained in specific pathogen-free conditions at the House for Experimental Therapy and the Institute for Experimental Immunology at the Hospital University Bonn, Germany and at the animal facility of the Department of Microbiology and Immunology, at The University of Melbourne, Australia.

Female C57BL/6 (B6) and PbT-I/uGFP (Lau et al., 2014) were used between 6-12 weeks of age. Animals used for the generation of the sporozoites were 4-5-week-old male Swiss Webster mice purchased from the Monash Animal Services (Melbourne, Victoria, Australia) and maintained at the School of Botany, The University of Melbourne, Australia.

*Anopheles stephensi* mosquitoes (strain STE2/MRA-128 from The Malaria Research and Reference Reagent Resource Center) were reared and infected with *P. berghei* ANKA (*P. berghei*) as described by (Benedict, 1997). Sporozoites were dissected from mosquito salivary glands, resuspended in cold PBS, and either left untreated for challenge experiments or irradiated with 20,000 rads using gamma 60<sup>Co</sup> source (Ramakrishnan et al., 2012). Sporozoites were kindly provided by our collaborators from the research group of Prof. Geoff McFadden, School of Biosciences, University of Melbourne; where Anton Cozijnsen and Vanessa Mollard bred, maintained and dissected the mosquitoes.



### 2.2.2 Naïve CD8 T cell enrichment and adoptive transfer

CD8 T cells were negatively enriched from naïve PbT-I/uGFP transgenic female mice (Lau et al., 2014). Cell suspensions were prepared from spleen and LN harvested in RPMI 2.5% through mechanical disruption and passaging through a 70 µm mesh. Cells were incubated for 2 min in filtered sterile red blood cell lysis (RBLIC) buffer at room temperature (RT) to lyse red blood cells and then washed (centrifuged at 596 G, 5min, 4°C) in RPMI 2.5%. To enrich for CD8 T cells, suspensions were incubated in a cocktail (1 µL per 10<sup>4</sup> cells) of rat mAbs specific for erythrocytes (Ter119), I-A/E (M5/114), CD4 (GK 1.5), Gr1 (RB6-8C5), Mac-1 (M1/70) and F4/80 (F4/80) and incubated for 30 minutes on ice. Cells were then washed, resuspended in 400 µL of RPMI 2.5% and incubated with an equal volume of 3 x washed BioMag<sup>®</sup> sheep anti-rat IgG-coupled magnetic beads (used at 10:1 bead : cell ratio) for 20 minutes on constant rotation at 4 °C. Tubes were then loaded onto a cold magnetic rack for bead separation. Negatively enriched CD8 T cells were collected in the supernatant and resuspended in RPMI 2.5% and kept on ice until use. Cells were counted with trypan blue and purity was determined by flow cytometry. For this, a small aliquot of 100 µL of cell suspension was stained with anti-CD8 and anti-Vα8.3. Purity was determined as the percentage of PI<sup>-</sup>, GFP<sup>+</sup>, CD8<sup>+</sup>, Vα8.3<sup>+</sup> cells of the total population. Cells were washed and resuspended in cold sterile PBS. Recipient naïve B6 mice were adoptively transferred with 50,000 PbT-I/uGFP cells by intravenous injection of 200 µL of cell suspension (Fernandez-Ruiz et al., 2016).

### 2.2.3 Mice immunisations:

#### 2.2.3.1 Prime immunisation with Clec9A mAb

Rat anti-mouse Clec9A monoclonal antibodies (mAb) (clone 24/04-10B4) were genetically coupled to the HSLSNVYDFNLLERD or STNVDFNLS peptides – containing the NVY mimic epitope or the PbRPL6<sub>120-127</sub> cognate epitope, respectively – via a 4 Alanine linker to generate the αClec9A-NVY (Fernandez-Ruiz et al., 2016) and αClec9A-NVF (in this thesis named: αClec9A-

Pb-1) mAb, respectively. The antibodies were kindly provided by our collaborators Prof. Irina Caminschi and Prof. Mirelle H. Lahoud, Monash University. B6 mice were injected i.v. with the indicated doses  $\alpha$ Clec9A-NVY and  $\alpha$ Clec9A-Pb-1 mAb plus indicated doses of adjuvants in a total volume of 200  $\mu$ L. All mice were injected via the tail vein using a 29G insulin syringes.

#### 2.2.3.2 Adjuvant stimulation and preparation

For the experiments described in chapter 3, B6 mice were i.v. injected with  $\alpha$ Clec9A-Pb-1 mAb plus an adjuvant: either 75  $\mu$ g of CpG-combo, generated by linking (5' to 3') CpG-2006 to CpG-21798 (Samulowitz et al., 2010), or 50  $\mu$ g of p I:C. For the experiments described in chapter 4, B6 mice were i.v. injected with 2  $\mu$ g of  $\alpha$ Clec9A-Pb-1 mAb in the presence or absence of one of the following adjuvants in indicated doses: CpG-combo, p I:C, CpG-1668 (Fernandez-Ruiz et al., 2016; Holz et al., 2018) or LPS.

For chapter 4, mice were also i.v. injected with adjuvants complexed in transfection reagents, as described next. Different doses of cGAS-L and RIG-I-L, generated by *in vitro* retro-transcription (Winfried Barchet, University Hospital Bonn), were complexed in the transfection reagent *In-vivo* Jet-PEI<sup>®</sup> according to manufacturer instructions. In detail, in a ratio of 0.12  $\mu$ L of *In-vivo* Jet-PEI<sup>®</sup> per 1  $\mu$ g of adjuvant, cGAS-L or RIG-I-L were mixed with Jet-PEI<sup>®</sup> in a total volume of 100  $\mu$ L of glucose (5% ultrapure) per mouse. This solution was mixed thoroughly and incubated at RT for 20 min. Additionally, CpG-1668 and two different doses of each, CpG-combo, CpG-2006 and TLR7-L were mixed thoroughly with DOTAP, in a ratio of 2.5  $\mu$ g DOTAP per 1  $\mu$ g of adjuvant and incubated for 20 min at RT (following manufacturer instructions). The complexed adjuvants were then mixed with 10 ng /  $\mu$ L of  $\alpha$ Clec9A-Pb-1 mAb, each mouse was injected with 200  $\mu$ L of priming solution containing a total of 2  $\mu$ g of mAb. Solution was kept on ice until injection.

### 2.2.3.3 Prime-and-trap vaccination

B6 mice were injected i.v. with the indicated doses  $\alpha$ Clec9A-Pb-1 mAb plus 75  $\mu$ g of CpG-combo as adjuvant and different doses of recombinant adeno-associated virus expressing the Pb-1 epitope (rAAV-Pb-1). The prime-and-trap vaccination strategy was developed by (Fernandez-Ruiz et al., 2016) and in this thesis is used to target the Pb-1 epitope. It differs from the previous publication in that all three components are administered simultaneously, whereas previously the adenovirus was given one day post immunisation with  $\alpha$ Clec9A mAb.

Viral particles were produced and kindly provided by our collaborators Prof. Patrick Bertolino and David G Bowen, University of Sydney. rAAV were prepared and purified in house at the Centenary Institute or by the Vector and Genome Engineering Facility (at the Children Medical Research Institute, Sydney, Australia) over caesium chloride (CsCl)-density gradient centrifugation followed by dialysis as previously described by (Tay et al., 2014). rAAV-NVF – in this thesis named rAAV-Pb-1 – vector was generated by replacing the sequence encoding for the natural SIINFEKL epitope of OVA (OVA<sub>257-264</sub>) contained in rAAV-OVA (Tay et al., 2014) with the sequence encoding for NVFDFNLL.

### 2.2.3.4 RAS vaccination

RAS were prepared by irradiating freshly isolated sporozoites with 20,000 rads using gamma 60<sup>Co</sup> source (Ramakrishnan et al., 2012). Immediately after this, B6 mice were injected i.v. with 50,000 RAS in the absence or presence of 75  $\mu$ g CpG-combo complexed or not in DOTAP, as previously described.

### 2.2.4 Challenge experiments with live PbA sporozoites

Vaccinated B6 mice were challenged i.v. with 200 freshly isolated PbA sporozoites per mouse resuspended in cold PBS. Mice were monitored daily for body condition until day 16 post infection (p.i.). Parasitemia was assessed according to (Fernandez-Ruiz et al., 2016). On days 6-8 and 10 p.i., 2  $\mu$ L of blood were taken from the tail vein and resuspended in 100  $\mu$ L of FACS buffer.

Parasitemia was assessed by incubating blood suspensions in 5 µg/mL Hoechst 33258 solution for 1 hour at 37°C. Parasites were detected from infected RBC by flow cytometry using a 405 violet laser and a 450/50 filter. Protection was considered sterile when no blood stage parasites had been detected within 10 days after challenge. Numbers over bars represent protected mice/total mice.

#### *2.2.5 Organ processing for T cell analysis*

Tissues were harvested in RPMI 2.5% at different time points after immunisation. Cell suspensions were generated from spleens and livers through mechanical disruption and passage through a 70 µm mesh and washed once with RPMI. Then, liver cell suspensions were resuspended in RT 35% isotonic Percoll and centrifuged at 500 G for 20 min at RT. (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Spleen and liver cell suspensions were incubated for 2 min in 3 mL of RBLIC at RT to lyse red blood cells, then washed and resuspended in cold FACS buffer (PBS 5% w/v BSA, 5% EDTA 5mM).

#### *2.2.6 Flow cytometry*

Cell surface immunostainings were performed according to standard protocols. After single cell suspensions were prepared from liver and spleen as described above, cells were stained with fluorophore-conjugated antibodies. H2-K<sup>b</sup>-PbRPL6<sub>120-127</sub> and H2-D<sup>b</sup>-PbTRAP<sub>130-138</sub> tetramers (Pb-1 and TRAP tetramer, respectively) were made in house. Tetramers were prepared by our collaborators Jie Lin and Prof. Andrew Brooks at the Peter Doherty Institute, University of Melbourne. Cell populations were identified based on forward-scatter area (FSC-A) and side-scatter area (SSC-A). Singlets were excluded based on FSC-A and forward-scatter height (FSC-H). Dead cells were excluded by propidium iodide (PI) staining. For the tetramer staining, B220<sup>+</sup>, NK1.1<sup>+</sup> and F4/80<sup>+</sup> cells were excluded before gating on tetramer<sup>+</sup> CD8 T cells. For the analysis of memory CD8 T cell populations in the spleen and the liver, tetramer<sup>+</sup> or PbT-I/GFP<sup>+</sup> CD8<sup>+</sup> CD44<sup>hi</sup> cells were subdivided into T<sub>CM</sub>, T<sub>EM</sub> or T<sub>RM</sub> based on CD69 and CD62L

expression ( $T_{CM}$  CD62L<sup>+</sup> CD69<sup>-</sup>,  $T_{EM}$  CD62L<sup>-</sup> CD69<sup>-</sup> and  $T_{RM}$  CD62L<sup>-</sup> CD69<sup>+</sup>). CXCR6 was included as an additional  $T_{RM}$  cell marker. In some instances, where indicated, cells were also stained with anti-PD1 and Tim-3 Ab, to determine an exhaustion-like phenotype. To enumerate cell numbers, Sphero blank calibrations beads (6-6.4 $\mu$ m) were diluted into FACS buffer containing PI and added to surface-stained cells immediately before acquisition of samples. The particles were distinguished from cells based on their differential size and granularity reflected by FSC-A, FSC-H and SSC-A representation. The number of cells detected by FACS was divided by the ratio of the number of particles added and the number of particles detected.

#### *2.2.7 Enumeration of antigen-specific CD8 T cells in the naïve repertoire*

Spleen, axillary, branchial, mesenteric, cervical and inguinal lymph nodes were harvested from naïve or immunised B6 mice (used as control). Single cell suspensions were produced by gently mashing the tissue through a 70  $\mu$ m mesh and resuspending cells in FACS buffer. Incubation with 50 nM dasatinib (Cell Signalling Technology, Danvers, MA, USA) for 30 min at 37°C was used to prevent TCR/tetramer internalization. Samples were stained with PE and APC-conjugated tetramer of the indicated specificity for 1 hour at room temperature. Solutions were then incubated with anti-PE and anti-APC conjugated magnetic microbeads, and tetramer-bound cells were enriched over a magnetic LS column MACS®. Enriched cells were then stained with fluorophore-conjugated antibodies specific for TCR $\beta$ , CD8, CD4, CD11b, CD11c, B220, F4/80, NK1.1 and CD62L. Dead cells were excluded by Live/Dead Aqua Blue or propidium iodide staining (La Gruta et al., 2010; Obar et al., 2008).

### 2.2.8 Enzyme-linked immunosorbent assay (ELISA)

#### 2.2.8.1 Plasma preparation

Blood samples were collected from live mice by cheek bleeding on BD Microtainer® blood collection tubes or in 1,5 mL tubes containing 100 µL of heparin solution (200 IU/mL in RPMI). After 6 h or 10h as indicated, mice were killed, and blood was collected immediately by cardiac puncture. Samples were mixed by inversion and plasma was separated by centrifugation (10,000 G) for 10 min at RT. Plasma samples were stored at -20 °C until further use.

#### 2.2.8.2 Organ processing

Spleens were cut in two and livers were perfused with cold PBS and two similar sized pieces of the main lobule were collected. The weight of each organ piece was determined for the normalization of protein content and levels of gene expression relative to organ weight. Tissues were collected in 1 mL tubes with 2.0 mm Zirconia beads, resuspended in RPMI supplemented with protease inhibitors or in TRIzol®, for ELISA assays or RNA extraction, respectively. Tissues were disrupted and homogenized in a Precellys 24 homogenizer. Tubes were shortly centrifuged at maximum speed and the supernatant was collected and stored at -20°C for ELISA assays or at -80°C until RNA extraction.

#### 2.2.8.3 ELISA to detect IFN $\alpha$

Murine IFN $\alpha$  (m IFN $\alpha$ ) ELISA was performed in serum and homogenized spleen and liver tissue. Nunc-Immuno™ MicroWell™ 96-well high protein binding plates were coated with 50 µL of PBS containing 250 ng/ mL of RMMA-1 antibody overnight (ON) at 4°C. Unbound protein was washed by soaking the plates 3 to 5 times with wash buffer (0.05% Tween20 in PBS) and blocked for 3 h with 50 µL of PBS containing 1% BSA at RT. After washing the plates 3 to 5 times, 50 µL of sera (diluted 1:5 in PRMI) and undiluted tissue samples were loaded on the plate. Two technical replicates were loaded per sample; additionally, a twofold serial dilution of recombinant IFN $\alpha$  was also loaded and used to calculate a standard curve. Plates with samples and standard were incubated at 4°C ON. For the

detection, washed plates were incubated at RT for 4 h with 50  $\mu$ L of antibody buffer (1% BSA, 0.05% Tween20 in PBS) containing rabbit polyclonal anti-mouse IFN $\alpha$  (1:1000) and after a subsequent washing, plates were incubated at RT for 3 h with donkey anti-rabbit HRP (diluted 1:2000 in antibody buffer). The wells were washed 6 to 12 times and HRP was detected by adding 100  $\mu$ L of TMB detection reagent (substance A + B in equal volumes) and incubated in darkness for 30 min before measurement. The chemical reaction was stopped with 100  $\mu$ L of 2N H $_2$ SO $_4$  or 100  $\mu$ L of 0.2 M HCl. Plates were analysed at 450/570 nm using Cytation 3 Imaging reader. The concentration of protein was calculated by linear regression of a standard curve.

### *2.2.9 Quantitative real-time PCR:*

#### 2.2.9.1 RNA isolation

Total RNA was isolated from homogenized liver and spleen tissue by phenol-chloroform extraction. To separate the aqueous phase containing RNA, 200  $\mu$ L of chloroform were added to samples in TRIzol<sup>®</sup>, mixed thoroughly, incubated for 10 min on ice and centrifuged at 12,000 G for 15 min at 4°C. The upper aqueous phase was transferred to a new tube and nucleic acids precipitated by adding 400  $\mu$ L isopropanol, mixed by inversion, incubated on ice for 10 min and centrifuged at 12,000 G for 10 min at 4°C. Supernatant was removed and pellet was washed (14,000 G for 5 min at 4°) twice with 70% ethanol. Supernatant was completely removed, and pellet was resuspended in 16  $\mu$ L of UltraPure<sup>™</sup> DEPC-treated water. RNA concentration was determined using Nanodrop 2000. DNA was eliminated by incubating RNA containing tubes with DNase 1 for 15 min at 37°C, followed by an inactivation step at 65°C for 5 min. Samples were stored at -80°C until use.

#### 2.2.9.2 cDNA synthesis

The cDNA was synthesized from the extracted RNA with a RevertAid RT Reverse Transcription kit. In detail, 200 ng of RNA were resuspended in 13  $\mu$ L of DEPC-treated water plus 1  $\mu$ L of Oligo (dT) $_{18}$  primer (0.5  $\mu$ g/ $\mu$ L) and incubated

for 10 min at 70°C. Samples were placed on ice and the reverse transcriptase mixture was added. This mixture contained 4 µL 5 x First Strand Buffer, 1 µL dNTP Mix (5nM each dNTP) and 1 µL RevertAid reverse transcriptase (200U/µL), per sample. cDNA synthesis was performed by incubating samples for 60 min at 42°C, then volume was adjusted to 60 µL in total and cDNA was stored at -20°C until use.

### 2.2.9.2 Real-time PCR

For real-time PCR a total of 2 µL of cDNA was mixed with 0.6 µL of 10 mM primer mix, 2 µL of Fast EvaGreen qPCR master mix in a total volume reaction of 10 µL. Reaction was performed in a Quant Studio 5 with the following cycle conditions: 95 °C for 2 minutes, 35 cycles at 95 °C for 15 seconds and 60 °C for 1 minute. The mRNA expression of each gene (*Ifi44* and *Irf7*) was normalized to the house keeping gene *Hprt* and the fold change relative expression was calculated using unstimulated controls as  $2^{-\Delta\Delta CT}$ .

Table 2.8 List of primers used for real-time PCR.

Name	Forward 5' → 3'	Reverse 5' → 3'
mHPRT	GCCCCAAAATGGTTAAGTT	CAAGGGCATATCCAACAACA
mIFI44	TCGATTCCATGAAACCAAT CAC	CAAATCGCAGAATGCCATGTT TT
mIrf7	CTTCAGCACTTTCTTCCGA GA	TGTAGTGTGGTGACCCTTGC

### 2.2.10 Statistical analyses

Figures were generated using GraphPad Prism 8. Data are shown as mean values ± standard error of the mean (SEM). Statistical analyses were performed using GraphPad Prism 8. The statistical test used is indicated in each figure legend.  $P < 0.05$  was considered to indicate statistical significance. \*,  $P < 0.05$ ; \*\*,  $P < 0.01$ ; \*\*\*,  $P < 0.001$ ; \*\*\*\*,  $P < 0.0001$ . Unless otherwise indicated, figures show



pooled data from all independent experiments performed. Statistical analyses of cell frequencies, percentage of iRBC and ratios were performed by comparing groups using a Kruskal- Wallis test with Dunn's multiple comparison post-test, as these data were considered non-parametric. Statistical comparisons of cell numbers in different groups were performed by log-transforming the data and using a Student's t-test (2 groups) or one-way ANOVA followed by Tukey's multiple comparisons test (>2 groups). Asterisks directly over groups denote statistical differences with the unvaccinated control group. Rates of sterile protection were compared using one-sided or two-sided Fisher's exact tests, for comparisons of immunized groups vs the unimmunized control group, or between immunized groups respectively.

## CHAPTER 3:

Characterization of the memory response induced by immunisation with an epitope from the ribosomal protein L6 (RPL6) of *P. berghei* ANKA.

### 3.1 Introduction:

In 2014, members of our group developed a C57Bl/6 (B6) T cell receptor (TCR) transgenic line able to produce MHC class I-restricted cells specific for PbA. This TCR transgenic line, termed PbT-I, produced cells that respond to blood-stage PbA, *P. yoelii* XNL and *P. chabaudi*, and to liver-stage parasites after infection with PbA radiation attenuated sporozoites (RAS). PbT-I cells were also shown to confer sterile protection against PbA sporozoite challenge (Lau et al., 2014).

To identify the cognate antigen for the PbT-I cells, an octamer combinatorial peptide library was used to determine ideal residues required for activation of PbT-I cells (Lau et al., 2014). These residues were then used to screen rodent malaria parasite genes for closely matching sequences that were then synthesised and tested for activation of PbT-I cells. After screening 151 potential peptides, the NCYDFNNI (NCY) epitope was described as the leading peptide candidate recognized by the PbT-I cells. This was shown by assessing *in vivo* cell proliferation and IFN- $\gamma$  production by PbT-I cells stimulated with this peptide (Lau et al., 2014). This epitope is part of a protein with unknown function, encoded on the genomes of PbA (PBANKA\_0714500) and *P. chabaudi* (PCHAS\_0723600). *In silico* analysis of the ortholog protein in *P. yoelii* (PY000568), however, revealed that this protein is truncated and does not contain the NCY epitope. This was later confirmed by sequencing the PY000568 gene within the *P. yoelii* genome (Valencia-Hernandez et al., 2020). Since PbT-I cells are able to respond to blood-antigen from *P. yoelii* infection, yet NCY was not present in this species genome, this indicated that NCY was not the authentic cognate peptide recognised PbT-I cells and that the authentic peptide antigen for the TCR remained to be identified.

From the initial screening of the octamer peptide library, another agonist epitope was identified, NVYDFNLL (NVY) (Fernandez-Ruiz et al., 2016). This is an antigen mimic epitope that is not present on the genome of *Plasmodium*

species but is able to induce the activation of the PbT-I cells. The NVY peptide was subsequently used as an antigen in a novel vaccine, termed “prime-and-trap” (described on Chapter 1) for the generation of PbT-I T<sub>RM</sub> cells in the liver, which were shown to be crucial for protection against the PbA sporozoite challenge (Fernandez-Ruiz et al., 2016).

Subsequently, analysis of the sequences from *Plasmodium* peptide candidates similar to NVY together with mass spectrometry of eluted peptides from dendritic cells presenting blood-stage PbA parasites identified the authentic cognate peptide of the PbT-I cells. In detail, the protein basic local alignment search tool (Blastp) within the PlasmoDB platform was used to identify peptides similar to NVY in the proteomes of PbA, *P. yoelii* and *P. chabaudi*. As a result, the NVFDFNNL epitope, in this document denominated as Pb-1, was identified. The Pb-1 epitope is present within the 60S ribosomal protein L6-2 of PbA (RPL6) (Valencia-Hernandez et al., 2020). Very similar epitope sequences were found in the ortholog proteins of *P. yoelii* XNL and *P. chabaudi chabaudi* (Table 1.).

Table 3.1. Amino acid sequence of the RPL6 protein. Alignment to the Pb-1 epitope. Same amino acids appeared in yellow, different in blue.

Strain	Gene code	Protein sequence
<i>P. berghei</i> ANKA	PBANKA_1351900	MAKNTKSGASADDKKKTLKHVVIKGQK KTLTPVRAKKTIAKKYYGKKLASKKKYIV QRKMRKSIQVGKVAIILTGHMGKRCIIT KVLKSGLLAVIGPYEVNGVPLKRVDPRY LIVTSTNVFDFNNLSQIKDKFIQAAERIN DEIFIKSIDIKKRQKKLLKNKNESLFMND VIQQIKEIRDSDPKMKRIKLLQKQLGDLL KPEISKDKMFRSYIKSKFTLRNMSFHN IKF
<i>P. yoelii</i> XNL	PY02722	MAKNNKNTKSVVSSDDKKSLLKHVVIK GQHKILTPIQAKKTISKKHVGRKIASKKK YIVQRKMRKSIQIGKVAIILTGHMGKRC

		IITKILNSGLLAVVGPYEVNGVPLKRVDP RYLIVTSTNVFDLNNLSQIKDKFIQTAEDI HDDIFIKSNDIKKRQKKLLKNKNESLFMN DVIQQIKEIRDSDPKMKKIKSLQKQLGDL LKPEMSKDKMLRSYLKSKFTLRNNMSF HTIKF
<i>P. chabaudi</i> <i>chabaudi</i>	PCHAS_1356500	MAKITKSTKSSKETKSVVPATGDKKNAL KHVVIKGQKKILTPVRAKKTISKKHVGRK LASKKKFVVQRKMRSSIQVGKVAIILTG KHMGRRCIIAKVLKSGLLAVVGPYEVNG VPLKRVDPRYLIVTSTNIFDFNNLSQVKE KFIQSAERINDEIFIKSMDVKKRQKKLLK NKNESLFMNDVIQKIKEIRDSDPKMKKIK LLQKELGDLLKPEISKDKMFRSYIKSKFT LRNNMSFHNMNF

The Pb-1 epitope as well as its orthologs from *P. yoelli* and *P. chabaudi* were tested for their capacity to stimulate PbT-I cells *in vitro*. Splenic dendritic cells (DC) coated with the different peptides were cultured with Cell-Trace Violet™ (CTV)-labelled PbT-I cells. All three peptides were able to induce proliferation of PbT-I cells and very importantly, the sensitivity to different peptide doses of the PbT-I cells to Pb-1 was equivalent to that of OT-I cells to OVA<sub>257-264</sub> (SIINFEKL) peptide (Valencia-Hernandez et al., 2020). This result supports the idea that Pb-1 is the authentic cognate antigen for the PbT-I cells.

To determine whether the Pb-1 epitope from PbA is naturally processed and presented by DC, mass spectrometry was utilised to identify epitopes eluted from H-2K<sup>b</sup> molecules of DC cells cultured with blood stage malaria parasites; an abundant source of antigens. First, eluted peptides were fractionated by *high-performance liquid chromatography* (HPLC); a small aliquot of each fraction was co-cultured with PbT-I cells and naive DC; then the activation and proliferation of PbT-I cells was determined by CD69 upregulation and CTV-labelling

respectively. Only one fraction triggered a PbT-I response (Valencia-Hernandez et al., 2020). Posteriorly, analysis by high-resolution liquid chromatography coupled to tandem mass spectrometry (LC-MS/MS) of the fraction of interest and the commercially synthesised Pb-1 peptide in presence of indexed retention time (iRT) peptides (Escher et al., 2012) demonstrated an overlap between the two retention time profiles (Valencia-Hernandez et al., 2020). These results indicate that Pb-1 is present in the fraction that activated PbT-I cells, implying that this peptide is a natural occurring parasite antigen presented on H-2K<sup>b</sup> molecules by DC during PbA infection (Valencia-Hernandez et al., 2020).

As mentioned earlier, PbT-I T<sub>RM</sub> cells in the liver are very effective at inducing immunity against PbA liver stage malaria. To this point, the generation of high numbers of liver T<sub>RM</sub> cells had only been achieved by prime-and-trap vaccination using the NVY mimic peptide in combination with transferred PbT-I cells, i.e. endogenous T cells had not been tested (Fernandez-Ruiz et al., 2016). Based on the success of prime and trap vaccination using PbT-I cells, together with recent identification of the cognate and natural occurring antigen recognised by these cells; the objective of the following study was to determine if the prime-and-trap vaccine strategy was able to use the Pb-1 epitope to evoke memory T cell responses in the liver and induce protection in mice with a normal endogenous T cell repertoire e.g. in the absence of transgenic PbT-I cells.

In this chapter, a series of different immunisation experiments utilising the Pb-1 epitope for priming were implemented and specific CD8 T cell responses were assessed. This study revealed the suitability of the novel Pb-1 epitope for induction of protection through vaccination. Importantly, when Pb-1 was targeted for the prime-and-trap protocol, high levels of sterile protection were achieved in B6 mice in absence of transgenic cells, supporting the potential for translation of this approach.

### 3.2 Results:

#### *3.2.1 Potency of $\alpha$ Clec9A-Pb-1 priming on PbT-I cell expansion and memory formation in spleen and liver:*

Rat anti-Clec9A monoclonal antibodies (mAb) genetically coupled to the Pb-1 epitope ( $\alpha$ Clec9A-Pb-1) were generated by our collaborators (Caminschi and Lahoud, Monash University). The expression of the Clec9A molecule is restricted to DC and is highly expressed by type 1 conventional dendritic cells (cDC1) (Caminschi et al., 2008; Desch et al., 2011; Huysamen et al., 2008; Sancho et al., 2008; Schreiber et al., 2012); which is the main antigen-presenting cell type responsible for stimulating T cells in malaria (Fernandez-Ruiz et al., 2017; Lau et al., 2014).  $\alpha$ Clec9A-Pb-1 was utilised to prime naive mice and study the memory CD8 T cell response to the Pb-1 peptide. Mechanistically, this reagent targets cDC1 delivering the Pb-1 peptide. In the spleen, these cDC1 process and cross-present the peptide to naive PbT-I or specific Pb-1-CD8 T cells; inducing their activation and proliferation. Details of the immune responses triggered by antigen DC-targeted immunization via Clec9A were reviewed in Chapter 1. It is important to mention that, for most of the experiments performed in this chapter, the response of PbT-I cells was examined first. This was done to get clear evidence that the priming reagents were working before examining their effectiveness at stimulating the responses from endogenous cells.

To determine the optimal dose of  $\alpha$ Clec9A-Pb-1 for induction of maximal expansion of specific CD8 T cells, a titration experiment was carried out. First, 50,000 PbT-I cells were adoptively transferred intravenously into naive B6 mice. The next day, mice were primed with increasing doses of  $\alpha$ Clec9A-Pb-1 or  $\alpha$ Clec9A-NVY, the last was used as control for comparative purposes between the authentic cognate epitope, i.e. Pb-1 and the mimic epitope, i.e. NVY. Additionally, CpG was co-injected with the  $\alpha$ Clec9A reagent as adjuvant, which is required for induction of CD8 T cell responses (Lahoud et al., 2011) and also favours liver T<sub>RM</sub> cell formation (Fernandez-Ruiz et al., 2016).

PbT-I cells from spleens were analysed on day 7 post priming, during the effector phase of the response, as these data provide a rapid indication of the effectiveness of the vaccine as well as an approximation of the proportion of memory cells that could be formed at a later time point. Results showed increasing expansion of PbT-I cells as the dose of the antibody increased (Fig. 3.1 B). In control unprimed mice (0  $\mu\text{g}$   $\alpha\text{Clec9A-Pb-1}$ ), the mean number of cells per spleen was approximately one thousand, while at the highest dose of  $\alpha\text{Clec9A-Pb-1}$  (16  $\mu\text{g}$  per mouse) the number of cells reached approximately 6 million (a six thousand-fold increase). When compared to  $\alpha\text{Clec9A-NVY}$ ,  $\alpha\text{Clec9A-Pb-1}$  was significantly better at inducing the expansion of PbT-I cells at all doses tested (Fig. 3.1 C). Additionally, no plateau of PbT-I cell expansion was evident; further experiments were performed using 2  $\mu\text{g}$  of  $\alpha\text{Clec9A-Pb-1}$  as it had one of the smallest deviations and was deemed sufficient to generate substantial expansion of PbT-I cells. This dose had been previously used for the OVA system; for these experiments mice have been adoptively transferred with OT-I cells, one day later primed with 2  $\mu\text{g}$  of  $\alpha\text{Clec9A-SIINFEKL}$  plus CpG; analyses at early and memory time points have revealed substantial expansion and memory cell formation of the OT-I cells (Holz et al., 2018).

The previous experiment revealed that PbT-I cells expand substantially in response to  $\alpha\text{Clec9A-Pb-1}$  priming at an early stage of the response. To explore the capacity of this reagent to induce long-term responses to Pb-1, the kinetics of the PbT-I response were examined at memory time points. Mice were transferred with naive PbT-I cells and, the next day, primed with 2  $\mu\text{g}$  of  $\alpha\text{Clec9A-Pb-1}$  plus CpG or poly I:C as adjuvants. The effect of these adjuvants and others will be evaluated in more detail on Chapter 4.

Blood was collected on day 7 to measure the percentage of the PbT-I cell expansion and study the potential correlation of this initial expansion with the numbers of memory cells at later points. Spleens and livers were harvest from day 3 to day 35. For the analysis of memory formation, cells were labelled for



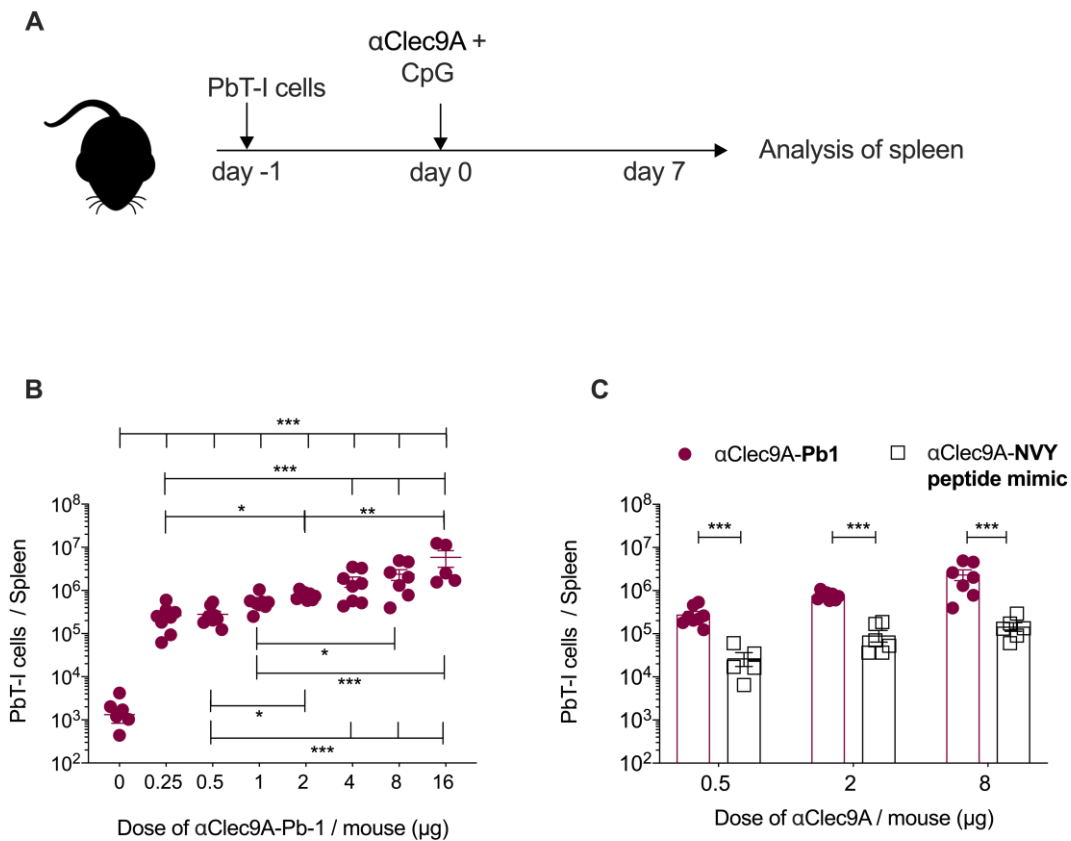


Figure 3.1. Effect of  $\alpha$ Clec9A-Pb-1 priming on the expansion of PbT-I cells in the spleen. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells. The next day, mice were immunised with increasing doses of  $\alpha$ Clec9A-Pb-1 or  $\alpha$ Clec9A-NVY plus 75 $\mu$ g of CpG. 7 days later, spleens were collected. Single cell suspensions were generated, and the number of PbT-I cells were determined by flow cytometry. A. Schematic of mice immunisation. B. Quantification of PbT-I cells (CD8+, GFP+ cells) in the spleen 7 days after priming with increasing doses of  $\alpha$ Clec9A-Pb-1. C. PbT-I cell expansion in the spleen 7 days after priming with 3 increasing doses of  $\alpha$ Clec9A-Pb-1 or  $\alpha$ Clec9A-NVY. The data on panel C. for the corresponding doses of  $\alpha$ Clec9A-Pb-1 are the same as shown in panel B. Results were pooled of two independent experiments. Data were log-transformed and compared by one-way ANOVA and Tuckey's multiple comparisons test.  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ . Two sets of statistical analysis were conducted. First, multiple comparisons were performed comparing all increasing doses. Second, the data of the three equivalent doses of  $\alpha$ Clec9A-Pb-1 vs.  $\alpha$ Clec9A-NVY were compared.

surface markers, including CD44, CD62L, CD69 and then analysed by flow cytometry; liver T<sub>RM</sub> like cells were defined as CD44<sup>high</sup>CD62L<sup>-</sup>CD69<sup>+</sup>. Since CD69 is also a marker for activation after stimulation (González-Amaro et al., 2013), CXCR6 was included as an additional T<sub>RM</sub> cell marker.

