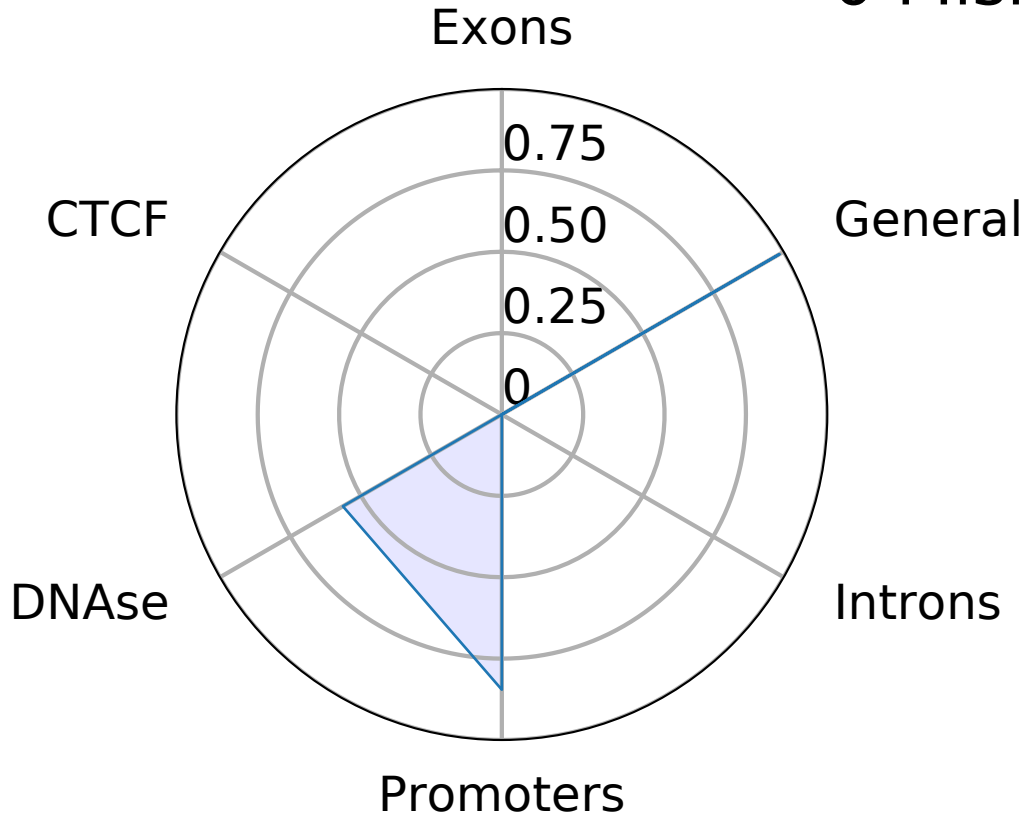


0 Mismatches



	Position	# Targets
General	0.98	7.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.85	1.0
DNase	0.56	1.0
CTCF	0.0	0.0

