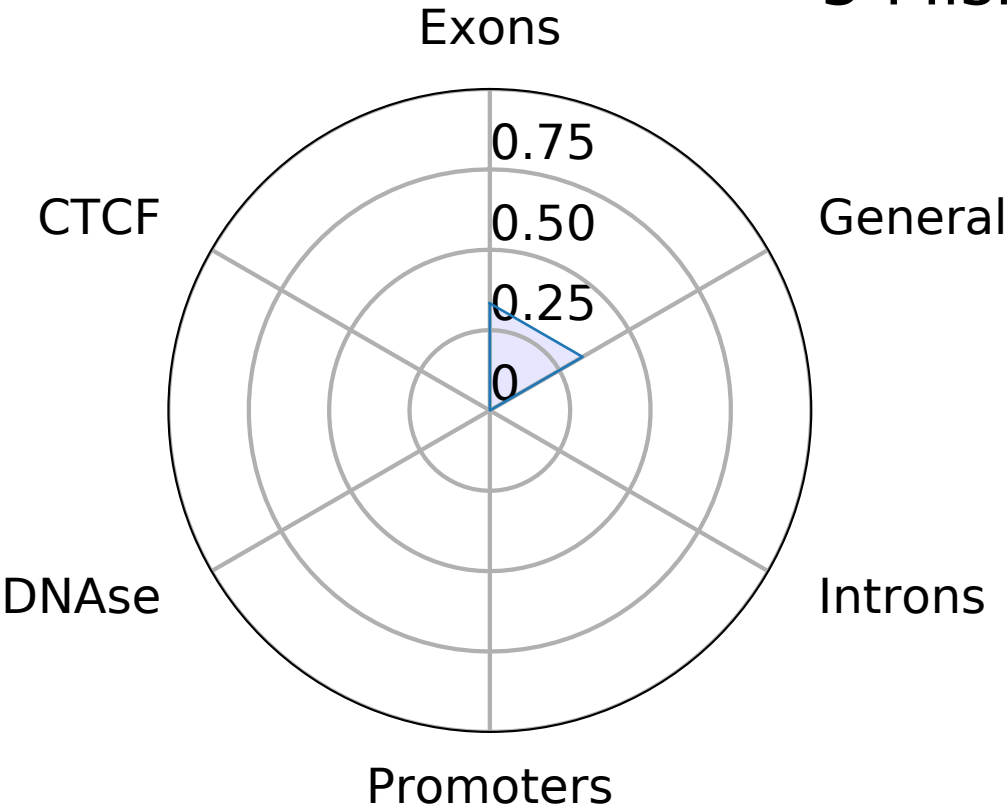


5 Mismatches



	Position	# Targets
General	0.33	28.0
Exons	0.33	1.0
Introns	0.0	0.0
Promoters	0.0	4.0
DNase	0.0	2.0
CTCF	0.0	0.0

