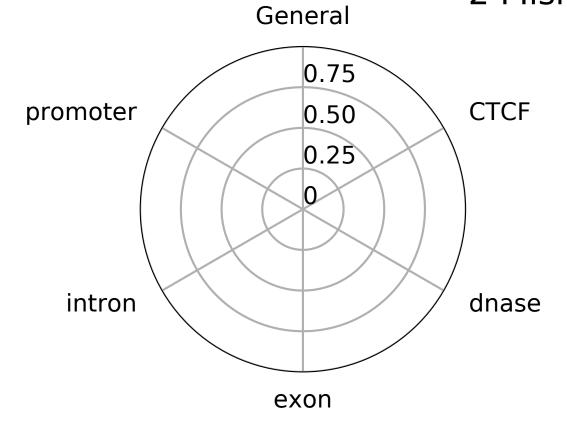
## 2 Mismatches



	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



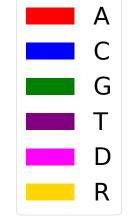
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



GCCGTTTG	G	T	G	G		С	A T C	CATC
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