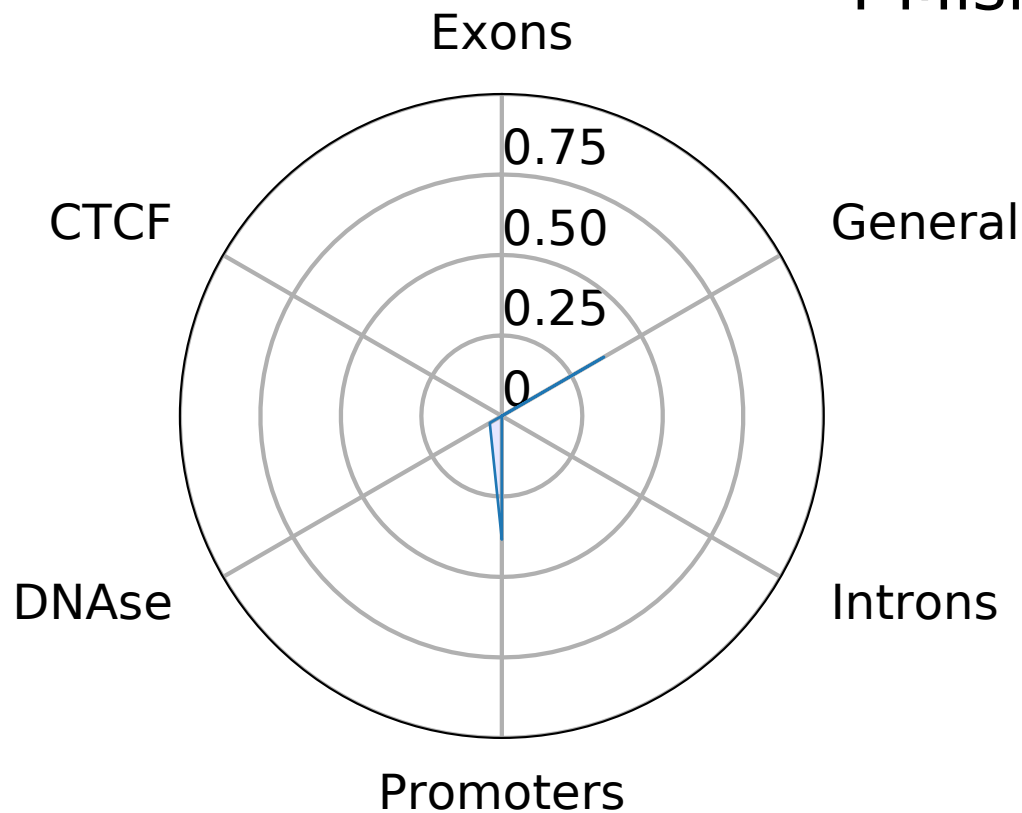


4 Mismatches



	Position	# Targets
General	0.37	66.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.38	11.0
DNase	0.04	7.0
CTCF	0.0	0.0

