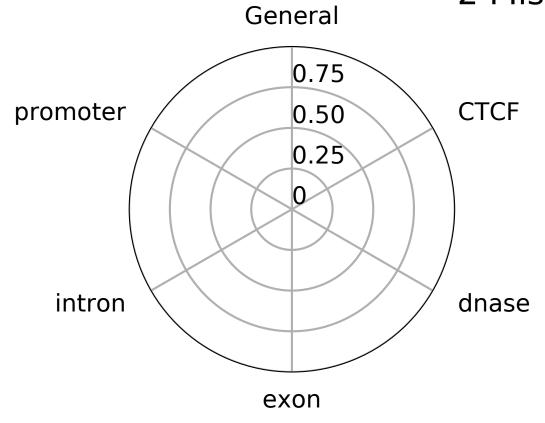
## 2 Mismatches



|          | Position | # Targets |
|----------|----------|-----------|
| General  | 0.0      | 0.0       |
| CTCF     | 0.0      | 0.0       |
| dnase    | 0.0      | 0.0       |
| exon     | 0.0      | 0.0       |
| intron   | 0.0      | 0.0       |
| promoter | 0.0      | 0.0       |



1.0 -

0.8 -

0.6 -

0.4 -

0.2 -

