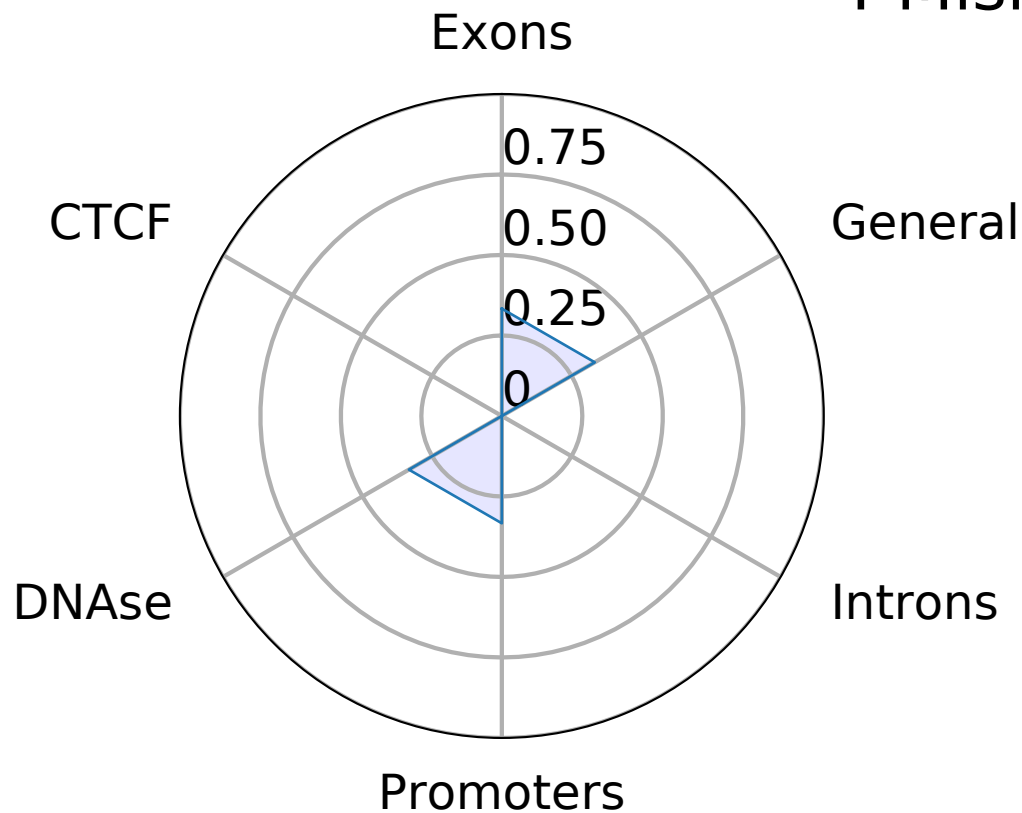


4 Mismatches



	Position	# Targets
General	0.33	286.0
Exons	0.33	6.0
Introns	0.0	0.0
Promoters	0.33	91.0
DNase	0.33	15.0
CTCF	0.0	0.0

