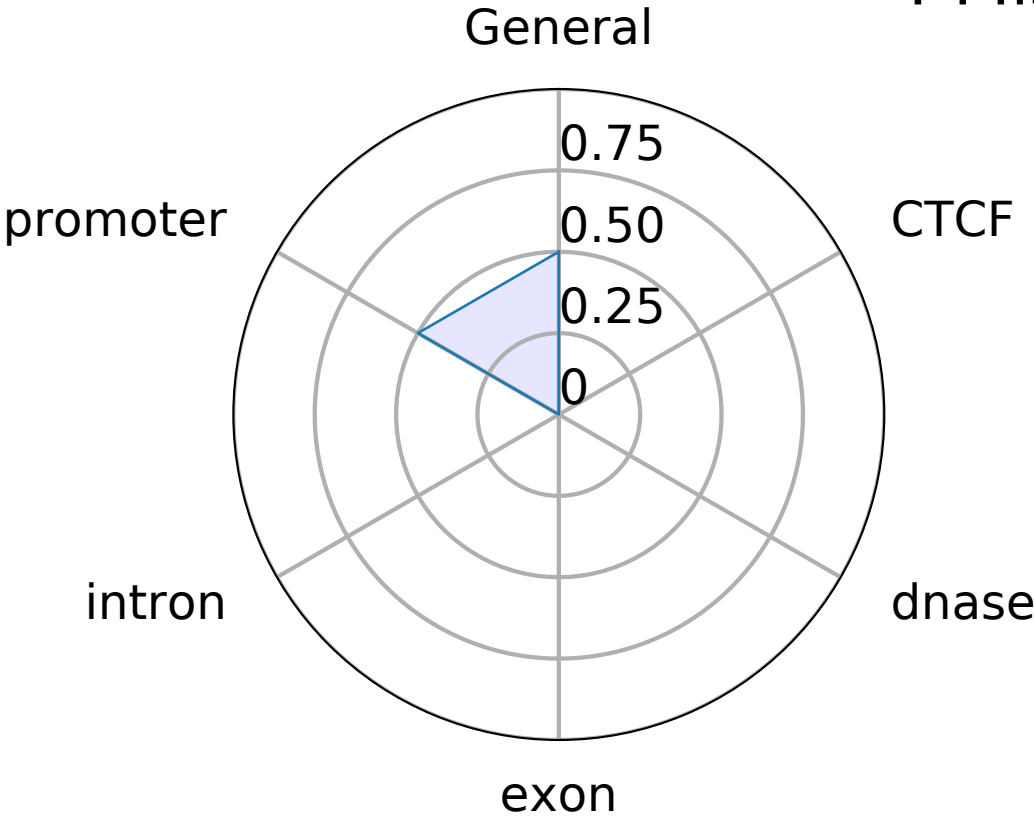


# 4 Mismatches



	Position	# Targets
General	0.5	423.0
CTCF	0.0	0.0
dnase	0.0	20.0
exon	0.0	11.0
intron	0.0	0.0
promoter	0.5	74.0