CpG and p I:C induced extensive expansion of PbT-I cells in blood on day 7 (Fig. 3.2 B). PbT-I cell expansion peaked on day 5 in the spleen, reaching in average of one million total PbT-I cells per spleen (Fig. 3.2 C). In the liver, p I:C induced a peak on day 5 with an average of  $1.7 \pm 0.2 \times 10^6$  cells while CpG on day 7 generated  $3.7 \pm 0.8 \times 10^6$  cells, which was significantly higher than p I:C on that day (Fig. 3.2 D). PbT-I cells slowly contracted over time in the spleen and liver. By day 35, the number of PbT-I cells in the liver was still significantly higher in mice primed with  $\alpha$ Clec9-Pb-1 mAb compared to naive mice, regardless of the adjuvant.

Moreover, CpG induced more T<sub>RM</sub>-like cells in the liver compared to p I:C. In fact, CpG showed a significant greater capacity at generating PbT-I T<sub>RM</sub> cells (CD44<sup>Hi</sup>, CD62L<sup>-</sup>, CD69<sup>+</sup>) on day 28 and 35 post immunisation compared with p I:C (Fig. 3.2 E). On day 35, mice immunised with CpG had approximately  $8.1 \pm 0.9 \times 10^4$  PbT-I T<sub>RM</sub> cells per liver. Interestingly, the numbers of T<sub>EM</sub> cells in the spleen was similar for the two adjuvants over time (Fig. 3.2 F).

As analyses in Figure 3.2 only used CD62L and CD69 expression to identify liver T<sub>RM</sub> cells, there was some possibility that CD69<sup>+</sup> cells represented effector T cells that were recognising antigen and upregulating CD69. To further substantiate the preliminary identification of CD69<sup>+</sup> cells in the liver as authentic T<sub>RM</sub> cells, the expression of the marker CXCR6 was also investigated in some of the experiments. The number of T<sub>RM</sub> like cells (CD62L<sup>-</sup>, CD69<sup>+</sup>) expressing CXCR6 was calculated for the different time points (Fig. 3.3 A).

These new analyses confirmed similar kinetics to those observed for CD69<sup>+</sup>, T<sub>RM</sub>-like cells. The expression profile of CXCR6 peaked on day 7 and

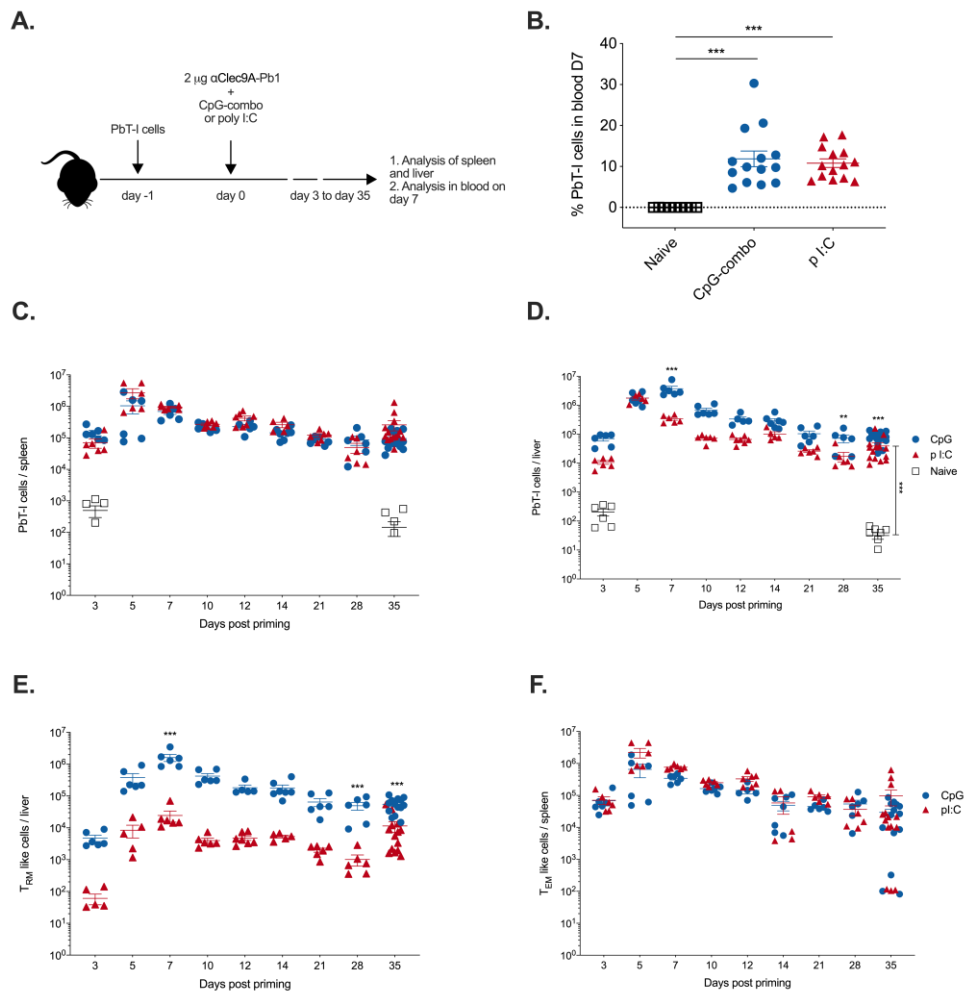


Figure 3.2. Kinetics of the PbT-I cell expansion and memory formation after  $\alpha$ Clec9A-Pb-1 priming. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells and immunised with 2  $\mu$ g of  $\alpha$ Clec9A-Pb-1 plus 75  $\mu$ g of CpG or 50  $\mu$ g of poly I:C. On day 7 after immunisation, mice were bled, and blood was analysed by flow cytometry. Spleens and livers were collected on various days up to day 35. Single cell suspensions were generated, and the number and memory status of PbT-I cells was determined by flow cytometry. A. Schematic of mice immunisation. B. Percentage of PbT-I cells (CD8+, GFP + cells) in the blood on day 7 post priming. C. and D. Total number of PbT-I cells in spleen and liver over time. E. and F. Kinetics of PbT-I  $T_{RM}$ -like cell numbers (CD8+, GFP+, CD44<sup>Hi</sup>, CD62L-, CD69+) per liver and PbT-I  $T_{EM}$  cells (CD8+, GFP+, CD44<sup>Hi</sup>, CD62L-, CD69-) per spleen, respectively. Results were pooled of two independent experiments, except for day 35 which corresponds to a pool of 4 independent experiments. Data were compared using a Kruskal- Wallis test with Dunn's multiple comparison post-test in (B.) or were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (D. and E.). P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001. Data was analysed separately for each corresponding day and the values for CpG vs. poly I:C were compared.

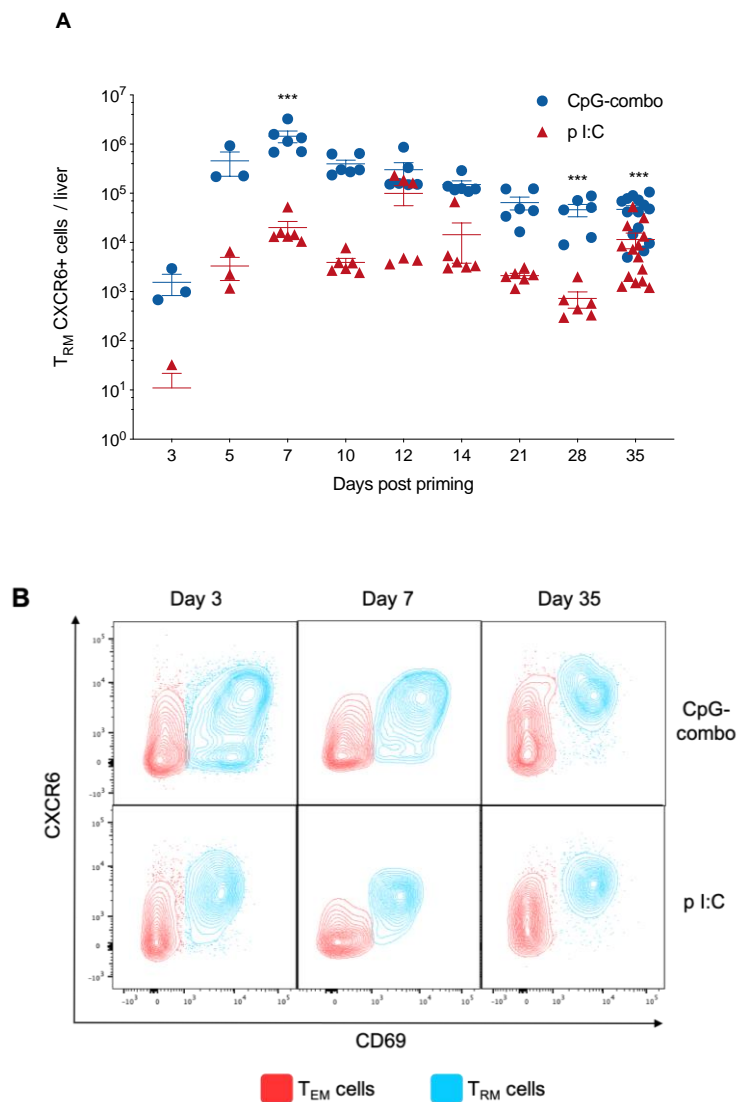


Figure 3.3. Kinetics of CXCR6+ PbT-I cells in  $\alpha$ Clec9-Pb-1 primed mice. Analysis of CXCR6 expression by flow cytometry from the previous experiment. A. Kinetics of the number of PbT-I  $T_{RM}$  cells expressing CXCR6 ( $CD8^+$ ,  $GFP^+$ ,  $CD44^{Hi}$ ,  $CD62L^-$ ,  $CD69^+$ ,  $CXCR6^+$ ) per liver. B. Representative flow cytometry plots for the expression of CD69 and CXCR6 in  $T_{RM}$  and  $T_{EM}$  like cells at three different time points. Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test.  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ . Data was analysed separately for each corresponding day and the values for CpG vs. poly I:C were compared.

slowly decreased until day 35. As expected, early after priming (day 3 – day 7) there was a population of T<sub>RM</sub> like cells negative for the expression of CXCR6, whereas on day 35 almost all T<sub>RM</sub> like cells express CXCR6 (Fig. 3.3 B). Again, CpG induced the highest number of T<sub>RM</sub> CXCR6+ cells in the liver compared to p I:C.

Together these results suggest that immunisation with  $\alpha$ Clec9-Pb-1 mAb is able to induce the formation of memory PbT-I cells and that CpG has a greater capacity to induce the generation of liver T<sub>RM</sub> cells than p I:C, as previously reported (Fernandez-Ruiz et al., 2016).

### *3.2.2 Endogenous response and protection induced by immunisation with the Pb-1 antigen:*

After demonstrating the efficacy of  $\alpha$ Clec9-Pb-1 mAb at inducing PbT-I cell expansion and memory formation, it was sought to assess the capacity of this reagent to evoke an endogenous response in B6 mice.

It has been shown that the frequency of peptide-specific naive T cell precursor determines to a certain extent the magnitude and expansion of the subsequent T cell response after exposure to the specific peptide (Jenkins and Moon, 2012; Moon et al., 2007; Obar et al., 2008). Thus, the number of naive CD8 T cell precursors harbouring a TCR specific for the Pb-1 epitope was first determined.

For this purpose, single cell suspensions of the spleen and major lymph nodes of naive B6 mice were generated and tetramer-associated magnetic enrichment was applied to those single cell suspensions. Tetramers for the Pb-1 and the TRAP<sub>130-138</sub> epitopes were used. PbTRAP<sub>130-138</sub> is an H-2D<sup>b</sup>-restricted epitope within the TRAP protein, identified as partially protective for B6 mice (Hafalla et al., 2013). Additional aspects of the TRAP protein

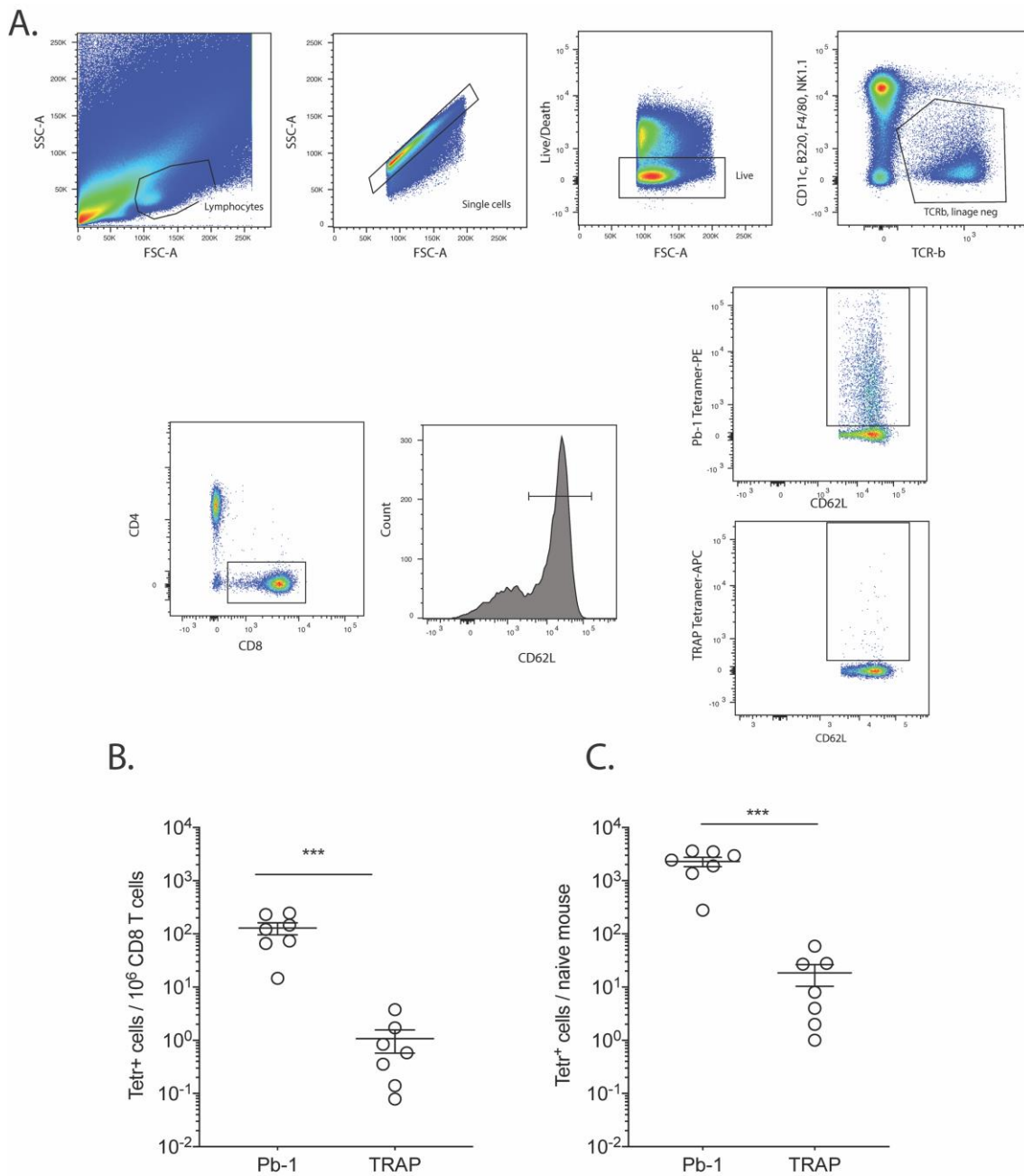


Figure 3.4. The naive Pb-1-specific CD8 T cell repertoire is very large. CD8 T cells specific for Pb-1 and the TRAP<sub>130-138</sub> epitopes were enriched from single cell suspensions of the spleen and major lymph nodes of naive B6 mice using specific tetramers. Cells were then stained for TCR $\beta$ , CD8, CD4, CD11c, B220, F4/80, NK1.1 and CD62L. A. Gating strategy on a representative sample: Naive tetramer-positive cells were defined as TCR $\beta$ <sup>+</sup>, lineage<sup>-</sup>, CD4<sup>-</sup>, CD8<sup>+</sup>, CD62L<sup>Hi</sup> and Pb-1 tetramer-positive. Number of tetramer-positive cells per million CD8 T cells (B) and per mouse (C) are displayed. Results were pooled from two independent experiments. Data were log-transformed and compared by unpaired T-test. P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001.

and this particular epitope were reviewed on Chapter 1. The number of TRAP<sub>130-138</sub> precursors has been reported (Van Braeckel-Budimir et al., 2017); and was used here as an internal control for the technique. Cells were stained for TCR $\beta$ , CD8, CD4, CD11c, B220, F4/80, NK1.1 and CD62L plus either the tetramer for Pb-1 or for TRAP<sub>130-138</sub>. Naive tetramer-positive cells were defined as TCR $\beta$ <sup>+</sup>, lineage<sup>-</sup>, CD4<sup>-</sup>, CD8<sup>+</sup>, CD62L<sup>Hi</sup> and tetramer-positive (Figure 3.4 A).

The frequency of TRAP-specific naive precursors ranged between 1 and 59 cells per mouse; with these numbers consistent with a previous report (Van Braeckel-Budimir et al., 2017). On the other hand, the average number of Pb-1-specific cells in naive B6 mice averaged  $2.3 \pm 0.4 \times 10^3$  cells per mouse, with variation ranging from just 278 to 3619 (Fig. 3.4 C). When compared to other well-known specificities such as OVA<sub>257-264</sub> or LCMV GP<sub>33-41</sub> with  $183 \pm 33$  and  $358 \pm 40$  cells per mouse, respectively (Jenkins and Moon, 2012), the number of Pb-1-specific naive precursors can be considered as extremely high. This compares well to GAP50<sub>40-48</sub>, another *P. berghei* epitope, recently reported to exhibit a large number of naïve precursors in the range of  $2.9 \pm 0.4 \times 10^3$  cells per mouse (Van Braeckel-Budimir et al., 2017).

Subsequently, the endogenous response of Pb-1-specific CD8 T cell response was studied in the context of natural infection. Data confirmed expansion of Pb1-specific precursors after blood-stage infection and memory formation in the liver after RAS immunisation. These experiments were performed by two postdoctoral researchers from our group, Maria Menezes and Nazanin Ghazanfari. For the blood infection experiment, naive B6 mice were infected with iRBC and 7 days later the number of tetramer-specific CD8 T cells was determined in spleen. For the RAS immunisation experiment, mice were injected with RAS and 25 days later, the number of specific memory cells was determined in the spleen and the liver. A large expansion of activated cells was observed in the spleen after blood stage compared to naive mice (Fig. 3.5 A), as was expansion of memory specific CD8 T cells in response to RAS immunisation (Fig. 3.5 B). The above findings showed that Pb-1-specific CD8 T cells could

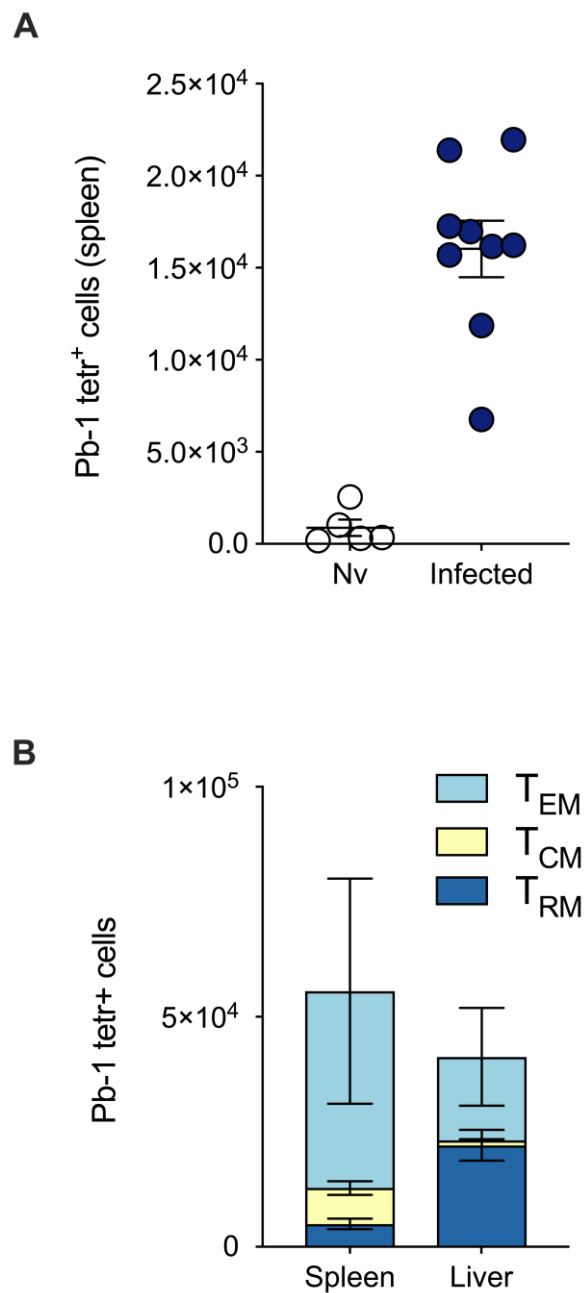


Figure 3.5. Expansion of Pb-1-specific CD8 T cells in response to blood stage and RAS infection. A. Mice were infected with  $10^4$  PbA iRBC, 7 days later spleens were dissected, and single cell suspensions were stained with Pb-1 tetramer and CD44. Activated Pb-1-specific CD8 T cells are displayed. B. Spleen and livers from mice primed with  $5 \times 10^4$  PbA RAS were collected 25 days post infection. Single cell suspensions were stained with Pb-1 tetramer, CD8, CD44, CD62L and CD69. Numbers of specific memory CD8 T cells subpopulations are displayed. Results were pooled from two independent experiments. Data were log-transformed and compared by unpaired T-test.  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ . Data obtained by Maria Menezes and Nazanin Ghazanfari modified from (Valencia-Hernandez et al., 2020).



respond to both blood-stage and RAS PbA in B6 mice but did not address the protective capacity of this response against liver-stage infection. The next series of experiments, therefore, addressed whether immunisation with  $\alpha$ Clec9-Pb-1 mAb could elicit a protective memory response against the liver stage of malaria through stimulation of the endogenous T cell repertoire.

Mice were immunised with 2  $\mu$ g or 16  $\mu$ g of  $\alpha$ Clec9-Pb-1 or 2  $\mu$ g of  $\alpha$ Clec9-NVY plus CpG; some mice were transferred with PbT-I cells and used as control for the  $\alpha$ Clec9 mAb priming, while other mice were left without cell transfer to monitor the response of the endogenous repertoire. 35-46 days later, when the memory cells had been formed, spleens and livers of some mice were analysed by flow cytometry for memory cell formation; and the remaining mice were challenged intravenously with live sporozoites on day 35 (Fig. 3.6 A).

After the challenge, blood samples were assessed for the presence of parasites up to day 10 post-infection. Mice were considered sterile protected when no blood stage parasites were detected 10 days after challenge.

Mice transferred with PbT-I cells and immunised with 2  $\mu$ g of  $\alpha$ Clec9-Pb1 + CpG generated similar numbers of liver  $T_{RM}$  cells to those that did not receive PbT-I cells and were immunized with 16  $\mu$ g of  $\alpha$ Clec9-Pb1 + CpG (Fig. 3.6 B). In absence of PbT-I cells, mice vaccinated with high dose of Ab generated on average  $1 \pm 0.2 \times 10^5$  tetramer-positive  $T_{RM}$  cells per liver. On the other hand, mice that did not receive PbT-I cells and were immunised with 2  $\mu$ g of  $\alpha$ Clec9-Pb1 generated significantly less liver  $T_{RM}$  cells. A similar pattern was also observed for the other memory populations of  $T_{EM}$  and  $T_{CM}$  cells in the spleen and the liver (Fig. 3.6 C, D).

Seven days after challenge all naive control mice were parasitemic; mice that received PbT-I cells and were primed with 2  $\mu$ g of  $\alpha$ Clec9-NVY as well as

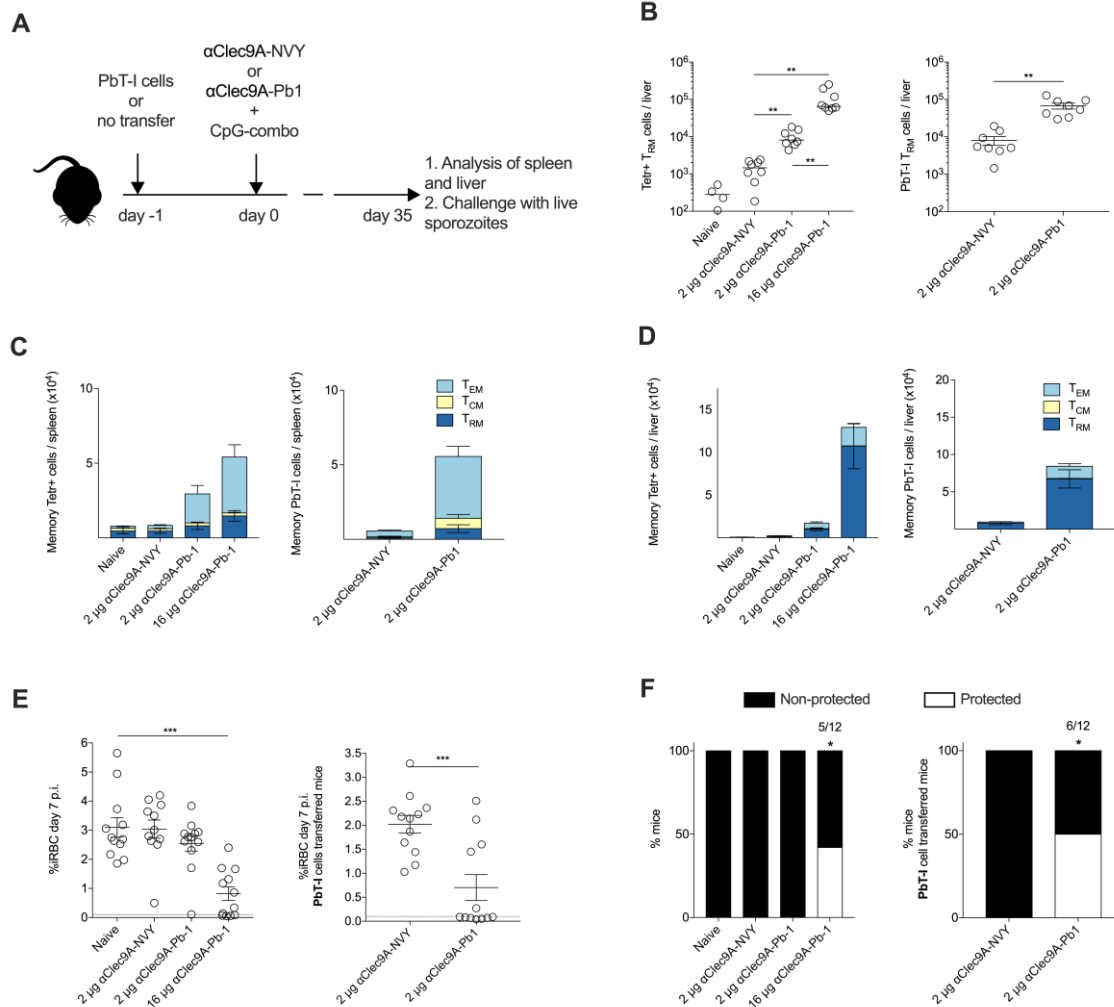


Figure 3.6 Immunisation with  $\alpha$ Clec9-Pb-1 mAb protects B6 mice against sporozoite challenge. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells or no cells and were immunised with  $\alpha$ Clec9-Pb-1 or  $\alpha$ Clec9-NVY plus 75  $\mu$ g of CpG on the next day. Spleens and livers from 4 mice were collected on day 35 or 46. Single cell suspensions were generated and the number of PbT-I or tetramer-Pb1 positive cells and memory status were determined by flow cytometry. The remaining mice (n=6) were challenged intravenously with 200 PbA sporozoites on day 35. Mice were monitored daily for parasitemia from day 6 until day 10 after challenge, when the percentage of sterile protection was determined. A. Schematic for the protection experiment. B. Number of  $T_{RM}$  cells per liver determined as CD8+ GFP+ or Tetr+, CD44Hi, CD62L-, CD69+. C. and D. Number of the CD8+ memory cells in spleen and liver. E. Percentage of infected red blood cells (iRBC) in blood on day 7 post challenge. F. Percentage of sterile protection determined by absence of blood stage infection by day 15 after challenge. Results were pooled of two independent experiments. Data were log-transformed and compared by one-way ANOVA with Tukey's multiple comparisons test in (B.), using a Kruskal-Wallis test with Dunn's multiple comparison post-test in (E.) or using Fisher's exact test (F.).  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ .

mice with no PbT-I cells immunised with either 2  $\mu\text{g}$  of  $\alpha\text{Clec9-NVY}$  or 2  $\mu\text{g}$   $\alpha\text{Clec9-Pb-1}$  were also positive for parasitemia on day 7 post infection. In contrast, 7 mice that received PbT-I cells and were primed with 2  $\mu\text{g}$   $\alpha\text{Clec9-Pb1}$  and 5 mice with no PbT-I cells injected with 16  $\mu\text{g}$   $\alpha\text{Clec9-Pb1}$  were negative for parasitemia (Fig. 3.6 E). Data from day 10, revealed that 41% of the mice with a normal endogenous T cell repertoire that were immunised with 16  $\mu\text{g}$  of  $\alpha\text{Clec9-Pb1}$  mAb plus CpG successfully controlled the liver infection and therefore were considered sterilely protected (Fig. 3.6 F).

This is the first time with this system that vaccinated B6 mice challenged with PbA liver-stage parasites become sterile protected in the absence of transgenic PbT-I cells. This suggests the Pb-1 epitope is an excellent antigen for the induction of protective immunity.

### *3.2.3 Induction of memory CD8 T cells using the Pb-1 epitope in prime-and-trap vaccination:*

“Prime-and-trap” is a three-component vaccine strategy used to induce liver  $T_{\text{RM}}$  cell formation through the priming of naive immune cells and the trapping of those cells in the liver (Fernandez-Ruiz et al., 2016). The first component is based on the use of an  $\alpha\text{Clec9}$  mAb coupled to a relevant epitope to achieve epitope delivery to cDC1 (Caminschi et al., 2008; Lahoud et al., 2011); which is the main antigen presenting cell stimulating T cells in malaria (Fernandez-Ruiz et al., 2017; Lau et al., 2014). The second component is a recombinant adeno-associated virus (rAAV) engineered to selectively infect hepatocytes (Tay et al., 2014), inducing the trapping of the T cells to the liver. This vector is also modified for the expression of the relevant epitope under the control of a hepatocyte-specific promoter. The third and last component, is a CpG adjuvant used for the licensing of DC involve in CD8 T cell priming (Lahoud et al., 2011) and to promote proinflammatory liver-microenvironment (Huang et al., 2013), which aids  $T_{\text{RM}}$  cell formation (Holz et al., 2018; Mackay et al., 2012) . In the past, it has been shown

that mice transferred with PbT-I cells and vaccinated with prime-and-trap using the mimic NVY epitope are sterilely protected against sporozoite challenge (Fernandez-Ruiz et al., 2016).

After confirming the efficacy of  $\alpha$ Clec9-Pb-1 mAb at promoting the expansion and memory formation of specific-endogenous CD8 T cells, the ability of the prime-and-trap vaccine to generate specific-endogenous liver T<sub>RM</sub> cells and induce protection against malaria was assessed using the Pb-1 epitope. To achieve this goal, reagents containing the Pb-1 epitope, including  $\alpha$ Clec9 mAb and rAAV particles expressing the Pb-1 epitope, were generated.

The priming capacity of the rAAV-Pb-1 alone (i.e. “trap” only) on the endogenous repertoire was first investigated. Mice with a normal endogenous T cell repertoire or that had been transferred with PbT-I cells were injected with increasing doses of rAAV-Pb-1. Livers and spleens were collected on day 35 post-injection when the viral particles has been cleared (Tay et al., 2014).

Analysis of the organs failed to identify specific T cells (data not shown). Spleens and livers were collected and analysed by flow cytometry for the presence of PbT-I cells or Pb-1 tetramer-positive cells. Neither GFP+ nor tetramer-positive cells were detected on day 35 in spleen or liver, indicating that the rAAV-Pb-1 on its own had no effect on the priming and or recruiting of PbT-I cells or Pb-1-specific cells. Endogenous non-specific cells were also analysed, to determine if this vaccine component in any way disturbed these populations. No differences in the number of memory cells in the liver or spleen were observed (including T<sub>RM</sub> cells) among groups injected with different doses of rAAV-Pb-1 or control naive mice.

While priming with rAAV-Pb-1 alone was ineffective, it was still possible that when combined with  $\alpha$ Clec9-Pb-1 mAb and CpG, liver T<sub>RM</sub> formation might be boosted. To test this possibility, mice were transferred with PbT-I cells and were immunised with  $\alpha$ Clec9-Pb-1 mAb, CpG and rAAV-Pb-1 on the next day.

