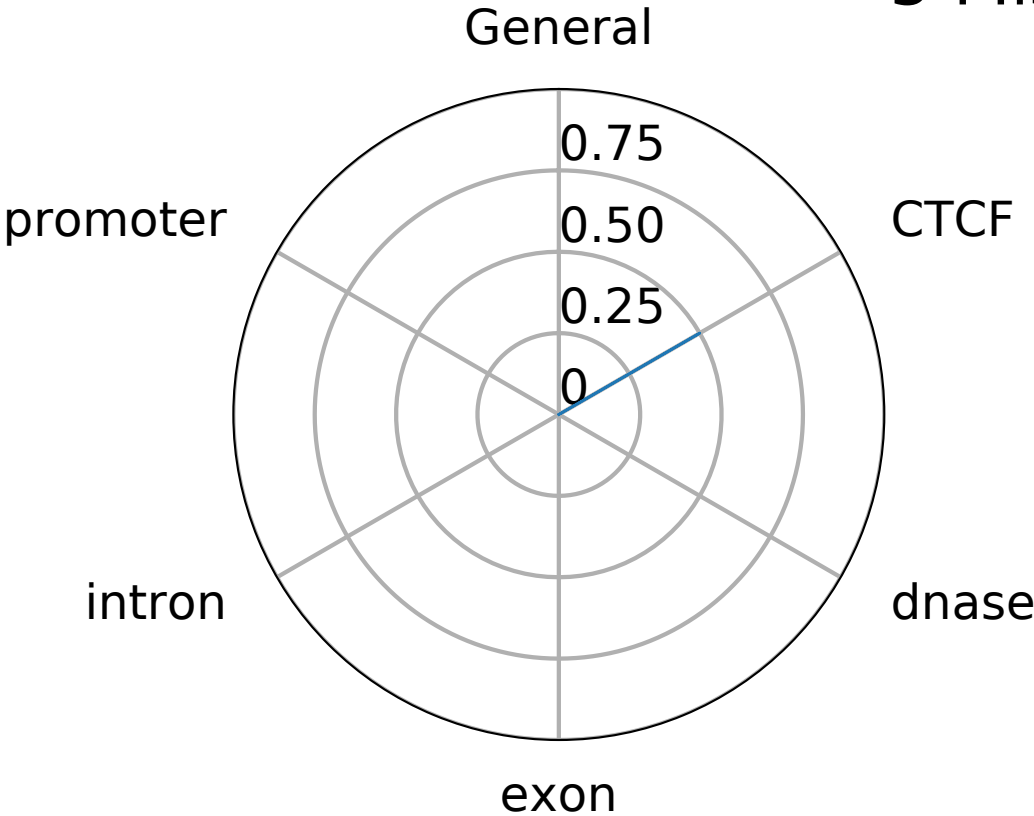


5 Mismatches



	Position	# Targets
General	0.0	93.0
CTCF	0.5	2.0
dnase	0.0	5.0
exon	0.0	5.0
intron	0.0	19.0
promoter	0.0	14.0

