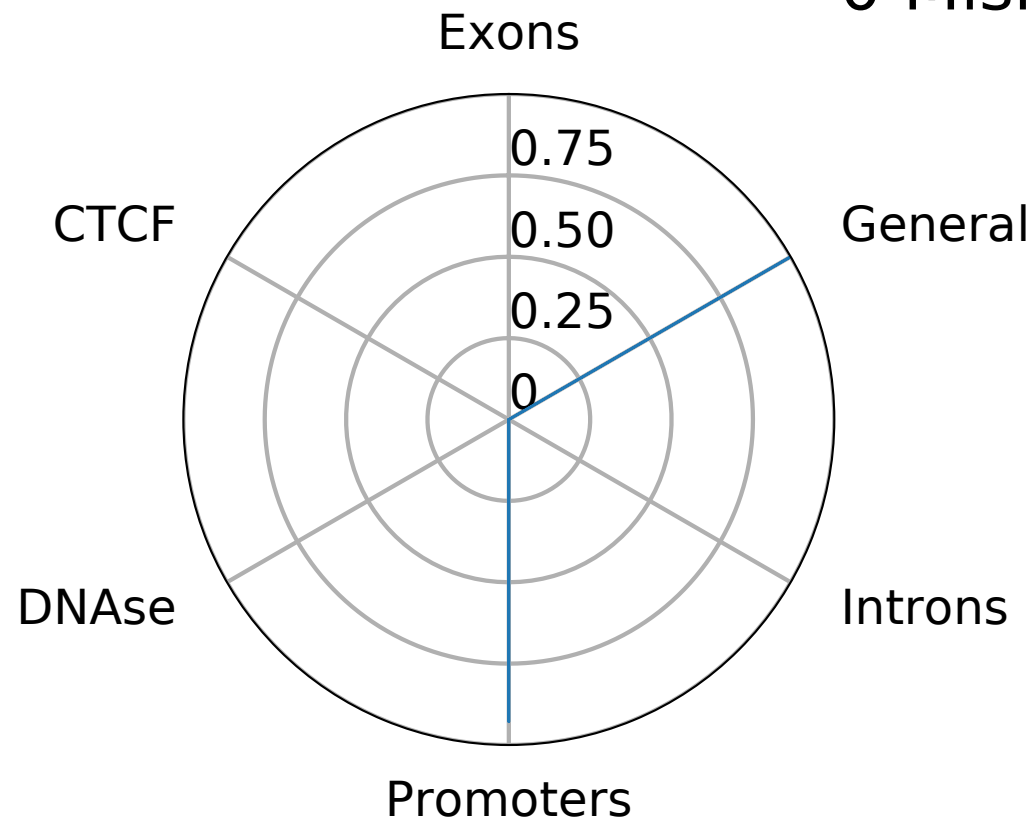


# 0 Mismatches



	Position	# Targets
General	1.0	14.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	0.93	2.0
DNase	0.0	0.0
CTCF	0.0	0.0

