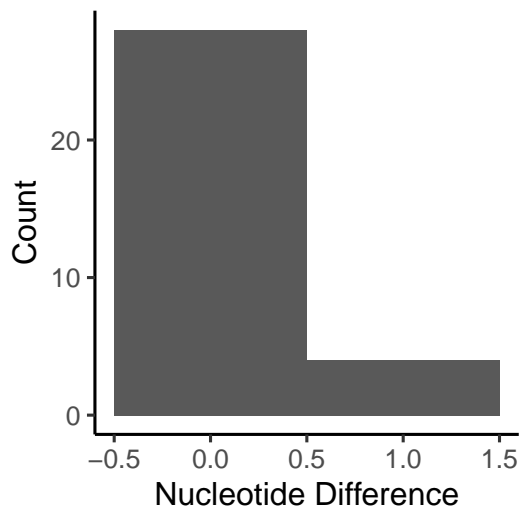


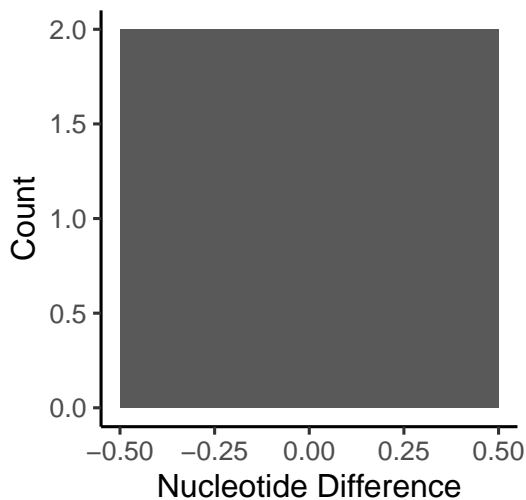
TRBV20-1*bp01

32 sequences assigned
28 (87.5%) exact matches, in which:
20 unique CDR3
9 unique J



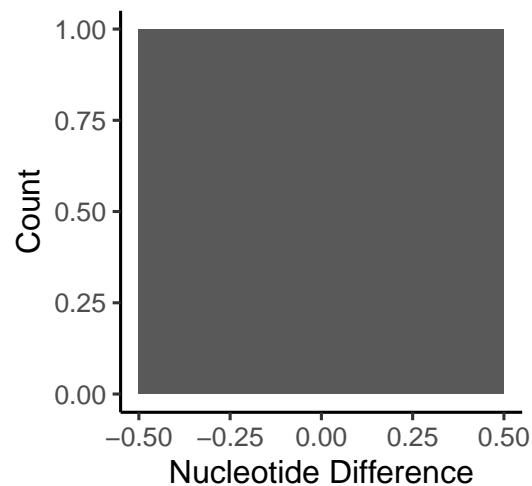
TRBV10-3*bp01

2 sequences assigned
2 (100%) exact matches, in which:
2 unique CDR3
2 unique J



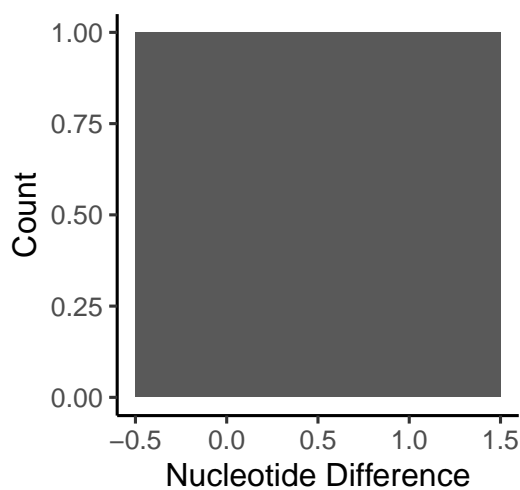
TRBV18*bp01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



