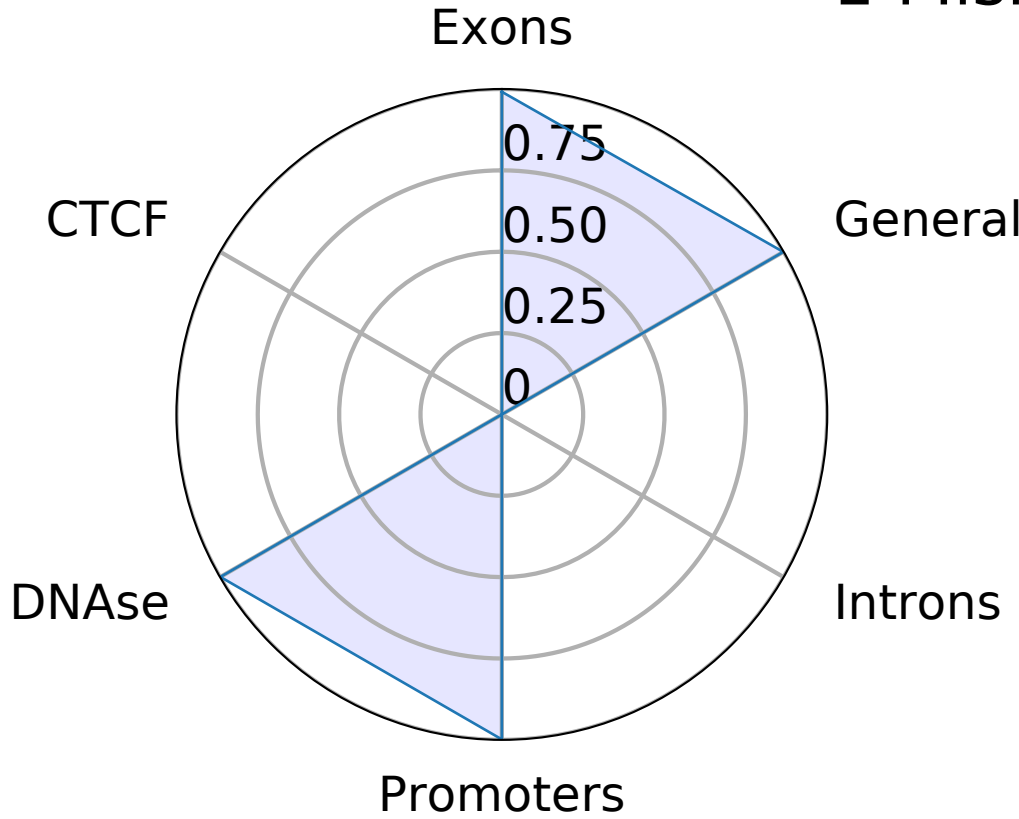


1 Mismatches



	Position	# Targets
General	1.0	354.0
Exons	0.99	6.0
Introns	0.0	0.0
Promoters	1.0	71.0
DNase	1.0	24.0
CTCF	0.0	0.0

