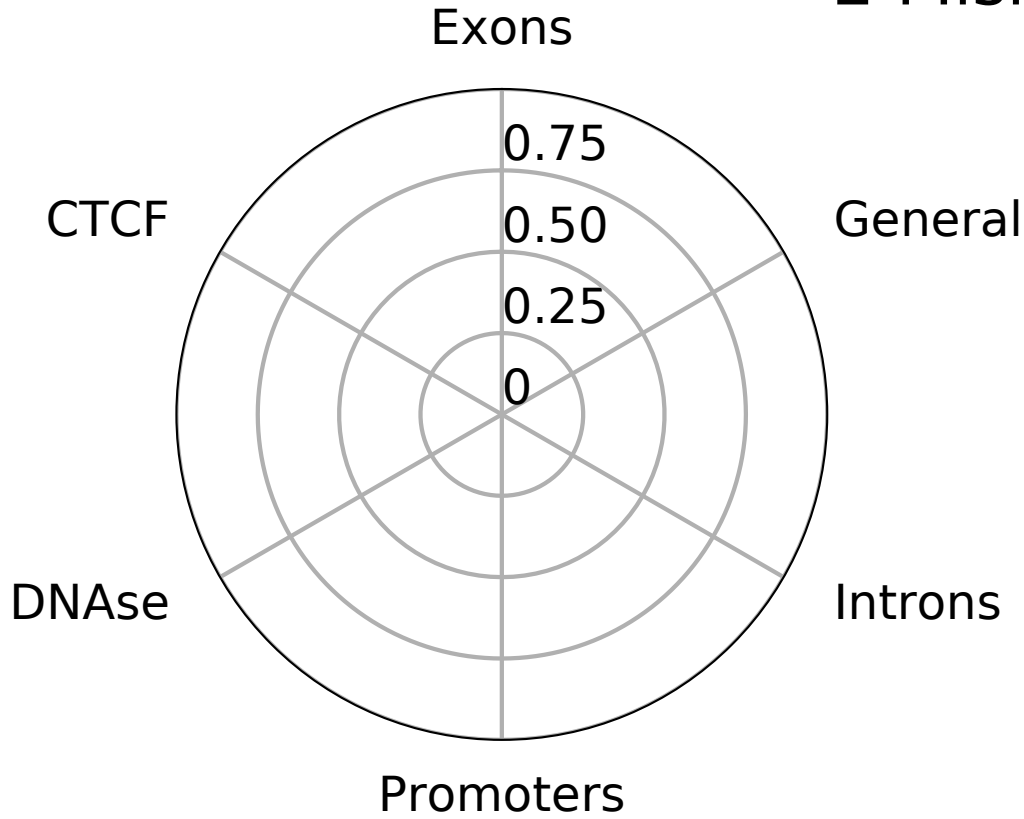


2 Mismatches



	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 -

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
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