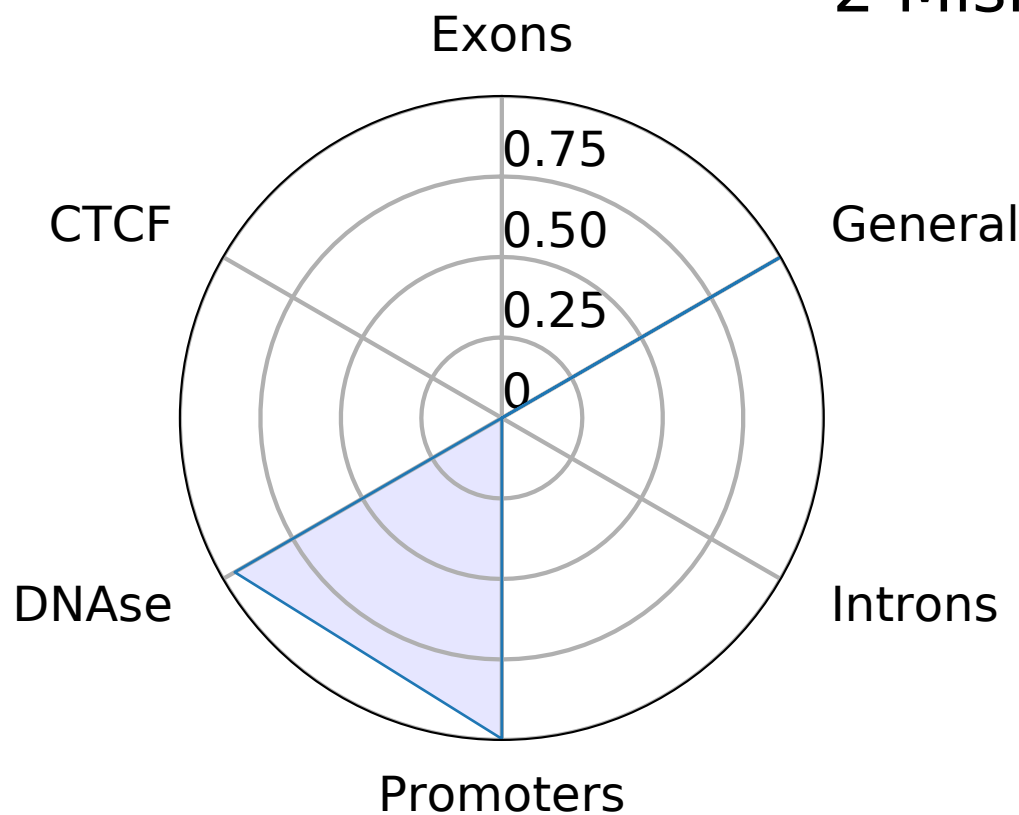


2 Mismatches



	Position	# Targets
General	1.0	48.0
Exons	0.0	0.0
Introns	0.0	0.0
Promoters	1.0	27.0
DNase	0.96	2.0
CTCF	0.0	0.0

