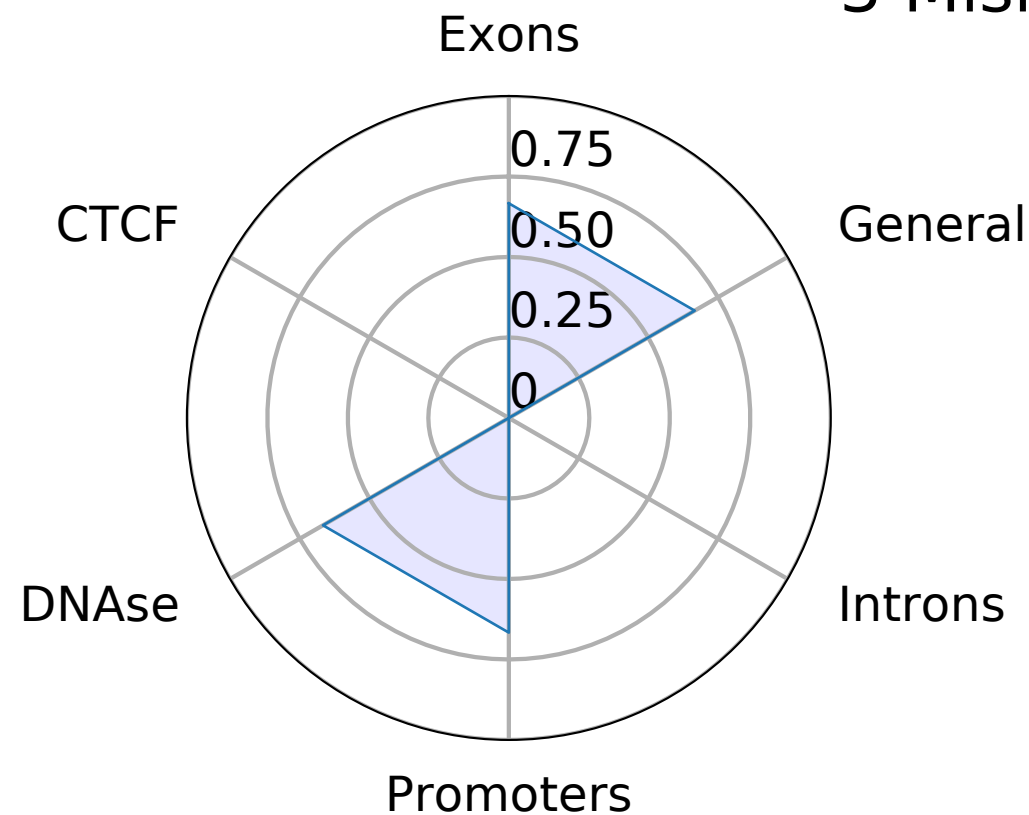


3 Mismatches



	Position	# Targets
General	0.67	312.0
Exons	0.67	5.0
Introns	0.0	0.0
Promoters	0.67	84.0
DNase	0.67	18.0
CTCF	0.0	0.0