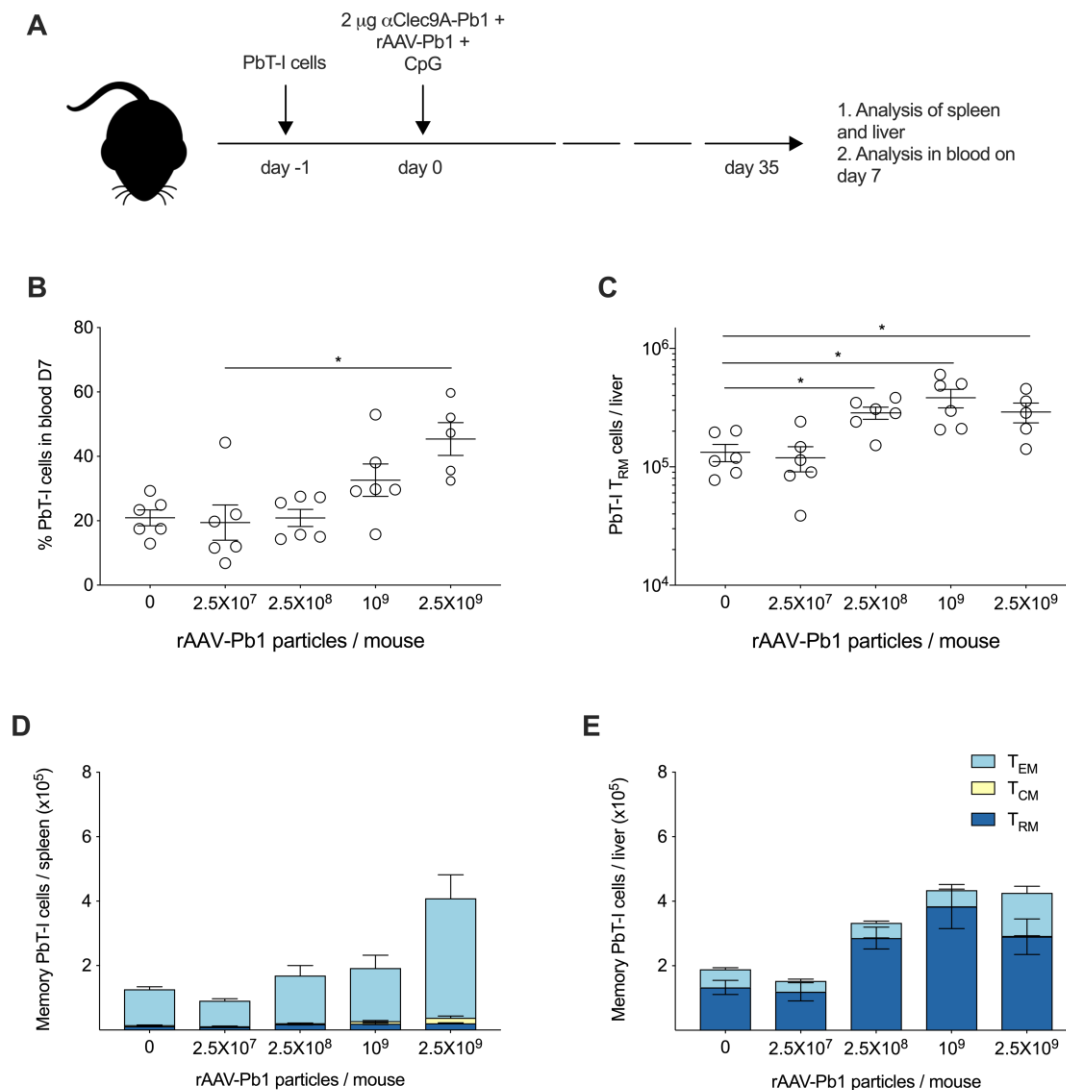


Figure 3.7. Prime-and-trap-Pb-1 induced high numbers of PbT-I  $T_{RM}$  cells in the liver. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells and next day immunised with 2  $\mu$ g of  $\alpha$ Clec9-Pb-1 plus 75  $\mu$ g of CpG plus different doses of rAAV-Pb-1. Mice were bled on day 7 and number of PbT-I cells was determined by flow cytometry. Spleens and livers were collected on day 35, for the analysis of memory formation. A. Schematic of the mice immunisation B. Percentage of PbT-I cells in blood on day 7, calculated as the frequency of total CD8+ cells. Data were compared by one-way ANOVA and Kruskal-Wallis test with Dunn's multiple comparison post-test. C. Number of  $T_{RM}$  cells per liver. Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test. C. and E. Number of the PbT-I CD8+ memory cells in spleen and liver, respectively. Results were pooled of two independent experiments.  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ .

Four different doses of rAAV-Pb-1 were tested, ranging from  $2.5 \times 10^7$  to  $2.5 \times 10^9$  viral particles per mouse; and one group were primed with  $\alpha$ Clec9-Pb-1 mAb and CpG without rAAV-Pb-1. Livers and spleens were analysed on day 35.

On day 7 after priming, 20% of total CD8 T cells corresponded to PbT-I cells in the blood of mice immunised with  $\alpha$ Clec9-Pb-1 mAb and CpG alone; this percentage increased up to 60% in mice that received prime-and-trap with a dose of  $2.5 \times 10^9$  rAAV-Pb-1 viral particles (Fig. 3.7 B). The percentage of PbT-I cells in blood correlated roughly with the number of PbT-I  $T_{RM}$  cells in the liver on day 35, though a slight decrease in liver  $T_{RM}$  numbers occurred at the highest viral dose (Fig. 3.7 C). The maximum number of PbT-I  $T_{RM}$  cells per liver was achieved in mice vaccinated with 2  $\mu$ g of  $\alpha$ Clec9-Pb-1 mAb, CpG and  $10^9$  rAAV-Pb-1, with a mean of  $3.8 \pm 0.7 \times 10^5$   $T_{RM}$  cells per liver. Mice vaccinated with a dose equal or higher than  $2.5 \times 10^8$  rAAV-Pb-1 formed significantly higher numbers of liver  $T_{RM}$  cells compared to mice that only got  $\alpha$ Clec9-Pb-1 mAb plus CpG (Fig. 3.7 C). Other memory subpopulations showed a similar pattern; in fact, increasing numbers of spleen  $T_{EM}$  cells formed in mice vaccinated with the highest doses of rAAV-Pb-1 (Fig 3.7 D, E).

These results demonstrated that indeed rAAV-Pb-1 is able to boost liver  $T_{RM}$  cell formation in the PbT-I system. Subsequently, the same effect was studied in mice with normal endogenous T cell repertoire i.e. no transferred PbT-I cells. This experiment assessed the expansion potential of Pb-1 specific precursor cells. In this case, mice were immunised with  $\alpha$ Clec9-Pb-1 mAb, CpG and increasing doses of rAAV-Pb-1. Stained cells from livers and spleens were analysed on day 35 (Fig. 3.8 A). Unlike the previous experiment, no differences were evident in the percentage of Pb-1 tetramer-positive cells in the blood on day 7 (Fig 3.8 B). However, on day 35 after immunisation, the number of liver  $T_{RM}$  cells increased with the amount of rAAV-Pb-1 virus administered (Fig. 3.8 C, E).

The maximum number of Pb-1 tetramer-positive  $T_{RM}$  cells per liver was achieved when mice were vaccinated with  $\alpha$ Clec9-Pb-1 mAb, CpG and  $2.5 \times 10^9$

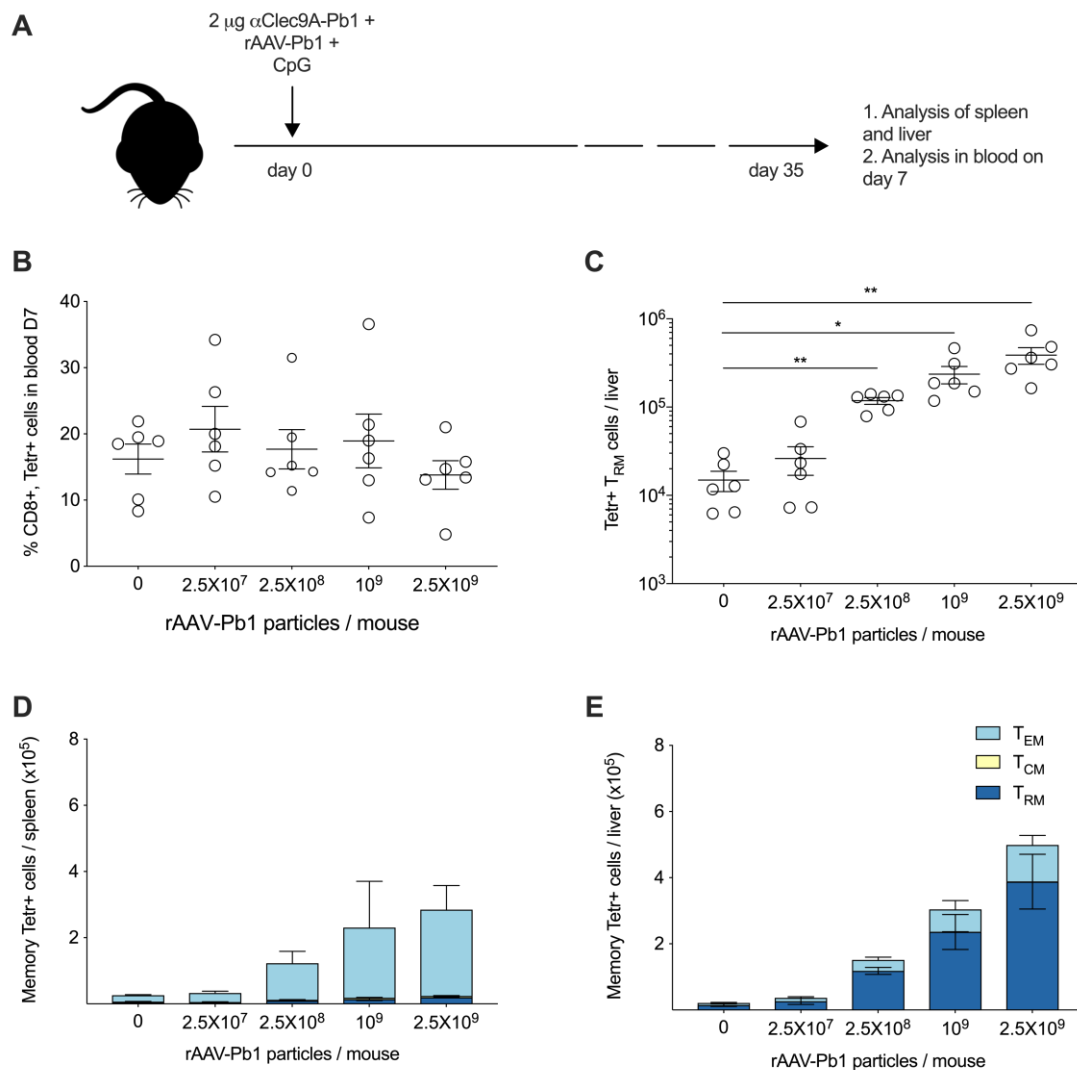


Figure 3.8. Prime-and-trap-Pb-1 induced high numbers of T<sub>RM</sub> cells in the liver of B6 mice with endogenous repertoire. B6 mice were immunised with 2 µg of  $\alpha$ Clec9-Pb-1 plus 75 µg of CpG plus different doses of rAAV-Pb-1. Mice were bled on day 7 and number of Pb-1 tetramer-positive cells was determined by flow cytometry. Spleens and livers were collected on day 35, single cell suspensions were generated and the number of Pb-1 tetramer-positive cells and memory cell subset were determined by flow cytometry. A. Schematic of the mice immunisation B. Percentage of tetramer-positive cells in blood on day 7. C. Number of T<sub>RM</sub> cells per liver. D. and E. Number of the Pb-1 tetramer-positive, CD8<sup>+</sup> memory cells in spleen and liver, respectively. Results were pooled of two independent experiments. Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test. P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001.

rAAV-Pb-1 (mean of  $3.8 \times 10^5$  T<sub>RM</sub> cells per liver). Mice vaccinated with a dose equal or higher than  $2.5 \times 10^8$  rAAV-Pb-1 developed significantly higher liver T<sub>RM</sub> cells compared to mice that only received  $\alpha$ Clec9-Pb-1 mAb plus CpG (Fig. 3.8 C).

Similar patterns were evident for the other memory sub-populations (Fig. 3.8 D, E). Together, these data confirmed the boosting capacity of rAAV-Pb-1 for the generation of liver T<sub>RM</sub> cells in the context of prime-and-trap vaccination in B6 mice with normal endogenous T cell repertoire.

Previous studies have shown that numbers of specific liver T<sub>RM</sub> cells correlate with levels of protection against sporozoite infection (Fernandez-Ruiz et al., 2016). Consequently, the next experiment sought to determine whether the number of cells could be increased by using even higher doses of the rAAV-Pb-1 virus. These experiments were performed only in mice with normal endogenous repertoire, in the absence of PbT-I cells. For this purpose, mice were immunised with  $\alpha$ Clec9-Pb-1 mAb, CpG and four different doses of rAAV-Pb-1, ranging from  $10^9$  to  $3.3 \times 10^{10}$  viral particles per mouse; and one group was primed only with  $\alpha$ Clec9-Pb-1 mAb and CpG. Livers and spleens were collected and processed for single cells staining and flow cytometry analysis on day 35 post immunisation.

Blood analysis on day 7 revealed a peak of Pb-1 specific CD8 T cells when injecting  $3.3 \times 10^9$  viral particles per mouse and lower numbers when using  $10^{10}$  or more viral particles per mouse (Fig. 3.9 B). The numbers of T<sub>RM</sub> cells in the livers mirrored this pattern, with a maximum mean of  $3.2 \times 10^5$  T<sub>RM</sub> cells per liver at  $3.3 \times 10^9$  viral particles and significantly reduced mean of  $5.7 \times 10^3$  T<sub>RM</sub> cells when injecting  $3.3 \times 10^{10}$  particles (Fig. 3.9 C). T<sub>EM</sub> and T<sub>CM</sub> cells in spleen and liver expanded and contracted in the same way as the liver T<sub>RM</sub> cells (Fig. 3.9 D, E).

In 2014, Tay and collaborators described how persisting high levels of antigen expression by rAAV promote the silencing of the CTL function (Tay et al.,



2014). In these studies, mice injected with high doses of rAAV-mOVA showed significantly less specific killing, high levels of programmed death-1 (PD-1) and T-cell immunoglobulin mucin 3 (Tim-3) expression in liver CD8 T cells and low expression of IFN- $\gamma$  upon *ex-vivo* restimulation (Tay et al., 2014). In fact, the expression of PD-1 has been associated with immunotolerance and CD8 T cell apoptosis mediated by PD-1 ligands in the context of chronic viral infections and persistence of antigen presentation (Golden-Mason et al., 2007; Okazaki and Honjo, 2006). Tim-3 is an inhibitory molecule induced during activation and involved in T-cell exhaustion and tolerance in cancers and chronic infections (Jones et al., 2008; Monney et al., 2002; Sabatos et al., 2003; Sakuishi et al., 2010; Sánchez-Fueyo et al., 2003). To investigate whether the reduction on the memory cell numbers that was observed at high doses of rAAV-Pb-1 is governed by a similar mechanism, the expression profile of Tim-3 and PD-1 was determined on day 35 after immunisation.

Indeed, activated specific CD8 T cells from the liver of mice vaccinated with high doses of rAAV-Pb-1 expressed significantly higher levels of PD-1 and Tim-3 (Fig 3.9 F, G). Very importantly, the expression of PD-1 was highly susceptible to small dose changes of the adenovirus. In fact, activated CD8 T cell from mice injected with  $3.3 \times 10^9$  rAAV-Pb-1 expressed significantly higher levels of PD-1 compared to mice that were injected with  $10^9$ . Higher levels of Tim-3 were observed on mice that received  $10^{10}$  or more viral particles. This result indicates that the levels of antigen expression greatly influence the numbers of liver  $T_{RM}$  cells generated, and therefore the doses of rAAV-Pb-1 should be considering carefully during the formulation of an efficient vaccine.

In summary, these findings revealed that prime-and-trap using the Pb-1 epitope as reagent induces very high numbers of liver specific  $T_{RM}$  cells in B6 mice with a normal endogenous repertoire or with transferred PbT-I cells. This number was maximized by immunisations with  $\alpha$ Clec9-Pb-1 mAb, CpG plus doses of rAAV-Pb-1 ranging from  $10^9$  to  $3.3 \times 10^9$  particles per mouse. However, raised levels of PD-1, a hallmark for T cell exhaustion (Okazaki and Honjo, 2006),

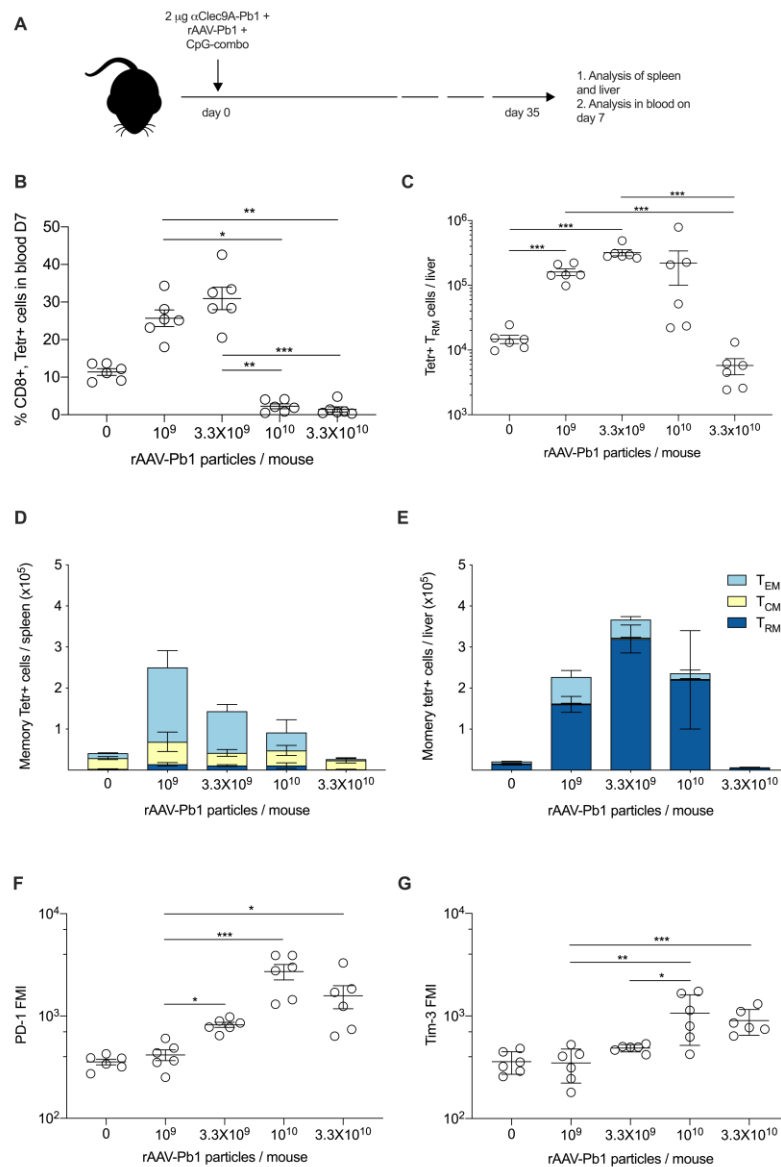


Figure 3.9. Adverse effect of very high doses of rAAV-Pb-1 on the liver T<sub>RM</sub> cell formation. B6 mice were immunised with 2  $\mu$ g of  $\alpha$ Clec9-Pb-1, 75  $\mu$ g of CpG and different high doses of rAAV-Pb-1. Mice were bled on day 7 and number of Pb-1 tetramer-positive cells was determined by flow cytometry. Spleens and livers were collected on day 35, single cell suspensions were generated and the number of Pb-1 tetramer-positive cells and memory status were determined by flow cytometry. A. Schematic of the mice immunisation B. Percentage of tetramer-positive cells in blood on day 7; data were compared by Kruskal-Wallis test with Dunn's multiple comparison post-test. C. Number of T<sub>RM</sub> cells per liver. D. and E. Number of the Pb-1 tetramer-positive, CD8<sup>+</sup> memory cells in spleen and liver, respectively. Analysis of PD-1 and Tim-3 expression on activated (CD44<sup>Hi</sup>) liver CD8 T cells from mice vaccinated with prime-and-trap and high doses of rAAV-Pb-1. Fluorescence mean intensity for PD-1 (F.) and Tim-3 (G.). Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test. P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001.

were observed on liver CD8 T cells from mice injected with  $3.3 \times 10^9$ . Consequently, it was determined the best dose of rAAV-Pb-1 for prime-and-trap to be between  $2.5 \times 10^9$  and  $10^9$  viral particles per mouse.

#### *3.2.4 Analysis of sterile protection against sporozoite infection by prime-and-trap vaccination using the Pb-1 antigen:*

Given the significant increase in liver T<sub>RM</sub> cells induced by the introduction of the rAAV-Pb-1 component, the capacity of prime-and-trap with the Pb-1 epitope to induce sterilizing immunity in B6 mice challenged with PbA sporozoites was assessed.

On previous experiments, the dose of Ab was 2 µg of αClec9-Pb-1 mAb per mouse and the optimal dose of rAAV-Pb-1 ranged between 1- $2.5 \times 10^9$  viral particles. For a first set of experiments we sought to investigate if higher doses of Ab could revert the upregulation of PD-1 and Tim-3 at high rAAV doses. For this, five different B6 mice groups were analysed. Two groups received 2 µg of αClec9-Pb-1 mAb plus CpG plus either  $2.5 \times 10^9$  or  $10^{10}$  of rAAV- Pb-1. The other two groups received 8 µg of αClec9-Pb-1 mAb plus CpG plus either  $2.5 \times 10^9$  or  $10^{10}$  of rAAV- Pb-1. The last group remained naive. On day 35, 3 mice per group were used for the enumeration of memory T cells populations in liver and spleen. The remaining mice (n=6) were challenged intravenously with 200 sporozoites and sterile protection was determined on day 18 post infection.

While there was a tendency towards higher numbers, there was no significant increase in liver T<sub>RM</sub> cells in mice vaccinated with the higher dose of αClec9-Pb-1 mAb (Fig. 3.10 C). On the other hand, the percentage of Pb-1 specific, circulating CD8 T cells on day 7 post immunisation was lower in mice injected with high doses of the viral particles compared to the mice injected with  $2.5 \times 10^9$  rAAV- Pb-1 (Fig. 3.10 B). A similar tendency was found on the number of liver T<sub>RM</sub> cells (Fig. 3.10 C), though this did not reach significance. PD-1 and Tim-3 expression was significantly higher on memory CD8 T cells of mice injected

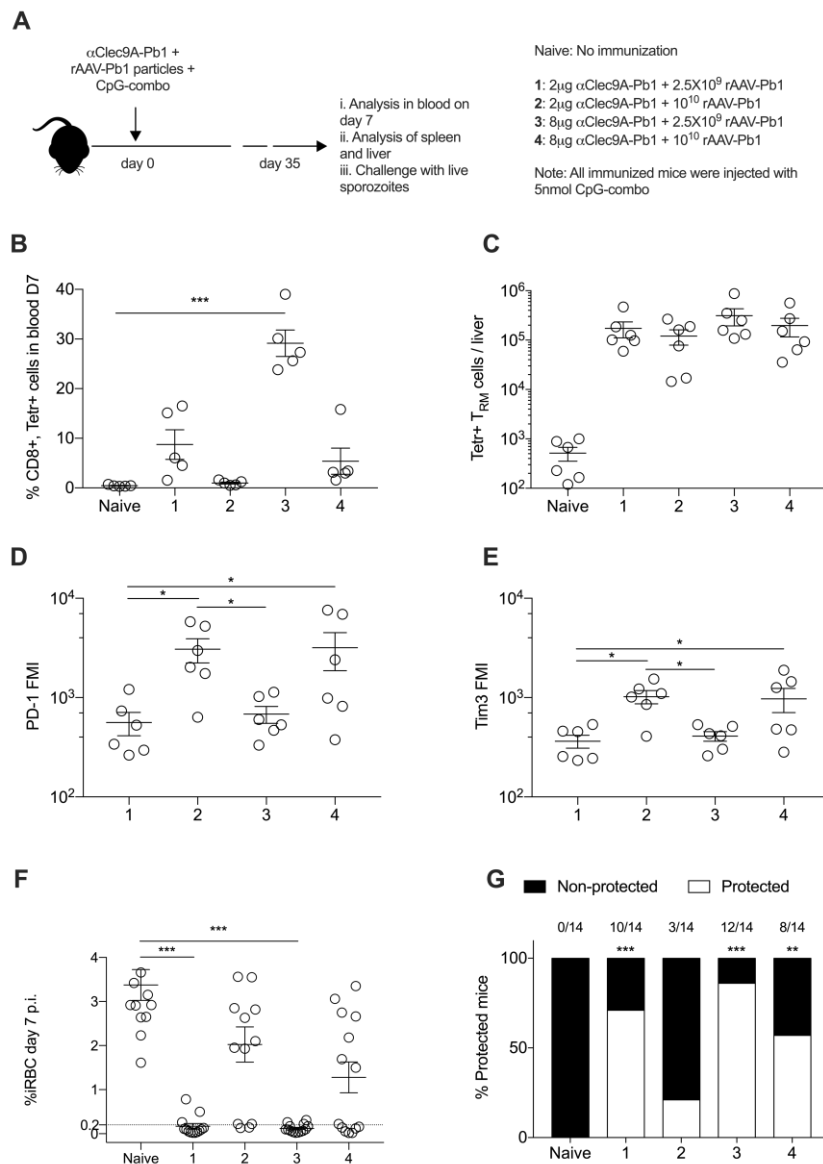


Figure 3.10 Effect of high doses of rAAV-Pb-1 on the induction of protection in mice vaccinated with Prime-and-trap-Pb-1. B6 mice were immunised with two different doses of  $\alpha$ Clec9-Pb-1, 75  $\mu$ g of CpG and two different doses of rAAV-Pb-1. One group of mice remained naive as a control for the viability of the sporozoites. Spleens and livers from 4 mice were collected on day 35 or 46. Single cell suspensions were generated. The remaining mice (n=6) were challenged intravenously with 200 living PbA sporozoites on day 35. A. Schematic of the mice immunisation B. Percentage of tetramer-positive cells in blood on day 7. C. Number of  $T_{RM}$  cells per liver D. and E. Fluorescence mean intensity for PD-1 and Tim-3. F. Percentage of infected red blood cells (iRBC) in blood on day 7 post challenge. G. Percentage of sterile protection determined by day 18 after challenge. Results are a pool of two independent experiments. Data were compared using Kruskal- Wallis test with Dunn's multiple comparison post-test in (B. and F.), log-transformed and compared using one-way ANOVA and Tukey's multiple comparisons test in (D. and E.) or using Fisher's exact test in (G.).  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ .

with  $10^{10}$  rAAV- Pb-1, compared to the lower dose of viral particles (Fig. 3.10 D, E). Protection was clearly affected by the use of high doses of the viral particles (Fig. 3.10 F, G). Mice injected with the lowest dose of viral particles showed a lower percentage of infected red blood cells on day 7 post infection and superior sterile protection. Importantly, approximately 80% sterile protection was achieved in mice vaccinated with  $8 \mu\text{g}$  of  $\alpha\text{Clec9- Pb-1}$  +  $2.5 \times 10^9$  rAAV- Pb-1 +  $75 \mu\text{g}$  CpG (Fig. 3.10 G).

These results together with the observation that memory cells from mice injected with  $3.3 \times 10^9$  rAAV-Pb-1 expressed significantly higher levels of PD-1 compared to those injected with  $10^9$ , led to the speculation that the optimal dose of viral particles for the for prime-and-trap-Pb-1 is  $10^9$  rAAV-Pb-1.

During the first round of experiments (Fig. 3.1 B and 3.6 B), it was observed that  $16 \mu\text{g}$  of  $\alpha\text{Clec9-Pb-1}$  mAb induced the highest number of liver  $T_{RM}$  cells when used alone, i.e. without rAAV-Pb-1 particles. Consequently, the role of this high dose of  $\alpha\text{Clec9-Pb-1}$  mAb was investigated in the context of a malaria infection in prime-and-trap vaccinated mice.

B6 were vaccinated with either  $2 \mu\text{g}$  or  $16 \mu\text{g}$  of  $\alpha\text{Clec9-Pb-1}$  mAb, CpG and  $10^9$  rAAV-Pb-1 (some mice were transferred with PbT-I cells as an experimental control). On day 35, when the virus has been cleared and thus memory cells have formed, some mice were used for the numeration of memory cells in spleen and liver, while the remaining mice were infected intravenously with 200 sporozoites as indicated on figure 3.11 A. The average number of liver  $T_{RM}$  cells generated on day 35 ranged between  $2.3$  to  $3.3 \times 10^5$  cells per liver among all the different groups (Fig. 3.11 B). Surprisingly, mice transferred with PbT-I cells prior vaccination did not form more liver  $T_{RM}$  cells than mice with a normal endogenous T cell repertoire. Furthermore, increasing the dose of  $\alpha\text{Clec9-Pb-1}$  from  $2 \mu\text{g}$  up to  $16 \mu\text{g}$  had no effect on the formation of  $T_{RM}$  cells. The number of effector memory cells in spleen and liver were similar among the

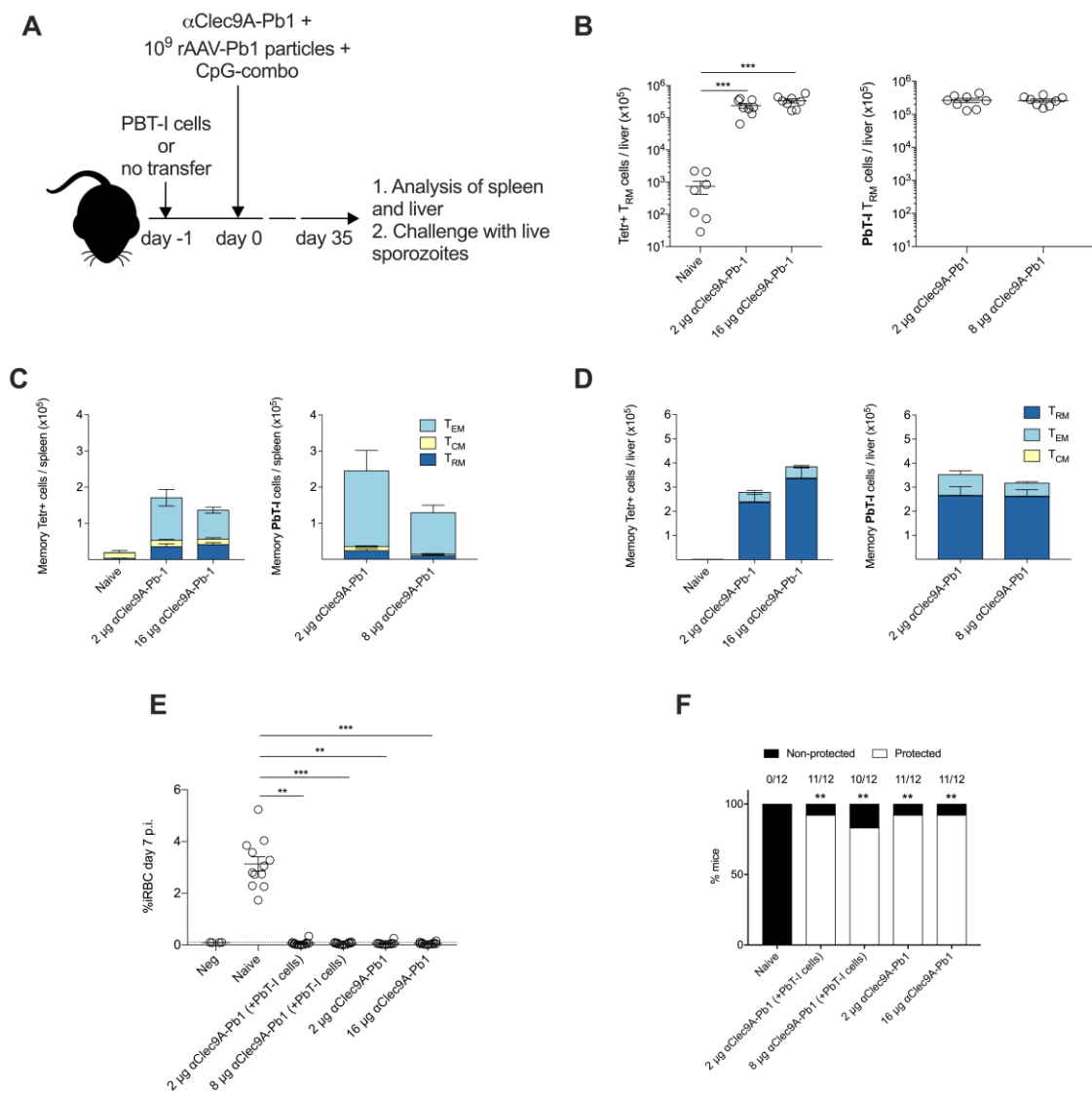


Figure 3.11 Prime-and-trap-Pb-1 induces more than 90% protection in B6 mice against sporozoite challenge. B6 mice were transferred with 50,000 PbT-I cells or no cells one day prior vaccination. Next day, mice immunised with different doses of  $\alpha\text{Clec9-Pb-1}$ , 75  $\mu\text{g}$  of CpG and  $10^9$  viral particles of rAAV-Pb-1. Spleens and livers from 4 mice were collected on day 35. Single cell suspensions were generated. The remaining mice ( $n=6$ ) were challenged intravenously with 200 living PbA sporozoites on day 35. A. Schematic of the mice immunisation B. Number of specific liver  $\text{T}_{\text{RM}}$  cells C. Number of memory CD8 T cells in spleen and livers of mice transferred with PbT-I cells D. Number of memory CD8 T cells in spleen and livers of mice with normal endogenous T cell repertoire E. Percentage of infected red blood cells (iRBC) in blood on day 7 post challenge. F. Percentage of sterile protection determined by day 18 after challenge. Results are a pool of two independent experiments. Data were log-transformed and compared using one-way ANOVA and Tukey's multiple comparisons test in (B.), using Kruskal-Wallis test with Dunn's multiple comparison post-test in (E.), or using Fisher's exact test in (F.).  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ .

different groups (Fig. 3.11 C and D). The levels of parasitemia on day 7 post infection were very low; additionally, more than 90% of the mice with a normal endogenous T cell repertoire or those transferred with PbT-I cells were sterilely protected against liver stage malaria (Fig. 3.11E and F).

In 2018 Gola and collaborators published a study using a similar vaccination principle to prime-and-trap, to protect B6 mice from *P. berghei* sporozoite infection, which they termed prime-and-target. For this strategy, they first primed the mice by injecting them intramuscularly with a modified adenovirus expressing an antigen of interest; and then two weeks later, they target the circulating specific CD8 T cells to the liver by injecting intravenously nanoparticles containing the specific peptide (Gola et al., 2018). With this technique they achieved high numbers of OVA-specific liver-associated CD8 T cells conferring 100% protection against transgenic *P. berghei* parasites expressing the OVA peptide (Gola et al., 2018).

These findings from Gola and collaborators motivated the following experiments, in which the effect of a “delayed trapping” for the generation of liver T<sub>RM</sub> cells was tested: B6 mice were immunised with  $\alpha$ Clec9-Pb-1 mAb plus CpG and 14 days later they were injected with  $10^9$  rAAV-Pb-1 viral particles. On day 35 after the last injection the number of liver T<sub>RM</sub> cells was determined and mice were challenged with sporozoites.

The number of liver T<sub>RM</sub> cells was not affected by the “delayed trapping” of two weeks, as similarly high numbers of these cells were obtained in mice vaccinated with the original prime-and-trap or the “delayed trapping” immunisation (Fig. 3.12 B and D). This was reflected on the high levels of protection (i.e. 100%) observed for all vaccinated groups (Fig. 3.12 E and F).

In summary, these data suggest that vaccination with prime-and-trap using the Pb-1 epitope is able to confer high levels of sterile protection in B6 mice either transferred with PbT-I cells or through their normal endogenous T cell repertoire.

