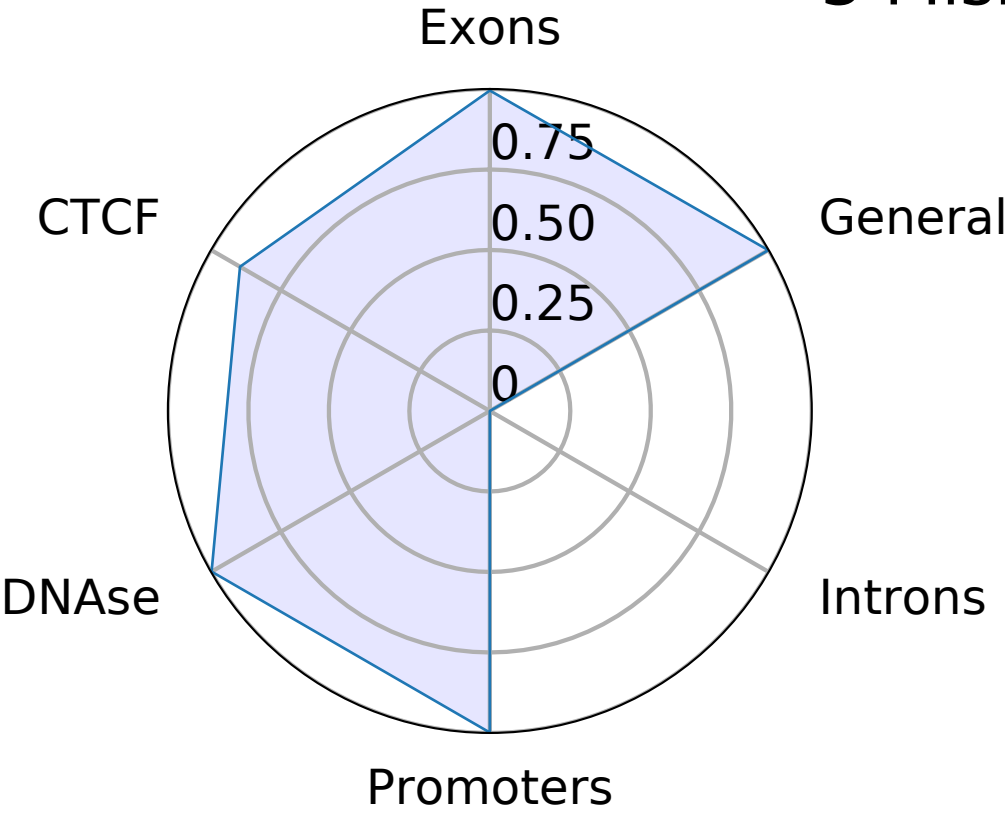


3 Mismatches



	Position	# Targets
General	1.0	4063.0
Exons	1.0	44.0
Introns	0.0	0.0
Promoters	1.0	1114.0
DNase	1.0	213.0
CTCF	0.9	1.0

