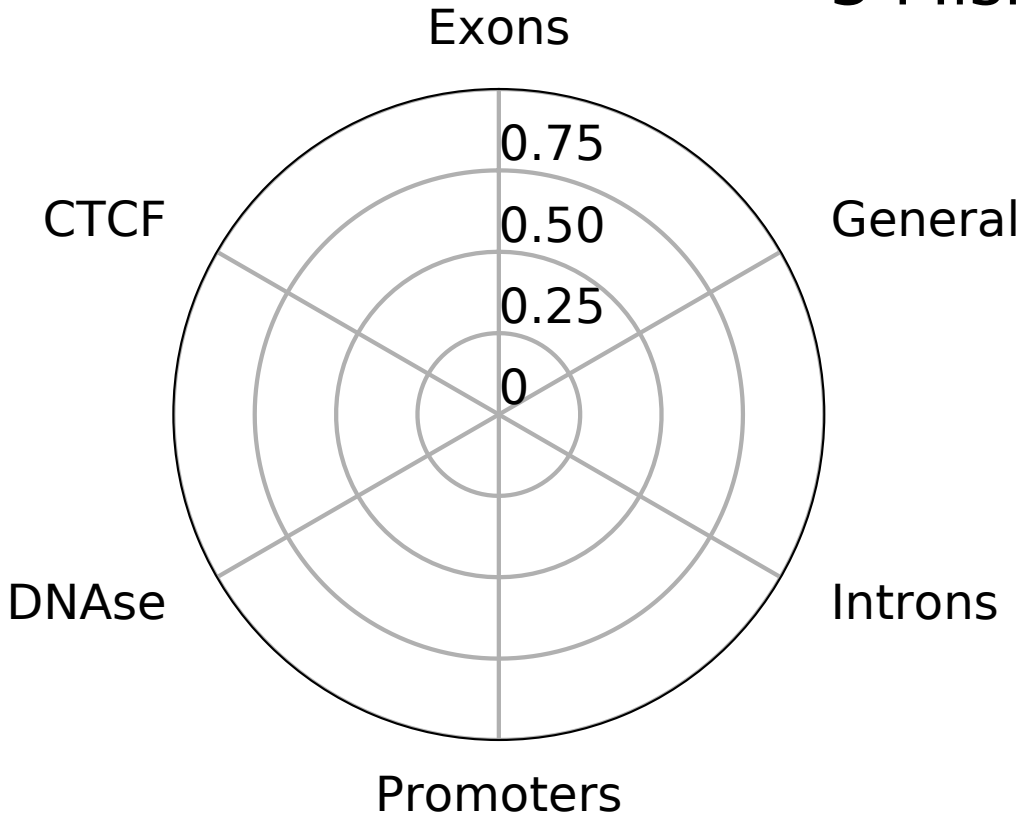


3 Mismatches



	Position	# Targets
General	0.0	1.0
Exons	0.0	64.0
Introns	0.0	0.0
Promoters	0.0	435.0
DNase	0.0	94.0
CTCF	0.0	0.0