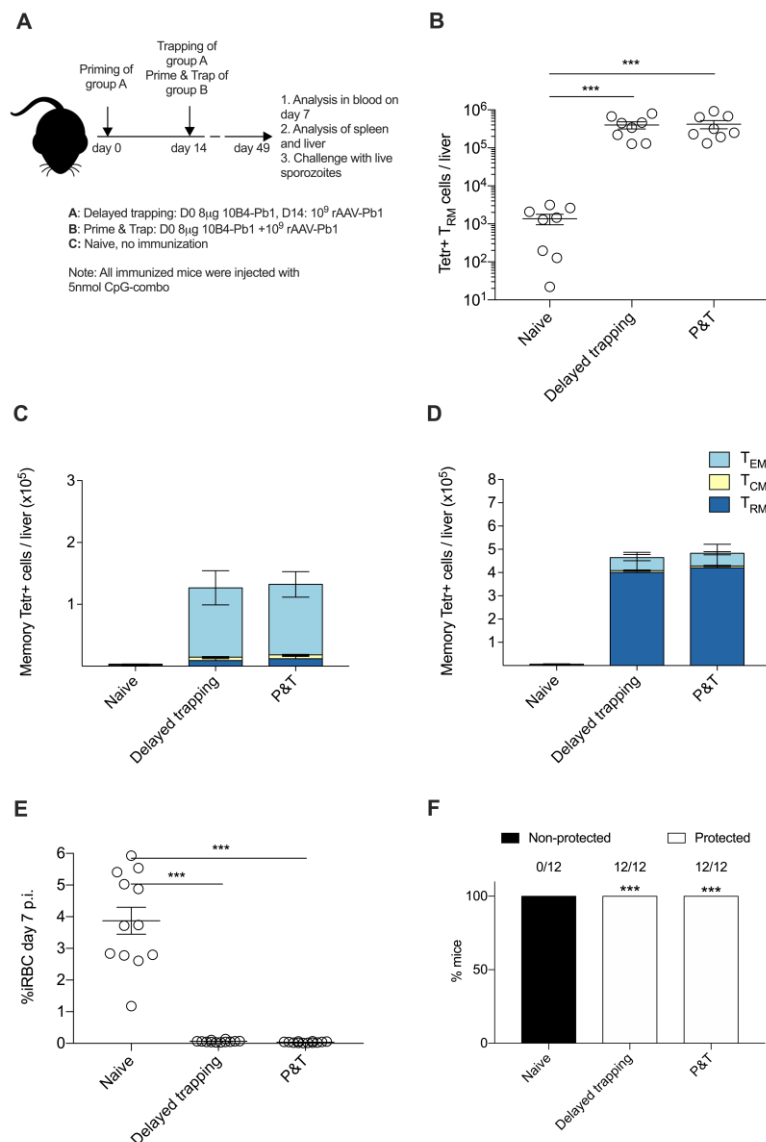


Figure 3.12 Delayed trapping with prime-and-trap-Pb-1. On day 0 a group of mice were injected with 8  $\mu$ g of  $\alpha$ Clec9- Pb-1 + 75  $\mu$ g CpG, 14 days later same group of mice were injected with 10<sup>9</sup> rAAV- Pb-1, while a second group of mice was vaccinated with 8  $\mu$ g of  $\alpha$ Clec9- Pb-1 + 10<sup>9</sup> rAAV- Pb-1 + 75  $\mu$ g CpG. Spleens and livers from 4 mice were collected on day 35. Single cell suspensions were generated. The remaining mice (n=6) were challenge intravenously with 200 living PbA sporozoites on day 35. A. Schematic of the mice immunisation B. Number of specific liver T<sub>RM</sub> cells C. and D. Number of memory CD8 T cells in spleen and liver E. Percentage of infected red blood cells (iRBC) in blood on day 7 post challenge. F. Percentage of sterile protection determined by day 18 after challenge. Results are a pool of two independent experiments. Data were log-transformed and compared using one-way ANOVA and Tukey's multiple comparisons test in (B.), using Kruskal- Wallis test with Dunn's multiple comparison post-test in (E.), or using Fisher's exact test in (F.). P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001..



Additionally, these results suggest that the optimal dose of prime-and-trap for the generation of liver T<sub>RM</sub> cells and induction of sterile protection ranged from 2 µg to 16 µg of αClec9- Pb-1 + 10<sup>9</sup> rAAV- Pb-1 + 75 µg CpG.

### 3.3 Discussion:

In this study, a series of different immunisation strategies targeting the Pb-1 epitope were implemented and specific CD8 T cell responses were assessed in B6 mice. Results demonstrated a large initial expansion of PbT-I cells in the spleen and the formation of relevant memory cell subpopulations in spleen and liver. Remarkably, analysis of the T cell repertoire of naive B6 mice revealed that the number of specific CD8 T cell precursors for the Pb-1 epitope was very large. These cells were successfully expanded using prime-and-trap with Pb-1 and large numbers of specific T<sub>RM</sub> cells were formed in the livers of immunised mice, which were efficiently protected from infection. In summary, the data presented in this chapter demonstrate the highly immunogenic nature of the Pb-1 epitope and its ability to confer high levels of sterile protection B6 mice with a normal endogenous T cell repertoire.

#### *3.3.1 Novel MHC-I restricted epitope derived from P. berghei ANKA:*

The critical role of CD8 T cell in protection against liver-stage malaria has been long confirmed by depletion and adoptive-transfer experiments in murine models (Jobe et al., 2007; Rodrigues et al., 1991; Schofield et al., 1987; Weiss et al., 1988). Despite many years of investigation, the quantitative and qualitative features of the protection and the effector functions mediated by CD8 T cells remain largely elusive. Only recently, liver-resident memory CD8 T cells have been implicated as an effective memory subpopulation for the induction of sterile protection (Fernandez-Ruiz et al., 2016; Tse et al., 2014). Other studies have also shown that γδ T cells play an important role in the memory response against liver-stage malaria (Ishizuka et al., 2016; Zaidi et al., 2017). The modest progress on revealing the nature of the of this CD8 T cell response, is partially explained

by the complexity of the *Plasmodium* genome and the limited number of defined MHC-I epitopes derived from the *Plasmodium* parasites (Schmidt et al., 2010). The identification and characterization of liver-stage derived antigens able to induce CD8 T cells responses through immunisation is critical for the development of new vaccine candidates (Van Braeckel-Budimir and Harty, 2014).

To date only two MHC-I restricted epitopes from PbANKA have been described to any level of protection against liver stage infection in B6 mice, a mouse strain that has been considered difficult to protect (Doolan and Hoffman, 2000; Schmidt et al., 2010). The first characterized epitope was the PbTRAP<sub>130-138</sub>. Hafalla and collaborators found that B6 mice vaccinated with a modified adenovirus expressing PbTRAP<sub>130-138</sub> showed 95% reduction in liver parasite load, but this vaccine did not induced sterile protection (Hafalla et al., 2013). The second epitope, PBANKA\_1031000<sub>631-639</sub> (Kb-17) is contained within the MIF4G-like protein of unknown function (Pichugin et al., 2018; Speake et al., 2016). Pichugin and colleagues showed that B6 mice primed intramuscularly with Kb-17 plasmid DNA and boosted intravenously with a recombinant adenovirus expressing Kb-17 had 7-fold less liver parasite burden compared to control mice (Pichugin et al., 2018). Nonetheless, this vaccination strategy using the Kb-17 epitope also failed to confer sterile protection in B6 mice.

In 2014, members of our group developed a B6 TCR transgenic line able to produce CD8 T cells for PbA. These cells were shown to respond to blood stage infection with different *Plasmodium*-species, to liver-stage parasites (PbA RAS) and to confer sterile protection against PbA sporozoite challenge (Lau et al., 2014). Very recently the authentic cognate epitope for the PbT-I cells, here termed Pb-1, was described. This MHC-I epitope is part of the ribosomal protein RPL6 of PbANKA (PBANKA\_1351900) with orthologs in *P. yoelli* (PY02722), and *P. chabaudi* (PCHAS\_1356500); and it is expressed during the liver and blood stage of the infection (Valencia-Hernandez et al., 2020).

In this chapter, the suitability of the Pb-1 epitope to evoke memory and induce protection mediated by two DC-targeted vaccination strategies, was investigated.

### *3.3.2 PbT-I cell expansion in response to immunisation targeting the Pb-1 epitope:*

Clec9A is DC surface protein specifically expressed by XCR1 and cDC1 subpopulations (Guilliams et al., 2014; Kroczeck and Henn, 2012; Shortman and Heath, 2010). Correspondingly, Clec9A is also selectively expressed on human BDCA3+ DC (Huysamen et al., 2008; Kerrigan and Brown, 2010; Poulin et al., 2010). Moreover, studies have demonstrated the important role of the cDC1 as the main antigen presenting cell stimulating T cells during blood-stage malaria infections (Fernandez-Ruiz et al., 2017; Lau et al., 2014).

Antigen targeting DC via Clec9A can enhance immune responses of CD8 T cell, CD4 T cell and B cells (Caminschi et al., 2008; Lahoud et al., 2011; Park et al., 2013). Other evidence, also showed that antigen targeting Clec9A in the absence of adjuvant, drives the development of long lasting follicular helper T cells and a potent antibody response (Kato et al., 2015; Park et al., 2013, 2017).

The nature of the CD8 T cell immune response induced by delivering peptides to DC via Clec9A has been investigated using model antigens such as OVA (Caminschi et al., 2008; Park et al., 2013; Sancho et al., 2008). These studies revealed the efficient formation of OVA-specific CTL responses in mice immunised with anti-Clec9A mAb together with an adjuvant. Furthermore, the induction of this specific CTL response at a memory time point accounted for the reduction of OVA-expressing lung tumour cells after or prior challenge (Sancho et al., 2008). These studies suggested that in principle delivering Ag via Clec9A could be exploited as an immunisation strategy to mediated CD8 T cell memory responses against intracellular infections.

For most of the experiments described in this chapter, the response of PbT-I cells was examined first, to test the effectiveness of priming reagents such as Clec9A mAb and the rAAV. Accordingly, mice adoptively transferred with PbA transgenic-specific CD8 T cells (i.e. PbT-I cells) and immunised with anti-Clec9A mAb targeting the Pb-1 epitope revealed a superior expansion of the PbT-I cells compared to mice immunised with the NVY mimic epitope. Relevant memory subpopulations, including T<sub>CM</sub>, T<sub>EM</sub> and T<sub>RM</sub>, were also formed in spleen and liver after immunisation with anti-Clec9A-Pb-1 mAb. A similar level of expansion has been achieved in mice adoptively transferred with OT-I cells and immunised with anti-Clec9A mAb targeting an OVA peptide (Holz et al., 2018). Supporting the idea that Pb-1 is the authentic cognate epitope for the PbT-I cells.

### *3.3.3 Endogenous response and protection induced by immunisation with the Pb-1 antigen:*

The magnitude and expansion of a given T cells response is partially determined by the frequency of the naive T cell precursor for a specific peptide (Jenkins and Moon, 2012; Moon et al., 2007; Obar et al., 2008). Therefore, to learn about the potential magnitude of the response to the Pb-1 epitope, the frequency of the naive T cell precursors for this epitope was determined. Results showed that the number of Pb-1 naive CD8 T cell precursors is extraordinarily high in naive B6 mice. In average, a female naive B6 mouse harbors approximately 2300 Pb-1 specific naive cells per mouse. On the other hand, the frequency of naive T cell precursor for PbTRAP<sub>130-138</sub> was less than 10 cells per mouse. Other well-known specificities like OVA<sub>257-264</sub> or LCMV GP<sub>33-41</sub> only report  $183 \pm 33$  and  $358 \pm 40$  cells per mouse, respectively (Van Braeckel-Budimir et al., 2017; Jenkins and Moon, 2012). It is not unprecedented to find large frequencies in B6 mice. In fact, a previous study reported that another PbA epitope, GAP50<sub>40-48</sub>, had an extremely large naive CD8 T cell repertoire of approximately 3000 cells per mouse (Van Braeckel-Budimir et al., 2017). In this study, they determined that more than 98% of the GAP50<sub>40-48</sub>-specific CD8 T cells express V $\beta$ 8.1 TCR and suggested that this extreme TCR bias is driven by

a strong thymic selection mediated by the peptide itself rather than MHC reactivity (Van Braeckel-Budimir et al., 2017). Future research investigating the expression of specific V $\alpha$  and V $\beta$  TCR genes on Pb-1 specific CD8 T cells as well as structural analysis of the TCR:Pb-1 epitope interaction can provide evidence for the mechanisms underlying the high frequency of Pb-1 naive CD8 T cell precursors.

The large number of Pb-1 naive CD8 T cell supports the idea that the magnitude of the endogenous CD8 T cell response to the Pb-1 epitope would be greater than the one for the PbTRAP epitope. Nonetheless, it has been suggested that epitopes with low precursor frequencies could be good vaccination targets if effective boosting strategies are applied (Doll et al., 2016). Hypothetically, the high frequency of Pb-1 naive CD8 T cell precursors could provide for a superior initial expansion and induction of memory formation mediated by DC-targeted vaccination. In fact, large memory populations, especially liver T<sub>RM</sub> cells, were formed in B6 mice immunised with anti-Clec9A mAb targeting the Pb-1 epitope in absence of PbT-I cells. In this regard, earlier studies have demonstrated the importance of the liver T<sub>RM</sub> cells for the protection against malaria liver-stage (Fernandez-Ruiz et al., 2016; Lau et al., 2014). Remarkably, 50% of B6 mice immunised with anti-Clec9A-Pb-1 were protected against liver-stage parasites in the absence of the transgenic PbT-I cells. Several murine studies have shown that immunisations with Clec9A targeting antigens drives potent humoral responses using model antigens such as OVA (Caminschi et al., 2008; Lahoud et al., 2011); and very recently in the context of infectious diseases, using antigens derived from pathogens like influenza and enterovirus 71 (Park et al., 2017). Other studies have demonstrated the CTL antitumor capacity of antigen targeting Clec9A using OVA-expressing tumour cells (Sancho et al., 2008). This is the first time that in the context of an infectious disease, like malaria, high levels of protection were achieved by inducing the formation of endogenous-specific memory CD8 T cells through Clec9A targeted vaccination.

### 3.3.4 Sterile protection against malaria induced by prime-and-trap vaccination using the Pb-1 epitope:

There are approximately 140 million hepatocytes/g liver in mice and humans (Sohlenius-Sternbeck, 2006) and the number of infected hepatocytes under a normal course of infection has been estimated to be only 1 in one million and one in 100 million cells in mice and humans, respectively (Van Braeckel-Budimir and Harty, 2014). Furthermore, the liver stage occurs in a very limited period of time, 2 days with *P. berghei* and 6–8 days with *P. falciparum* in mice and humans, respectively (Sturm et al., 2006; Todryk and Hill, 2007). Accordingly, the number of memory CD8 T cells required for the surveillance of the whole liver in that very small time window in order to induce sterile protection is massive. In the context of other infectious diseases, such as *L. monocytogenes*, the transfer of specific memory CD8 T cells induced a significant reduction on the bacterial load in the liver, leading to a sublethal infection that can be cleared from all the mice (Badovinac et al., 2002). In the case of B6 mice infected with *P. berghei* all infected hepatocytes have to be cleared before any parasite progresses to blood stage and the mouse succumb to a lethal malaria.

Early studies published in 2008 and 2010 by Schmidt and collaborators found that sterilizing immunity induced after several i.v. injections with RAS in mice was correlated with the number of specific CD8 T cells in the blood and liver (Schmidt et al., 2010, 2008). They estimated that the threshold of specific cells is one million in the spleen and 20,000 in the liver of BALB/c mice to achieve sterile protection against *P. berghei* (Schmidt et al., 2008). This threshold is even higher when protecting B6 mice from *P. berghei* sporozoites (Schmidt et al., 2010). These studies addressed mainly the requirement of circulating memory CD8 T cells.

Data have revealed that liver T<sub>RM</sub> cells are very effective at inducing sterilizing immunity against *P. berghei* in B6 mice. These cells constantly survey

for liver infection by patrolling through the liver sinusoids (Fernandez-Ruiz et al., 2016). The main advantage of T<sub>RM</sub> cells over conventional circulating memory CD8 T cells is that they are already present and resident in the liver when parasites start infecting hepatocytes; and no extra time is invested in recruiting these cells. Liver T<sub>RM</sub> cells, as effector memory cells, are also equipped for the elimination of malaria infected hepatocytes (Fernandez-Ruiz et al., 2016). Given the high efficacy of the liver T<sub>RM</sub> cells at inducing sterile protection, the next goal was to maximize the formation of endogenous-specific liver T<sub>RM</sub> cells able to confer protection.

Previous reports have shown that the presence of local antigen and inflammatory stimuli provide by adjuvants enhanced the liver T<sub>RM</sub> cell formation through vaccination strategies like prime-and-trap (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Data here produced revealed that more than 90% of the mice vaccinated with prime-and-trap using the Pb-1 epitope were sterile protected against infection with *P. berghei* sporozoites.

During the optimisation experiments for the different doses of the prime-and-trap components, it was observed that slight changes in the dose of the recombinant adenovirus (rAAV-Pb-1) led to dramatic changes in the number of liver T<sub>RM</sub> cells. In fact, beyond the optimal dose of the virus, significant reduction of liver T<sub>RM</sub> cells was observed, accompanied by a T cell exhaustion phenotype, which translated in poor sterile protection. Elegant studies have demonstrated that CD8 T cell exhaustion, in the context of chronic viral infections, develops in a hierarchy manner (McLane et al., 2019; Wherry, 2011). The model suggest that initially during a chronic infection, CD8 T cells are highly functional and they progressively lose function and their ability to produce proinflammatory cytokines if they are constantly exposed to specific antigen. In this hierarchical model, it was also demonstrated that ultimately specific-CD8 T cell populations were physical deleted when exposed to high antigen loads, as oppose to other specific populations exposed lower antigen loads (Wherry et al., 2013). For the prime-and-trap system, high doses of rAAV-Pb-1 appeared to induce loss of function by

specific-CD8 T cells inducing to poor control of infection; and may have even led to the physical deletion of some specific-CD8 T cells.

In summary, this data revealed the highly immunogenic nature of the Pb-1 epitope; and the capacity of the prime-and-trap vaccine using Pb-1 to induce the formation of endogenous-specific liver T<sub>RM</sub> cells which were effective at conferring high levels of sterile protection. This study also demonstrate that vaccination strategies that achieve the formation of T<sub>RM</sub> cells have the potential to induce effective protection, even if they stimulate cells specific for a single antigen.

The development of sub-unit vaccines aiming for the formation of liver T<sub>RM</sub> cells represents a novel and potentially effective strategy to be implemented in clinical trials. One of the biggest limitations for the development of effective malaria vaccines mediated by CD8 T cell immunity is the lack of tools that allow the monitoring of liver T<sub>RM</sub> cell formation. Currently, clinical trials study circulating T cell responses through cytokine production assays performed under *ex vivo* settings and the local immune response in the tissue remain unknown. Therefore, there is an urgent need for the identification of biomarkers in blood that could effectively predict the formation of T<sub>RM</sub> cells. Moreover, there is high demand for the discovery of immunogenic epitopes derived from human malaria parasites expressed during liver-stage infection. The major challenge for the identification of these antigens is the lack of a suitable and sufficient source of infected hepatocytes. These concepts together with other translational approaches will be further discussed on Chapter 5.



## CHAPTER 4

Analysis of nucleic acid-based adjuvants for liver T<sub>RM</sub> cell formation induced by Clec9A targeted immunisation.

#### 4.1 Introduction:

Tissue-resident memory T cells ( $T_{RM}$  cells) are part of a memory T cell subpopulation that are often localised to the tissue where the initial infection occurred. In case of secondary infections, the strategic location of the  $T_{RM}$  cells allows them to mediate effector responses more rapidly than their counterparts, the circulating memory cells (Mueller and Mackay, 2016).  $T_{RM}$  cells exert effective killing of infected cells through many effector mechanisms, including the expression of granzyme B (Casey et al., 2012; Mackay et al., 2012b; Schenkel et al., 2014) and the production of proinflammatory cytokines such as IFN- $\gamma$  (Glennie et al., 2015; Schenkel et al., 2013), the latter of which also promotes the recruitment of circulating memory T cells. A detailed description of the  $T_{RM}$  cell effector functions is found in Chapter 1.

Several studies have revealed the critical role of  $T_{RM}$  cell-mediated protection in the context of different pathogen models, such as herpes simplex, vaccinia and influenza virus infections (Ariotti et al., 2012; Gebhardt et al., 2009; Jiang et al., 2012; Mackay et al., 2012a; Shin and Iwasaki, 2012; Teijaro et al., 2011). Other studies have analysed the role of  $T_{RM}$  cells in the context of viral hepatitis in humans. Pallett *et al.* described that  $T_{RM}$  cells are enriched in samples of liver tissues from patients with chronic hepatitis B virus (HBV) infection as compared with healthy controls. Importantly, they observed an inverse correlation between  $T_{RM}$  frequency and HBV viral load, and was associated with the *ex vivo* capacity of these cells to produce IFN $\gamma$  and TNF $\alpha$  in response to antigen stimulation (Pallett et al., 2017). Liver  $T_{RM}$  cells have also been reported to be effective at inducing sterile immunity against *P. berghei* infection in mice (Fernandez-Ruiz et al., 2016). These  $T_{RM}$  cells patrol the liver sinusoids, scanning for infected hepatocytes. Antibody depletion of liver  $T_{RM}$  cells from vaccinated mice abrogated protection against sporozoite challenge, indicating the critical role of these cells in vaccine-mediated protection (Fernandez-Ruiz et al., 2016).

Previous reports and data collected in Chapter 3 revealed that the number of liver T<sub>RM</sub> cells correlates with the level of sterile protection against malaria infection (Fernandez-Ruiz et al., 2016). Given the importance of liver T<sub>RM</sub> cells in the context of protection against malaria and other liver-associated pathogens (Fernandez-Ruiz et al., 2016; Pallett et al., 2017; Stelma et al., 2017), it may be valuable to study how to enhance their formation.

It has been recently reported that liver T<sub>RM</sub> cell formation is increased by TLR agonists, used as adjuvants, in the absence of antigenic stimuli *in vivo* (Holz et al., 2018). Moreover, studies applying the prime-and-trap vaccination strategy provide evidence that the nature of the adjuvant influences liver T<sub>RM</sub> cell formation; justifying a greater investigation of adjuvant options (Fernandez-Ruiz et al., 2016; Chapter 3 in this thesis). It is the aim of this chapter to assess various nucleic acid agonists as adjuvants for liver T<sub>RM</sub> cell formation induced by Clec9A targeted immunisation *in vivo*.

#### 4.1.1 The adjuvants

Historically, TLR9 agonists, such as synthetic CpG oligonucleotides (ODN), are used as vaccine adjuvants (Bode et al., 2011). Moreover, model antigen studies have reported the use of synthetic CpG ODN, as the preferred adjuvant, for the induction of CD8 T cell and cytotoxic responses mediated by immunisations with mAb targeting antigen to Clec9A (Caminschi et al., 2008; Lahoud et al., 2011).

There are four major types of synthetic CpG ODN, (A-, B-, C- and P-Class) each with distinct base pair composition and biological properties (Bode et al., 2011). Detailed properties for the different CpG ODN were reviewed in Chapter 1.

B-class ODN have been shown to be required for optimal cytokine production and DC activation by binding to DEC-205, a surface molecule highly expressed on cross-presenting DC (Lahoud et al., 2012). Also, it has been

reported that signalling via TLR9 with B-class ODN induces the formation of intrahepatic myeloid cell aggregates (iMATEs) that facilitate the accumulation of CD8 T cells in the liver (Huang et al. 2013). These two factors might explain why CpG is a suitable adjuvant for the generation of effective liver TRM cells mediated by prime-and-trap vaccination (Fernandez-Ruiz et al. 2016).

Polyinosinic–polycytidylic acid (p I:C), a TLR3 agonist, has also been used as an adjuvant to induce CTL responses mediated by targeting antigen to Clec9A *in vivo* (Caminschi et al., 2012b, 2008; Lahoud et al., 2011). Recently, it has been shown in vaccinated mice and cell transfer experiments that stimulation of TLR3 through p I:C increases the formation of liver T<sub>RM</sub> cells and yet is less potent than CpG (Fernandez-Ruiz et al., 2016; Holz et al., 2018; Chapter 3 in this thesis).

Both synthetic CpG ODN and p I:C mediate type I IFN responses through TLR engagement. Other adjuvants investigated in this chapter also mediate type I IFN responses by binding to specific nucleic acid receptors. Based on location, there are two groups of nucleic acid receptors described at the moment; the endosomal and the cytosolic sensors (Junt and Barchet, 2015). While those members of the TLR family that recognise nucleic acids, which are generally localised in the endosome, are primarily expressed by cell types of the innate immune system, cytosolic receptors are broadly expressed by most cell types (Roers et al., 2016). The fact that TLR and cytosolic receptors are differentially expressed, suggests that their ligands might be used as strategic adjuvants to specifically activate different cell types depending on the desired immune response.

To cover a range of different receptors, two endosomal and two cytosolic receptors were selected and their agonists studied as adjuvants for the induction of liver T<sub>RM</sub> cells. Agonists were selected for TLR7 and TLR9, receptors located in the endosome, and for RIG-I and cGAS, located in the cytoplasm.

TLR7 detects ssRNA, short dsRNA and short synthetic GU-rich ODN (Zhang et al., 2016). Retinoic acid-inducible gene I (RIG-I), recognizes short double stranded RNA bearing 5' triphosphorylated or 5' diphosphorylated ends (Goubau et al., 2014; Hornung et al., 2006). In a recent study, it was observed that RIG-I ligands promoted cross-presentation enhancing CTL responses in the context of an antiviral vaccine (Hochheiser et al., 2016). Cyclic GMP–AMP synthase (cGAS) recognizes dsDNA, which upon activation produces a second messenger that signals through the stimulator of IFN genes (STING) (Sun et al., 2013). The activation of endosomal and cytosolic sensors leads to the production of type-I IFN and NF- $\kappa$ B-dependent pro-inflammatory cytokines, the signature of innate immune activation during infection (Junt and Barchet, 2015). Further information about nucleic acid receptors, their agonists and mediated immune responses was given in Chapter 1.

## 4.2 Results:

### *4.2.1 TLR9 ligands are potent at inducing liver T<sub>RM</sub> cell formation*

In Chapter 3, the highly immunogenic nature of the novel Pb-1 epitope derived from PbA was revealed. It was demonstrated that vaccinations targeting Pb-1 to Clec9A induced the formation of effective liver T<sub>RM</sub> cells able to confer high levels of sterilizing protection in B6 mice.

In this chapter, immunisation targeting the novel Pb-1 epitope to Clec9A was used as a platform for the assessment of the different agonists as adjuvants for the formation of liver T<sub>RM</sub> cells.

For the first round of experiments, only TLR agonists were assessed. The TLR4 agonist lipopolysaccharide (LPS) was included as a non-nucleic acid ligand. To control for the initial number of responding cells and to facilitate the analysis of these experiments, mice were transferred with PbT-I GFP expressing cells one day prior to the priming. The next day, mice were injected with 2 µg of αClec9A-Pb-1 mAb – which is a dose that induces significant expansion of PbT-I cells (see Chapter 3) – plus one of the selected adjuvants in a single injection. A small blood sample was taken at day 7 as these data provide a rapid indication of the effectiveness of the priming as well as an approximation of the proportion of memory cells that could be formed at a later time point. At day 28, post immunisation spleens and livers were collected, and the formation of memory cells was investigated (Figure 4.1A).

Table 4.1. Characteristics of the adjuvants tested during the first round of experiments:

Name	Receptor	Class	Dose per mouse
CpG-combo	TLR9	CpG class B (ODN2006)	75 µg

		linked to a CpG class P (21798)	
CpG 1668	TLR9	CpG class B	75 µg
Poly I:C (p I:C)	TLR3, RIG-I / MDA5	Synthetic analog of dsRNA	50 µg
Lipopolysaccharide (LPS)	TLR4	Non-nucleic acid	1 µg

On day 7 post immunization, CpG-combo showed a significant superior capacity at inducing expansion of PbT-I cells compared to the naive control mice, though p I:C was capable of inducing similar PbT-I cell expansion (Fig. 4.1 B). On day 28, the four adjuvants, CpG-combo, CpG-1668, p I:C and LPS induced significantly more liver T<sub>RM</sub> cells compared to naive control mice, with CpG-combo the most potent adjuvant for the induction of liver T<sub>RM</sub> cells (Fig. 4.1 C and E). Mice primed with  $\alpha$ Clec9A-Pb-1 mAb plus CpG-combo formed on average  $1.1 \times 10^5$  liver T<sub>RM</sub> cells, while CpG-1668 induced 5 times less cells. P I:C and LPS were poor at inducing T<sub>RM</sub> cell formation compared to CpG-combo. Interestingly, CpG-1668 and p I:C induced approximately  $1.4 \times 10^5$  T<sub>EM</sub> cells in spleen while CpG-combo induced approximately  $7.7 \times 10^4$  cells (Fig. 4.1 D).

It is important to mention that the doses of the adjuvants differed somewhat. LPS was used at a relatively low mass as higher doses cause toxicity. The doses of CpG-combo, CpG-1668 and p I:C corresponded to values used previously in murine studies were they were combined with  $\alpha$ Clec9A mAb to provide adjuvant activity (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Results from these studies revealed that these doses are well tolerated and effective at inducing memory T cell formation (Fernandez-Ruiz et al., 2016; Holz et al., 2018).

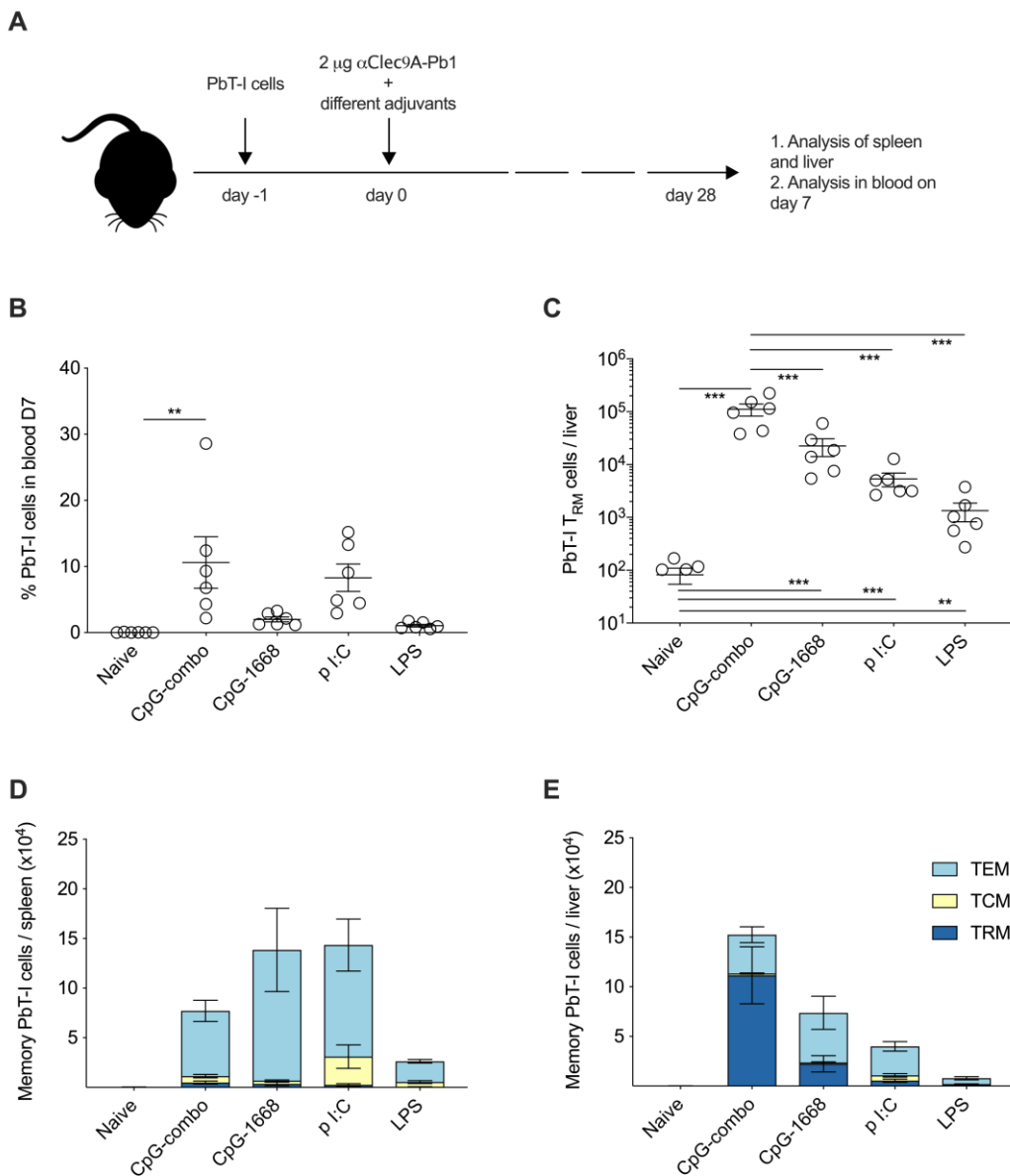


Figure 4.1 CpG-combo is a potent adjuvant for the induction of liver memory cells. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells. The next day, mice were immunized with 2 µg of αClec9-Pb-1 plus 75 µg of CpG-combo, 75 µg of CpG-1668, 50 µg of p I:C or 1 µg of LPS. One group of mice remain naive. Spleens and livers were collected on day 28. Single cell suspensions were generated, and the number of PbT-I cells and memory status were determined by flow cytometry. A. Schematic for the protection experiment. B. Percentage of PbT-I cells (CD8+, GFP + cells) in blood on day 7 post immunization. C Number of T<sub>RM</sub> cells per liver determined as CD8+, GFP+, CD44Hi, CD62L-, CD69+. D. and E. Total number of PbT-I cells in spleen and liver, respectively. Results are a pool of two independent experiments. Results are a pool of two independent experiments. Data were compared using a Kruskal-Wallis test with Dunn's multiple comparison post-test in (B.) or were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (C.). P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001..



For a second series of experiments, agonists for the cytosolic receptors cGAS and RIG-I, as well as ligands of the endosomal receptors TLR7 and TLR9, were included on the assessment and compared to CpG-combo. Moreover, these additional ligands were mixed and complexed in liposomal transfection reagents, which protects them from extracellular degradation and promotes the cellular uptake of nucleic acids (Smyth Templeton, 2002).

CpG-ODN 2006 (CpG-2006) and a GU-rich ssRNA TLR7 ligand (TLR7-L) were complexed in DOTAP, which is a cationic lipid that mediates the delivery of CpG-ODN and TLR7 ligands to the endosomal compartment (Heil et al., 2004; Honda et al., 2005; Yasuda et al., 2005; Yotsumoto et al., 2008). RIG-I ligand (RIG-I-L) and cGAS ligand (cGAS-L) were complexed in in vivo-jetPEI®, a polyethyleneimine that enhances *in vivo* delivery of both DNA and RNA (Yin et al., 2014). The nucleic acid sequences of each agonist were listed in Chapter 2, as well as the protocol used to complex them with the liposomal transfection reagents.

