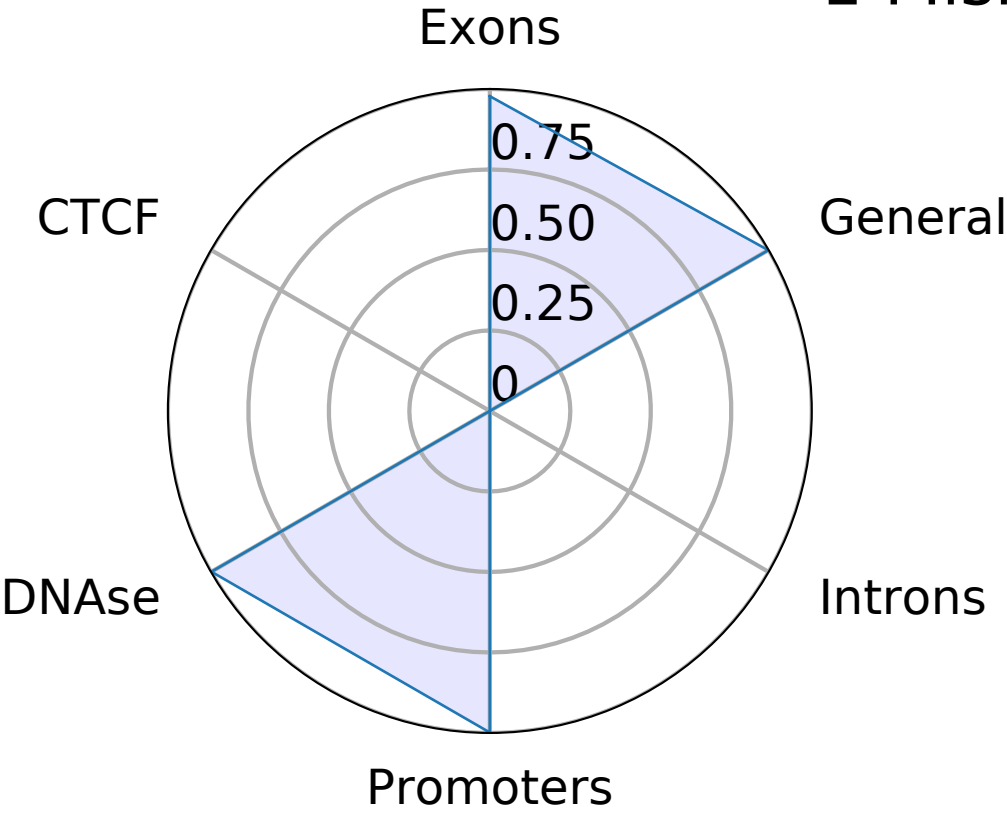


1 Mismatches



	Position	# Targets
General	1.0	322.0
Exons	0.98	2.0
Introns	0.0	0.0
Promoters	1.0	98.0
DNase	1.0	19.0
CTCF	0.0	0.0

