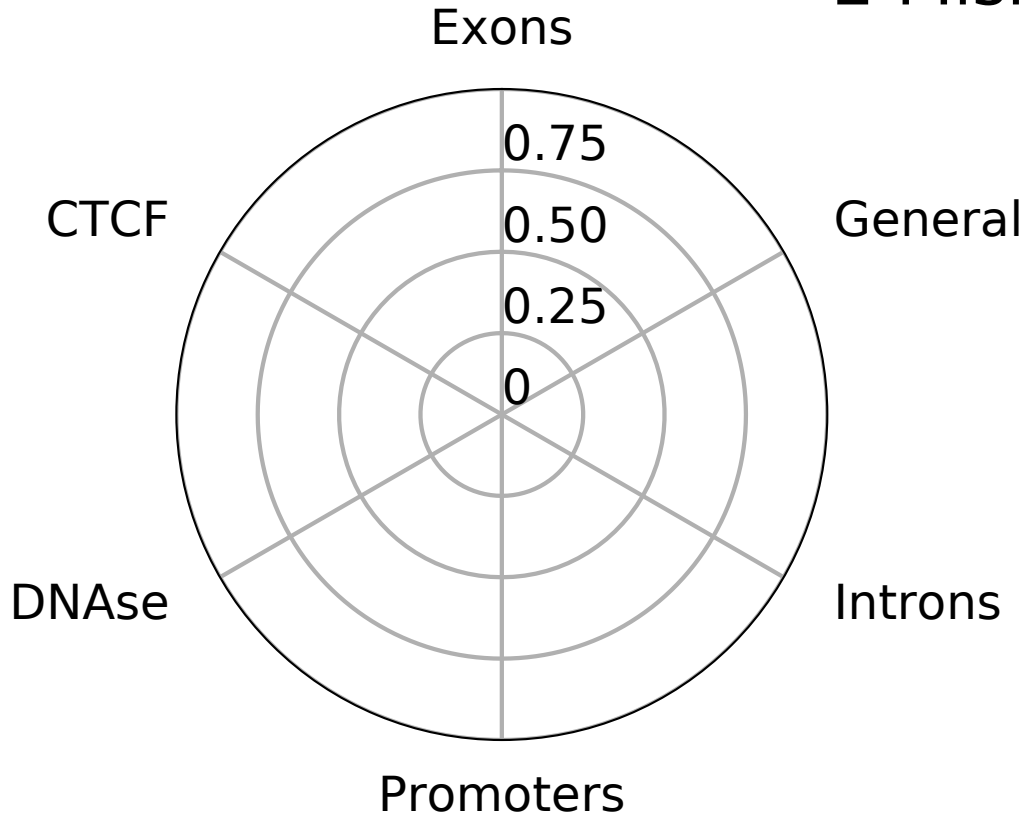
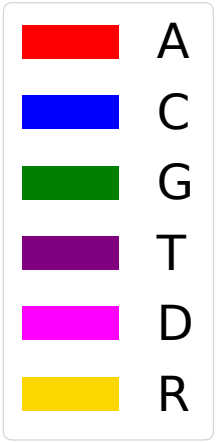


2 Mismatches



	Position	# Targets
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

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



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