Table 4.2. Characteristics of the adjuvants that were complexed in transfection reagents:

Name	Receptor	Class	Transfection reagent	Dose per mouse
CpG-combo	TLR9	CpG class B (ODN2006) linked to a CpG class P (21798)	No	20 µg or 75 µg
CpG-2006	TLR9	CpG class B (ODN2006)	DOTAP	20 µg
RIG-I-L	RIG-I	5' 3p dsRNA cytosol ligand	jet-PEI®	20 µg
TLR7-L	TLR7	ssRNA endosome ligand	DOTAP	30 µg

cGAS-L	cGAS	dsDNA cytosol ligand	jet-PEI®	20 µg
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The doses for the different adjuvants were selected based on previous research performed at the Institute for Clinical Chemistry and Clinical Pharmacology at the University Hospital Bonn, Germany (unpublished).

B6 mice were transferred with PbT-I cells and primed the next day with  $\alpha$ Clec9A-Pb-1 plus one of the selected adjuvants. Two groups of mice received the CpG-combo, one with a lower dose than usual (20 µg instead of 75 µg per mouse). Analysis from these results allowed for the direct comparison of a potential dose impact on the expansion at the effector phase and the formation of memory cells. Another group of mice were only injected with the  $\alpha$ Clec9A-Pb-1 as a control to account only for the adjuvant effect. Similar to the previous experiment, a small blood sample was taken on day 7, and on day 28 memory cell formation was assessed in the spleen and liver.

Analysis on day 7 revealed that RIG-I-L induced the highest percentage of PbT-I cells in blood (26%) as compared to the other agonists including CpG-combo-high dose (10%) (Fig. 4.2 B). Mice injected with a low dose of CpG-combo (i.e. 20 µg) expanded less PbT-I cells than mice primed with the high dose. Additionally, 20 µg of CpG-2006 complexed in DOTAP induced a significantly higher percentage of PbT-I cells in the blood as compared to mice primed only with  $\alpha$ Clec9A-Pb-1. TLR7-L and cGAS-L did not induce significant expansion.

On day 28 post priming, stimulation with 75 µg of CpG-combo per mouse induced the largest number of T<sub>RM</sub> cells, reaching approximately 10<sup>5</sup> cells per liver (Fig. 4.3 C and E). Mice that were treated with 20 µg of CpG-combo generated significantly lower numbers of liver T<sub>RM</sub> cells (9.3 x 10<sup>3</sup> cells). CpG-2006 and RIG-I ligand induced significantly higher numbers of T<sub>RM</sub> cells compared to prime no-adjuvant treated mice, whereas TLR7-L and cGAS-L failed to induce liver T<sub>RM</sub> cell formation (Fig. 4.3 C and E), consistent with their poor

responses on day 7. In the spleen, CpG-2006 prompted the formation of large numbers of T<sub>EM</sub> cells, closely followed by RIG-I-L and CpG-combo-high dose with similar numbers (Fig. 4.3 D).

All together, these results suggest that CpG-combo is the best adjuvant, given that mice treated with  $\alpha$ Clec9A-Pb-1 mAb plus 75  $\mu$ g of CpG-combo generated the highest liver T<sub>RM</sub> cell numbers, with an average of  $10^5$  T<sub>RM</sub> cells per liver. CpG-1668 and CpG-2006, the last complexed in DOTAP, also induced high numbers of liver memory cells. For most of the group treatments, the memory data from day 28 mirrored the results of the early expansion on day 7. RIG-I-L induced a large number of spleen T<sub>EM</sub> cells, in similar proportion to CpG-combo and CpG-2006, and low numbers of liver T<sub>RM</sub> cells. An important finding is the dramatic dose effect observed in mice that received 20  $\mu$ g instead of 75  $\mu$ g of CpG-combo. Mice from the last group formed on average 10 times more liver T<sub>RM</sub> cells than the mice in the group of the lower dose of CpG-combo. In a later section on this chapter the comparison between liver T<sub>RM</sub> cells and circulating memory T cells will be evaluated.

#### *4.2.2 High dose of agonists enhanced liver T<sub>RM</sub> cells formation*

Given that the number of liver T<sub>RM</sub> cells was significantly influenced by the dose of CpG-combo, it was important to determine whether a similar effect could be observed for the other adjuvants. For this, the dose of adjuvants was increased to 75  $\mu$ g per mouse, which corresponds to the dose of CpG-combo used in a previous published study (Fernandez-Ruiz et al., 2016) and for the experiments carried out in Chapter 3.

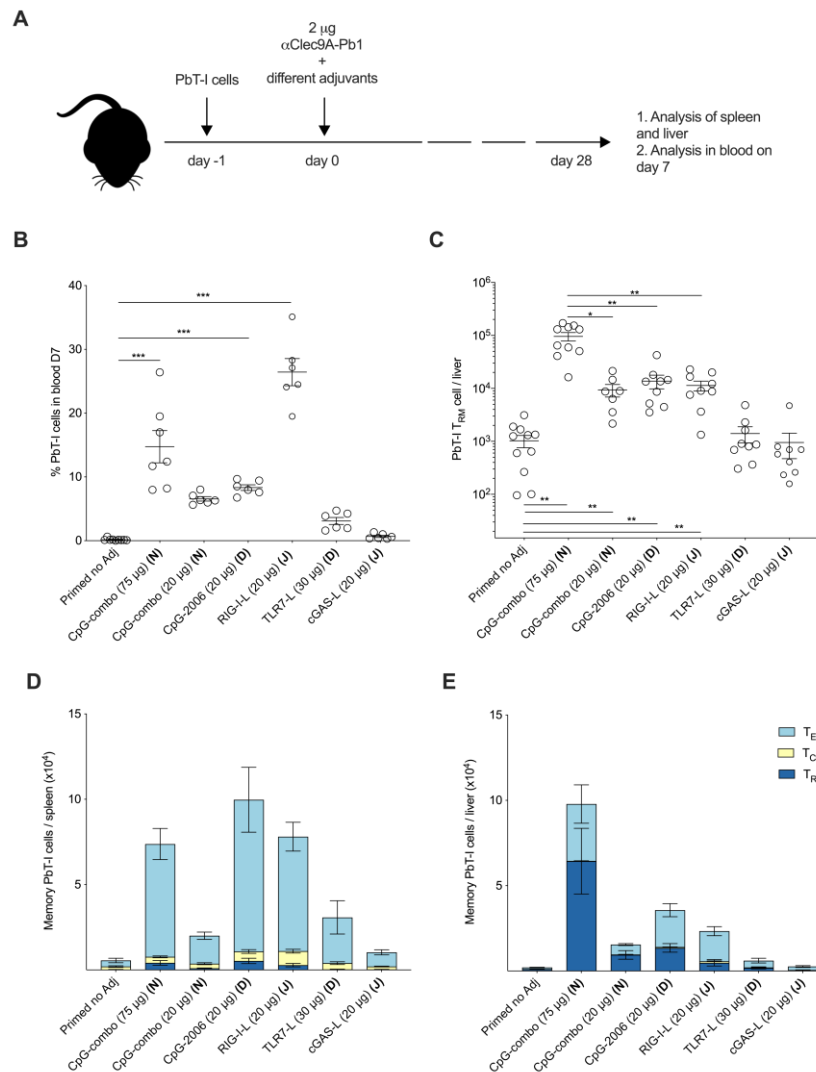


Figure 4.2 RIG-I ligand and CpG 2006 also induce great numbers of memory cells in liver. A. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells. The next day, mice were immunized with 2  $\mu$ g of  $\alpha$ Clec9A-Pb-1 plus: 75  $\mu$ g or 20  $\mu$ g of CpG-combo, 20  $\mu$ g of CpG-2006 complexed in DOTAP, 20  $\mu$ g of RIG-I-L complexed in *in vivo* jet-PEI®, 30  $\mu$ g of TLR7-L complexed in DOTAP or 20  $\mu$ g of cGAS-L complexed in *in vivo* jet-PEI®. On day 7, mouse blood was analyzed by flow cytometry. Spleens and livers were collected at day 28, single cell suspensions were generated, and the number of PbT-I cells and memory status were determined by flow cytometry. B. Percentage of PbT-I cells (CD8+, GFP + cells) in blood on day 7 post immunization. C Number of T<sub>RM</sub> cells per liver determined as CD8+, GFP+, CD44<sup>Hi</sup>, CD62L<sup>-</sup>, CD69<sup>+</sup>. D. and E. Total number of PbT-I cells in spleen and liver, respectively. Results are a pool of two independent experiments. Data were compared using a Kruskal- Wallis test with Dunn's multiple comparison post-test in (B.) or were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (C.). (N) refers to naked or uncomplexed adjuvants, (D) for the ligands complexed in DOTAP and (J) for the agonists complexed in *in vivo* jet-PEI®. P>0.05, \*, P<0.05; \*\*, P<0.01, \*\*\*, P<0.001.

Mice were transferred with PbT-I cells and primed the next day with  $\alpha$ Clec9A-Pb-1 mAb plus a high dose of one of the selected adjuvants. High doses of CpG-2006, TLR7-L, RIG-I-L and cGAS-L were assessed and compared to the CpG-combo. To control for toxicity, during the first two days, mice were monitored twice per day and their body condition was assessed. Body weight was recorded daily for 7 days. If body condition did not comply to healthy standards or 15% body weight was lost, mice were euthanized.

Records of the body weight are presented in figure 4.3. Each plot shows the weight fluctuation over 7 days after stimulation with each individual adjuvant. The dash line represents 15% of body weight loss of the lightest mouse at day 0. In general, a high dose of the adjuvants was well tolerated. The exception was for the RIG-I-L, which was not tolerated at 75  $\mu$ g per mouse and was changed for a tolerable dose of 40  $\mu$ g per mouse (Fig. 4.3). In addition, in the group of mice injected with 75  $\mu$ g of TLR7-L, one mouse registered a weight loss of 15% and was euthanized. As expected, no fluctuation was evident in mice that received no adjuvant. Minor body weight loss was observed in mice primed with CpG-combo, CpG-2006 and 40  $\mu$ g of RIG-I-L but this weight was recovered by day 7 (Fig. 4.3). Mice immunised with 75  $\mu$ g of TLR7-L or cGAS-L did not experience fluctuations in their weight (Fig. 4.3).

After testing the mouse tolerability to a high dose of the adjuvants, agonists were assessed for their capacity to induce PbT-I cell expansion and liver T<sub>RM</sub> cell formation. Blood samples from the surviving mice were taken on day 7 after immunisation and on day 28 spleen and liver were harvested and analysed for the formation of memory T cells.

CpG-combo, CpG-2006 and RIG-I-L promoted significant expansion of PbT-I cells on day 7, while no expansion was evident in mice stimulated with TLR7-L and cGAS-L, compared to mice that received no adjuvant (Fig. 4.4 B).

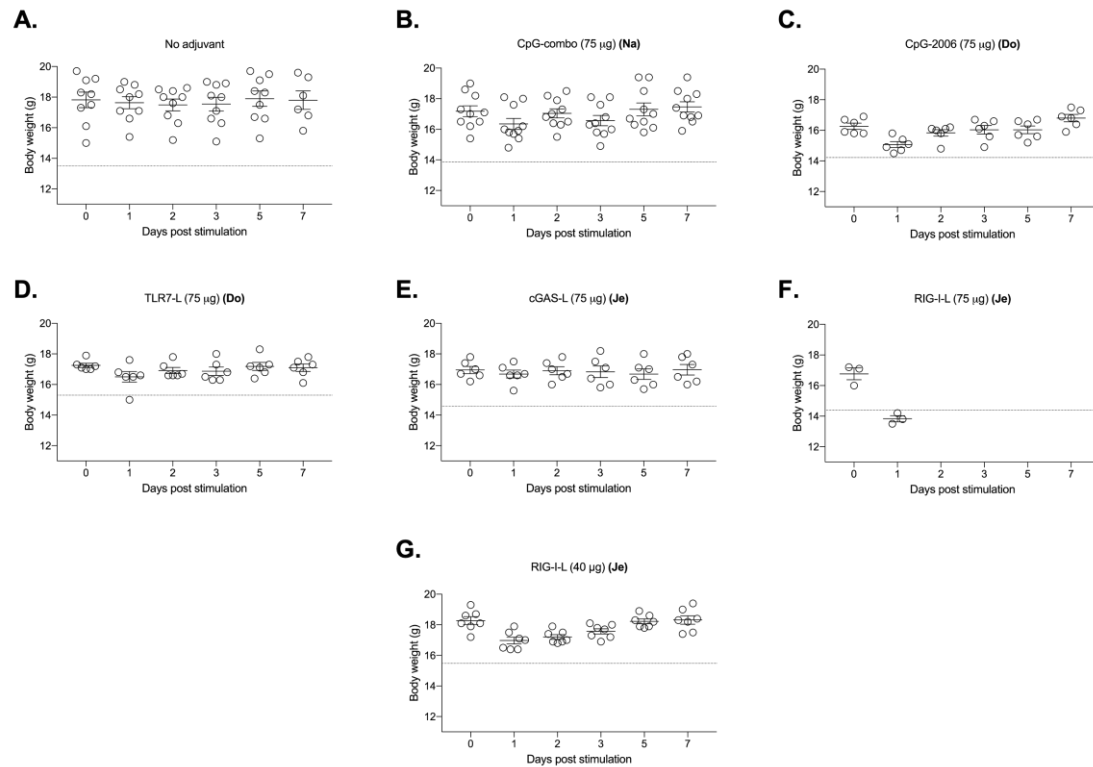


Figure 4.3 Mice body weight after priming with high dose of adjuvants. Female B6 mice, ranging from 6 to 8 weeks of age, were transferred intravenously with 50,000 PbT-I/uGFP cells. The next day, body weight was recorded and mice were immunized with 2  $\mu$ g of  $\alpha$ Clec9A-Pb-1 plus: 75  $\mu$ g of CpG-combo (B.), 75  $\mu$ g of CpG-2006 complexed in DOTAP (C.), 75  $\mu$ g of TLR7-L complexed in DOTAP (D.), 75  $\mu$ g of cGAS-L complexed in *in vivo* jet-PEI® (E.), 75  $\mu$ g (F.) or 40  $\mu$ g of RIG-I-L complexed in *in vivo* jet-PEI® (G.). (Na) refers to naked or uncomplexed adjuvants, (Do) for the ligands complexed in DOTAP and (Je) for the agonists complexed in *in vivo* jet-PEI®. During the first two days, mice were monitored twice per day and the body condition was assessed. Body weight was recorded daily for 7 days. If body condition did not comply to healthy standards or 15% body weight was lost, mice were euthanized. Open circles represent each individual mouse and the dash line represent 15% body weight loss relative to day 0 per group. Data were pooled from 2 to 3 independent experiments.

Analysis of memory cells on day 28 revealed that CpG-combo, CpG-2006, TLR7-L and RIG-I-L induced significantly high numbers of liver  $T_{RM}$  cells compared to mice that did not receive an adjuvant. cGAS-L was poor at inducing liver  $T_{RM}$  cell formation (Fig. 4.4 C). CpG-combo, CpG-2006 and RIG-I-L induced significantly higher numbers of liver  $T_{RM}$  cells compared to TLR7-L and cGAS-L. Moreover, CpG-combo and CpG-2006 also induced higher number of cells compared to RIG-I-L (Fig. 4.4 C). CpG-2006 (75  $\mu$ g) in DOTAP induced twice the number of liver  $T_{RM}$  cells compared to uncomplexed CpG-combo, although non-significant difference was observed. Mice that received CpG-2006 (75  $\mu$ g) formed in average 234,344 liver  $T_{RM}$  cells, while CpG-combo (75  $\mu$ g) induced 125,508 cells (Fig. 4.4 E). CpG-2006 also induced a slightly higher number of  $T_{EM}$  cells in the spleen as compared to CpG-combo but this was not a significant difference (Fig. 4.4 D).

When comparing with previous results described on Fig. 4.2, mice that received 75  $\mu$ g of CpG-combo formed 13 times more liver  $T_{RM}$  cells than mice receiving 20  $\mu$ g. Similarly, though not directly compared in the same experiment, 75  $\mu$ g of CpG-2006 induced a 17-fold increase in liver  $T_{RM}$  cells compared to 20  $\mu$ g of CpG-2006; for RIG-I-L and TLR7-L the increment was 2- and 1.5-fold, respectively (Fig. 4.4 C and Fig. 4.2 C).

These data indicate that the liver  $T_{RM}$  cell formation is influenced by the dose of the adjuvant for most of the cases, except for cGAS-L agonist which seems to fail to provide an adjuvant effect in this system.

#### *4.2.3 CpG-ligands preferentially induce liver $T_{RM}$ cell formation*

As discussed in Chapter 1, the agonists investigated here stimulate a broad range of different cell types by activating diverse pathways expressed differentially on the target cells (Junt and Barchet, 2015; Roers et al., 2016). Additionally, results depicted in this chapter demonstrated that the dose of the adjuvants influenced the formation of liver  $T_{RM}$  cells (Fig. 4.2).

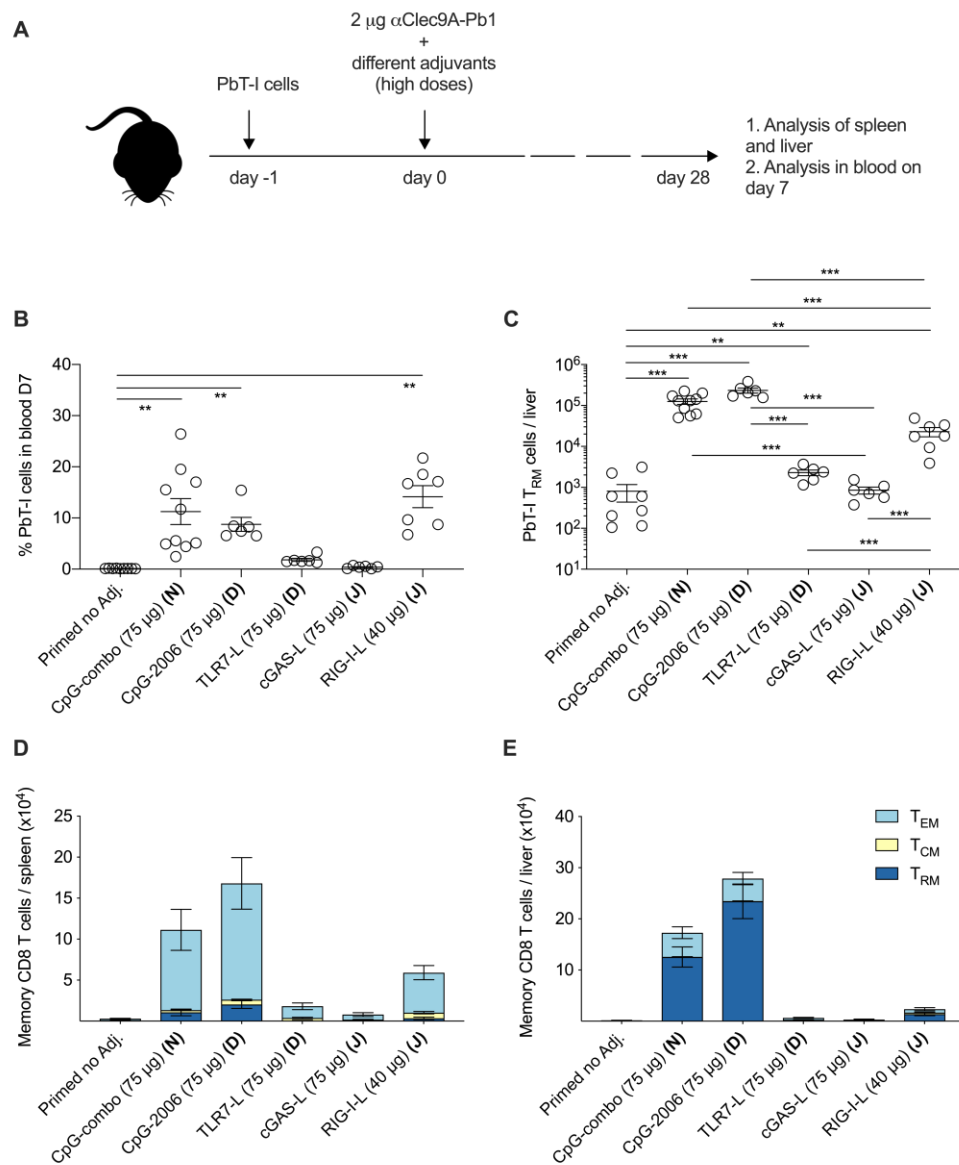


Figure 4.4 High dose of CpG-2006 is very potent at inducing liver  $T_{RM}$  cell formation. A. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells and primed the next day with 2  $\mu\text{g}$  of  $\alpha\text{Clec9A-Pb-1}$  plus: 75  $\mu\text{g}$  of CpG-combo, 75  $\mu\text{g}$  of CpG-2006 complexed in DOTAP, 75  $\mu\text{g}$  of TLR7-L complexed in DOTAP, 75  $\mu\text{g}$  of cGAS-L complexed in *in vivo* jet-PEI®, or 40  $\mu\text{g}$  of RIG-I-L complexed in *in vivo* jet-PEI®. B. Percentage of PbT-I cells (CD8+, GFP + cells) in blood on day 7 post immunization. C Number of  $T_{RM}$  cells per liver determined as CD8+, GFP+, CD44Hi, CD62L-, CD69+. D. and E. Total number of PbT-I cells in spleen and liver, respectively. Results are a pool of two independent experiments. Data were compared using a Kruskal- Wallis test with Dunn's multiple comparison post-test in (B.) or were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (C.). (N) refers to naked or uncomplexed adjuvants, (D) for the ligands complexed in DOTAP and (J) for the agonists complexed in *in vivo* jet-PEI®.  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ ..



Since the aim of this study was to identify an adjuvant that favoured the induction of liver T<sub>RM</sub> cells, the data collected from previous experiments were assessed for the ability to induce T<sub>RM</sub> cells relative to the induction of circulating memory cells. To do this in a simple way, the ratio of liver T<sub>RM</sub> cells over circulating memory cells – T<sub>EM</sub> plus T<sub>CM</sub> cells from the spleen – was calculated. Analysis of this ratio allows a comparison of the capacity of different vaccination strategies (e.g. different adjuvants) to promote T<sub>RM</sub> cell differentiation from a pool of circulating memory T cells (Holz et al., 2018).

Examination of the ratio of T<sub>RM</sub> cells to circulating memory T cells revealed that a high dose of CpG-2006 and CpG-combo (e.g. 75 µg instead of 20 µg per mouse) had the best adjuvant effect for the induction of liver T<sub>RM</sub> cell differentiation, as compared to the primed no-adjuvanted mice (Fig. 4.5). There was a tendency for lower doses to generate less liver T<sub>RM</sub> cells as compared to higher doses of same adjuvant, although this difference was not significant (Fig. 4.5). Remarkably, all other adjuvants, including CpG-1668 (uncomplexed) and high doses of RIG-I-L, TLR7-L and cGAS-L had similarly poor capacity to preferentially induce liver T<sub>RM</sub> cells in this particular system (Fig. 4.5).

All together, these data support previous evidence demonstrating that CpG ODN enhance the formation of resident memory T cells in the liver (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Potential mechanisms behind this effect will be explored in the discussion of this chapter.

#### *4.2.4 CpG-combo has no significant effect on the liver T<sub>RM</sub> cell formation induced by RAS*

Published studies have demonstrated that vaccination with radiation-attenuated sporozoites (RAS) induces liver T<sub>RM</sub> cells and provides a certain level of protection against sporozoite challenge

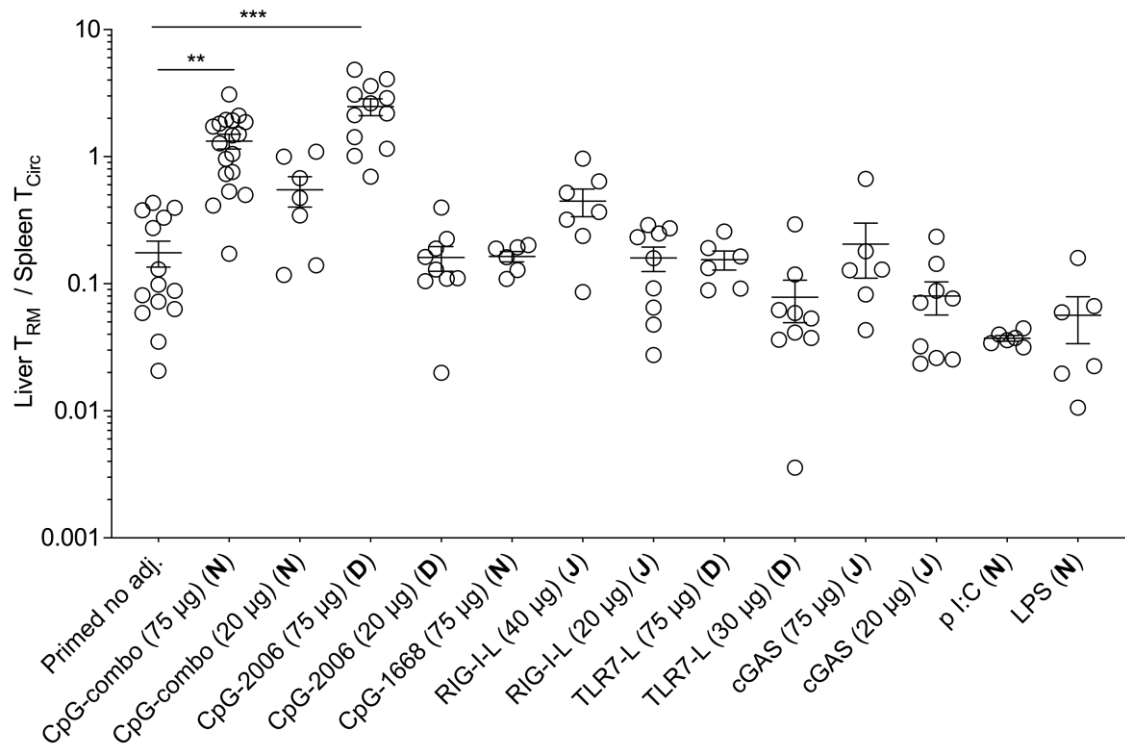


Figure 4.5 CpG-ligands are the best at inducing T<sub>RM</sub> cell formation. The number of liver T<sub>RM</sub> cells was divided by the number of circulating T cells (T<sub>Circ</sub>) in the spleen (T<sub>EM</sub> cells plus T<sub>CM</sub> cells) to calculate the relative proportion of T<sub>RM</sub> cells in the liver. This figure compiles results from figures 4.1, 4.2 and 4.4, which are results pooled from 2 to 4 independent experiments. (N) refers to naked or uncomplexed adjuvants, (D) for the ligands complexed in DOTAP and (J) for the agonists complexed in *in vivo* jet-PEI®. Groups were compared using a Kruskal-Wallis test with Dunn's multiple comparison post-test.

(Epstein et al., 2011; Fernandez-Ruiz et al., 2016). As CpG-based adjuvants were the best adjuvants for inducing liver T<sub>RM</sub> cells in the context of Clec9A targeted immunisation, the role of CpG as adjuvant for RAS vaccination was investigated. For this experiment, mice were adoptively transferred with PbT-I cells, as these cells are known to expand in response to RAS vaccination and can be easily monitored (Fernandez-Ruiz et al., 2016; Lau et al., 2014). The next day, mice were vaccinated with RAS or RAS plus CpG-combo. On day 28, when memory cells had been formed, spleens and livers were harvested from 4 mice per group and the memory T cell profile was assessed by flow cytometry. Remaining mice (n=6) were infected with live sporozoites and monitored for the occurrence of parasites in blood.

Data from day 28 determined that CpG-combo had no significant effect on the formation of liver T<sub>RM</sub> cells (Fig. 4.6 B, D). Mice vaccinated with RAS generated on average  $1.8 \pm 0.1 \times 10^5$  T<sub>RM</sub> cells per liver, and mice that received RAS plus CpG-combo formed a similar number of  $1.3 \pm 0.1 \times 10^5$  cells (Fig. 4.6 B, D). Likewise, other memory populations in the spleen or liver were unaffected by the addition of CpG-comb, although mice that received only RAS formed slightly higher numbers of T<sub>EM</sub> cells in the spleen and liver compared to RAS plus CpG-combo vaccinated mice (Fig. 4.6 C, D). It was previously reported in Chapter 3 and in a recent publication, that relatively high numbers of PbT-I T<sub>RM</sub> cells in the liver effectively protect against sporozoite challenge (Fernandez-Ruiz et al., 2016). As a consequence, mice vaccinated here with both, RAS or RAS plus CpG-combo had significantly less parasitemia as compared to naive mice (Fig. 4.6 E). Additionally, 78% of the mice were sterilely protected for both treatment groups (Fig. 4.6 F).

These data indicate that CpG-combo does not enhance the formation of liver T<sub>RM</sub> cell induced by one dose of RAS vaccination in B6 mice and has no effect on the protection induced by this immunisation.

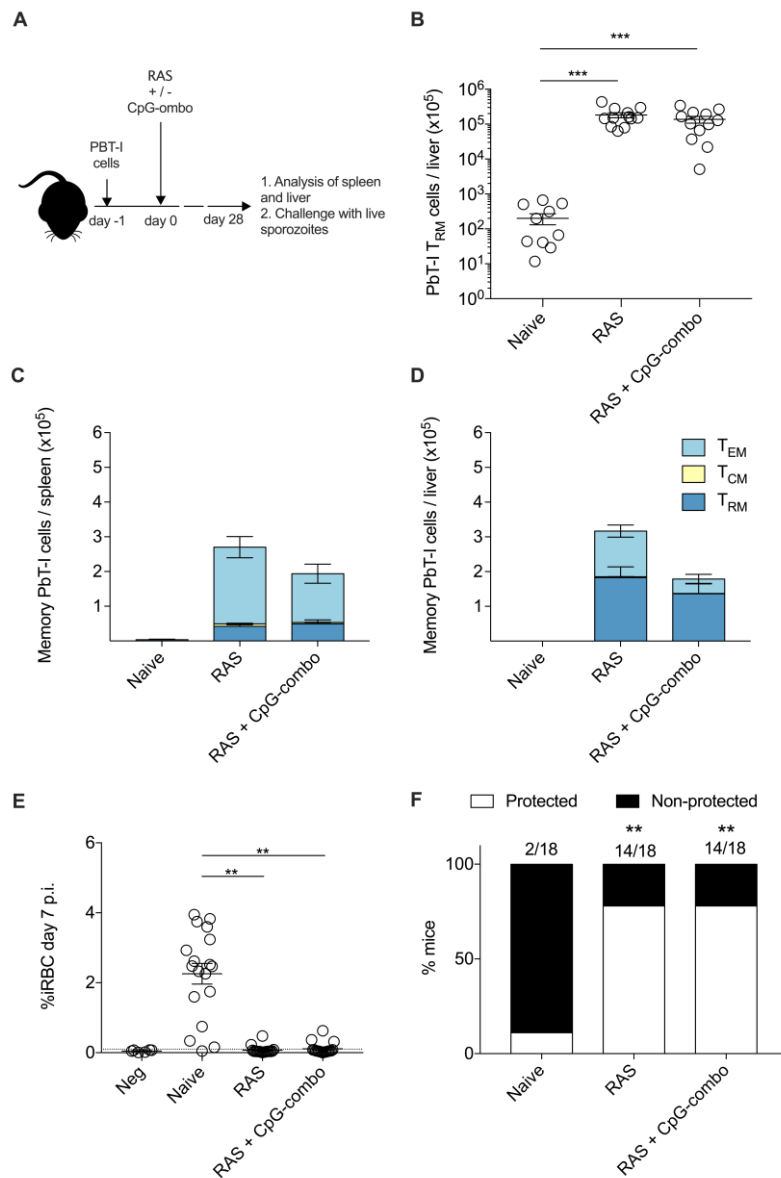


Figure 4.6 CpG-combo has no significant effect on the liver  $T_{RM}$  cell formation induced by RAS vaccination. A. B6 mice were transferred intravenously with 50,000 PbT-I/ $\mu$ GFP cells and vaccinated the next day with 50,000 RAS with or without 75  $\mu$ g of CpG-combo. Spleens and livers from 4 mice were collected on day 28, single cell suspensions were prepared, and memory T cell formation was assessed by flow cytometry. The remaining mice ( $n=6$ ) were challenged intravenously with 200 live sporozoites. Parasitemia was monitored in blood until day 10 post infection. B. Number of  $T_{RM}$  cells per liver on day 28. D. and E. Total number of PbT-I cells in spleen and liver, respectively. E. Percentage of infected red blood cells (iRBC) in blood on day 7 post F. Percentage of sterile protection. Results are a pool of two or three independent experiments. Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (E.), compared using a Kruskal-Wallis test with Dunn's multiple comparison post-test in (E.) or using Fisher's exact test (F.).  $P>0.05$ , \*,  $P<0.05$ ; \*\*,  $P<0.01$  \*\*\*,  $P<0.001$ .

#### *4.2.5 DOTAP enhances the effect of the CpG-ligands on the formation of liver T<sub>RM</sub> cells*

The above studies revealed that TLR9 ligands were the best adjuvants at inducing liver T<sub>RM</sub> cell formation and yet each one induced different numbers of cells. In fact, CpG-2006 induced twice as many T<sub>RM</sub> cells as CpG-combo (though this was not a significant difference; Fig. 4.4 C). As previously mentioned, some of the adjuvants were mixed and complexed in transfection reagents to enhance the differential delivery of agonists within the cell to the endosome or cytosol. Importantly, CpG-2006 was complexed in the transfection reagent DOTAP, whereas CpG-combo was used naked, i.e. without a transfection reagent. To understand the role of a transfection reagent on the induction of T<sub>RM</sub> cells, the effect of DOTAP on the three CpG tested here (CpG-2006, CpG-combo and CpG-1668) was investigated.

For this, mice were transferred with PbT-I cells and primed the next day with  $\alpha$ Clec9A-Pb-1 mAb plus CpG-2006, CpG-combo or CpG-1668 naked or complexed in DOTAP (Figure 4A). On day 7, a small blood sample was taken to investigate early cell expansion, and on day 28 the analysis on the memory cell formation in the spleen and liver was assessed.

