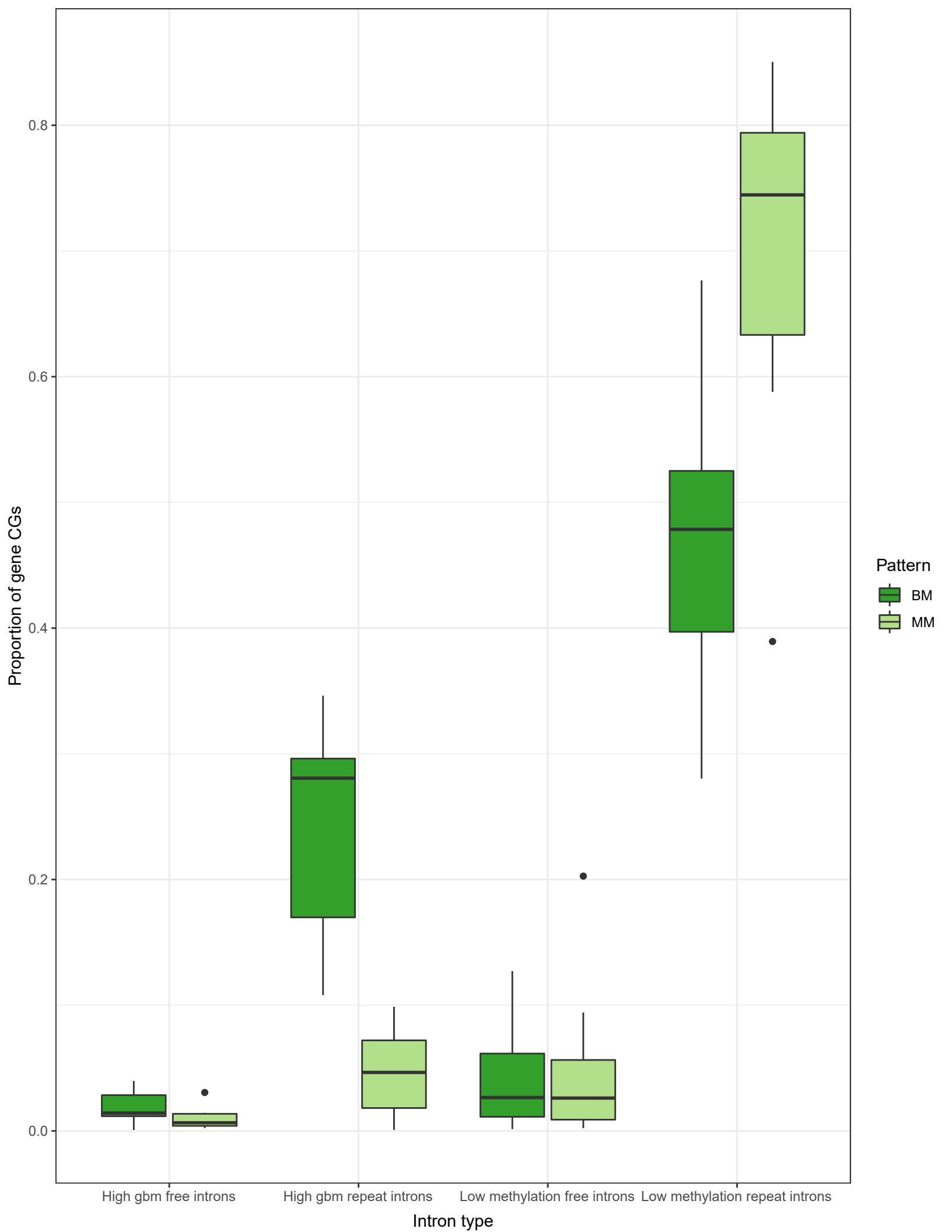
Proportion of gene CGs
0.0
0.2
0.4
0.6
0.8
High gbm free introns
High gbm repeat introns
Low methylation free introns
Low methylation repeat introns
Intron type
Pattern
BM
MM