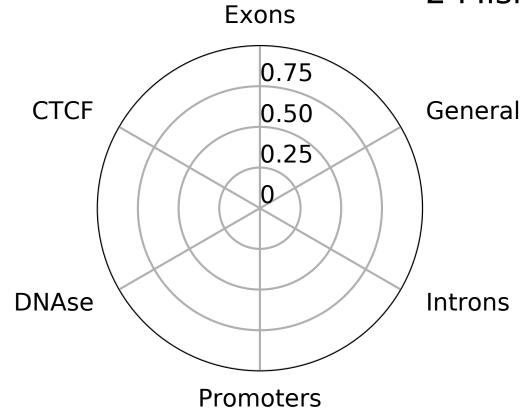
## 2 Mismatches



	Position	# Targets
General	0.0	0.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.0	0.0
DNAse	0.0	0.0
CTCF	0.0	0.0

D

R



1.0 -

0.8 -

0.6 -

0.4 -

0.2 -

0.0 T G G G T T T A G A G C C C A G A T