Results from day 7 bleeds revealed that mice injected with CpG-2006 or CpG-1668 complexed in DOTAP expanded significantly more PbT-I cells as compared to mice injected with the same CpG in absence of transfection reagent (Fig. 4.7B). CpG-combo complexed in DOTAP induced slightly more PbT-I cells in blood as compared to naked CpG-combo, but a significant difference was not evident. Data from day 28 mirrored the results from day 7. CpG-2006 complexed in DOTAP induced a 14-fold increase liver T<sub>RM</sub> cells as compared to CpG-2006 in absence of DOTAP. In a similar fashion, CpG-1668 in DOTAP prompted the formation of 7-fold more T<sub>RM</sub> cells as compared to naked CpG-1668 (Fig. 4.7C, E). CpG-combo in DOTAP induced 3 times more T<sub>RM</sub> cells compared to CpG-combo in the absence of a transfection reagent, although this difference was not

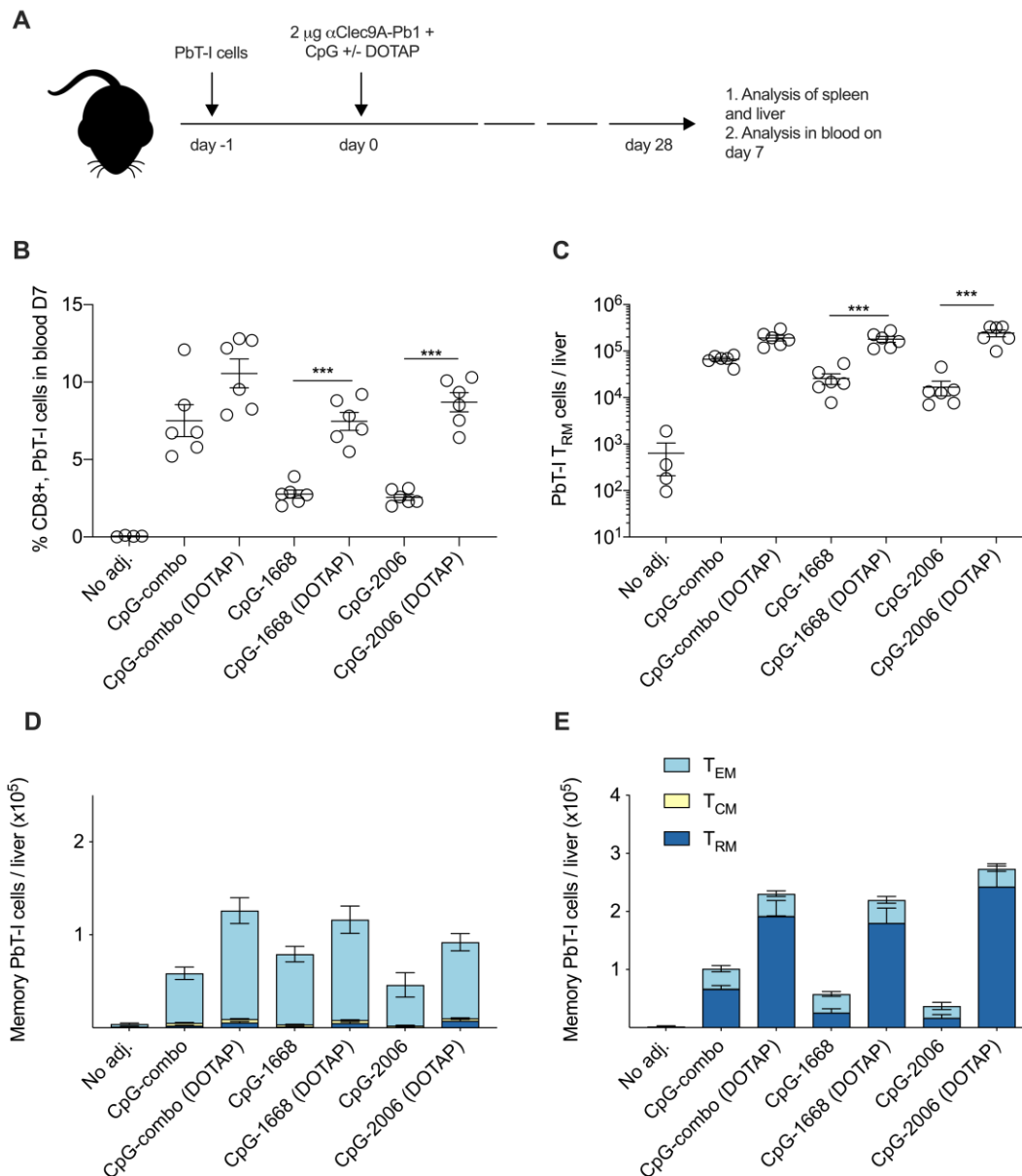


Figure 4.7 DOTAP enhances the effect of the CpG-ligands on the formation of liver T<sub>RM</sub> cells. A. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells and primed next day with 2 µg of  $\alpha$ Clec9A-Pb-1 plus: 75 µg of CpG-combo, 75 µg of CpG-2006, 75 µg of CpG-1668 complexed or not in DOTAP. On day 7 mice blood was analyzed by flow cytometry. Spleens and livers were collected at day 28, single cell suspensions were generated, and the number of PbT-I cells and memory status were determined by flow cytometry. B. Percentage of PbT-I cells (CD8<sup>+</sup>, GFP + cells) in blood on day 7 post immunization. C Number of T<sub>RM</sub> cells per liver determined as CD8<sup>+</sup>, GFP<sup>+</sup>, CD44<sup>Hi</sup>, CD62L<sup>-</sup>, CD69<sup>+</sup>. D. and E. Total number of PbT-I cells in spleen and liver, respectively. Results are a pool of two independent experiments. Data were compared using a Kruskal- Wallis test with Dunn's multiple comparison post-test in (B.) or were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (C.) P>0.05, \*, P<0.05; \*\*, P<0.01 \*\*\*, P<0.001.

significant (Fig. 4.7C, E). A similar pattern was observed for T<sub>EM</sub> cells in the spleen, as DOTAP enhances the number of these cells compared to CpG in absence of the transfection reagent (Fig. 4.7D).

After demonstrating that DOTAP enhances the ability of CpG to induce liver T<sub>RM</sub> cells during Clec9A vaccination, the role of this transfection reagent was investigated in the context of RAS vaccination (Fig. 4.8). For this experiment, mice were transferred with PbT-I cells, then on the next day were vaccinated with RAS plus CpG-combo complexed or not in DOTAP (Fig. 4A). Blood was collected on day 7 to measure the percentage of the PbT-I cells and monitor their expansion (Fig. 4B). On day 28, spleens and livers were harvested, and the memory T cell profile was assessed (Fig. 4C). Remaining mice were infected with live sporozoites and monitored for sterile immunity (Fig. 4D, E). Of note, only one experiment was performed due to time constraints. Analysis of bloods on day 7 showed PbT-I cell expansion in mice that were vaccinated with RAS or RAS + naked CpG-combo, while PbT-I cells expanded poorly in mice receiving RAS + CpG-combo complexed in DOTAP (Fig. 4.8 B). Similarly, on day 28, mice generated high numbers of liver T<sub>RM</sub> cells when vaccinated with RAS or RAS + naked CpG-combo, but not in those immunised with RAS + CpG complexed in DOTAP formed (Fig. 4.8 C). As a consequence of the poor induction of liver T<sub>RM</sub> cell formation, mice vaccinated with RAS + CpG-combo in DOTAP developed levels of parasitemia comparable to naïve untreated mice after 7 days of challenge (Fig. 4.8 D) and none were sterilely protected (Fig. 4.8 E). These preliminary data suggest that CpG-combo complexed in DOTAP has an adverse effect on the formation of liver T<sub>RM</sub> cells mediated by RAS.

While complexing CpG-combo in DOTAP had an adverse effect on T<sub>RM</sub> cell formation induced by RAS, it was still possible that when used for prime-and-trap vaccination, memory cell formation might be boosted. To test this possibility, mice were vaccinated with prime-and-trap using the Pb-1 epitope. One group received CpG-combo complexed in DOTAP, while another group received

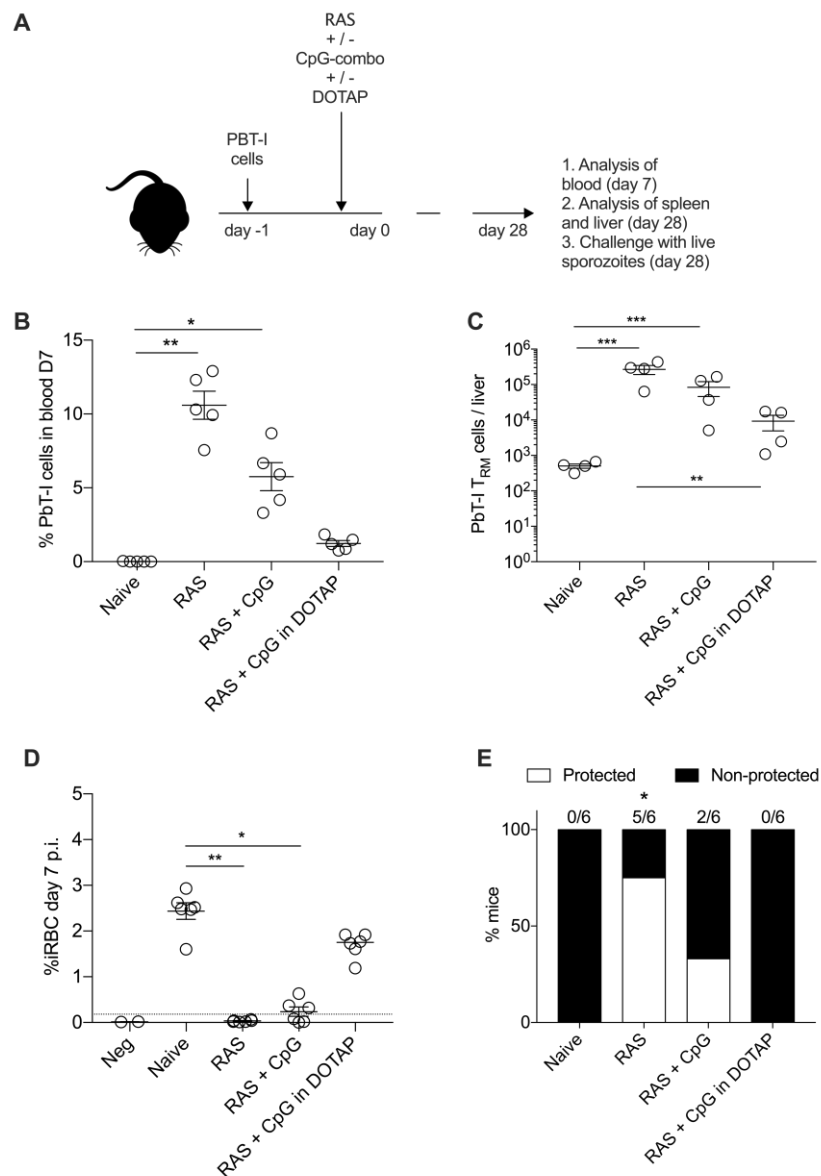


Figure 4.8 DOTAP has an adverse effect on the formation of  $T_{RM}$  cells induced by RAS (preliminary results). A. B6 mice were transferred intravenously with 50,000 PbT-I/uGFP cells and vaccinated the next day with 50,000 RAS with or without 75  $\mu$ g of CpG-combo complexed or not in DOTAP. On day 7, mice blood was analyzed by flow cytometry. Spleens and livers from 4 mice were collected on day 28 and memory T cell formation was assessed by flow cytometry. The remaining mice (n=6) were challenged intravenously with 200 live sporozoites. Parasitemia was monitored in blood until day 10 post infection. B. Percentage of PbT-I cells in blood on day 7 post immunization. C Number of  $T_{RM}$  cells per liver. D. Percentage of infected red blood cells (iRBC) in blood on day 7 post F. Percentage of sterile protection. Results are from a single experiment. Data were compared using a Kruskal- Wallis test with Dunn's multiple comparison post-test in (B. and D.) or were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (C.) or compared by using Fisher's exact test (E.).  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ .



CpG-combo in the absence of DOTAP. On day 35, 4 mice per group were used for the enumeration of memory T cells populations in the liver and spleen. The remaining mice (n=6) were challenged intravenously with 200 sporozoites and sterile protection was determined. Surviving mice were challenged a second time on day 104 after immunisation, since this data provides insights for the role of DOTAP in a long-term set up.

Results from day 35 indicated that the number of liver T<sub>RM</sub> cells induced in mice vaccinated with prime-and-trap was not significantly altered by the presence of the transfection reagent DOTAP, as similar high numbers of T<sub>RM</sub> cells were formed in mice from both groups (Fig. 4.9B). As previously reported in Chapter 3, mice challenged on day 35 that have been vaccinated with prime-and-trap targeting Pb-1 developed significantly less parasitemia as compared to naive mice, which was reflected on the induction of high levels of sterile protection (Fig. 4.9 C, D). These observations were also true for mice vaccinated with prime-and-trap using CpG-combo complexed in DOTAP; in fact, the level of sterile protection was the same as compared to the original prime-and-trap vaccine.

It is important to mention that data from day 104 were based on a single experiment due to time constraints. On day 104, some of the surviving mice from both the prime-and-trap vaccine with or without DOTAP were infected with 200 live sporozoites. Results on day 7 p.i. indicated a tendency in both groups of vaccinated mice to control parasitemia as compared to naive mice, although this difference was not significant (Fig. 4.9 E). A similar tendency was also observed in the level of protection (Fig. 4.9 F).

Together, these data indicate that even though DOTAP was able to enhance liver T<sub>RM</sub> cell formation mediated by CpG when vaccinating with Clec9A-targeted antigen, it did not have a significant effect in the context of prime-and-trap vaccination.

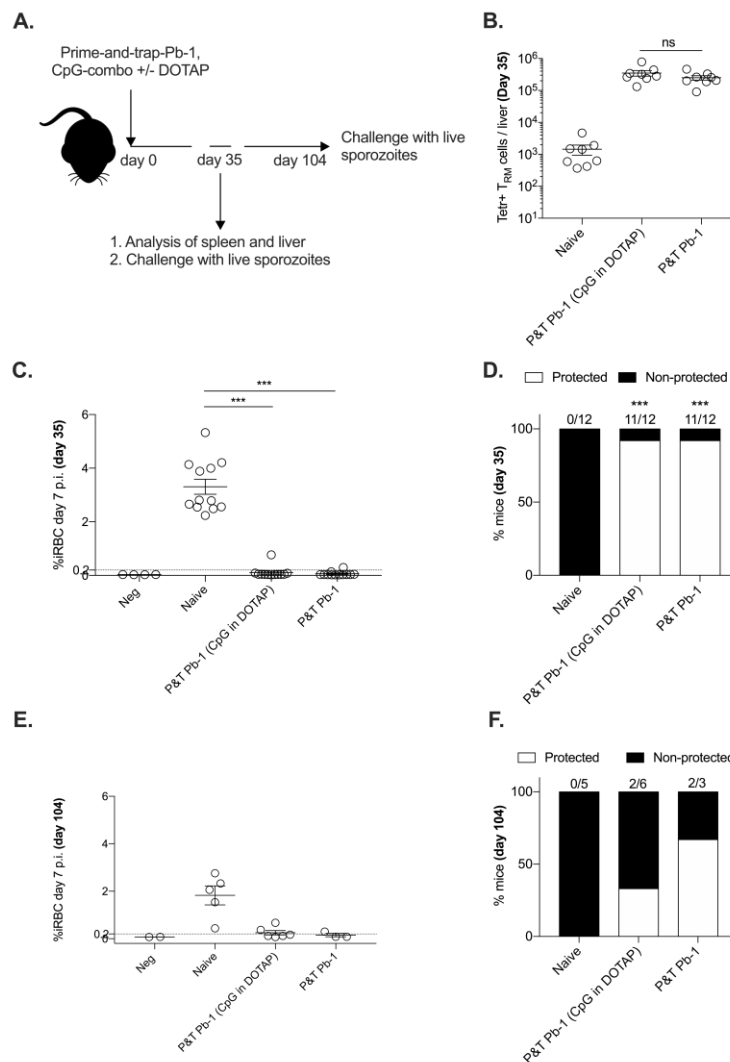


Figure 4.9 DOTAP has no significant effect on the sterile protection induced by prime-and-trap targeting the Pb-1 epitope. A. B6 mice were vaccinated with 8  $\mu\text{g}$  of  $\alpha\text{Clec9A-Pb-1}$ ,  $10^9$  rAAV-Pb-1 and 75  $\mu\text{g}$  of CpG-combo complexed or not in DOTAP. Spleens and livers from 4 mice were collected on day 35 and single cell suspensions were prepared, and memory T cell formation was assessed by flow cytometry. The remaining mice ( $n=6$ ) were challenged intravenously with 200 live sporozoites. Parasitemia was monitored in the blood up to day 10 post infection. Sterilely protected mice were challenged a second time with 200 living PbA sporozoites on day 104 B. Number of  $T_{RM}$  cells per liver on day 35. C. Percentage of infected red blood cells (iRBC) in blood on day 7 post challenge in mice infected on day 35. D. Percentage of sterile protection in mice challenged on day 35. E. Percentage of iRBC on day 7 post challenge in mice infected on day 104. F. Percentage of sterile protection in mice challenged on day 104. Results are a pool of two independent experiments, for data of day 35 and a single experiment for data of day 104. Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test in (B.) or compared using a Kruskal-Wallis test with Dunn's multiple comparison post-test in (C.) and (E.), or compared by using Fisher's exact test (D.) and (F.).  $P>0.05$ , \*,  $P<0.05$ ; \*\*,  $P<0.01$  \*\*\* ,  $P<0.001$ .

#### 4.2.6 Assessment of adjuvants for type I IFN production: Control experiments

In the above studies, CpG ODN performed best for induction of T<sub>RM</sub> cells while most other adjuvants favoured induction of circulating T cells. Two adjuvants, TLR7-L and cGAS-L had particular poor capacity to induce the formation of memory cells in these Clec9A priming studies, prompting the question the whether or not they were functional.

In preliminary studies prior to the assessment with the Clec9A Ab, adjuvants were tested for the stimulation of type I IFN response. These experiments were carried out in Germany before undertaking the Clec9A priming studies in Australia. For this, the synthetically produced CpG-2006, TLR7-L, cGAS-L and RIG-I-L were tested *in vivo* for the production of IFN- $\alpha$ , a well-known type I IFN cytokine. Additionally, the expression of several interferon stimulated genes (ISG), a hallmark for type I IFN response (Schneider et al., 2014), was investigated in the spleen and liver. Since these experiments were performed before assessing the adjuvants for the formation of liver T<sub>RM</sub> cells, only low doses of the adjuvants were assessed.

Mice were grouped into independent experiments for endosomal delivery and cytosolic delivery. The first group was stimulated with CpG-2006 and TLR7-L complexed in DOTAP, while the second group received cGAS-L and RIG-I-L complexed in Jet-PEI. A few control mice received a non-stimulating nucleic acid (NA) complexed in DOTAP or Jet-PEI for the endosomal and cytosolic delivery, respectively. Four h after stimulation, mice were cheek bled and serum was collected. At 6 h, mice were sacrificed and blood, the spleen and the liver were collected. The serum was stored and RNA was isolated from the homogenized organs. Production of IFN $\alpha$  was measured in serum by ELISA and the expression of several ISG was investigated by qPCR.

Results from these preliminary experiments revealed extensive variation between experiments. Importantly, substantial background for the production of IFN- $\alpha$  was observed in mice that received non-stimulating NA, potentially masking the stimulation induced by the adjuvants. There was a tendency towards higher production of IFN- $\alpha$  induced by CpG-2006 and TLR7-L as compared to control mice (Fig. 4.10 B). A similar tendency was found in mice stimulated with cGAS-L and RIG-I-L (Fig. 4.10 C).

The relative gene expression of representative ISG *Irf 7* and *Irf 44* is shown in Fig. 4.10 D, E. Cytosolic stimulation of cGAS and RIG-I leads to the activation of the transcription factor IRF7, which promotes the production of IFN and other ISG (Schneider et al., 2014; Schoggins and Rice, 2012). Preliminary results from two mice revealed that while cGAS and RIG-I induced higher expression of *Irf 7* and *Irf 44* in the spleen and liver compared to CpG-2006 and TLR7-L, these last two also induced a 10-fold increase of *Irf 7* expression in liver (Fig. 4.10 D, E). Data from these pilot experiments suggested these adjuvants were functional in their capacity to induce type IFN.

CpG-2006, TLR7-L, cGAS-L and RIG-I-L were shipped from Germany to Australia prior the Clec9A priming studies. Given the limited effect of some of those adjuvants, it was necessary to confirm their function once again in Australia. For this purpose, mice were stimulated with the different adjuvants and the production of IFN $\alpha$  was assessed in sera. Mice were grouped in two different functional analysis: i) they received high or low doses of TLR7-L, cGAS-L or RIG-I-L and ii) they were stimulated with different CpG in the presence or absence of DOTAP. Control mice received only DOTAP or remained naive. Four and 10 h later, mice were bled and serum collected.

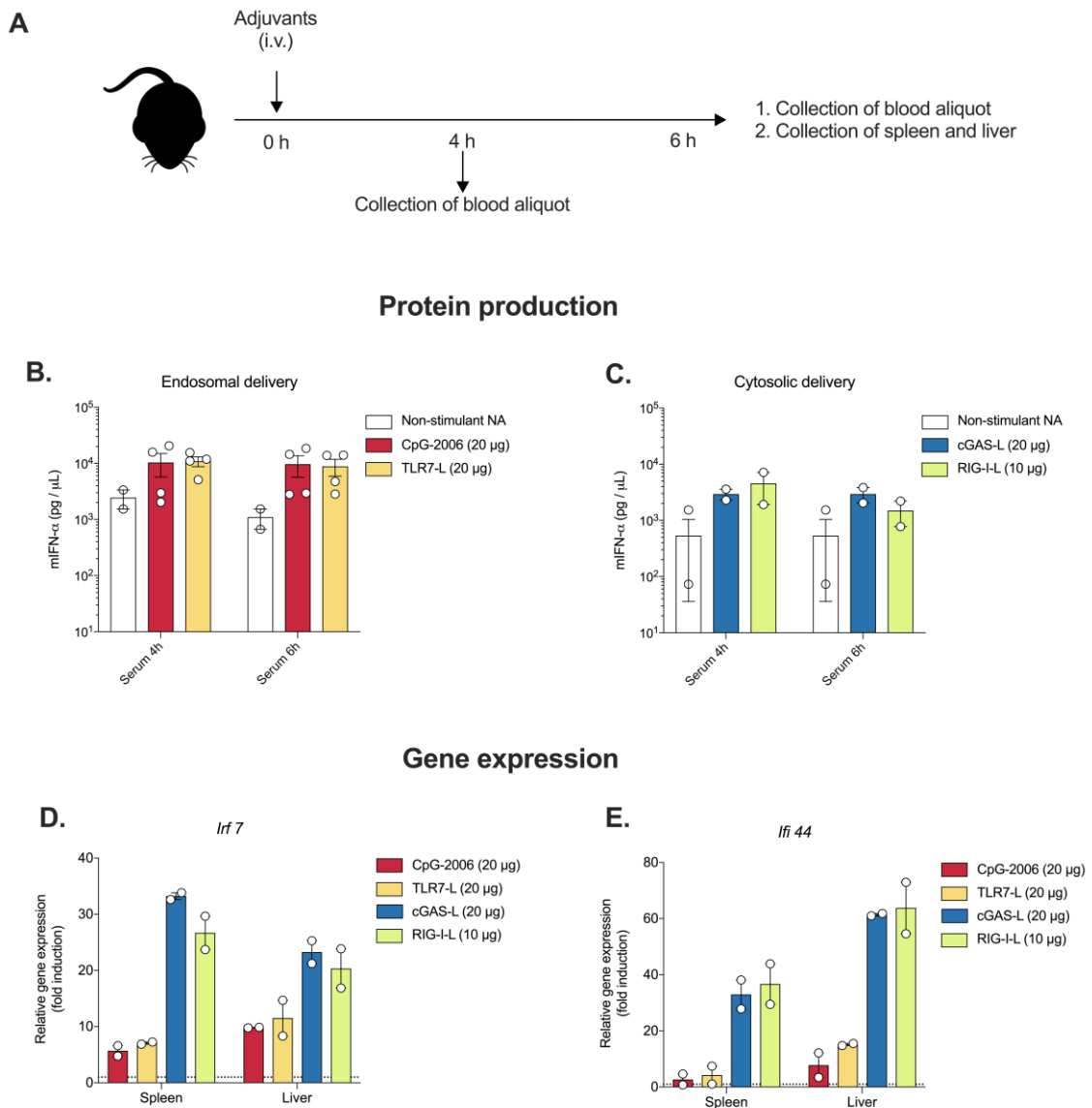


Figure 4.10 Adjuvants induce type I IFN production, preliminary experiment. A. B6 mice were stimulated i.v. with 20  $\mu$ g of CpG-2006, 20  $\mu$ g of TLR7-L or 20  $\mu$ g of non-stimulating DNA complexed in DOTAP; or with 20  $\mu$ g of cGAS-L, 10  $\mu$ g of RIG-I-L or 10  $\mu$ g of non-stimulating RNA complexed in Jet-PEI. 4 h later mice were cheek bled and serum collected. 6 h later mice were sacrificed, blood, spleen and liver were collected. Serum was collected and RNA was isolated from the homogenized organs. Production of mIFN- $\alpha$  was measured in serum by ELISA and the expression of several ISG was investigated in spleen and liver by qPCR. Production of mIFN- $\alpha$  in sera of mice stimulated with CpG-2006 or TLR7-L (B.) and cGAS-L or RIG-I-L (C.). Relative gene expression of *Irf 7* (D.) and *Ifi 44* (E.) expressed by fold induction in the spleen and liver.

These data are from a single experiment and require confirmation. After 4 h, DOTAP substantially enhances the early production of IFN $\alpha$  induced by CpG-combo and CpG-2006 and it had no effect when complexed with CpG-1668 (Fig. 4.11 A). Despite the limited production of IFN $\alpha$  induced by CpG-1668, this adjuvant was clearly functional as it induced the formation of liver T<sub>RM</sub> cells in the absence or presence of DOTAP (Fig. 4.1 and Fig. 4.7). All three CpG in the absence of DOTAP failed to induce the production of this cytokine as compared to control mice (Fig. 4.11 A). After 10 h of stimulation, the levels of IFN- $\alpha$  in sera were all similar to control mice (Fig. 4.11 B).

One of the major reasons driving this experiment was the poor performance of the cGAS-L and TLR7-L to induce expansion and formation of liver T<sub>RM</sub> cells. Importantly, results from this experiment showed that these two adjuvants (at both high and low doses) had the capacity to induce the early production of IFN $\alpha$  (Fig. 4.11 C). Observations indicated that there was no dose effect for RIG-I-L, TLR7-L or cGAS-L. Interestingly, high levels of the cytokine were still detected after 10 h in mice stimulated with RIG-I-L independently of the dose (Fig. 4.10 D). Preliminary data from this experiment suggested that all adjuvants transported from Germany to Australia were still active and able to induce the production of IFN- $\alpha$  *in vivo*.

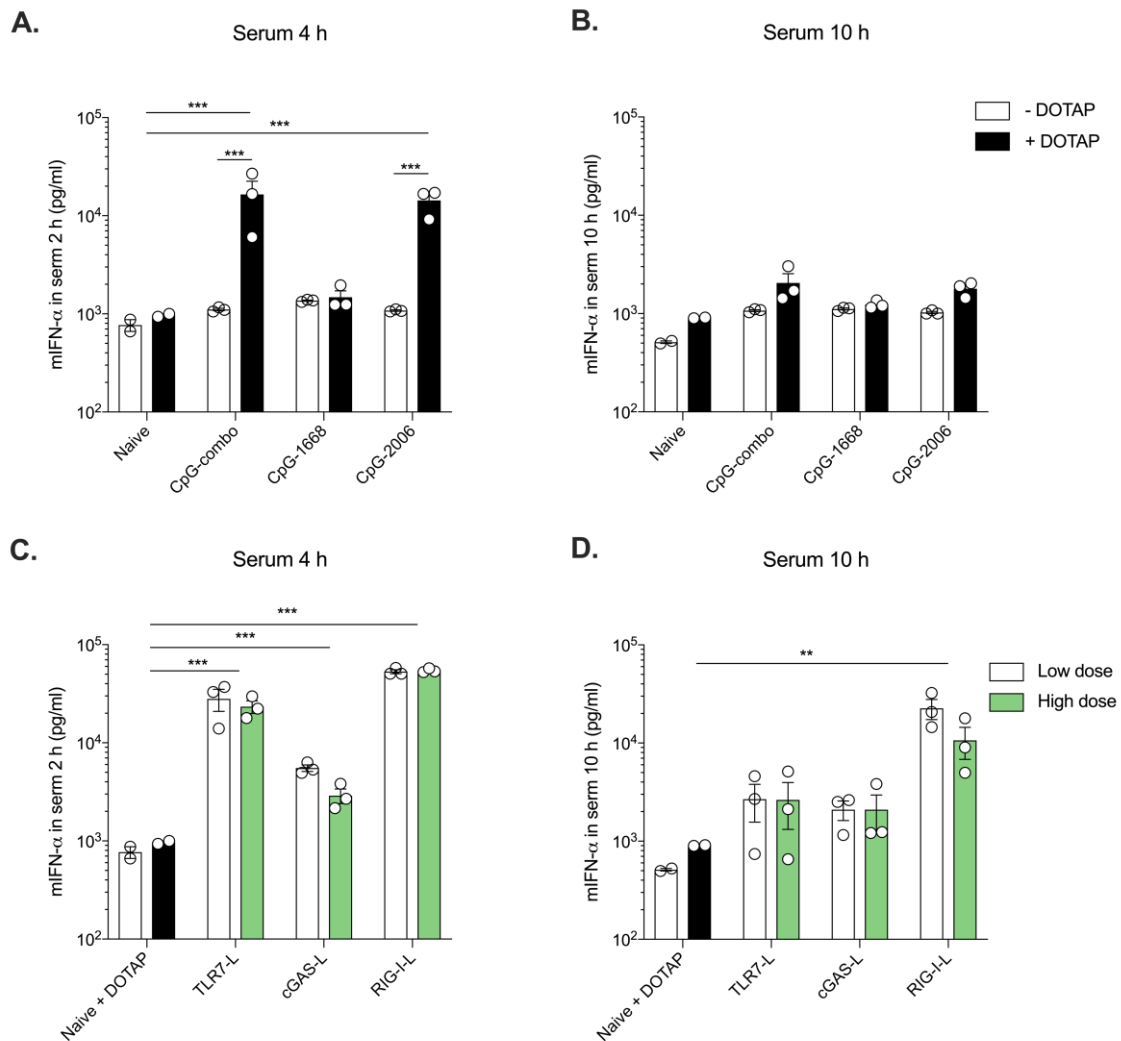


Figure 4.11 Adjuvants induce type I IFN production, late control experiment. A. B6 mice were grouped in 2. The first group received 75  $\mu$ g of CpG-combo, or CpG-1668 or CpG-combo complexed or not in DOTAP. The second group were stimulated with 30 or 75  $\mu$ g of TLR7-L complexed in DOTAP, or 20 or 75  $\mu$ g of cGAS-L complexed in Jet-PEI®, or 20 or 40  $\mu$ g of RIG-I-L complexed in Jet-PEI®. Control mice received only DOTAP or remained naive. Three mice were stimulated per each treatment. Later, at 4 and 10 h mice were bled and serum collected. Production of mIFN $\alpha$  was measured in serum by ELISA. Production of IFN $\alpha$  in mice stimulated with three different CpG at 4 h (A.) and 10 h (B.). Level of IFN $\alpha$  in mice receiving different doses of cGAS-L or RIG-I-L or TLR7-L at 4 h (C.) and 10 h (D.). Results are from a single experiment. Data were log-transformed and compared by one-way ANOVA and Tukey's multiple comparisons test.  $P > 0.05$ , \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  \*\*\*,  $P < 0.001$ .

### 4.3 Discussion:

In this study, a series of different nucleic acids were assessed as novel adjuvants for the induction of liver T<sub>RM</sub> cell formation induced by Clec9A targeted immunisation. The agonists investigated here included 3 types of CpG ODN, which are TLR9 ligands; p I:C, a TLR3 ligand; LPS, a TLR4 ligand, a synthetically produced ssRNA as TLR7 ligand, a dsDNA for cGAS stimulation and a 5' 3p dsRNA as RIG-I agonist. Results demonstrated that CpG ODN were the best adjuvant to promote formation of T<sub>RM</sub> cells in the liver, as well as large numbers of T<sub>EM</sub> cells in the spleen. Importantly, the number of liver T<sub>RM</sub> cells was heavily influenced by the amount of the adjuvant per mouse, as a high dose induced the formation of significantly and higher proportions of memory T cells in the liver. The results also showed that the transfection reagent DOTAP enhanced the adjuvant effect mediated by several CpG ODN, at least in their ability to form liver T<sub>RM</sub> cells; however, it did not improve immunity mediated by the prime-and-trap vaccine. Moreover, CpG ODN had no significant effect on the formation of liver T<sub>RM</sub> cells induced by RAS vaccination nor on the protection mediated by this vaccine. In summary, the data presented in this chapter demonstrate the superior capacity of CpG ODN to promote liver T<sub>RM</sub> cell formation in B6 mice mediated by Clec9A targeted immunisation.

#### *4.3.1 CpG ODN as an adjuvant for CD8 T cell expansion*

The specific mechanisms that govern the early expansion and the formation of memory CD8 T cells mediated by CpG ODN remain to be fully elucidated. Some of the factors that may influence these effects will be discussed below.

Intracellularly, the stimulation of TLR9 in APC by CpG ODN activates the adaptor protein myeloid differentiation primary response protein 88 (MYD88) (Häcker et al., 2000); which ultimately leads to the expression of type I IFN stimulated genes (ISG) or the production of proinflammatory cytokines via the



NF- $\kappa$ B pathway (Junt and Barchet, 2015). There are 4 types of CpG ODN described, and although all of them bind to TLR9, they have different immunological effects (Bode et al., 2011).

A-class CpG ODN induces strong production of type I IFN by pDC, leading to the activation of cellular antiviral state in human primary cells and murine splenic DC (Guiducci et al., 2006; Hemmi et al., 2003). B-class ODN, triggers pDC differentiation and the production of TNF- $\alpha$ , which in turn leads to T cell activation. Additionally, B-class ODN induces B cells to proliferate and secrete IgM (Hartmann and Krieg, 2000; Hemmi et al., 2003; Rothenfusser et al., 2001; Verthelyi et al., 2001). Studies have demonstrated that the distinct effects mediated by A-class versus B-class relies on differences in the retention time in early or late endosomes (Guiducci et al., 2006; Honda et al., 2005). A-class ODN are able to form complexed structures, allowing for a longer retention into early endosomes. This promotes the signalling through the transcription factor IRF7, which leads to IFN- $\alpha$  expression (Bode et al., 2011; Guiducci et al., 2006; Honda et al., 2005). In contrast, the B-class ODN single strand structure enables rapid transport to late endosomes for signalling via NF- $\kappa$ B, for the production of proinflammatory cytokines, such as TNF- $\alpha$  (Bode et al., 2011; Guiducci et al., 2006; Honda et al., 2005). C-class combines features from both A and B-class, as it stimulates B cells to secrete IL-6 and pDC to produce IFN- $\alpha$  (Hartmann et al., 2003; Marshall et al., 2003). P-class ODN is the most recently described ODN, shown to produce more IFN- $\alpha$ , IL-6 and the CXCR3 ligand CXCL10 than B- or C-class ODN (Samulowitz et al., 2010).

The diversity of immune responses mediated by CpG ODN and their relatively simple structure have triggered investigation of these compounds as adjuvants. In fact, several vaccine clinical trials have been performed using CpG ODN for the generation of antibodies for hepatitis B (Cooper et al., 2004) and malaria (Mullen et al., 2008); and for the induction of CD8 T cell responses against melanoma, breast and ovarian cancer (Valmori et al., 2007).

In this chapter and previous published studies, data revealed that CpG-combo, which is a B-class ODN linked to a P-class ODN, was one of the best adjuvants at inducing early expansion of CD8 T cells and liver T<sub>RM</sub> cell formation (Fernandez-Ruiz et al., 2016; Holz et al., 2018). B-class ODN not only promotes pDC differentiation and production of TNF- $\alpha$ , but also binds to DEC205, a molecule highly expressed on CD8+ DC, improving their maturation, production of IL-12 and stimulation of CTL (Caminschi et al., 2013; Lahoud et al., 2012). Therefore, by linking a B-class to a P-class ODN, the superior capacity for cytokine induction of the P-class ODN is targeted to CD8+ DC. Importantly, the Clec9A mAb used in this study for the antigen immunisation also target CD8+ DC, thus both the antigenic peptide and this potent adjuvant could converge in the same DC. These features make CpG-combo a superior adjuvant for priming of CD8 T cells mediated by Clec9A targeted immunisation.

