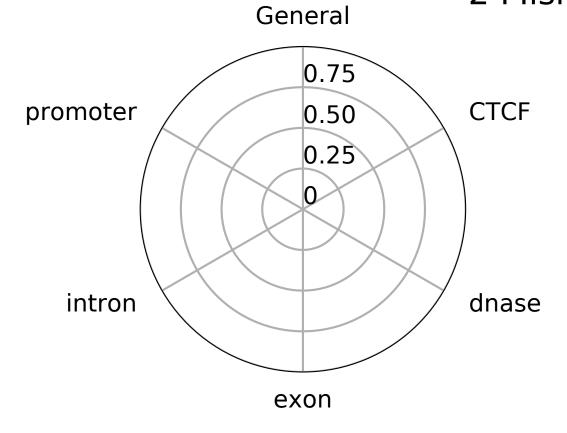
2 Mismatches



	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				



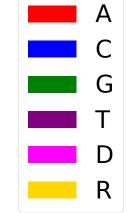
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



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0.0 -		Λ	T			T					T	T	T			
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