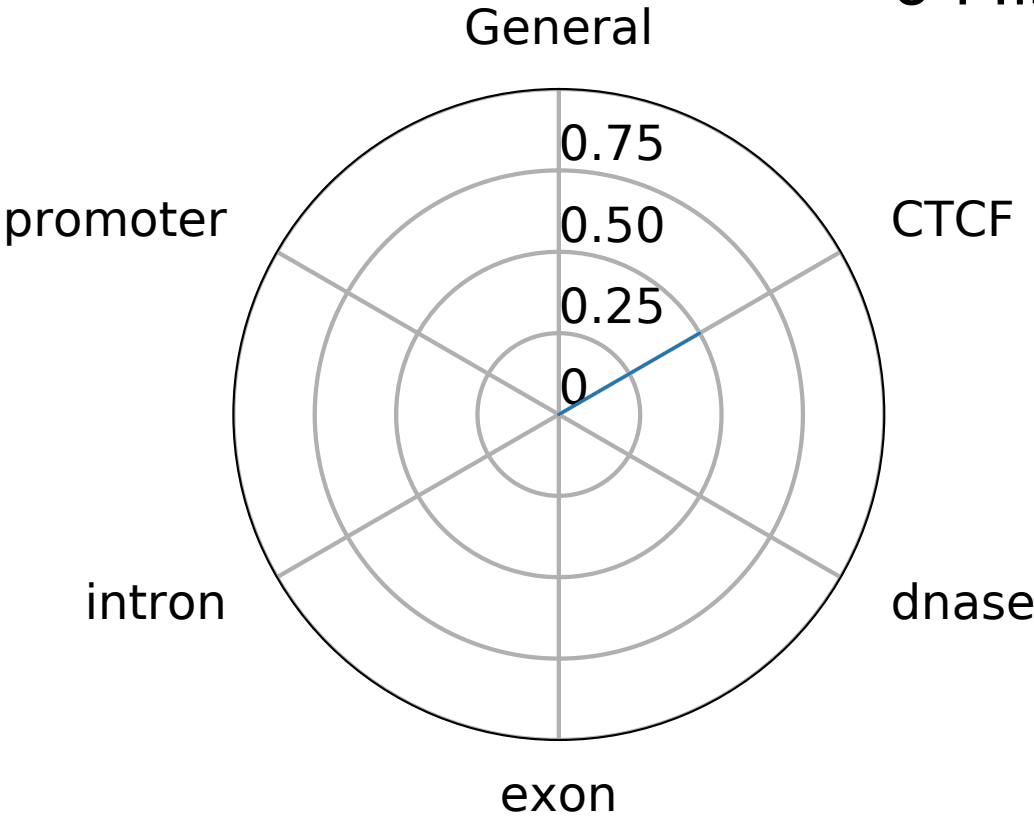


6 Mismatches



	Position	# Targets
General	0.0	723.0
CTCF	0.5	3.0
dnase	0.0	74.0
exon	0.0	55.0
intron	0.0	155.0
promoter	0.0	265.0

