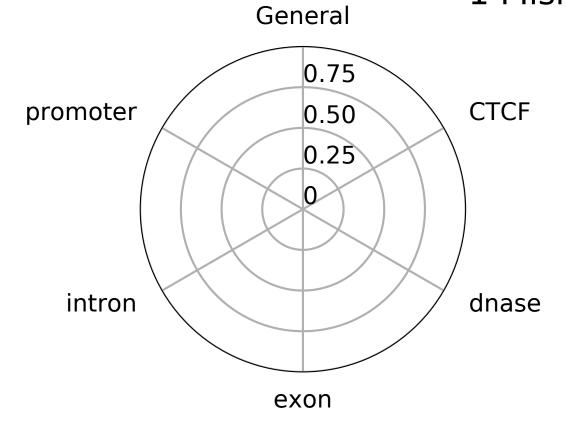
1 Mismatches



	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					



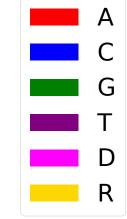
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



0.0^{-1}	G	Δ	G	T		G	Δ	G	Δ	G	Δ	Δ	G	Δ	Δ	G	Δ	Δ
			U	I		U		U		J			U			J		