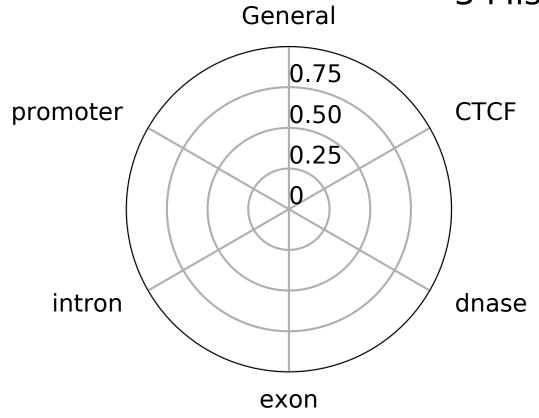
## 3 Mismatches



	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				



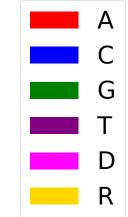
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



$\cap \cap$																				
0.0 -	G	Α	G	T	С	С	G	Α	G	C	Α	G	Α	Α	G	Α	Α	G	Α	Α