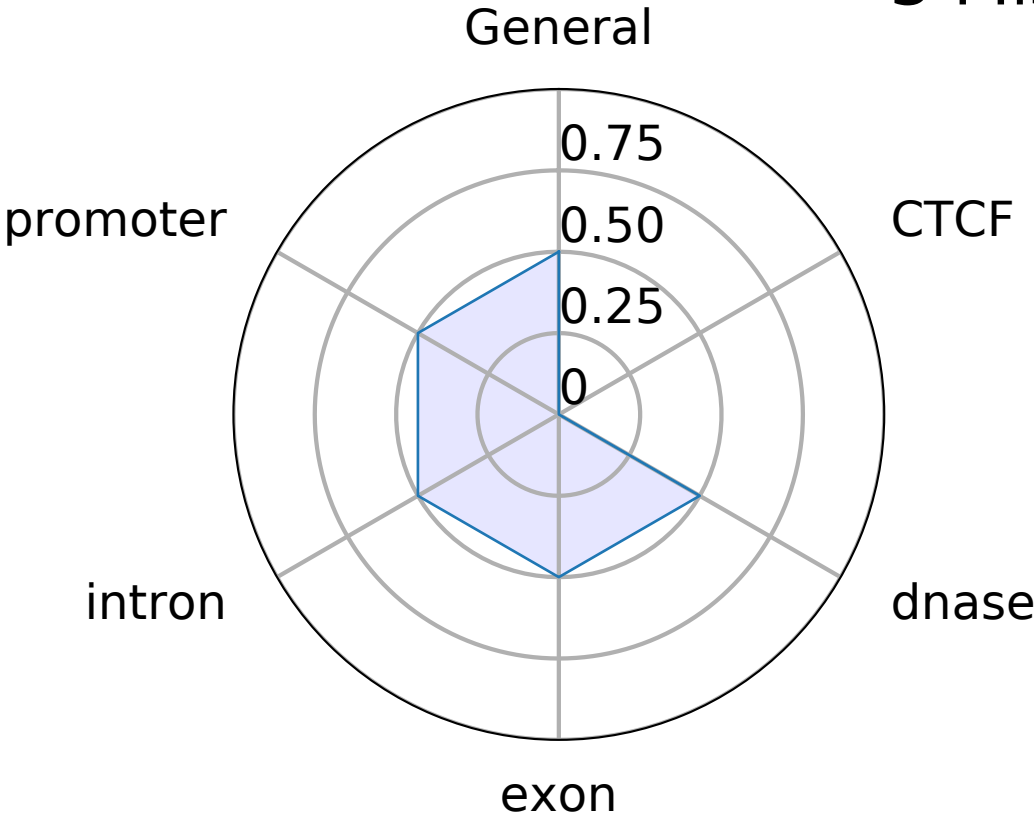


3 Mismatches



| | Position | # Targets |
|----------|----------|-----------|
| General | 0.5 | 35.0 |
| CTCF | 0.0 | 0.0 |
| dnase | 0.5 | 4.0 |
| exon | 0.5 | 2.0 |
| intron | 0.5 | 11.0 |
| promoter | 0.5 | 13.0 |

