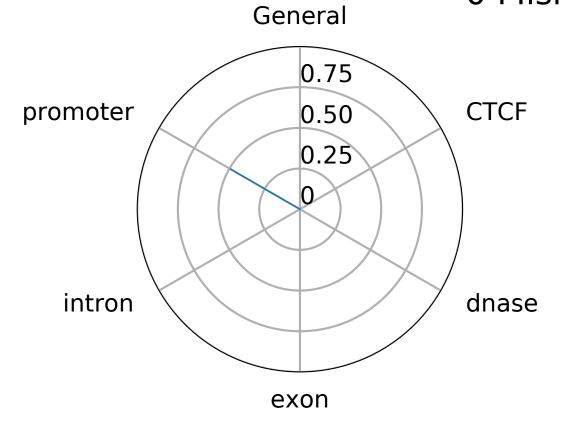
0 Mismatches



	Position	# Targets				
General	0.0	1.0				
CTCF	0.0	0.0				
dnase	0.0	0.0				
exon	0.0	0.0				
intron	0.0	1.0				
promoter	0.5	1.0				



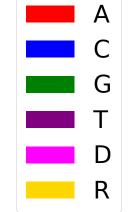
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



$\cap \cap$	_															
0.0 -		Δ	T	G	G	T	G	C		G	T	T	T	G		
		/ / \		_)			•		•			