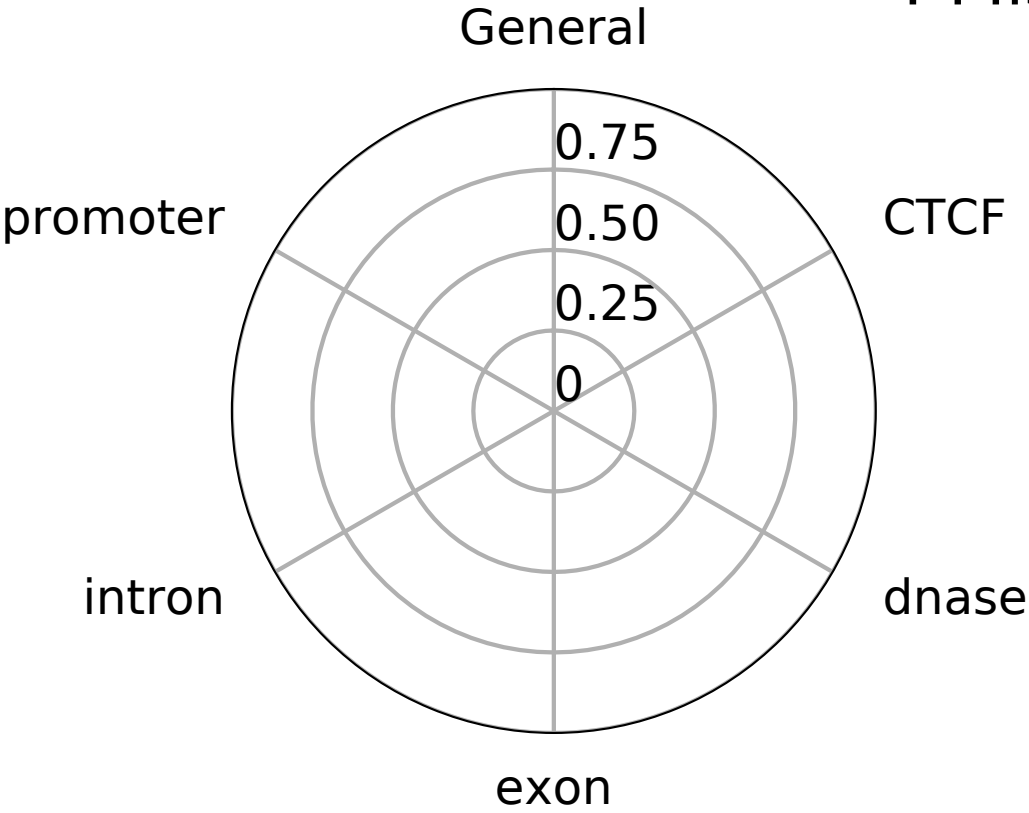


# 4 Mismatches



	Position	# Targets
General	0.0	8.0
CTCF	0.0	0.0
dnase	0.0	1.0
exon	0.0	0.0
intron	0.0	5.0
promoter	0.0	1.0

