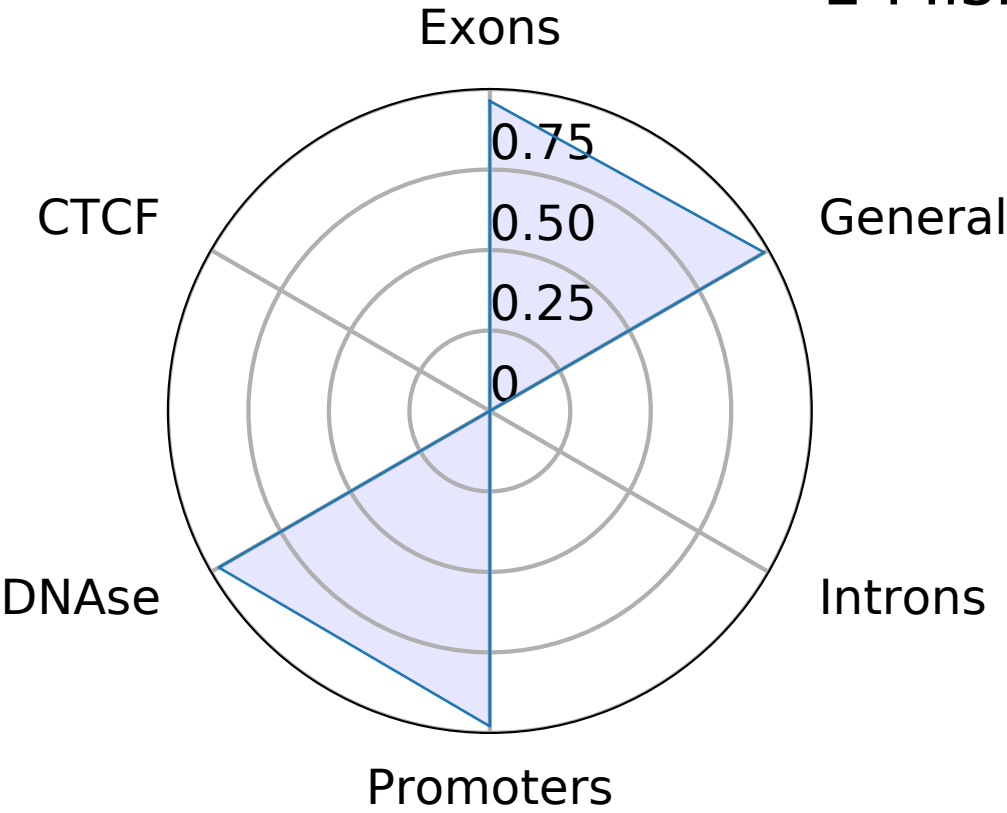


1 Mismatches



	Position	# Targets
General	0.98	4.0
Exons	0.96	1.0
Introns	0.0	0.0
Promoters	0.98	1.0
DNase	0.97	1.0
CTCF	0.0	0.0

