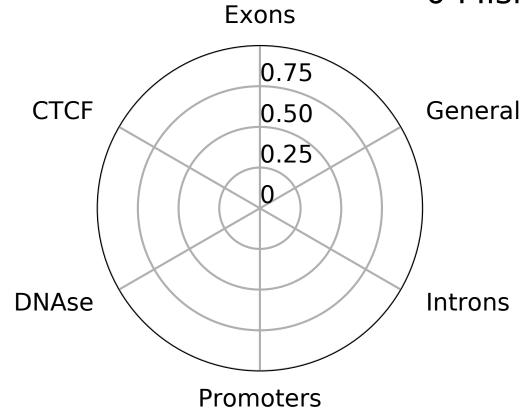
0 Mismatches



		Position	# Targets
I	General	0.0	0.0
	Exons	0.0	0.0
	Introns	0.0	0.0
	Promoters	0.0	0.0
	DNAse	0.0	0.0
	CTCF	0.0	0.0



1.0 -

0.8 -

0.6 -

0.4 -

0.2 -

