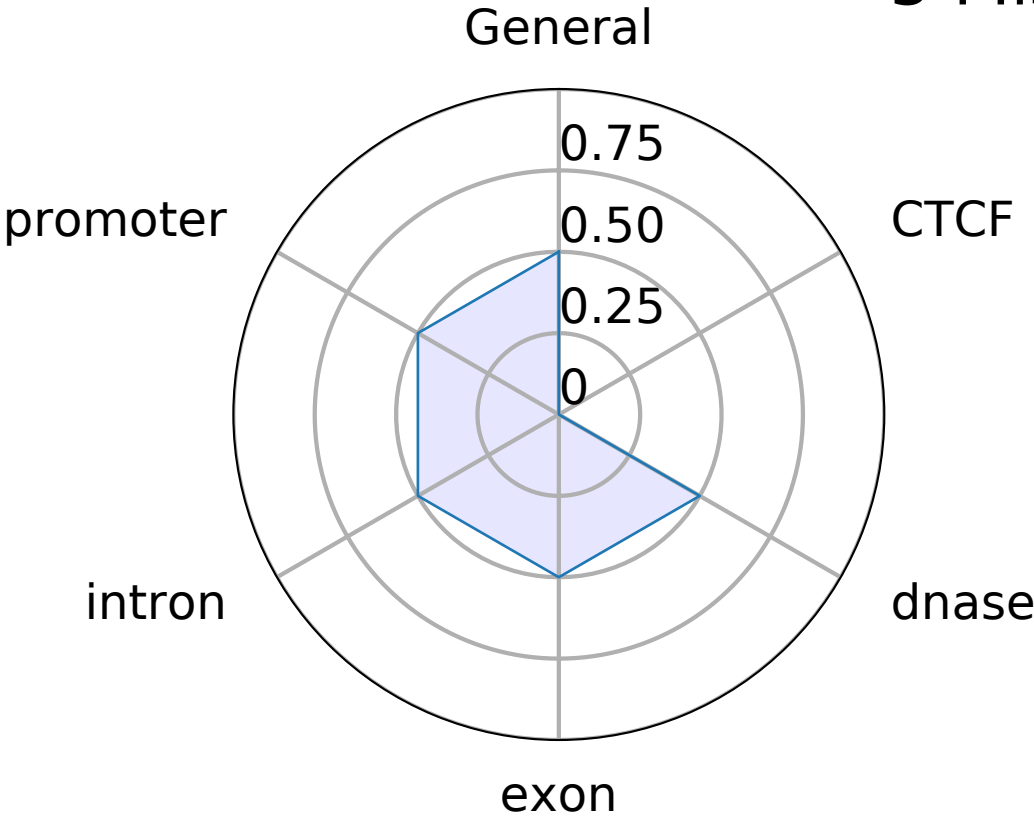


5 Mismatches



	Position	# Targets
General	0.5	289.0
CTCF	0.0	0.0
dnase	0.5	19.0
exon	0.5	18.0
intron	0.5	57.0
promoter	0.5	68.0

