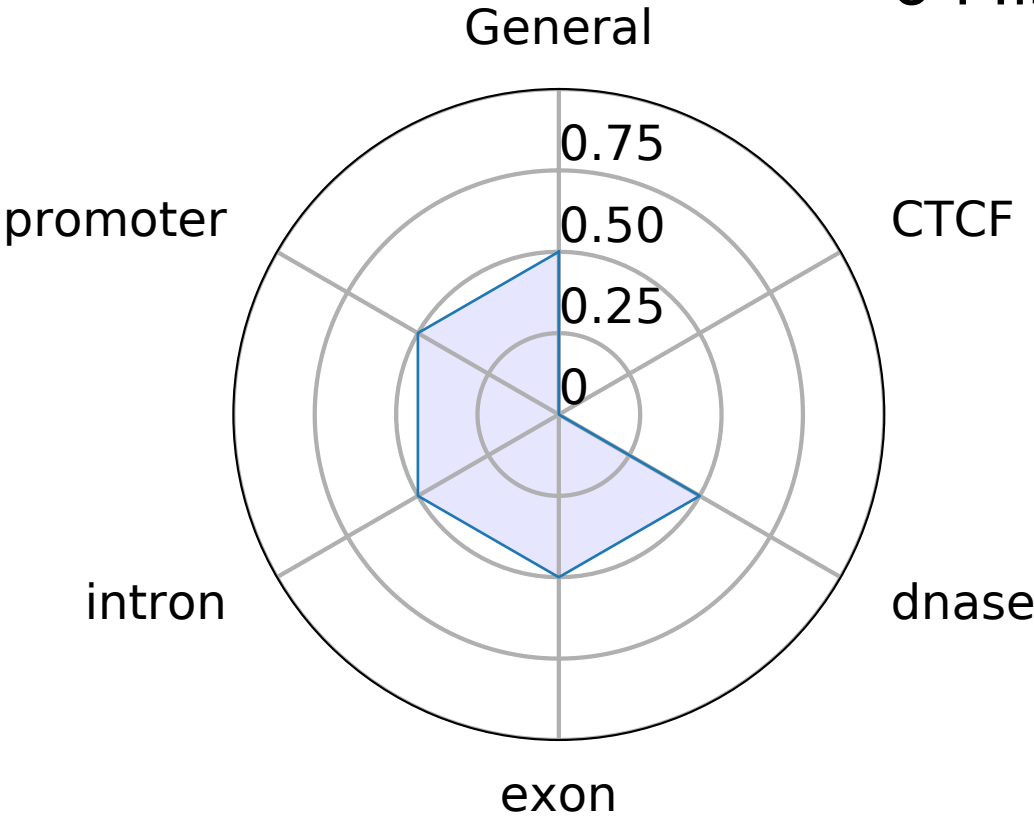


6 Mismatches



	Position	# Targets
General	0.5	1973.0
CTCF	0.0	1.0
dnase	0.5	168.0
exon	0.5	119.0
intron	0.5	451.0
promoter	0.5	461.0

