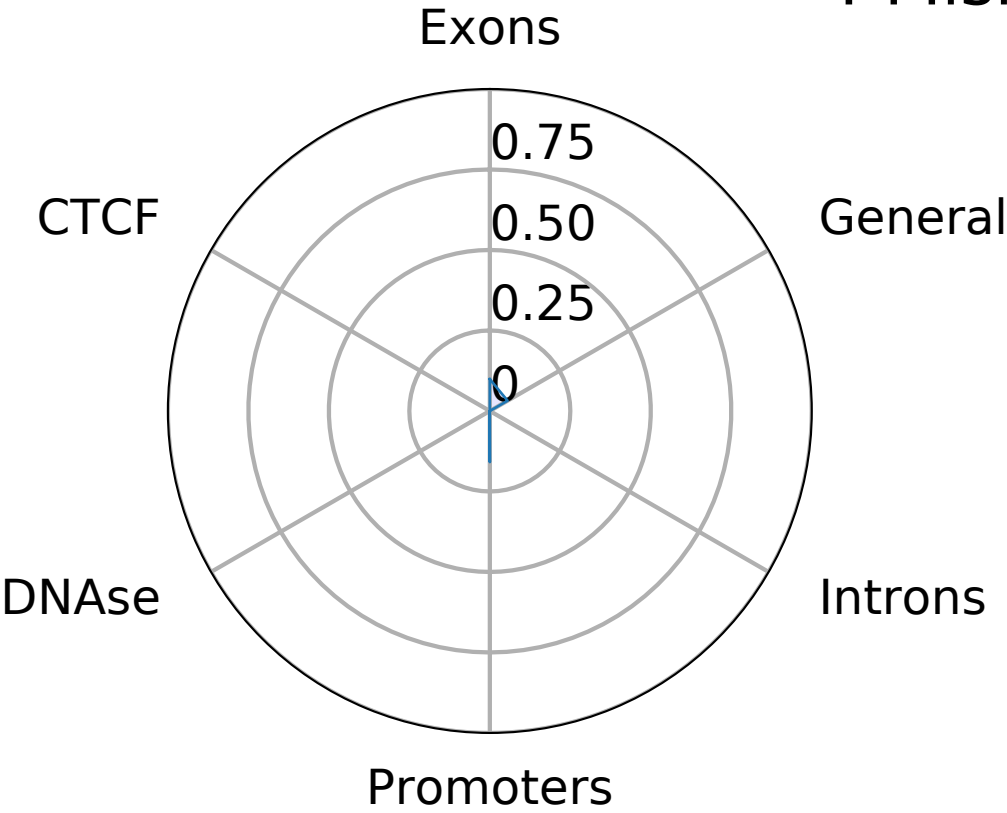


4 Mismatches



	Position	# Targets
General	0.06	23.0
Exons	0.1	2.0
Introns	0.0	0.0
Promoters	0.16	5.0
DNase	0.0	1.0
CTCF	0.0	0.0

