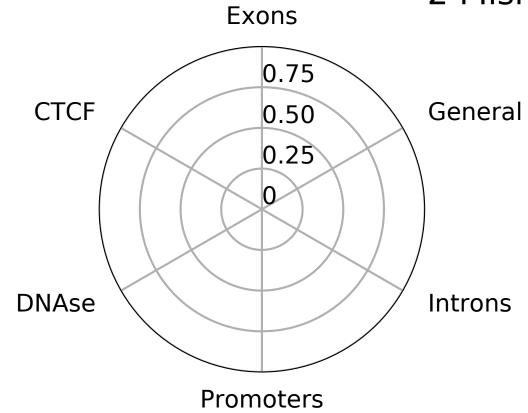
2 Mismatches



	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					



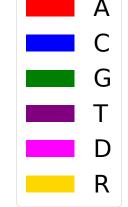
1.0 -

0.8 -

0.6 -

0.4 -

0.2 -



0.0 -	_																			
0.0	T	Α	G	T	T	G	T	Α	Α	Α	Α	С	T	T	Т	G	Α	T	T	T