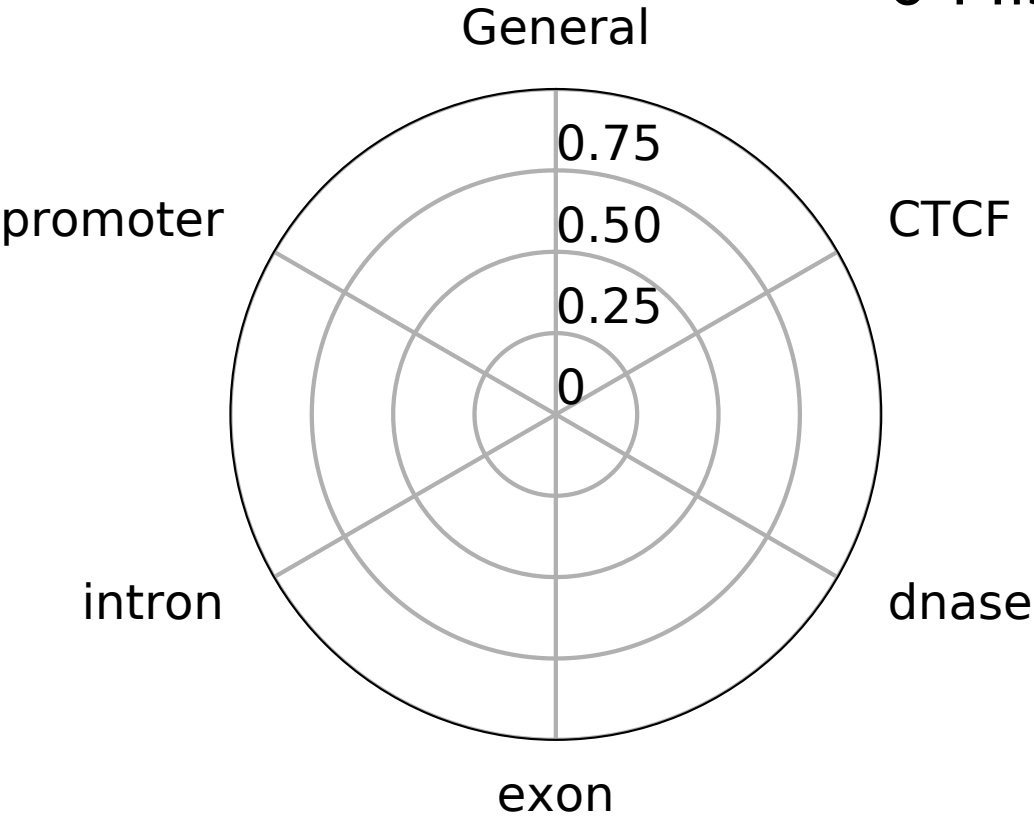


0 Mismatches



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

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

A

C

G

T

D

R

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