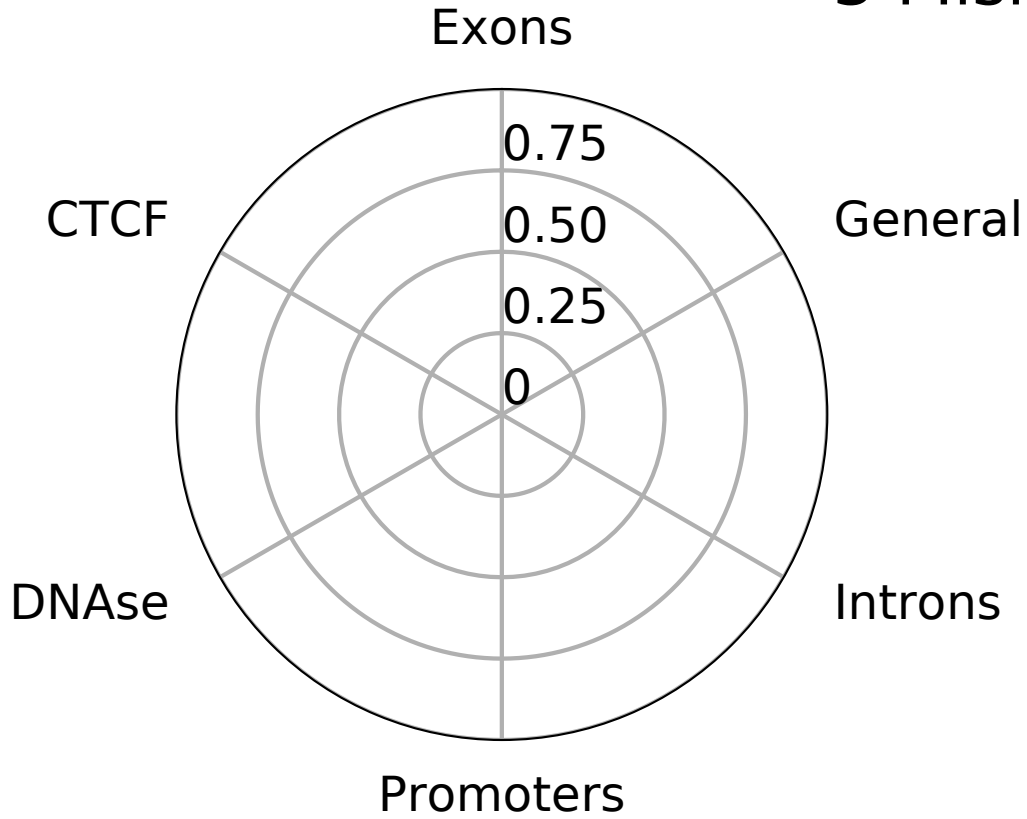


3 Mismatches



	Position	# Targets
General	0.0	0.0
Exons	0.0	2.0
Introns	0.0	0.0
Promoters	0.0	18.0
DNase	0.0	4.0
CTCF	0.0	0.0

