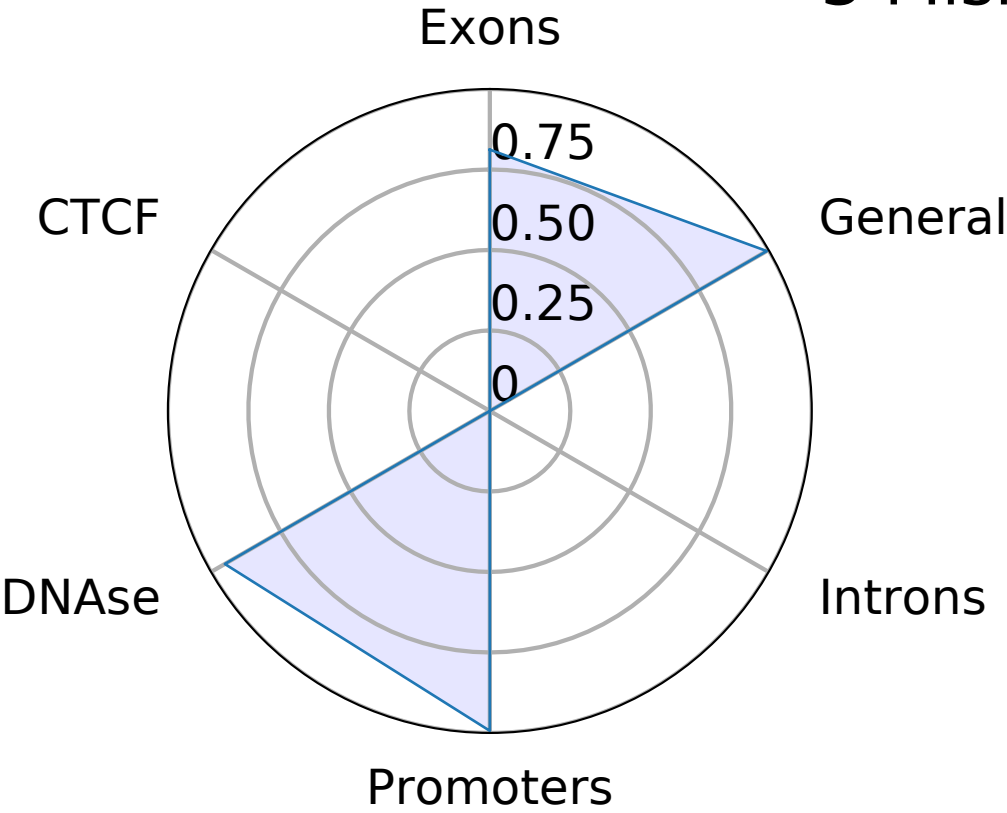


3 Mismatches



	Position	# Targets
General	0.99	133.0
Exons	0.81	2.0
Introns	0.0	0.0
Promoters	0.99	48.0
DNase	0.95	11.0
CTCF	0.0	0.0

