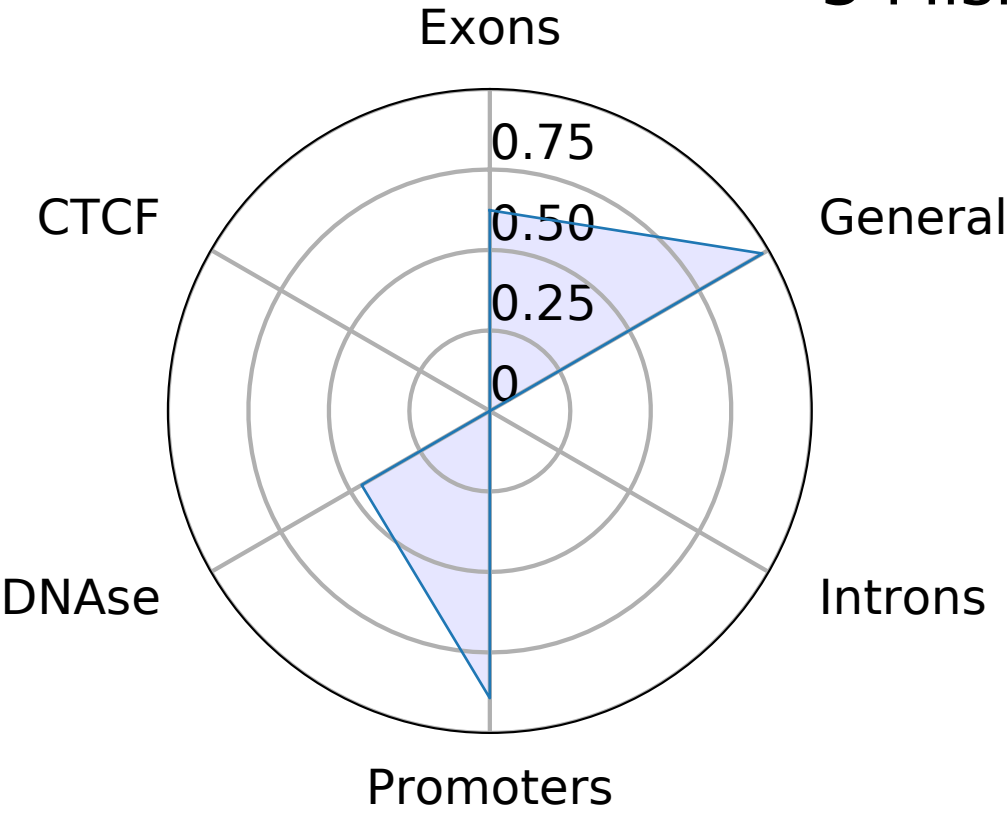


3 Mismatches



	Position	# Targets
General	0.98	47.0
Exons	0.62	1.0
Introns	0.0	0.0
Promoters	0.89	5.0
DNase	0.46	2.0
CTCF	0.0	0.0

