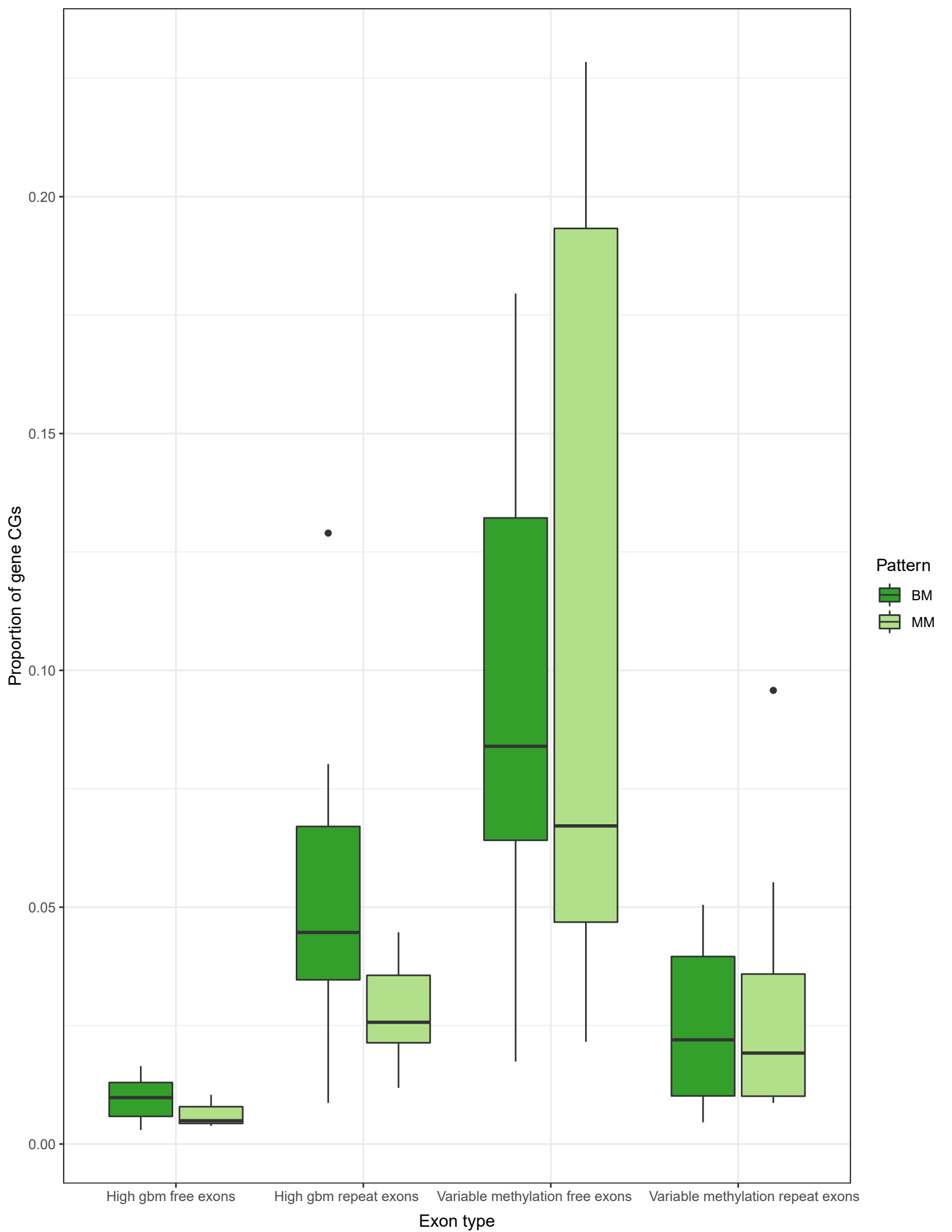
Proportion of gene CGs
0.20
0.15
0.10
0.05
0.00
High gbm free exons
High gbm repeat exons
Variable methylation free exons
Variable methylation repeat exons
Exon type
Pattern
BM
MM