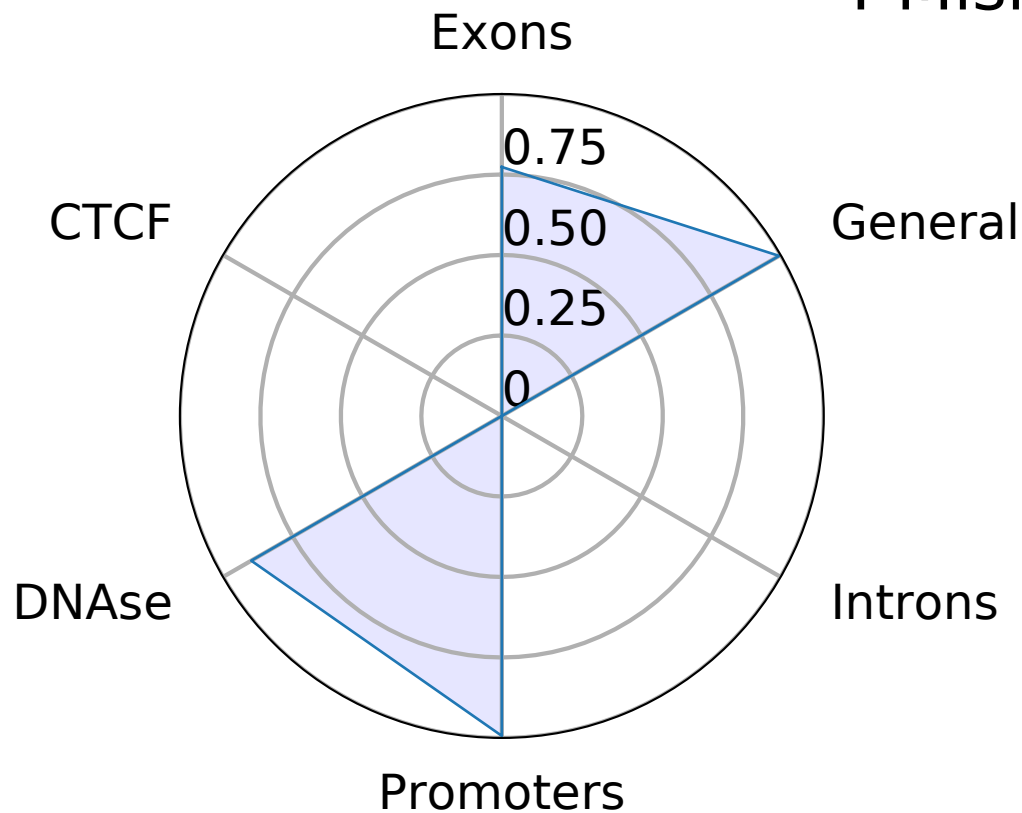


# 4 Mismatches



	Position	# Targets
General	0.99	1201.0
Exons	0.77	13.0
Introns	0.0	0.0
Promoters	0.99	304.0
DNase	0.9	71.0
CTCF	0.0	0.0

