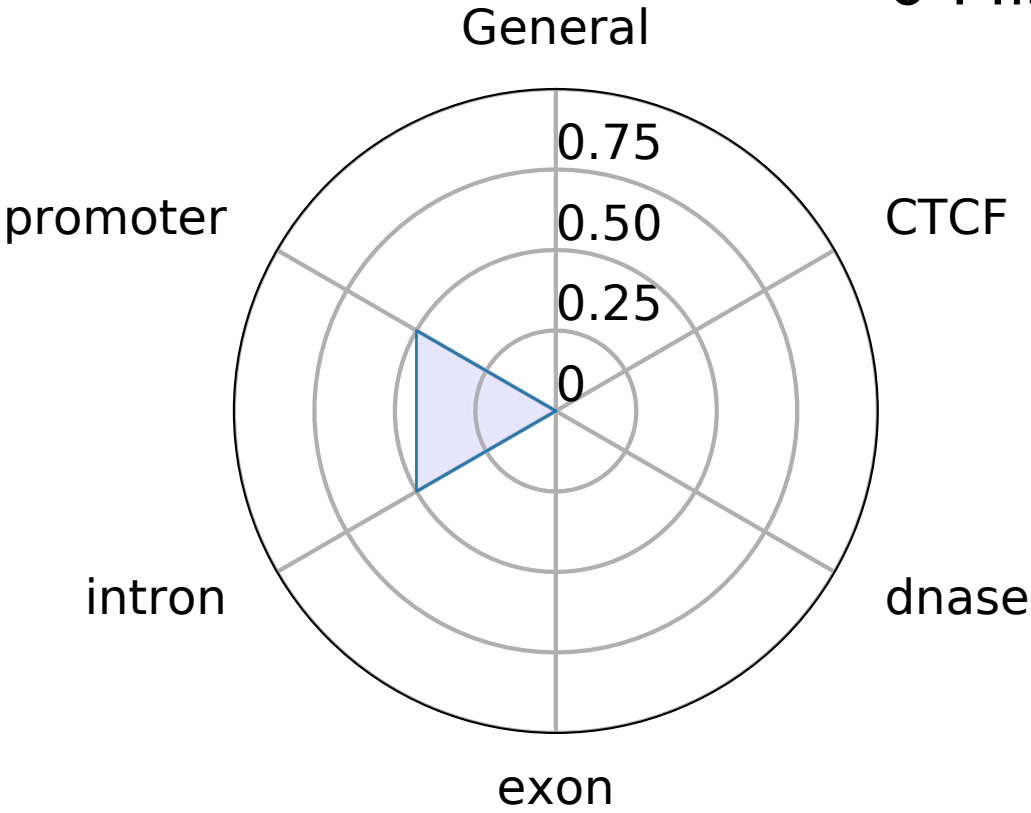
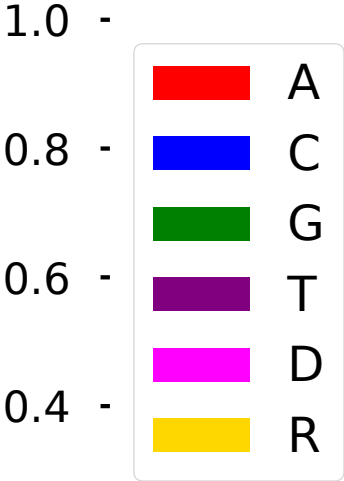


0 Mismatches



	Position	# Targets
General	0.0	1.0
CTCF	0.0	0.0
dnase	0.0	0.0
exon	0.0	0.0
intron	0.5	2.0
promoter	0.5	2.0



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

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

