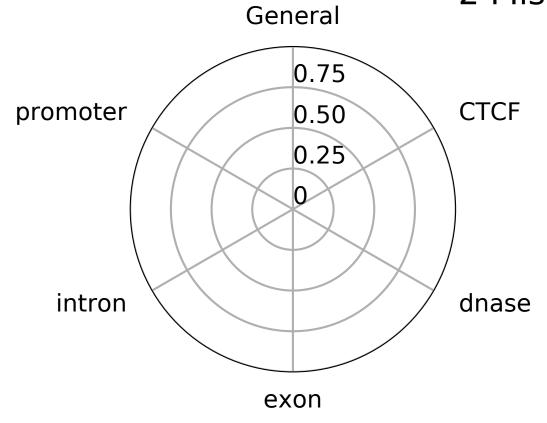
2 Mismatches



	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



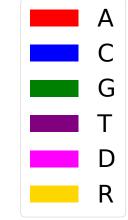
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



Α
Α
G
Α
Α
G
Α
Α
G
Α
С
G
Α
G
С
С
Т
G
Α
G
0.0