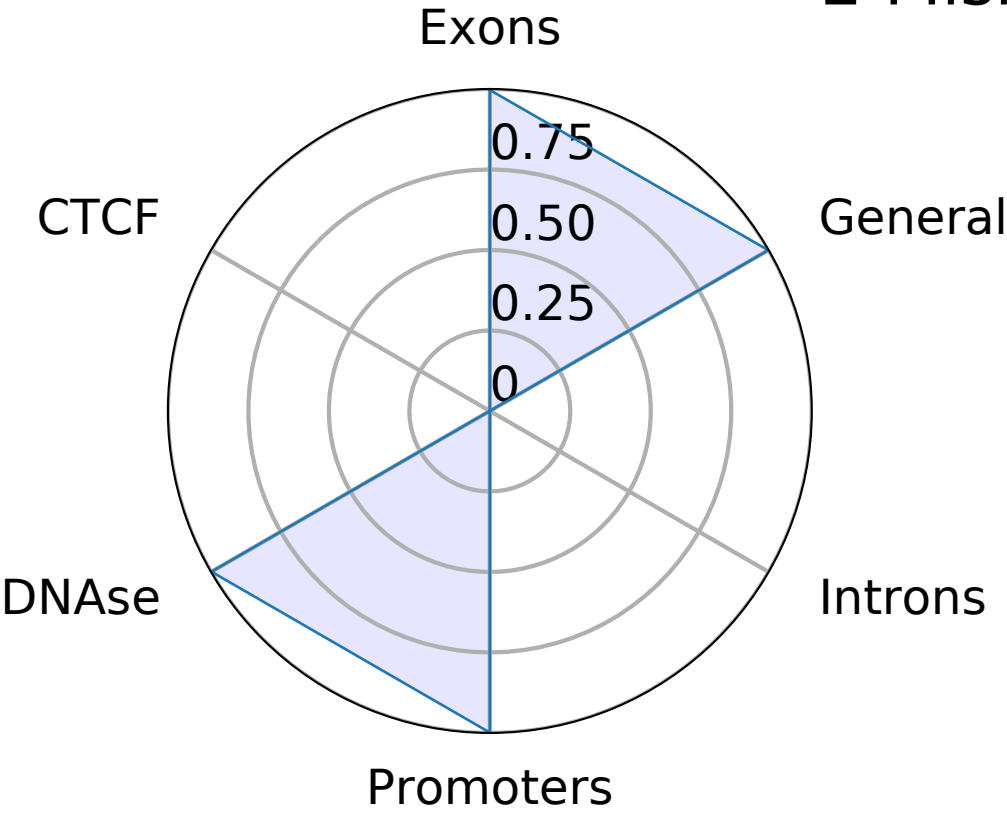


2 Mismatches



	Position	# Targets
General	1.0	1618.0
Exons	1.0	26.0
Introns	0.0	0.0
Promoters	1.0	466.0
DNase	1.0	95.0
CTCF	0.0	0.0

