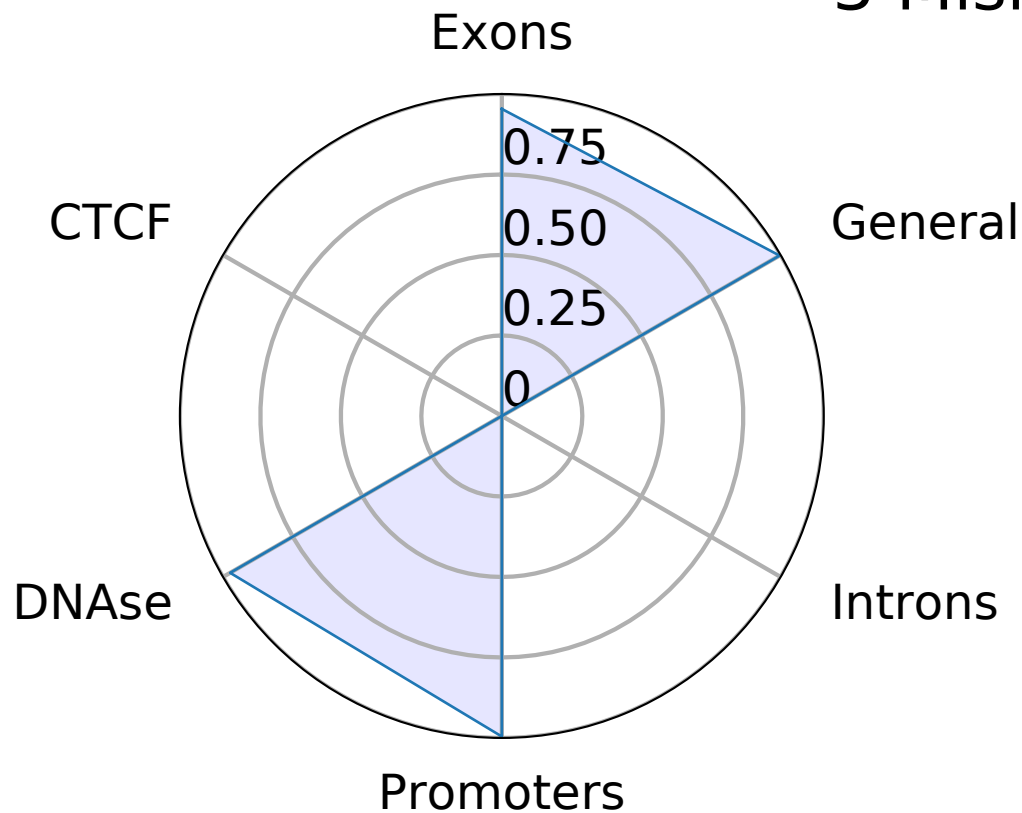


3 Mismatches



	Position	# Targets
General	1.0	312.0
Exons	0.95	5.0
Introns	0.0	0.0
Promoters	1.0	84.0
DNase	0.97	18.0
CTCF	0.0	0.0

