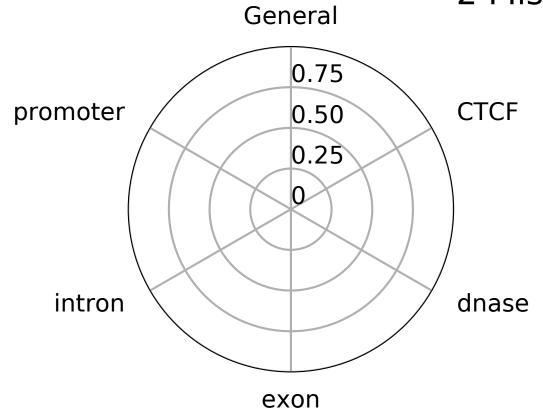
2 Mismatches



	Position	# Targets
General	0.0	0.0
CTCF	0.0	0.0
dnase	0.0	0.0
exon	0.0	0.0
intron	0.0	0.0
promoter	0.0	0.0



1.0 -

0.8 -

0.6 -

0.4 -

0.2 -

C G T D R