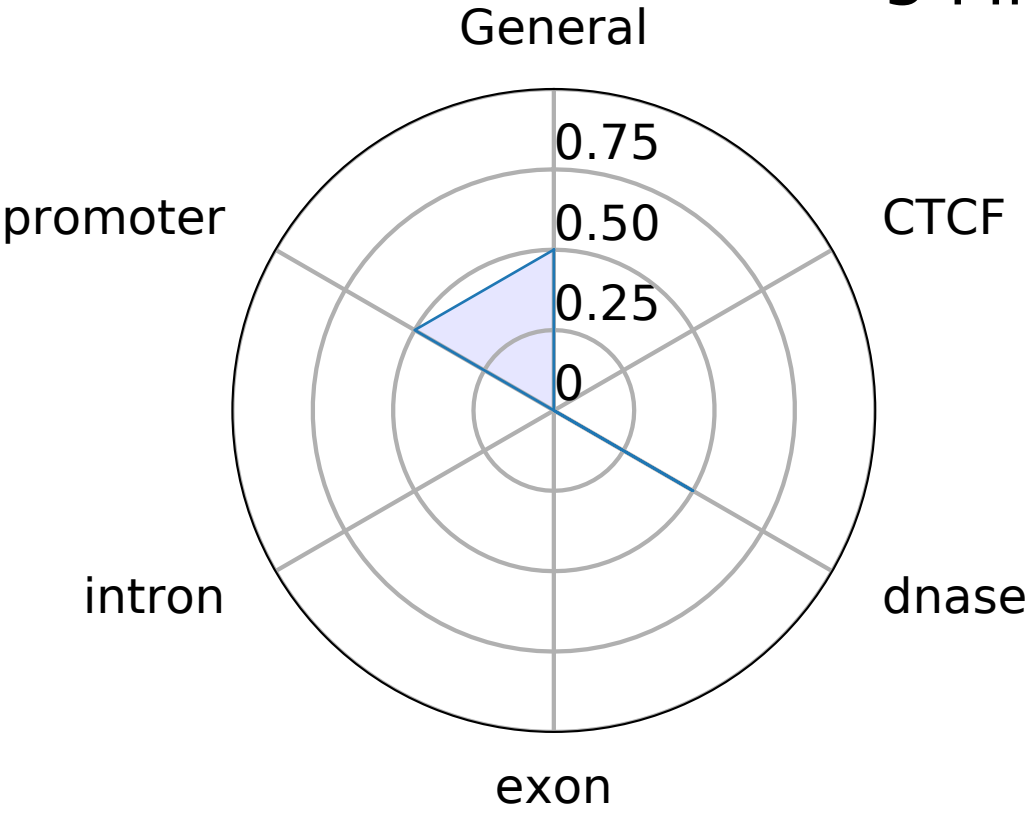


5 Mismatches



	Position	# Targets
General	0.5	34.0
CTCF	0.0	0.0
dnase	0.5	4.0
exon	0.0	0.0
intron	0.0	0.0
promoter	0.5	5.0

