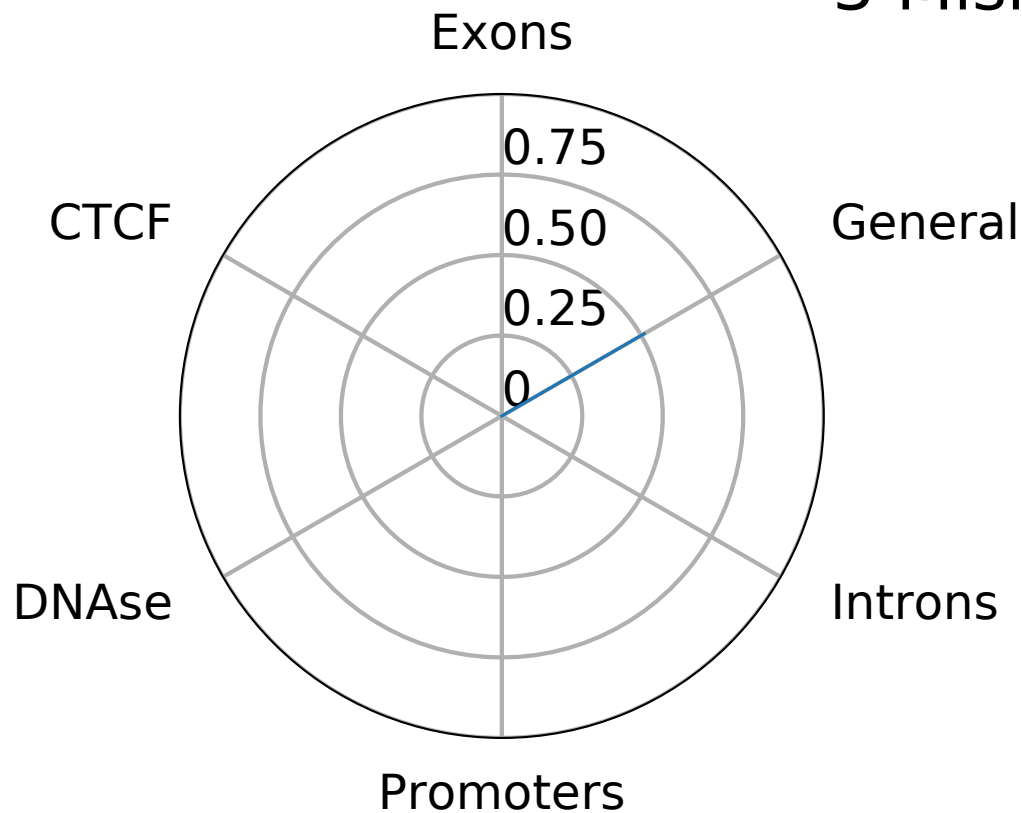


# 3 Mismatches



	Position	# Targets
General	0.51	6.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.0	0.0
DNase	0.0	0.0
CTCF	0.0	0.0