In the current study, it was demonstrated that as long as CpG-2006 and CpG-1668 were complexed in DOTAP, they were as potent as CpG-combo for the induction of T<sub>RM</sub> cells in the liver. CpG-2006 and CpG-1668 are both B-class ODN characterised by a simple structure that makes them susceptible to the activity of DNases present in sera (Hanagata, 2012). If they successfully reach the endosome, they are rapidly transported to lysosomes where they signal TLR9 through the NF- $\kappa$ B pathway (Honda et al., 2005) before being degraded. The P-class ODN, present in CpG-combo, contains palindromic sequences that enable the formation of complex structures, protecting them from extracellular degradation (Samulowitz et al., 2010). Potentially, this feature also promotes the retention of the ODN within early endosomes for the activation of type I IFN. Forming a complex of B-class ODN with a cationic lipid such as DOTAP provides several advantages. First, encapsulation within a cationic lipid protects DNA from extracellular degradation (Capaccioli et al., 1993). Second, the now positively charged DNA-lipid complex enhances uptake by the negatively charged cellular membrane (Chesnoy and Huang, 2000). Finally, this complex is more likely to be retained into both the early and late endosome, as compared to a “naked” B-class

ODN, triggering type I IFN as well as proinflammatory cytokines mediated by NF- $\kappa$ B (de Jong et al., 2010; Khalil et al., 2006).

The superior capacity of liposomes to enhance CpG-mediated CD8 T cell responses has been previously demonstrated by several studies. Vaccination studies of a breast cancer model, using a HER-2/neu epitope, revealed that encapsulation of the peptide together with CpG ODN in liposomes substantially enhanced the peptide-specific CD8 T cell response as measured by IFN- $\gamma$  production (Li et al., 2003). In a B16 melanoma model, Jérôme and collaborators showed that co-administration of liposome-encapsulated antigen and a B-class CpG-ODN elicited a potent anti-tumor CTL response (Jérôme et al., 2006). For the OVA antigen model, it was also described that delivery by liposomal-nanoparticles of CpG-ODN in the presence of OVA peptide, enhanced the specific killing mediated by CD8 T cells (Chikh et al., 2009).

Despite the potent adjuvant effect of CpG-combo, it failed to enhance the formation of liver T<sub>RM</sub> cells mediated by RAS vaccination. Similarly, it was previously reported that B-class CpG ODN do not enhance the expansion of malaria specific-CD8 T cells (i.e. PbT-I cells) mediated by RAS immunisation (Lau et al., 2014). Interestingly, a preliminary observation reported in this chapter indicated that mice vaccinated with RAS plus CpG-combo complexed in DOTAP generated less liver T<sub>RM</sub> cells than mice receiving only RAS and failed to control infection. This may be explained by the fact that non-specific-protection against sporozoite challenge is induced by CpG-combo alone, at least for a short time (Gramzinski et al., 2002, unpublished data Heath lab). It is possible that interference with hepatocyte invasion by RAS reduces its capacity to induce the formation of memory cells. Further experiments are required to unveil the mechanisms that govern this effect. For example, if the adjuvant CpG is administered 24 h after RAS vaccination, sporozoites may have more time to infect hepatocytes to ensure local antigen presentation and liver T<sub>RM</sub> cell formation.

In summary, the data presented here are in line with several previous publications, demonstrating the superior capacity of CpG to induce CD8 T cell expansion and for the cationic liposome DOTAP to enhance this adjuvant effect.

#### *4.3.2 Other agonists*

In addition to early CD8 T cell expansion, CpG ODN also enhanced the formation of liver T<sub>RM</sub> cells mediated by Clec9A targeted immunisation. All other adjuvants tested, i.e. p I:C, LPS, a TLR7, cGAS and RIG-I ligand; failed to promote the formation of liver T<sub>RM</sub> cells as effectively as CpG ODN.

As briefly mentioned before, CpG ODN has the capacity to induce inflammation in the liver and the formation of iMATEs, which in turn, supports proliferation and accumulation of CD8 T cells in the liver (Huang et al. 2013). This feature is of special interest for the design of strategies designed to induce CD8 T cell responses in the liver. Importantly, Huang and collaborators also tested MPLA a TLR4 ligand less toxic than LPS, p I:C and a TLR7 ligand. They showed that MPLA induced a moderate number of CTL in the liver and that neither p I:C nor TLR7 ligand had the capacity to induce the formation of iMATEs. Consequently, the proliferation of CD8 T cells was reduced (Huang et al., 2013). Data generated here support the idea that LPS, p I:C and TLR7-L are poor adjuvants for the formation of memory CD8 T cells in the liver.

Whereas, TLR4 activation enhances Th1 immune responses and ligands of TLR4 have been exploited as potential adjuvants (Steinhagen et al., 2011), LPS, a well-known TLR4 agonist, is also known to induce tolerance. Excessive MyD88 activation by large amounts of free LPS causes the production of pro-inflammatory cytokines that can lead to endotoxin shock when overproduced systemically (Biswas and Lopez-Collazo, 2009). Endotoxin tolerance is a host protection mechanism intended to counteract the effects of extreme inflammation. In fact, mice exposed to non-lethal doses of LPS are protected from a secondary otherwise lethal dose of LPS (Cavaillon and Adib-Conquy, 2006;

Freudenberg and Galanos, 1988). In the clinic, patients who survive sepsis and have undergone “cytokine storm”, usually exhibit an immunocompromised phase correlated with high risk of secondary infection and mortality (Biswas and Lopez-Collazo, 2009). *In vitro* studies in mouse macrophages and human monocytes have demonstrated a reduced ability to produce pro-inflammatory cytokines such as TNF- $\alpha$ , IL-6 and IL-12 after a second administration of LPS (Akira and Takeda, 2004). Importantly, *in vivo* stimulation with LPS leads to a substantial loss of DC and further impairment in IL-12 production (Wysocka et al., 2001) – possibly influencing subsequent priming of CD8 T cells and their recruitment to the liver. Novel biosynthetically engineered TLR4 ligands have been designed to overcome the induction of toxicity and currently are being tested as adjuvants for vaccines against bacterial infections, such as *Bordetella pertussis*, *Shigella flexneri* and *Vibrio cholerae* (Zariri and Van Der Ley, 2015).

TLR3 signalling induces strong type I IFN response and promotes cross-presentation leading to potent CD8 T cell responses (Schulz et al., 2005; Sistigu et al., 2014). In this regard, poly I:C, a synthetic TLR3 agonist has been shown to have substantial adjuvant capacity for an influenza vaccine and strong antitumor activity in murine tumour models (Currie et al., 2008; Ichinohe et al., 2005; Pulko et al., 2009; Salem et al., 2005). These studies differ from the experiments described in this chapter by two major aspects. First, the route of administration is different. While mice here received p I:C systemically through intravenous injection, for the influenza vaccine and the solid tumour studies, p I:C was locally administered, respectively, by intranasal or intra-tumoral application (Currie et al., 2008; Ichinohe et al., 2005). Second, the time frame of the CD8 T cell response is different. These published studies investigated the therapeutic properties of effector CD8 T cells mediated by p I:C stimulation (Pulko et al., 2009; Salem et al., 2005) rather than the formation of memory CD8 T cells, which is the main topic of investigation in this chapter.

Studies have demonstrated the capacity of TLR7 agonists as adjuvants for the induction of CTL antitumor responses in murine models (Drobits et al.,

2012; J. Wu et al., 2017). Particularly, for these two studies, the TLR7 ligands were applied locally either topically in the case of the melanoma model, or intratumorally in case of the breast cancer model (Drobits et al., 2012; J. Wu et al., 2017). Similarly, pulmonary administration of a synthetically modified TLR7 ligand, termed 1V270, was shown to increase resistance to lethal pulmonary viral infection in mice (Wu et al., 2014). These data may indicate that the cell-mediated immunity induced by stimulation with TLR7 agonists is better achieved by a local rather than the systemic administration route used in this chapter. In fact, some studies have reported *in vivo* tolerance induced by the systemic and repeated administration of TLR7 agonists (Bourquin et al., 2011; Gunzer et al., 2005; Hayashi et al., 2009). Pre-treatment of pDC and myeloid DC with TLR7 ligands leads to a reduced production of the proinflammatory cytokines IFN- $\alpha$ , IL-6 and IL-12p70 after TLR restimulation (Bourquin et al., 2011). Moreover, *in vitro* cultures of pDC with TLR7 ligand showed increased secretion of the anti-inflammatory cytokine IL-10 after a second stimulation with the ligand (Bourquin et al., 2011), and high expression of inhibitory molecule PD-L1 after a single TLR7 stimulation (Wu et al., 2019).

Despite promising data suggesting RIG-I agonists have the potential to be used as pan-antivirals, vaccine adjuvants or mediators of antitumor activity (Y. Wu et al., 2017; Yong and Luo, 2018), only a few studies have demonstrated the effect of RIG-I ligands as vaccine adjuvants (Kasumba and Grandvaux, 2019). For a lethal influenza murine model, a sendai virus-derived RIG-I ligand was shown to increase specific IgG titres when co-administered with inactivated virus vaccine, leading to reduced morbidity and mortality after challenge (Martínez-Gil et al., 2013). Moreover, the effect of a 3pRNA RIG-I agonist as an adjuvant was confirmed for an OVA murine model, in which mice vaccinated with OVA peptide plus 3pRNA exhibited high specific cytotoxicity correlated to viral clearance after challenge with a modified adenovirus expressing OVA (Hochheiser et al., 2016). This last study supports the idea that RIG-I agonists could be exploited as adjuvants for hepatic infections, like hepatitis and malaria liver-stage. However, in this chapter, the RIG-I ligand used as adjuvant had a limited effect on the

induction of liver T<sub>RM</sub> cells. Importantly, Hochheiser *et al.* complexed the adjuvant together with the antigen, ensuring simultaneous delivery of adjuvant and antigen to the same APC (Hochheiser *et al.*, 2016). For the experiments described here, Clec9A mAb coupled to the antigen was co-administered in the same injection as the complexed RIG-I-L, but this does not ensure simultaneous delivery to the same cell. This factor may explain why RIG-I-L and other agonists failed to induce the formation of substantial liver T<sub>RM</sub> cells.

Recent studies have identified cGAS as a sensor of cytosolic DNA able to induce interferons and cytokines. Upon activation, cGAS produces a cyclic dinucleotide GMP-AMP (cGAMP) (Ablasser *et al.*, 2013; Diner *et al.*, 2013; Gao *et al.*, 2013). cGAMP functions as a second messenger and activates the adaptor protein STING (Burdette *et al.*, 2011; Burdette and Vance, 2013; Danilchanka and Mekalanos, 2013). In turn, STING activates the transcription factors NF- $\kappa$ B and IRF3 to induce type I IFN and cytokines (Tanaka and Chen, 2012; Wu *et al.*, 2013). In recent reports, cGAMP rather than ligands for cGAS has been used as an antiviral or adjuvant in different murine models. Li and collaborators demonstrated that cGAMP functions as an adjuvant for T cell activation and antibody production in an OVA *in vivo* model (Li *et al.*, 2013). Additionally, a study in healthy volunteers confirmed that cGAMP enhances the antigen-specific expansion, activation and killing of melanoma cell lines mediated by CD8 T cells isolated from PBMC (Gutjahr *et al.*, 2019). Two important factors may influence the fact that cGAS ligand had a limited effect for the induction of liver T<sub>RM</sub> cells. First and as already mentioned, the adjuvant may not reach the same APC as the antigen, failing to provide the licensing signals for an adequate CD8 T cell priming. Second, if the cGAS ligand successfully reaches the cytoplasm, this does not ensure the production of cGAMP and further activation of the STING pathway.

The last observation in this chapter showed early production of IFN- $\alpha$  detected in serum of mice stimulated with TLR-7-L, cGAS-L and RIG-I-L. Importantly, it was also established that these three agonists did not promote

substantial formation of liver T<sub>RM</sub> cells. Further experiments are required to confirm these findings, but as it stands this may indicate that level of IFN- $\alpha$  in serum does not correlate with T<sub>RM</sub> cell formation in liver.

Studies have demonstrated that licensing of DC for optimal stimulation of CD8 T cells requires cis-activation by danger signals rather than indirect stimulation provide by inflammatory cytokines produced by adjacent cells (Desch et al., 2014). Accordingly, vaccine formulations that link antigen and danger signals, which act as adjuvants showed an enhanced capacity to induce humoral and cellular immunity (Demento et al., 2010; Hou et al., 2011; Moon et al., 2011; Steers et al., 2009). Thus, further experiments exploring the possibility of linking the here tested adjuvants to the antigen represent a strategic variation to enhance the expansion of CD8 T cells and the potential formation of liver T<sub>RM</sub> cells. Translational implications will be further discussed on Chapter 5.



## CHAPTER 5

General discussion

## 5.1 Context

In 2014, members of our group developed a B6 TCR transgenic line able to produce CD8 T cells specific for PbA. This TCR transgenic line, termed PbT-I, produced cells that responded to blood-stage PbA, *P. yoelii* XNL and *P. chabaudi*, and to liver stage PbA parasites (RAS) (Lau et al., 2014). Soon after, these cells were used to develop the novel vaccination strategy “prime-and-trap” and generate liver PbT-I T<sub>RM</sub> cells, which were shown to be crucial for protection against PbA sporozoite challenge (Fernandez-Ruiz et al., 2016). T<sub>RM</sub> cells are part of a memory T cell subpopulation that are localised in tissues and provide potent immune surveillance. The strategic location of T<sub>RM</sub> cells allows these cells to mediate effector responses against secondary infections more rapidly than their counterparts, the circulating memory cells, which need to be recruited to the infected organ (Mueller and Mackay, 2016). Given the importance of liver T<sub>RM</sub> cells in the context of protection against malaria and other liver-associated pathogens (Fernandez-Ruiz et al., 2016; Pallett et al., 2017; Stelma et al., 2017), studies on how to enhance their formation are highly relevant, and thus the experiments conducted and presented in this thesis aimed to investigate this goal. Recently, the cognate epitope for the PbT-I cells, called the Pb-1 epitope in this thesis, has been identified as part of the RPL6 protein of PbA (Valencia-Hernandez et al., 2020). Here we addressed the suitability of this epitope for the induction of endogenous liver T<sub>RM</sub> cells that confer sterilizing protection through vaccination. Additionally, it is known that inflammation favours the formation T<sub>RM</sub> cells (Mackay et al., 2012a) and that adjuvants can affect their numbers (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Recent data showed that CpG induced liver inflammation (Huang et al., 2013) and supports the formation of T<sub>RM</sub> cells in this organ (Fernandez-Ruiz et al., 2016; Holz et al., 2018). Based on these findings, in this thesis, nucleic acid sensor agonists were assessed as novel adjuvants for the formation of liver T<sub>RM</sub> cells.

## 5.2 Summary of main findings

In chapter 3, a series of different immunisation strategies targeting the Pb-1 epitope were implemented and specific CD8 T cell responses were assessed in B6 mice. Analysis of the T cell repertoire of naive B6 mice revealed that the number of specific CD8 T cell precursors for the Pb-1 epitope was very large. These cells were successfully expanded by two different immunisation strategies: i) Single injection with Clec9A mAb coupled to Pb-1 plus adjuvant (vaccine 1) and ii) Prime-and-trap targeting the Pb-1 epitope. Large numbers of specific T<sub>RM</sub> cells were formed in the livers of vaccinated mice. While mice developed a certain level of protection when vaccinated with vaccine 1, higher levels of protection were achieved in mice vaccinated with prime-and-trap.

In Chapter 4, various pattern recognition receptor (PRR) agonists were assessed as novel adjuvants for the induction of liver T<sub>RM</sub> cell formation induced by Clec9A targeted immunisation. The agonists investigated included: i) LPS, a TLR4 ligand, which is a plasma membrane receptor; ii) agonists of the endosomal receptors TLR3, TLR7 and 3 types of TLR9 ligands; and iii) ligands of the cytosolic receptors cGAS and RIG-I. The data revealed that CpG ODN was the best adjuvant to promote formation of T<sub>RM</sub> cells in the liver. Also, the number of liver T<sub>RM</sub> cells was influenced by the amount of adjuvant administered; that is, a high dose induced the formation of a higher proportion of memory T cells in the liver. It was also determined that the transfection reagent DOTAP enhanced the adjuvant effect mediated by several CpG ODN, at least in their ability to form liver T<sub>RM</sub> cells. However, use of DOTAP with the prime-and-trap vaccine did not improve protection. Finally, CpG ODN did not enhance formation of liver T<sub>RM</sub> cells when used in combination with RAS, nor improve protection mediated by this vaccine.

### 5.3 A protective malaria epitope

Malaria still claims more than 400,000 lives per year, mostly among African children (WHO, 2019). Naturally acquired immunity (NAI) to clinical malaria only develops with age in areas with high parasite transmission. In adults, NAI protects against clinical malaria by maintaining low parasitemia levels (Baird et al., 2003; Owusu-Agyei et al., 2001). Sterilizing immunity, i.e. complete absence of blood stage parasites, never develops and clinical immunity is not long lasting (Doolan et al., 2009; Keegan and Dushoff, 2013). Hence, there is an urgent need for the development of an effective vaccine. However, despite decades of intense research, an effective malaria vaccine for application in endemic areas remains elusive (Coelho et al., 2017; Draper et al., 2018). Vaccines targeting pre-erythrocytic (PE) parasite stages are ideal because they avoid the occurrence of symptoms and block transmission. PE vaccines based on whole-sporozoite administration have proven to be effective at conferring protection in murine models and in malaria naïve humans (Clyde, 1990; Mordmüller et al., 2017; Nussenzweig et al., 1967; Seder et al., 2013). However, these approaches face major limitations that question their suitability for routine use in endemic areas (Draper et al., 2018). Some of those challenges are associated with technical and logistical issues of large scale sporozoite production (mosquito breeding and manual dissection) and distribution in endemic areas (maintenance of the cold chain). However, a greater concern is that while RAS vaccination induces protection in malaria naïve humans, it has failed to protect with a comparable efficacy in malaria-experienced adults in Mali (Sissoko et al., 2017).

It is known that the sterilizing immunity induced by PE vaccines is substantially mediated by CD8 T cells (Cockburn et al., 2014; Doll et al., 2016; Epstein et al., 2011; Ishizuka et al., 2016; Nganou-Makamdop et al., 2012; Schofield et al., 1987; Weiss et al., 1988), with liver-associated memory cells having been recently shown to play a prominent role (Fernandez-Ruiz et al., 2016; Tse et al., 2014). In this regard, vaccine strategies aimed at generating

CD8 T cell responses against specific PE epitopes have been developed recently. These liver-stage subunit vaccines overcome the manufacturing challenges of whole-parasite approaches, but also come with the limitation that highly protective *Plasmodium* epitopes are largely unknown.

Prime and boost immunisations with heterologous viral vectors encoding for PE antigens have been exploited for their capacity to elicit CD8 T cell responses (Draper et al., 2018). Murine studies demonstrated high levels of protection associated with CD8 T cells induced by prime and boost immunisations with viral vectors encoding several PE antigens – contained in a construct termed ME-TRAP (Gilbert et al., 2002; Li et al., 1993; Reyes-Sandoval et al., 2010; Schneider et al., 1998). In contrast, mixed results in vaccine efficacy were obtained in humans depending on malaria pre-exposure. That is, in malaria naïve volunteers, vaccine efficacy (determined as sterilizing immunity or delay in the onset of parasitemia) was partial ranging from 46-57%, whereas clinical trials in malaria endemic countries revealed negligible efficacy (Bliss et al., 2017; Ewer et al., 2013; Hodgson et al., 2015; Mensah et al., 2016; Ogwang et al., 2015; Tiono et al., 2018).

Given these mixed results, efforts have been made to improve the immunogenicity and improve protection levels of liver-stage subunit vaccines. Viral vectored immunisations with other PE antigens not included in ME-TRAP have failed to protect with a comparable efficacy to those induced by the ME-TRAP construct (Hodgson et al., 2015). Immunisations with viral vectors encoding for blood-stage antigens failed to confer protection against sporozoite challenge in malaria naïve volunteers (Sheehy et al., 2012). Even the co-administration of viral vectors encoding for ME-TRAP together with the blood-stage epitope MSP1 did not improve vaccine efficacy (Sheehy et al., 2012). In another attempt to increase the efficacy of the viral vectored ME-TRAP vaccine, malaria naïve volunteers were immunised with RTS,S, which induces Ab against CSP, plus viral vectors encoding for ME-TRAP simultaneously. The groups who received RTS,S plus the viral vectors had lower protection levels compared to

the groups immunised with RTS,S only, although differences were not significant due to sample size (Rampling et al., 2018).

To date, CSP and TRAP have been extensively targeted by several vaccination strategies, since they have been considered important immunogenic antigens found in *P. falciparum* (Müller et al., 1993; Rogers et al., 1992), because both are essential for sporozoite motility and liver-stage infection (Nussenzweig and Nussenzweig, 1985; Sultan et al., 1997). However, as mentioned before, these strategies have failed to provide robust protection. Therefore, there is need for the discovery of novel and highly immunogenic CD8 T cell epitopes. One of the major challenges for the design of liver-stage sub-unit vaccines is the limited number of known protective MHC-I epitopes derived from liver stage *Plasmodium* parasites (Draper et al., 2018; Schmidt et al., 2010). The lack of a suitable and sufficient source of epitopes eluted from MHC molecules of infected hepatocytes hampers the efforts to identify immunogenic CD8 T cell epitopes. In fact, the identification of the Pb-1 epitope, described in chapter 3, took more than 6 years. Several strategies have been recently developed to tackle this problem.

Transcriptomic studies revising gene expression patterns across different parasite stages and *Plasmodium* species provided valuable information on proteins differentially expressed during liver-stage infection (Howick et al., 2019; Sacci et al., 2005; Speake et al., 2016; Tarun et al., 2008). This information together with epitope prediction software (Vita et al., 2019) support the identification of immunogenic peptides that can be potentially targeted by liver-stage subunit vaccines (Hafalla et al., 2013). Furthermore, humanized liver-chimeric mouse models and new *in vitro* liver-stage cell culture methods are currently exploited as a way to yield sufficient epitopes from infected hepatocytes for analysis by mass spectrometry (Longley et al., 2015). In a recent study, human liver chimera FAH<sup>-/-</sup> NOD Rag1<sup>-/-</sup> IL2R $\gamma$ <sup>NULL</sup> (FNRG) mice were infected with *P. falciparum* sporozoites and then peptides from the liver were isolated for sequencing. The collected peptides were derived from the proteasomal degradation of proteins and potentially have the right size to be presented by

MHC molecules (Winer et al., 2020). Deviating from several other studies, the authors used a top down mass spectrometry approach, a technique that avoids the digestion of peptides and the loss of post-translational modifications. This approach allows for the identification of peptides in their native form (Cui et al., 2011). In this preliminary study, a series of CD8 T cell *P. falciparum*-derived antigens were identified as potential targets for a malaria vaccines (Winer et al., 2020).

As well as identifying potential immunogenic epitopes, it is important to understand the features of protective antigens to allow application of this knowledge to the selection of new antigenic targets for use in subunits vaccines. Some factors that may influence immunogenicity of CD8 T cell epitopes include: i) the number of naïve CD8 T cell precursors (Jenkins and Moon, 2012), ii) the expression pattern during liver-stage infection (Butler et al., 2011; Speake et al., 2016), iii) the presence of the protein in the hepatocyte cytosol for presentation on MHC I (Boddey et al., 2016; Cockburn et al., 2011; Montagna et al., 2014), and iv) protein polymorphism.

As mentioned before, the TRAP protein has been an antigen of choice for liver-stage subunit vaccines and, as such, has been tested in several clinical trials. However, robust immunity has not been achieved (Bliss et al., 2017; Ewer et al., 2013; Hodgson et al., 2015; Mensah et al., 2016; Ogwang et al., 2015; Tiono et al., 2018). Moreover, in B6 mice – a strain that is considered difficult to protect (Doolan and Hoffman, 2000; Schmidt et al., 2010) – vaccination targeting TRAP induced a reduction in liver parasite load, but failed to induce sterile protection (Hafalla et al., 2013). Recently published data generated by members of our group revealed that B6 mice vaccinated with prime-and-trap targeting a TRAP epitope generated large numbers of specific liver T<sub>RM</sub> cells, but only low levels of sterile protection (Valencia-Hernandez et al., 2020). On the contrary, results from chapter 3 demonstrated that B6 mice immunised with prime-and-trap targeting the RPL6 protein developed both, high numbers of endogenous liver T<sub>RM</sub> cells as well as high levels of sterile protection.

We hypothesize that two major factors may contribute to the different immunogenicity of RPL6 and TRAP in B6 mice. The first is the relatively high number of Pb-1-specific precursors compared to the small number specific for the TRAP epitope (Chapter 3). It has been demonstrated that the frequency of peptide-specific naive T cell precursor determines to a certain extent the magnitude and expansion of the subsequent T cell response after exposure to the specific peptide (Jenkins and Moon, 2012; Moon et al., 2007; Joshua J Obar et al., 2008). The second factor is the differences in the expression patterns of the proteins. While TRAP is mainly expressed by sporozoites and not liver-stage parasites (Amino et al., 2006), RPL6 is expressed throughout the liver and blood stages of infection by *P. berghei* and *P. yoelii* (Howick et al., 2019; Sacci et al., 2005; Tarun et al., 2008). This feature of RPL6 increases its possibility to be presented by hepatocytes and mediate CD8 T cell recognition and activation of effector functions.

An important characteristic of RPL6 that might be beneficial for future translation of a vaccine that targets this protein is its highly conserved nature (Valencia-Hernandez et al., 2020). Analysis on a dataset of more than 2,000 *P. falciparum* genomes from 15 countries determine that more than 99% of the sequences share same haplotype for PfRPL6 (Valencia-Hernandez et al., 2020). By contrast, TRAP, which is a surface protein and likely subjected to Ab selection pressure, is highly polymorphic, as confirmed by sequencing of isolates from endemic areas (Kosuwin et al., 2014; Ohashi et al., 2014; Robson et al., 1990; Srisutham et al., 2018).

When compared to a single injection of RAS, prime-and-trap targeting PbRPL6 induced higher levels of protection in B6 mice (data reported in Chapter 3 and 4). Data presented here indicated that T<sub>RM</sub> cell-based subunit vaccines have the potential to induce sterile protection in B6 mice, even when using a single epitope. Moreover, the fact that *P. falciparum* RPL6 is highly conserved, makes of this protein a potential target for the development of a human malaria vaccine. Notably, our research group has also now identified an immunogenic



HLA\*A02:01-restricted epitope in *P. falciparum* RPL6 (Valencia-Hernandez et al., 2020).

#### 5.4 Adjuvants for liver T<sub>RM</sub> cell formation

T<sub>RM</sub> cells provide effective tissue surveillance and can respond rapidly to infection by providing local effector functions and the recruitment of immune cells leading to the elimination of pathogens (Schenkel et al., 2014, 2013). Mouse studies have revealed that T<sub>RM</sub> cells play important roles in fighting infections against several types of pathogens, including viruses (Gebhardt et al., 2009; Teijaro et al., 2011), bacteria (Benoun et al., 2018; Sheridan et al., 2014; Wilk et al., 2017), fungi (Park et al., 2018), and parasites (Fernandez-Ruiz et al., 2016; Glennie et al., 2015).

Recently, efforts to develop vaccination strategies aimed at generating T<sub>RM</sub> cells have been increasing. Animal vaccine studies have shown promising results targeting T<sub>RM</sub> responses at barrier sites such as lung (Perdomo et al., 2016; Wakim et al., 2015; Zens et al., 2016), female reproductive tract (Çuburu et al., 2014; Shin et al., 2016; Tan et al., 2018), and liver (Fernandez-Ruiz et al., 2016; Gola et al., 2018). However, only a few studies have investigated the effect of adjuvants on the formation of T<sub>RM</sub> cells induced by those vaccination strategies.

In an influenza virus model, mice receiving *in vitro* activated effector T cells were immunised intranasally with DC-targeted mAb coupled to OVA plus LPS as adjuvant (Wakim et al., 2015). These mice generated lung T<sub>RM</sub> cells that contributed to high levels of protection against a lethal influenza virus challenge, but the effect of immunisation with the mAb in absence of LPS was not examined (Wakim et al., 2015).

Fernandez-Ruiz and collaborators explored the role of adjuvants on the induction of liver T<sub>RM</sub> cells in vaccinated mice. Using prime-and-trap vaccination, they observed that the adjuvants poly I:C and CpG both enabled the formation of

liver T<sub>RM</sub> cells, but CpG was superior and, in contrast to poly I:C, was able to induce strong protection against malaria liver-stage infection (Fernandez-Ruiz et al., 2016). A later study also showed that in the absence of antigen presentation, *in vitro* activated CD8 T cells formed liver T<sub>RM</sub> cells and that their formation was increased by poly I:C and CpG (Holz et al., 2018). This effect may be in part explained by the fact that i.v. injection of CpG induces intrahepatic myeloid cell aggregates that supports the accumulation of CD8 T cells in the livers of stimulated mice (Huang et al., 2013). Another group also investigated the role of adjuvants on the formation of liver T<sub>RM</sub> cells mediated by vaccination (Gola et al., 2018). For this study, liver T<sub>RM</sub> cells were generated in mice by priming with a modified adenovirus and boosting with nanoparticles both containing OVA antigen. They showed that the presence of R848-MPL (ligand for TLR7 and TLR4, respectively) on nanoparticles that also contained OVA antigen did not enhance the formation of liver T<sub>RM</sub> cells (Gola et al., 2018).

Given that inflammation favours T<sub>RM</sub> cell formation (Mackay et al., 2012b), adjuvants that induce inflammation in particular organs can be used to enhance numbers of T<sub>RM</sub> cells in those organs. In this thesis, a wide variety of adjuvants were assessed for the induction of liver T<sub>RM</sub> cells in the context of antigen targeted immunisation. The studied agonists included 3 types of CpG ODN, ligands for TLR3, TLR4, TLR7, as well as agonists for cGAS and RIG-I. The above 8 ligands bind a diverse range of pattern recognition receptors located in the plasma membrane, the endosomal compartment or the cytoplasm, and have the ability to induce the release of proinflammatory cytokines and the upregulation genes involve in type I IFN response. We hypothesized that some of the proinflammatory signals triggered by this type of adjuvant could promote the formation of liver T<sub>RM</sub> cells. However, the liver has particular features that may pose a challenge for the generation of a certain type of inflammation that could enhance T<sub>RM</sub> cell formation without causing tissue damage. These features will be discussed in the following paragraphs.

### 5.5 Challenges for the induction of liver inflammation

The liver receives blood from the gastrointestinal track through the portal vein, meaning it is constantly exposed to PAMPS derived from microbes present in the gut lumen. While the liver is constantly exposed to molecules from microorganisms and diet, it maintains homeostasis via tolerogenic immune mechanisms (Bigorgne and Crispe, 2010; Kesar and Odin, 2014; Schwabe et al., 2006). Liver tolerance has been associated with a tight regulation of the expression levels of pattern recognition receptors. For instance, the expression levels of TLR, as well as the signalling adaptor molecule MyD88 in liver cells are lower as compared to the levels observed in cells from other organs (De Creus et al., 2005; Faure-Dupuy et al., 2018; Lichtman et al., 1998; Zarembler and Godowski, 2002).

Hepatic immune regulation can also be promoted by liver resident immune cells. Upon activation, human and murine Kupfer cells (hepatic macrophages) release proinflammatory cytokines such as IL-6, TNF $\alpha$ , as well as the anti-inflammatory cytokine IL-10 (Frink et al., 2007; Knolle et al., 1995; Overland et al., 2005; Seki et al., 2001; Wu et al., 2009). It has been shown by *in vitro* assays that murine freshly isolated Kupffer cells respond less strongly to LPS if they have been pre-exposed to LPS, whereas pre-exposure to CpG does not affect the response of Kupffer cells to LPS stimulation (Schuchmann et al., 2004). Moreover, one study suggests that hepatic DC express lower mRNA levels of TLR4 relative to their splenic counterparts and that this phenotype may contribute to the tolerogenic environment in the liver (De Creus et al., 2005).

Dysregulation of the liver tolerance environment, in particular of the TLR stimulation pathway, has been associated with inflammatory liver diseases, such as alcoholic liver disease and hepatocellular carcinoma (Kiziltas, 2016; Mencin et al., 2009; Pimentel-Nunes et al., 2010; Soares et al., 2010). For example, it has been shown *in vivo* that chronic alcohol intake leads to the upregulation of the mRNA levels of TLR2, TLR4, TLR7, TLR8 and TLR9 in the liver, which is

associated with hepatitis (Gustot et al., 2006). Remarkably, in a murine model of alcoholic liver disease, TLR4 deficient mice were protected from the liver tissue damage induced by the prolonged alcohol intake (Hritz et al., 2008; Uesugi et al., 2001).

In a model of chemical-induced liver cancer, the LPS levels in sera of treated mice were higher compared to healthy control mice; and additionally, animals lacking TLR4 or microbiota developed significantly less hepatocellular carcinoma (Dapito et al., 2012; Yu et al., 2010). In liver samples of patients with non-alcoholic fatty liver disease the protein levels of TLR4 and TLR9 were higher as compared to healthy volunteers (Mridha et al., 2017). In the same study, authors showed in a murine model of non-alcoholic fatty liver disease that mice lacking TLR9 had reduced chronic inflammation and consequently, were protected from developing the disease (Mridha et al., 2017).

Based on the information described above, the liver is an organ known for its immunotolerance microenvironment that upon dysregulation can lead to severe chronic diseases. Under these circumstances, generating beneficial inflammation, one that could enhance T<sub>RM</sub> cell formation (Mackay et al., 2012a) without inducing liver tissue damage, is a difficult task to achieve. On the one hand, the threshold for TLR activation in the liver is higher given the downregulation of TLR expression compared to other organs (De Creus et al., 2005; Faure-Dupuy et al., 2018; Lichtman et al., 1998; Zarembek and Godowski, 2002), thus potent immune stimulators should be used to generate intrahepatic inflammation. On the other hand, hyperstimulation of TLR could lead hepatocyte apoptosis, to tissue damage (Dapito et al., 2012; Yu et al., 2010), and as a consequence, poor liver T cell accumulation.

In this thesis, mice were immunised with anti-Clec9A antibodies targeting the RPL6 *P. berghei* protein, in the presence of several nucleic acid-based adjuvants. The hypothesized mechanism is that these antibodies target cDC1 which when activated by the adjuvant, prime naïve CD8 T cells in secondary

lymphoid organs (Caminschi et al., 2008; Sancho et al., 2008). After priming, CD8 T cells would then migrate to the liver recruited by the inflammation induced locally by the adjuvant and ultimately generate T<sub>RM</sub> cells. Results from experiments performed in this thesis showed that mice receiving CpG-based adjuvants developed substantially more liver T<sub>RM</sub> cells, when compared to mice that received the other agonists (TLR3, -4, -7, RIG-I and cGAS ligands). These results support previous findings showing that CpG ODN are better at promoting T cell liver residency, when compared to TLR3 and TLR4 ligands (Fernandez-Ruiz et al., 2016; Holz et al., 2018). CpG ODN has the capacity to induce the formation of intrahepatic myeloid cell aggregates that promote proliferation and accumulation of CD8 T cells in the liver (Huang et al. 2013); and possibly, this is the mechanism by which CpG favours the formation of liver T<sub>RM</sub> cells. Huang and collaborators also showed in this study that TLR3, TLR4 and TLR7 ligands failed to induce the formation of such myeloid cell aggregates and subsequently, the proliferation of CD8 T cells was reduced (Huang et al. 2013). Accordingly, data generated in this thesis showed that TLR3, TLR4 and TLR7 ligands are poor adjuvants for the formation of memory CD8 T cells in the liver.

