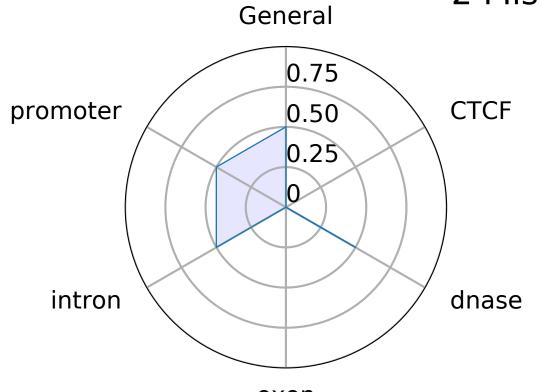
2 Mismatches

Α

G



	Position	# Targets
General	0.5	2.0
CTCF	0.0	0.0
dnase	0.5	1.0
exon	0.0	0.0
intron	0.5	1.0
promoter	0.5	2.0

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