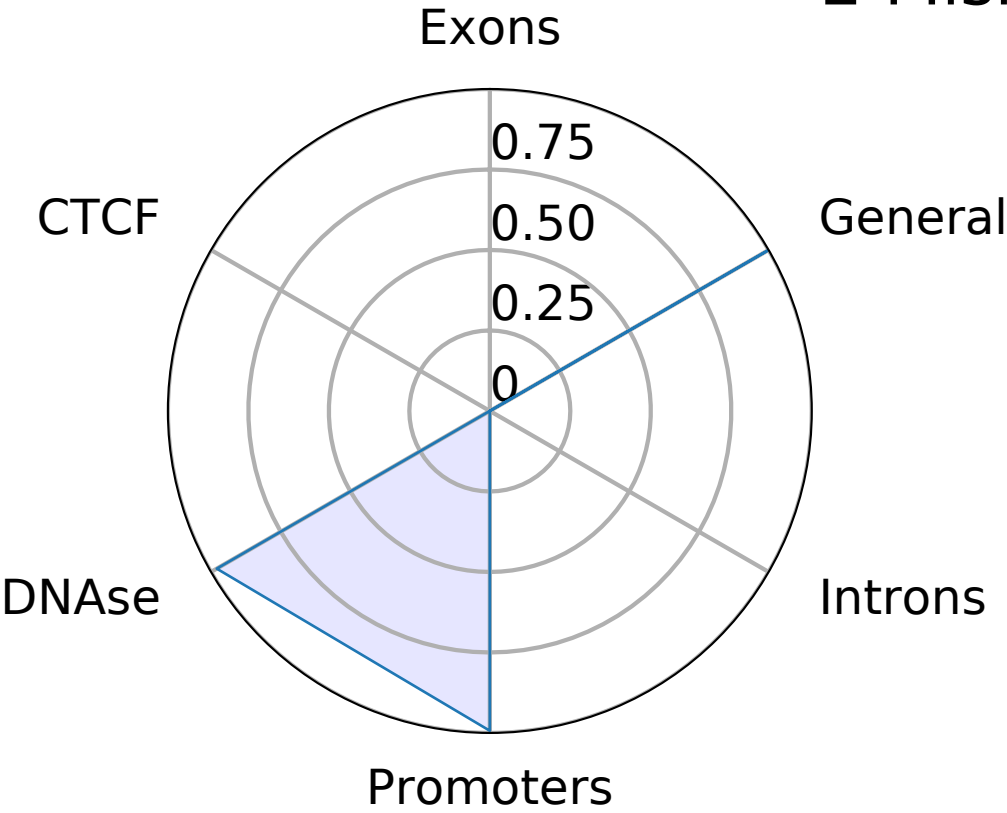


2 Mismatches



	Position	# Targets
General	1.0	45.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.99	11.0
DNase	0.98	3.0
CTCF	0.0	0.0

