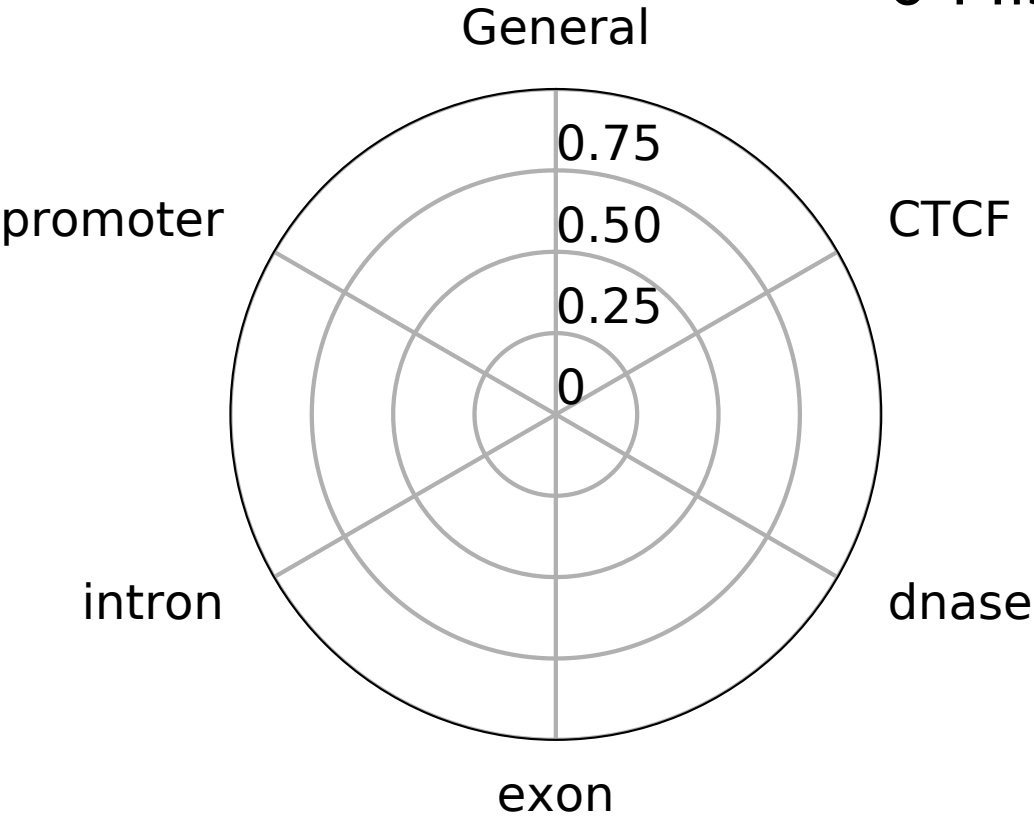
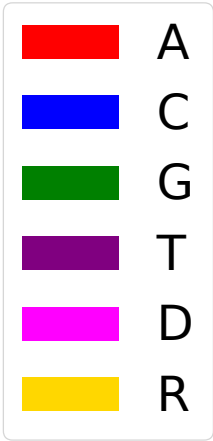


# 0 Mismatches



	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

1.2 -  
1.0 -  
0.8 -  
0.6 -  
0.4 -  
0.2 -  
0.0 -



G	A	G	T	C	C	G	A	G	C	A	G	A	A	G	A	A	G	A	A
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