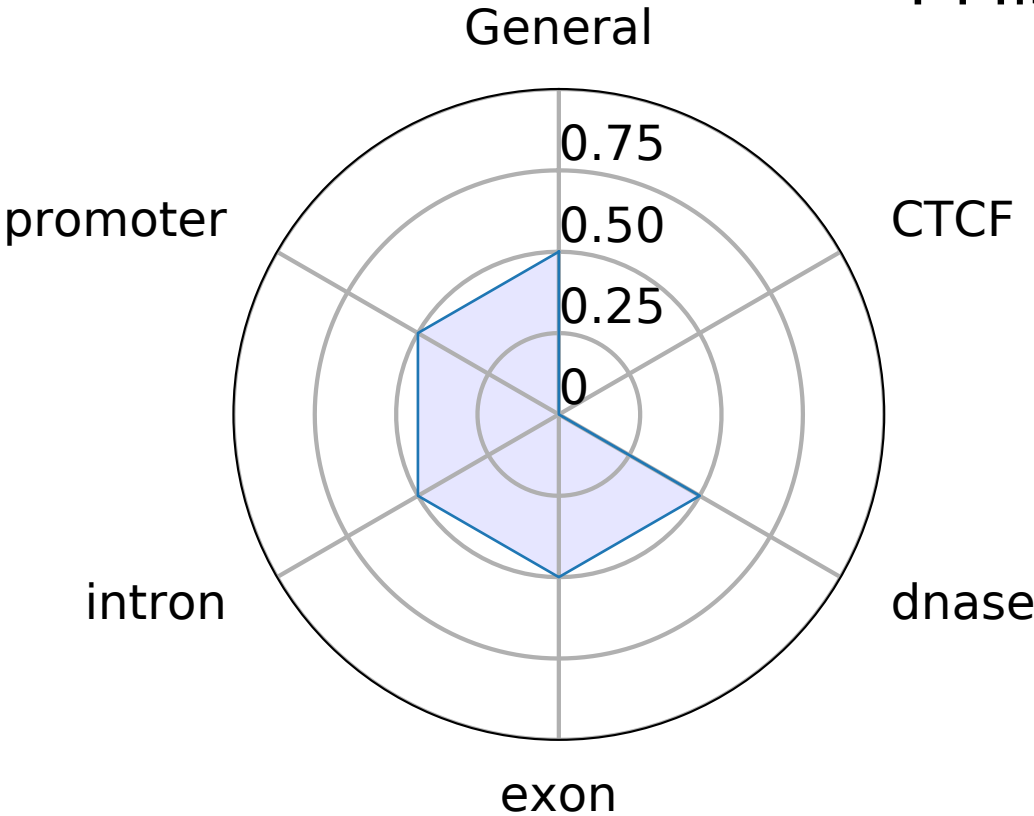


4 Mismatches



	Position	# Targets
General	0.5	354.0
CTCF	0.0	0.0
dnase	0.5	32.0
exon	0.5	10.0
intron	0.5	127.0
promoter	0.5	69.0

