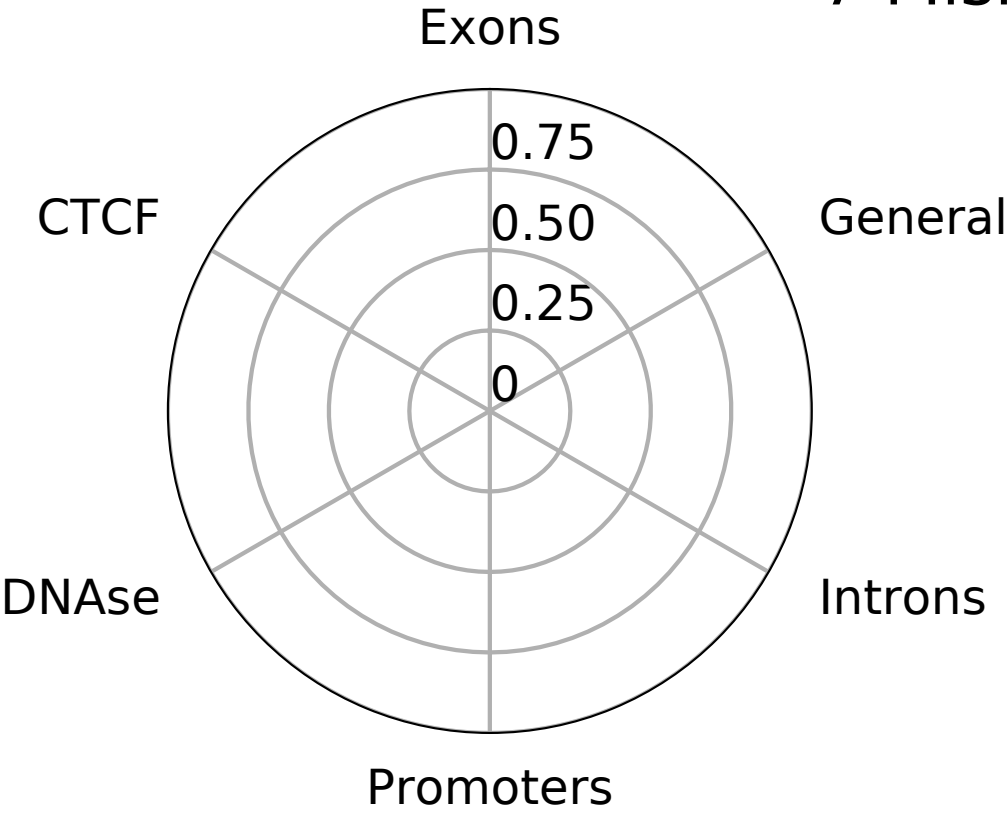


7 Mismatches



	Position	# Targets
General	0.0	310.0
Exons	0.0	23.0
Introns	0.0	0.0
Promoters	0.0	98.0
DNase	0.0	127.0
CTCF	0.0	1.0

