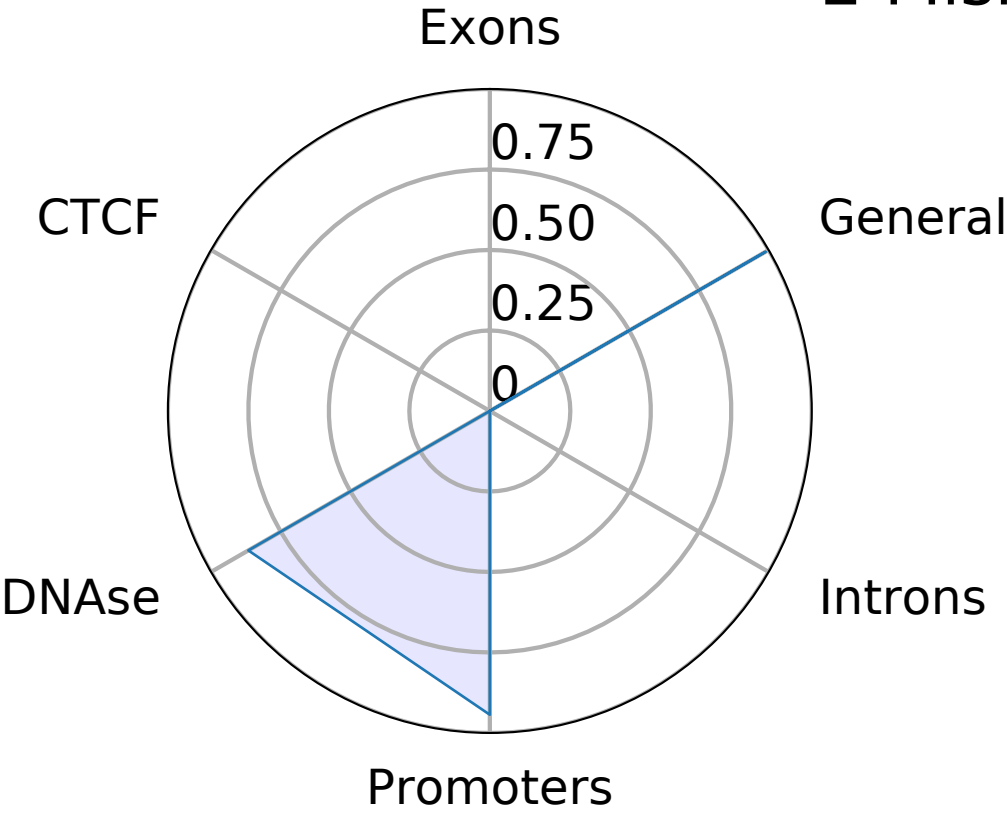


2 Mismatches



	Position	# Targets
General	0.99	11.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.94	1.0
DNase	0.87	1.0
CTCF	0.0	0.0

