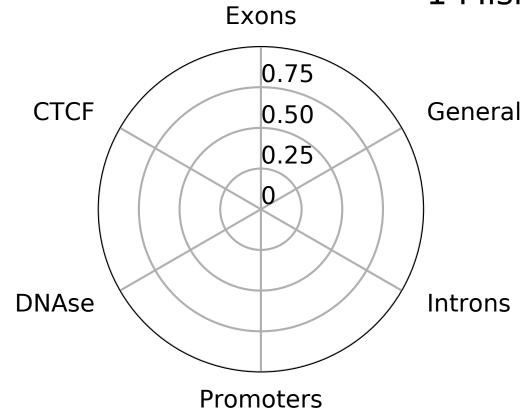
1 Mismatches



	Position	# Targets				
General	0.0	0.0				
Exons	0.0	0.0				
Introns	0.0	0.0				
Promoters	0.0	0.0				
DNAse	0.0	0.0				
CTCF	0.0	0.0				



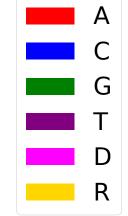
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



INN.																				
0.0	T	G	\cup	T	T	G	G	T	C	G	G	C	Α	C	T	G	Α	T	Α	G