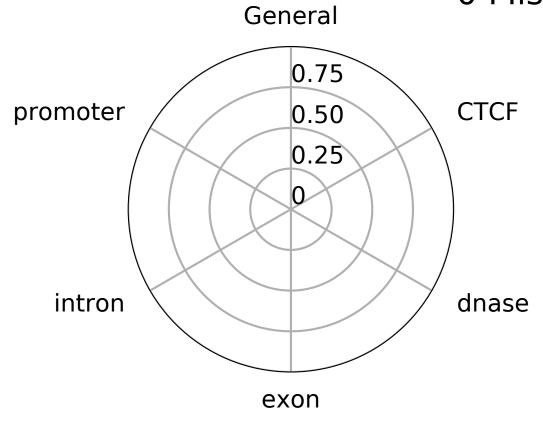
0 Mismatches



	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 -

G G T D R

\cap \cap																				
0.0	G	Α	G	T	С	С	G	Α	G	С	Α	G	Α	Α	G	Α	Α	G	Α	Α