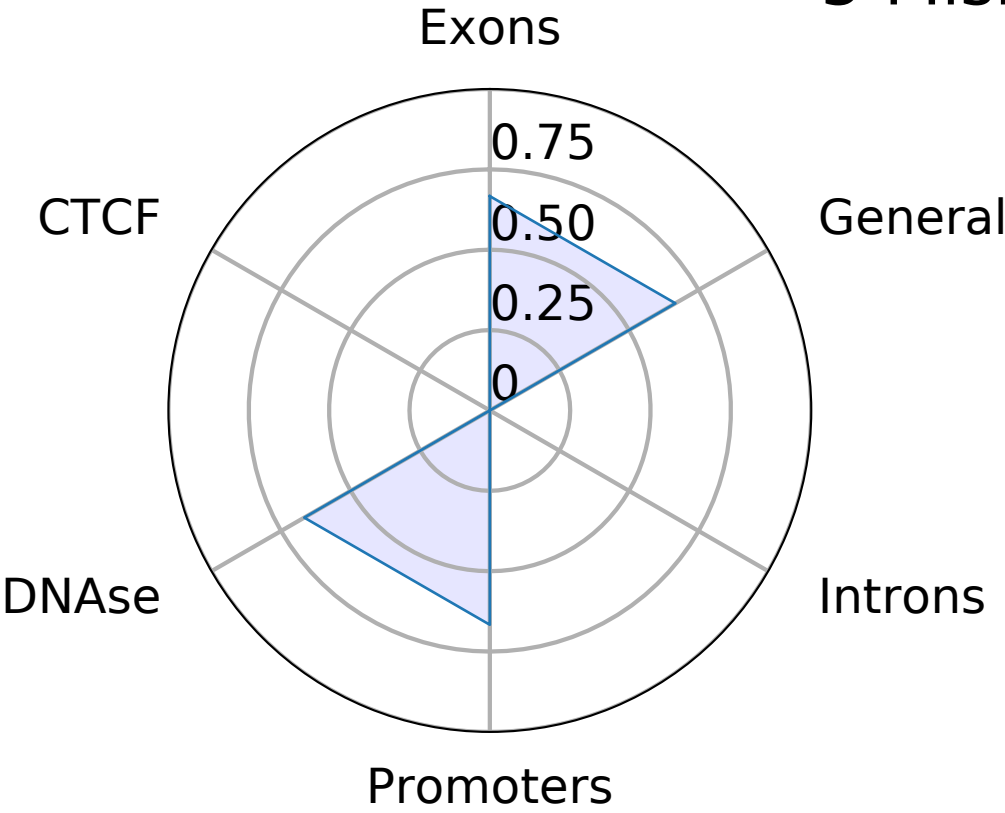


5 Mismatches



	Position	# Targets
General	0.67	113.0
Exons	0.67	8.0
Introns	0.0	0.0
Promoters	0.67	35.0
DNase	0.67	9.0
CTCF	0.0	0.0

