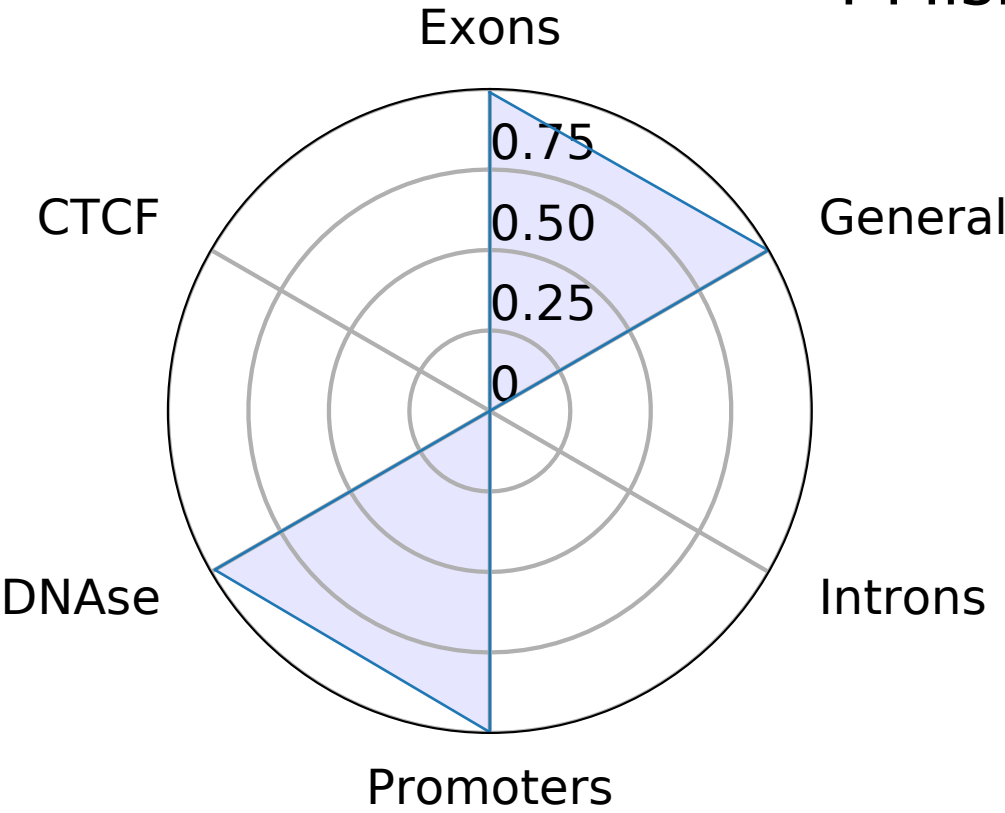


4 Mismatches



	Position	# Targets
General	1.0	4657.0
Exons	0.99	84.0
Introns	0.0	0.0
Promoters	1.0	1417.0
DNase	0.99	254.0
CTCF	0.0	0.0

