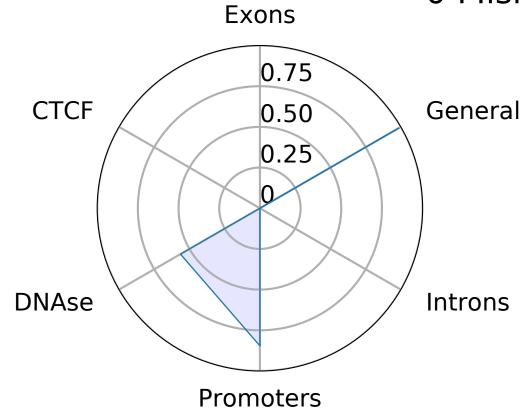
## 0 Mismatches



	Position	# Targets
General	0.98	7.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.85	1.0
DNAse	0.56	1.0
CTCF	0.0	0.0

D

R



1.0 -

0.8 -

0.6 -

0.4 -

0.2 -

O.0 C T C A C C A T T G C A C T C C A G C C