TLR3 synthetic ligands, such as poly I:C and TLR7 agonists have been used as adjuvants to promote potent CD8 T cells responses in the context of vaccines against viral infection and tumour therapy (Currie et al., 2008; Drobits et al., 2012; Ichinohe et al., 2005; Pulko et al., 2009; Salem et al., 2005; J. Wu et al., 2017). These published studies differ from the experiments described in this thesis by two major features, the route of administration and the triggered CD8 T cell response.

In this thesis, mice received the adjuvants systemically through intravenous injection, whereas on the published studies the TLR3 and TLR7 ligands were administered locally (intranasal, intrapulmonary, topically or intratumoral) depending on the model (Currie et al., 2008; Drobits et al., 2012; Ichinohe et al., 2005; J. Wu et al., 2017). These data may indicate that the cell-mediated immunity induced by stimulation with TLR3 and TLR7 agonists is better

achieved by a local rather than a systemic administration. Importantly, the systemic and repeated administration of TLR7 ligands has been shown to induce tolerance in *in vivo* studies (Bourquin et al., 2011; Gunzer et al., 2005; Hayashi et al., 2009). Moreover, the above published reports investigated the therapeutic effects of effector CD8 T cells rather than differentiated memory T cells (Currie et al., 2008; Drobits et al., 2012; Ichinohe et al., 2005; Pulko et al., 2009; Salem et al., 2005; J. Wu et al., 2017), thus, information about these adjuvants for the generation of potent CD8 T cell memory responses, particularly liver T<sub>rm</sub> cells, is missing from the published data.

Activation of TLR4 enhances Th1 immune responses and accordingly, ligands of TLR4 have been exploited as potential adjuvants (Steinhagen et al., 2011). However, not all TLR4 agonists can be used as adjuvants. In fact, LPS, a well-known TLR4 agonist, induce excessive activation and can generate tolerance and endotoxin shock (Biswas and Lopez-Collazo, 2009). Therefore, alternative TLR4 ligands have been designed to avoid toxicity and tolerance while keeping the benefits of TLR4 activation.

Monophosphoryl lipid A (MPLA) is a detoxified derivative of LPS isolated from *Salmonella minnesota* that activates TLR4 (Baldrige et al., 2004; Casella and Mitchell, 2008; Ribi et al., 1979). Inspection of the literature indicates in most cases MPLA, and not LPS, has been used in combination with other adjuvants in animal vaccination studies and clinical trials. For example, MacLeod and colleagues (2011) conducted an influenza vaccine study in which mice that received antigen plus MPL together with alum (a well-known adjuvant in human vaccines) generated higher CTL differentiation and long-lasting memory CD8 T cells, as compared with mice that received antigen plus only one of the two adjuvants. In the same study, mice vaccinated with influenza antigen with MPL plus alum developed higher levels of protection against challenge as compared to mice vaccinated with antigen plus only one adjuvant (MacLeod et al., 2011). These data indicate that signals from both adjuvants (MPL and alum) are required to enhance vaccine efficacy and drive potent CD8 T cell responses. To date, at

least two adjuvants based on TLR4 ligands (AS01 and AS04) are used in human licensed vaccines against virus, including herpes zoster, hepatitis B and human papillomavirus (Del Giudice et al., 2018). Both AS01 and AS04 are composed of MPL plus a second adjuvant, saponin QS-21 and alum hydroxide, respectively (Didierlaurent et al., 2017, 2009).

As mentioned, data generated in this thesis revealed the poor adjuvant capacity of LPS to induce liver T<sub>RM</sub> cell formation mediated by Clec9A targeted immunisation (Chapter 4). Future studies investigating the adjuvant effects on T<sub>RM</sub> cells would benefit from exploring MPL instead of LPS together with other adjuvants such alum salts and the saponin SQ-21.

Another ligand that showed a poor adjuvant capacity for the induction of liver memory CD8 T cells in experiments performed in this thesis was the cGAS ligand (Chapter 4). Upon activation of cGAS by cytoplasmic DNA, it produces a messenger molecule, known as cGAMP, which in turn activates STING that ultimately leads to the production of pro-inflammatory cytokines and type I IFN (Ablasser et al., 2013; Burdette et al., 2011; Diner et al., 2013; Sun et al., 2013; Tanaka and Chen, 2012). STING agonists have been extensively investigated in murine models as vaccine adjuvants against bacterial and viral infections and tumour malignancies (Blaauboer et al., 2015; Corrales et al., 2015; Ebensen et al., 2007; Karaolis et al., 2007b; Ogunniyi et al., 2008; Wang et al., 2016). Interestingly, published data suggests that WT murine hepatocytes express undetectable levels of STING and cGAS when compared to spleenocytes (Thomsen et al., 2016). This has important implications for hepatotropic infections with DNA viruses including hepatitis B virus (HBV) and cytomegalovirus (CMV). In fact, two recent studies have shown that ablation of STING or cGAS has no effect on the *in vivo* immune response to HBV and MCMV as compared to WT mice (Tegtmeyer et al., 2019; Thomsen et al., 2016). Importantly, upon induction of interferon stimulated genes by poly I:C (TLR3 ligand) or by forcing the expression of STING in hepatocytes, the STING deficient mice gained the ability to reduce the HBV viral load after infection (Thomsen et al., 2016). These findings

suggest that cGAS or STING ligands are probably poor inducers of inflammation mediated by hepatocytes because of the low levels of expression in these cells. In this sense, combining cGAS ligands with other adjuvants might provide signals important for CD8 T cell memory formation in the liver.

RIG-I was first discovered as a host sensor for viral RNA and upon activation leads to type I IFN response (Rothenfusser et al., 2005; Yoneyama et al., 2004). Since then, RIG-I ligands have been effectively used in animal models to induce antiviral and antitumoral responses (Chiang et al., 2015; Goulet et al., 2013; Kasumba and Grandvaux, 2019; Linehan et al., 2018; Y. Wu et al., 2017; Yong and Luo, 2018). Moreover, two studies have confirmed the use of RIG-I agonists as adjuvant in murine vaccines against infections with influenza virus and an adenovirus expressing OVA (Hochheiser et al., 2016; Martínez-Gil et al., 2013). This last study supports the idea that RIG-I agonists could be exploited as adjuvants for hepatotropic infections, like hepatitis and malaria liver-stage. However, in this thesis, the RIG-I ligand had a limited effect on the induction of liver T<sub>RM</sub> cells. Importantly, Hochheiser and colleagues used a transfection reagent to complexed the adjuvant together with the antigen, aiming for a simultaneous delivery of adjuvant and antigen to the same APC (Hochheiser et al., 2016). For the experiments described in this thesis, Clec9A mAb coupled to the antigen was co-administered in the same injection as the RIG-I ligand, but this does not ensure simultaneous delivery to the same cell. This factor may explain why RIG-I ligand and other agonists failed to induce an adequate initial priming of the CD8 T cells.

### 5.6 CpG favours liver T<sub>RM</sub> cell formation

Data generated in this thesis demonstrate the superior capacity of CpG (compared to the other tested adjuvants) to promote liver T<sub>RM</sub> cell formation mediated by Clec9A targeted immunisation. As mentioned before, the superiority of CpG may be related to the capacity of CpG to generate intrahepatic myeloid cell aggregates that support proliferation of CD8 T cells in the liver (Huang et al.,



2013). Remarkably, the transfection reagent DOTAP enhanced the formation of liver T<sub>RM</sub> cells mediated by different types of CpG (Chapter 4). DOTAP is a cationic lipid that mediates intracellular delivery and enhances DC activation (Honda et al., 2005; Yasuda et al., 2005; Yotsumoto et al., 2008).

In the current study, two B-class CpG and a special CpG, named CpG-combo (a B-class linked to a P-class CpG), were assessed for the induction of liver T<sub>RM</sub> cell formation. B-class CpG are known to promote pDC differentiation and the production of TNF- $\alpha$  (Hartmann and Krieg, 2000; Hemmi et al., 2003; Rothenfusser et al., 2001; Verthelyi et al., 2001), which in turn leads to T cell activation. P-class CpG, on the other hand, is known to promote a strong production of IFN- $\alpha$ , IL-6 and the CXCR3 ligand CXCL10 *in vivo* (Samulowitz et al., 2010).

Results from chapter 4 showed that mice receiving CpG-combo developed much higher numbers of liver T<sub>RM</sub> cells compared to mice that received B-class CpG. However, when B-class CpG were complexed in DOTAP they induced as many T<sub>RM</sub> cells as CpG-combo, regardless of whether or not CpG-combo was also complexed in DOTAP. Structural differences between the two CpG types might explain this finding. B-class CpG are composed by single strand structures, which are susceptible to degradation by DNases present in sera (Hanagata, 2012). On the contrary, the P-class ODN, present in the CpG-combo, contains palindromic sequences that enable the formation of complex structures (Samulowitz et al., 2010), protecting it from extracellular degradation. It is possible that B-class CpG and DOTAP complexes are more stable than free B-class CpG and therefore more effective at inducing DC activation and/or inflammation in the liver. These complexes are less prone to extracellular degradation (Capaccioli et al., 1993) and as they are positively charged, they could be easily drawn-in by the negatively charged cellular membrane (Chesnoy and Huang, 2000). Existing evidence from vaccination studies suggest that transfection reagents based on liposomes enhance the CpG-mediated CD8 T cells responses. In fact, the encapsulation of relevant antigen together with CpG

in liposomes, increases CTL responses *in vivo* for the OVA antigen model and tumour models for breast cancer and melanoma (Chikh et al., 2009; Jérôme et al., 2006; Li et al., 2003).

### 5.7 Too much of a good thing?

Pre-erythrocytic malaria vaccines based on the administration of whole sporozoites have demonstrated high efficacy against infection in mice and humans. From these approaches, immunisation with radiation-attenuated sporozoites (RAS) is the most clinically tested strategy, and it has been demonstrated to induce sterile protection against sporozoite challenge in animal models and humans (Clyde, 1990; Gwadz et al., 1979; Ishizuka et al., 2016; Nussenzweig et al., 1967; Seder et al., 2013). Recently, murine studies showed that vaccination with RAS induces liver T<sub>RM</sub> cells providing protection against sporozoite challenge (Epstein et al., 2011; Fernandez-Ruiz et al., 2016).

Given the superior capacity of CpG-combo to favour the formation of liver T<sub>RM</sub> cells mediated by Clec9A immunisation (chapter 4), this compound was assessed as an adjuvant for RAS vaccination. Surprisingly, CpG-combo had no enhancing effect on the generation of liver T<sub>RM</sub> cells nor on the protection induced by RAS. Additionally, preliminary results indicate that mice receiving RAS together with CpG-combo complexed in the transfection reagent DOTAP generated less T<sub>RM</sub> cells than mice immunised with RAS alone and therefore failed to control infection (chapter 4). These findings question the capacity of CpG-combo to adjuvant RAS vaccination, but it will be important to repeat these last experiments to confirm our findings.

Data from chapter 4 also indicate that CpG not only failed to enhance the formation of memory cells, it also failed to promote the early expansion of circulating CD8 T cells 7 days after priming with RAS. Similarly, published data show that CpG does not enhance early expansion of malaria transgenic-CD8 T cells (i.e. PbT-I cells) mediated by RAS immunisation (Lau et al., 2014). Notably,

preliminary data suggest that mice vaccinated with RAS and CpG in DOTAP had fewer specific cells in blood at day 7 post priming than mice receiving only RAS (chapter 4). Altogether, these findings indicate that administration of CpG, in particular CpG in DOTAP, reduces not only the initial CD8 T cell expansion and the later formation of liver memory cells, but also the protection against sporozoite challenge normally induced by immunisation with RAS.

It is known that insufficient early expansion of CD8 T cells may result from an inadequate priming process. For instance, it has been recently reported that suboptimal protection induced by intradermal RAS vaccination results from insufficient early CD8 T cell activation, which in turn is a consequence of a suboptimal priming by cDC1 (Haeberlein et al., 2017). In light of the results from chapter 4, it is possible that the administration of CpG (in the presence or absence of DOTAP) somehow alters the CD8 T cell priming induced by RAS.

Early studies have shown that immunisation with non-viable sporozoites, e.g. heat-killed and freeze-thawed parasites, provide much less protection against sporozoite challenge as compared to RAS (Alger and Harant, 1976; Hafalla et al., 2006). This indicates that certain characteristics of viable parasites, in this case of RAS, are important for the induction of effective protection. Indeed, the protective immunity induced by RAS is abrogated by treatment of hosts with primaquine to eliminate the remaining liver parasites (Klotz et al., 1995; Scheller and Azad, 1995). It is possible that if attenuated parasites fail to establish a hepatocyte infection, the probability of antigen acquisition and presentation reduces and with that the chances to prime T cell responses.

We and others have observed that CpG alone induces a short (1-2 days) non-specific-protection against sporozoite infection (Gramzinski et al., 2002, unpublished data Heath lab). Importantly, these authors have also shown that this effect mediated by CpG is abrogated by treatment with monoclonal antibodies against IL-12 or IFN- $\gamma$  (Gramzinski et al., 2002). In line with these findings, early studies in mice and primates have demonstrated that pre-

treatment (1-2 days prior infection) with recombinant IFN- $\gamma$  or IL-12 inhibits the development of liver-stage infection by both human and mouse *Plasmodium* species (Ferreira et al., 1986; Hoffman et al., 1997; Sedegah et al., 1994). Additionally, the administration of IFN- $\gamma$  induces the production of nitric oxide and reactive nitrogen intermediates inside hepatocytes, which in turn mediates the elimination of intracellular parasites (Klotz et al., 1995; Nahrevanian and Dascombe, 2001).

Shortly after systemic administration, CpG complexed in DOTAP induces significantly higher amounts of IL-12 and IFN- $\gamma$  than CpG alone (Suzuki et al., 2004; Yotsumoto et al., 2008). Based on these previous studies describing the role of IL-12 and IFN- $\gamma$  on the elimination of intra-hepatic parasites, we hypothesize that CpG in DOTAP induces “too much of a good thing” and hinders the protective immunity induced by RAS. Hypothetically, high levels of IL-12 and IFN- $\gamma$  induced by the systemic administration of CpG in DOTAP contribute to the early elimination of intrahepatic RAS parasites, reducing the availability of liver-stage antigens for the priming and activation of CD8 T cells. Ultimately, this results in a reduction of liver T<sub>RM</sub> cells and therefore, a loss of protection against sporozoite infection. Further experiments are required to unveil the mechanisms that govern this effect. For example, if the adjuvant CpG is administered days later after RAS vaccination, sporozoites may have more time to infect hepatocytes to ensure antigen presentation and liver T<sub>RM</sub> cell formation.

### 5.8 The translational potential of ‘prime-and-trap’

Prime-and-trap vaccination successfully induces the formation of liver T<sub>RM</sub> cells that protect against sporozoite challenge in B6 mice transferred with *Plasmodium*-specific transgenic cells (Fernandez-Ruiz et al., 2016), or in mice with a normal endogenous T cell repertoire (chapter 3) (Valencia-Hernandez et al., 2020). Ideally, this vaccine could be engineered for human use, but prime-and-trap is a complex 3-component vaccine that currently faces some challenges before translation is actually feasible. These complications are not necessarily

insurmountable and in principle a vaccine like prime-and-trap can be produced for clinical trials.

The first component of the vaccine consists of a monoclonal antibody anti-Clec9A (a cDC1 receptor) coupled to a highly immunogenic liver-stage epitope (Fernandez-Ruiz et al., 2016; Valencia-Hernandez et al., 2020), designed to achieve epitope delivery to cDC1 (Caminschi et al., 2008; Lahoud et al., 2011). Antigen targeting using anti-Clec9A antibodies together with adjuvants has shown to be well tolerated and to induce effective humoral responses in non-human primates (Li et al., 2015). To date, no clinical studies have demonstrated the safety of anti-Clec9A antibodies administration. However, the first human studies using antibodies targeting another DC receptor, the DEC-205 (also expressed by cDC1), coupled to a tumour antigen, have demonstrated the feasibility, safety and biological activity of this vaccine (Dhodapkar et al., 2014; Griffiths et al., 2018). Hence, in principle, protein DC-targeted vaccines are safe for use in humans, as well in non-human primates. Moreover, current advances in technologies for large-scale production of plant-derived antibodies for immunotherapy (Buyel et al., 2017), can lower the cost of production increasing feasibility.

The second component is a recombinant adeno-associated virus (rAAV) engineered to selectively infect hepatocytes (Tay et al., 2014), which expresses the targeted antigen under the control of a hepatocyte-specific promoter, to induce the trapping of the T cells to the liver (Fernandez-Ruiz et al., 2016; Valencia-Hernandez et al., 2020). Clinical trials based on adenovirus vector-vaccines expressing antigens against several infectious diseases including malaria, HIV, HCV and Ebola have increased rapidly during the last decade (Ewer et al., 2016; Lauer et al., 2017; Naso et al., 2017; Rauch et al., 2018). These studies not only demonstrate the safety of this approach but also report promising vaccine efficacies (Hodgson et al., 2015; Mensah et al., 2016; Ogwang et al., 2015; Tiono et al., 2018). Currently, human malaria vaccine studies use a chimpanzee adenovirus (ChAd63) and a modified vaccinia virus (MVA) to

express different *P. falciparum* peptides (Ewer et al., 2016). In mice, intravenous administration of ChAd63 or MVA induces liver-tropism and hepatic expression of antigen, facilitating the formation of liver T<sub>RM</sub> cells (Gola et al., 2018). Given the existing clinical data for ChAd64 administration, it will be beneficial to design a ChAd63 viral vector expressing the RPL6 epitope and use it for the prime-and-trap vaccine instead of the original adeno-associated virus.

The third and last component is a CpG adjuvant used for the licensing of DC involve in CD8 T cell priming (Lahoud et al., 2011) and to induce liver inflammation (Huang et al., 2013), known to support T<sub>RM</sub> cell formation (Holz et al., 2018; Mackay et al., 2012a). CpG has been used extensively in clinical trials as an adjuvant for vaccines against several infection diseases and tumours and was recently approved as part of a licensed vaccine against HBV (Food and Drug Administration, 2020; Del Giudice et al., 2018; Scheiermann and Klinman, 2014). This indicates that CpG is well tolerated and actually improves vaccine efficacy in humans. Tolerability to different types of CpG, in particular of “CpG-combo” and administration in liposomal transfection reagents, such as DOTAP, are yet to be determined.

In summary, these clinical studies indicate that the development of the prime-and-trap components for human use is feasible in the near future. Some components could be changed, for example, instead of using anti-Clec9A to target DC, anti-DEC205 antibodies could be used, given their ability to induce potent CD8 T cell responses (Lahoud et al., 2011) and safety administration in humans (Dhodapkar et al., 2014; Griffiths et al., 2018). To induce the trapping of primed T cells to the liver, the ChAd63 viral vector could be used instead of adeno-associated virus, since the former has been tested in clinical trials and it is known to favour the formation of liver T<sub>RM</sub> cells (Gola et al., 2018; Hodgson et al., 2015; Mensah et al., 2016). Despite the need to simplify and modify the actual components, the main idea behind this vaccination strategy, to achieve formation of liver T<sub>RM</sub> cells, should remain.

Currently, all our experimental data for the prime-and-trap vaccine are based on intravenous administration in mice and no other application routes have been investigated (Fernandez-Ruiz et al., 2016; Valencia-Hernandez et al., 2020). Conventional vaccination uses oral and parental administration, which includes intradermal, intramuscular and subcutaneous delivery (WHO, 2020; Zhang et al., 2015). In spite of the reluctance to employ intravenous vaccines, recent work suggests they are safe and immunogenic. A recent study in macaques showed that intravenous vaccination with BCG increased the number of T cells in lung expressing T<sub>RM</sub> markers and that these animals developed higher levels of protection against *Mycobacterium tuberculosis* as compared to macaques that received the BCG intradermally (Darrah et al., 2020). RAS immunisation, which protects against malaria, is also delivered through intravenous injections in humans (Seder et al., 2013). Importantly, a small phase I clinical trial revealed that intravenous injection with a viral vector expressing malaria epitopes (ChAd63-ME-TRAP) caused only moderate adverse events and induced immunogenicity measured by *ex vivo* assays (Gola et al., 2018). Altogether, these studies point to promising translational prospects and an increase feasibility of the prime-and-trap vaccine and similar strategies.

### 5.9 T<sub>RM</sub>-based vaccines for humans

An important limitation of the approaches favouring the generation of T<sub>RM</sub> cells is a lack of tools to monitor T<sub>RM</sub> cell formation. The identification of bona fide human T<sub>RM</sub> cells is controversial because gold standard tests (e.g. parabiosis experiments and intravascular cell labelling) cannot be performed in humans. Currently, clinical trials assess vaccine immunogenicity of T cell responses through cytokine production assays performed on circulating cells and hence, the local immune response in the tissue remains unknown. Data presented in this thesis suggests that, for most cases, the initial expansion of specific CD8 T cells measured in blood at day 7 post immunisation reflected the numbers of liver T<sub>RM</sub> cells at a later time point. However, this did not hold true for all treatments e.g. for mice receiving RIG-I-L as adjuvant. Therefore, there is an urgent need for the

identification of biomarkers in blood that could effectively predict the formation of T<sub>RM</sub> cells.

Efforts have been made to characterize human liver T<sub>RM</sub> cells, by several means including extensive studies of protein expression and gene transcriptional profiling. These studies have identified a unique CD8 T cell subpopulation in the liver, absent in PBMC, of healthy and HBV infected humans that matched core signature features of T<sub>RM</sub> cells (Pallett et al., 2017; Swadling et al., 2020). These recent studies realized there are important differences in how human and murine liver T<sub>RM</sub> cells are formed and maintained. For example, while murine liver T<sub>RM</sub> cells lack the expression of CD103 (Fernandez-Ruiz et al., 2016; McNamara et al., 2017), an integrin expressed by T<sub>RM</sub> cells located in epithelial and neuronal tissues (Mueller and Mackay, 2016), this molecule is apparently found in the liver T<sub>RM</sub> cells of healthy and HBV infected humans (Pallett et al., 2017). At the transcriptional level, whereas Blimp-1 and Hobit regulate the maintenance of murine liver T<sub>RM</sub> cells (Mackay et al., 2016), it has been suggested that human T<sub>RM</sub> cells express low levels of Blimp-1 and Hobit (Pallett et al., 2017). Functionally, liver T<sub>RM</sub> cells have been found to express high and low levels of granzyme B in mice and humans, respectively (Fernandez-Ruiz et al., 2016; Pallett et al., 2017). There are also shared features between the human and murine liver T<sub>RM</sub> cells. For instance, for both cases, liver T<sub>RM</sub> cells express high levels of CD69, CXCR6, CXCR3 and require IL-15 exposure (Fernandez-Ruiz et al., 2016; Holz et al., 2018; Pallett et al., 2017). Additionally, data presented in this thesis support previous findings suggesting that the generation of large numbers of cells is boosted by hepatocyte antigen presentation, even though local antigen exposure is not required for the formation of liver T<sub>RM</sub> cells in mice and humans (Fernandez-Ruiz et al., 2016; Holz et al., 2018; Pallett et al., 2017).

Differences similar to those described between human and mice T<sub>RM</sub> cells in the liver (see above) have also been found for other tissues, including the lung, intestine and skin (Kumar et al., 2017; Szabo et al., 2019). This situation raises concerns about the translational potential of inbred mouse models to investigate



human T<sub>RM</sub> cell responses. Therefore, there is a necessity for developing models that better recapitulate the human immunology within tissues. It has been recently proposed that outbred mice from pet stores, which have been exposed to pathogens and have an enriched microbiota compared to the “clean” lab mice, have T<sub>RM</sub> cells that more closely resemble those from humans (Beura et al., 2018, 2016; Kumar et al., 2017). Extensive investigation on “dirty” mice is still required to determine their translational potential as compared to inbred mice.

In the context of vaccines for infectious diseases, several mouse studies have demonstrated that protective T<sub>RM</sub> responses can be generated through different immunisation strategies. For instance, administration of live-attenuated pathogens locally or intravenously in mice generates T<sub>RM</sub> cells that mediate protection against influenza (Zens et al., 2016), tuberculosis (Perdomo et al., 2016) and malaria (Fernandez-Ruiz et al., 2016; Tse et al., 2013). Intravaginal administration of viral vectors to deliver targeted antigens to the vaginal mucosa of mice successfully induces protective T<sub>RM</sub> cells against HIV (Tan et al., 2018) and genital herpes virus 2 (Çuburu et al., 2014). Another particularly successful vaccination strategy for the induction of protective T<sub>RM</sub> cells is known as “prime and pull”. Those vaccines and their variations incorporate a priming step, with a relevant antigen, together with a pull or target step for the recruitment of T<sub>RM</sub> precursors to a specific tissue. With this approach, protective T<sub>RM</sub> responses have been proven against HSV-2 challenge (Bernstein et al., 2019; Shin et al., 2016; Shin and Iwasaki, 2012) and *Plasmodium* sporozoite infection (Fernandez-Ruiz et al., 2016; Gola et al., 2018; Valencia-Hernandez et al., 2020). Altogether, these studies provide evidence of the potential of T<sub>RM</sub>-based vaccines to protect against infections. Whether or not these strategies can be applied to humans will likely be determined in the coming years.

#### 5.10 Concluding remarks

The data presented in this thesis demonstrated the highly immunogenic nature of an epitope within the RPL6 protein of PbA and its ability to evoke high

levels of sterile protection against liver stage infection in vaccinated mice with a normal endogenous T cell repertoire. Additionally, the results showed the superior capacity of CpG ODN to promote liver T<sub>RM</sub> cell formation in mice vaccinated with antigen targeted immunisation. These findings provide insights into elements that favour the generation of protective liver T<sub>RM</sub> cells; information that can be used for the design of T<sub>RM</sub> cell-based subunit vaccines against *Plasmodium* infection. While the 'prime-and-trap' vaccine currently faces challenges, they are not insurmountable and cutting-edge scientific and technological developments are increasing the translational prospects of T<sub>RM</sub>-based vaccines.

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## CURRICULUM VITAE

### Key experience

- Experience with mouse *in vivo* experiments: Equivalent course on animal welfare FELASA B1. During my PhD I have been performing mouse *in vivo* experiments on C57BL/6 mice, especially intravenous injections and monitoring of infected mice with experimental cerebral malaria. Processing of livers, lymph nodes, blood and spleens for the generation of single cell solutions and flow cytometry analysis.
- Cellular biology: Cell culture in Bio Safety Level 2; flow cytometry for cell death analysis; immunofluorescence microscopy; electron microscopy; cell culture of cells in suspension; growth curves; transfections in *Trypanosoma brucei* and cell culture of *Plasmodium falciparum* in human erythrocytes.
- Molecular biology: Development of CRISPR-Cas system in a model parasite, chromatin immuno-precipitation (ChIP), phage display assays, preparation of libraries for next generation sequencing, qPCRs, western blots, SDS-PAGE, cloning, PCR, plasmid and genomic DNA isolation, RNA isolation, synthesis of cDNA, ligation, transformation.
- Training in software such as “FlowJo” for data analysis of flow cytometry, “Graphpad Prism 5” for statistical analysis of data and “CLC Workbench” for DNA, RNA, and protein sequence data analysis.
- Analytical skills for problem solving, objectivity, leadership, attention to detail and ability to work in team.

### Formal Education

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|----------------|--|
| 2016 - Present | <p>PhD student of joint program in Immunology<br/>The University of Melbourne, Australia &amp; University of Bonn, Germany.<br/>Thesis title: “Characterization of the memory response induce by immunisation with a novel antigen and adjuvants for a dendritic cell-targeted malaria vaccine”.<br/>Supervisor (University of Melbourne): Prof Dr William R Heath<br/>Supervisor (University of Bonn): Dr. rer. Nat. Winfried Barchet<br/>Scholarship from: University of Melbourne</p> |
| 2013 - 2015    | <p>MSc FOKUS Life Sciences.<br/>University Wuerzburg, Germany<br/>Thesis title: “Role of histone acetylation and bromodomain factors in the targeted deposition of H2AZ in <i>Trypanosoma brucei</i>”.<br/>Supervisor: Prof Dr Nicolai Siegel</p>  |
| 2007 – 2011    | <p>BsC in Microbiology<br/>Universidad de los Andes, Bogotá, Colombia.<br/>Thesis title: “Importance of the cell death type in the acquisition of adaptive immunity against <i>Mycobacterium Tuberculosis</i>”.<br/>Supervisor: Prof. Dr. Carlos Alberto Parra Lopez</p>   |

### Complementary Education

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|------|---|
| 2016 | DGFI Autumn School, Current Concepts in Immunology (Hours: 40h) |
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## Professional Experience

- 2016 – 2019      University Hospital Bonn (Germany)  
Institute for Clinical Chemistry and Clinical Pharmacology.  
Scientific employee, Position: PhD Student, Working hours: 65% (to full time).
- Coordination of a research collaborative project in Malaria vaccination with the participation of different institutes of the University Hospital Bonn, University Hospital Heidelberg in Germany and The Doherty Institute, Melbourne.
  - Investigating the role of nucleic acids as adjuvants for a malaria vaccine against liver stage using the mouse model of C57BL/6 and *Plasmodium berghei*.
- 2015 - 2016      Fraunhofer IME, Aachen (Germany)  
Division of Pharmaceutical product development,  
Scientific employee, Position: Assistant research scientist (HIWI)
- Performing phage display assays for a phage library containing Fab fragments of antibodies specific against *P. falciparum*.
- 2013 – 2014      University Wuerzburg (Germany)  
Institute of Molecular Infection Biology (IMIB)  
Scientific employee, Position: Assistant research scientist (HIWI)
- Studying the presence of proteins involved in autophagy processes during infection of murine macrophages with *L. major*.
  - Embedding electron microscope images of murine infected-dendritic cells with promastigotes of *L. major*.
- 2012 – 2013      University Wuerzburg (Germany)  
Institute of Molecular Infection Biology (IMIB)  
Scientific employee, Position: Assistant research scientist (HIWI)
- Investigating the mechanisms of transcription termination in *T. brucei*.

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Presentations in Events

1. Valencia-Hernandez, AM., Menezes, MN., Lahoud, M., Cozijinzen, A., McFadden GI., La Gruta, N., Bertolino, P., Caminschi, I., Fernandez-Ruiz, D., Barchet, W., & Heath WR. Subunit vaccination targeting novel malaria liver-stage epitope confers protection in mice with natural endogenous repertoire. Oral presentation at the 48th Annual Scientific Meeting of the Australasian Society for Immunology (ASI), 2019.
2. Valencia-Hernandez, AM., Lahoud, M., Cozijinzen, McFadden GI., La Gruta, N., Caminschi, I., Fernandez-Ruiz, D., Barchet, W., & Heath WR. Effect of nucleic acids as adjuvants for the induction of *P. berghei*-specific liver tissue resident memory CD8 T cells. Poster presentation at the Malaria in Melbourne 2019 conference.
3. Semi-finalist 3 Minutes Thesis (3MT), "An army of Immune cells". The University of Melbourne, 2019.
4. Valencia-Hernandez, AM., Fernandez-Ruiz, D., Caminschi, I., Lahoud, M., Cozijinzen, A., Mollard, V., Barchet, W., & Heath WR. Effect of adjuvants on the induction

of protective CD8+ T cell immunity against liver stage malaria parasites. Oral presentation (Science communication session) at the 47th Annual Scientific Meeting of the Australasian Society for Immunology (ASI), 2018.

5. Valencia-Hernandez, Ana M., Fernandez-Ruiz, D., Müller, AK., Frank, R., Caminschi, I., Lahoud, M., Barchet, W. & Heath WR. Increasing Plasmodium based vaccine immunogenicity to induce protective CD8+ T cell immunity against Malaria liver stage. Poster presentation at the 1st Malaria World Congress, 2018.

6. Valencia-Hernandez, Ana M., Fernandez-Ruiz, D., Müller, AK., Frank, R., Caminschi, I., Lahoud, M., Barchet, W. & Heath WR. Increasing Plasmodium based vaccine immunogenicity to induce protective CD8+ T cell immunity against Malaria liver stage. Poster presentation at the 7th Australasian Vaccines & Immunotherapeutics Development Meeting, 2018.

7. Valencia-Hernandez, Ana M., Fernandez-Ruiz, D., Schumak, B., Müller, A., Hörauf, A., Heath W., Hartmann G. and Barchet, W. Increasing *Plasmodium* based vaccine immunogenicity to induce protective CD8+ T cell immunity against Malaria liver stage. Poster presentation at the Annual Meeting of German Center for Infection Research, 2016.

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#### Awards

Scholarship "Programa Quiero Estudiar" (100%) awarded by the Universidad de Los Andes (Colombia) to students with outstanding results in their applications, it covered the full tuition and other education related expenses along the 4 years of my undergraduate studies.

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#### Articles Published in Scientific Journals

1. Ghilas S., Valencia-Hernandez, A. M., Enders, M. H., Heath, W. R & Fernandez-Ruiz, D. (2020). Resident memory T cells and their role within the liver. *International Journal of Molecular Sciences*, 1-15. <https://doi.org/10.3390/ijms21228565>
2. Valencia-Hernandez, A. M., Ng, W. Y., Ghazanfari, N., Ghilas, S., de Menezes, M. N., Holz, L. E., Huang, C., English, K., Naung, M., Tan, P. S., Tullett, K. M., Steiner, T. M., Enders, M. H., Beattie, L., Chua, Y. C., Jones, C. M., Cozijnsen, A., Mollard, V., Cai, Y., ... Fernandez-Ruiz, D. (2020). A Natural Peptide Antigen within the Plasmodium Ribosomal Protein RPL6 Confers Liver TRM Cell-Mediated Immunity against Malaria in Mice. *Cell Host & Microbe*, 1-13. <https://doi.org/10.1016/j.chom.2020.04.010>
3. Holz, L. E., Chua, Y. C., de Menezes, M. N., Anderson, R. J., Draper, S. L., Compton, B. J., Chan, S. T. S., Mathew, J., Li, J., Kedzierski, L., Wang, Z., Beattie, L., Enders, M. H., Ghilas, S., May, R., Steiner, T. M., Lange, J., Fernandez-Ruiz, D., Valencia-Hernandez, A. M., ... Heath, W. R. (2020). Glycolipid-peptide vaccination induces liver-resident memory CD8+ T cells that protect against rodent malaria. *Science Immunology*, 5(48), 1-14. <https://doi.org/10.1126/sciimmunol.aaz8035>.

4. Masic, A., Valencia-Hernandez, A. M., Hazra, S., Glaser, J., Holzgrabe, U., Hazra, B., & Schurigt, U. (2015). Cinnamic acid bornyl ester derivatives from *valeriana wallichii* exhibit antileishmanial in vivo activity in leishmania major-infected BALB/c Mice. *PLoS ONE*, *10*(11), 1–20. <https://doi.org/10.1371/journal.pone.0142386>

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## References

Available on request