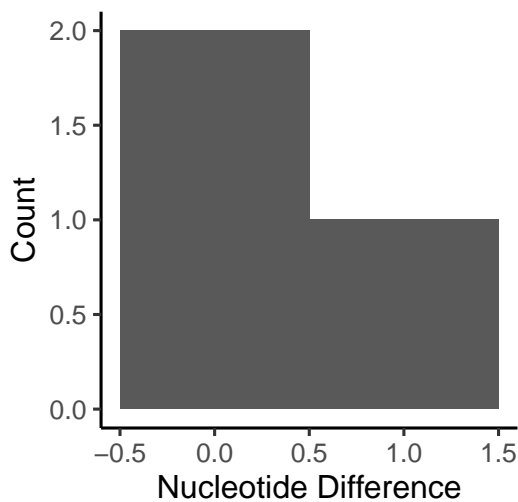
TRBV10-1*bp01

2 sequences assigned
1 (50%) exact matches, in which:
1 unique CDR3
1 unique J



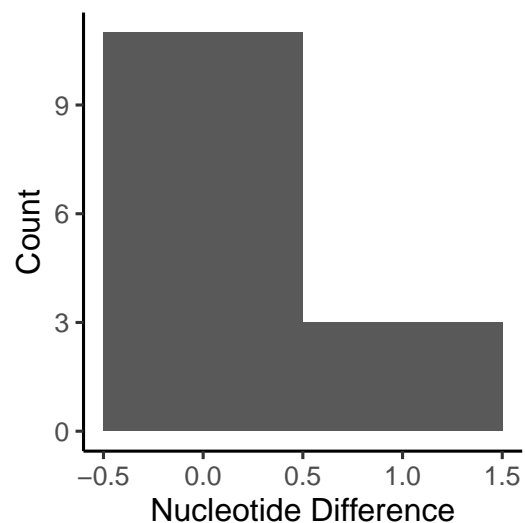
TRBV13*bp01

3 sequences assigned
2 (66.7%) exact matches, in which:
2 unique CDR3
1 unique J



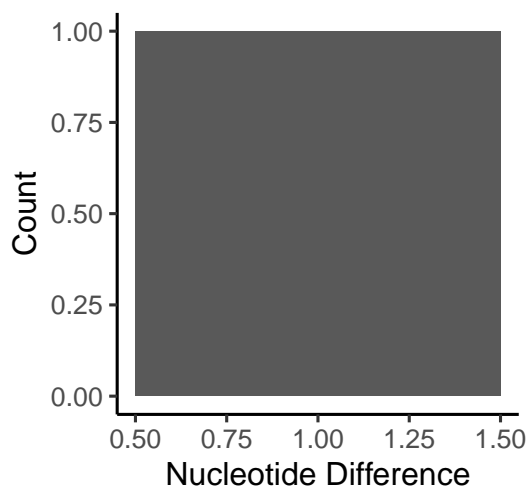
TRBV19*bp01

14 sequences assigned
11 (78.6%) exact matches, in which:
9 unique CDR3
7 unique J



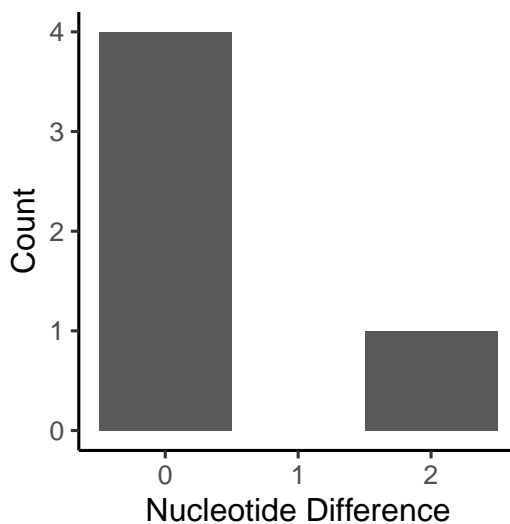
TRBV10-2*bp01

1 sequences assigned
No exact matches.



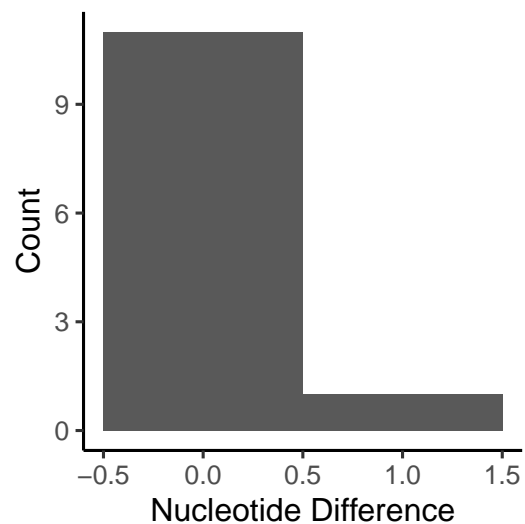
TRBV15*bp02

5 sequences assigned
4 (80%) exact matches, in which:
3 unique CDR3
3 unique J



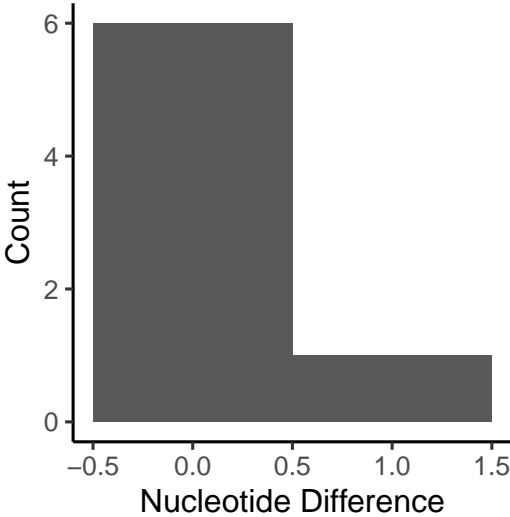
TRBV27*bp01

12 sequences assigned
11 (91.7%) exact matches, in which:
8 unique CDR3
5 unique J



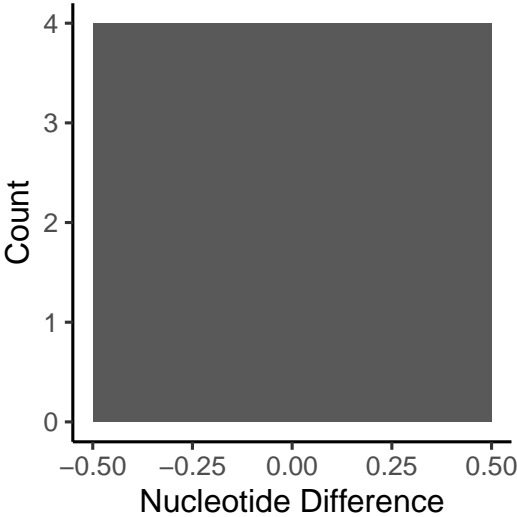
TRBV28*bp01

7 sequences assigned
6 (85.7%) exact matches, in which:
6 unique CDR3
4 unique J



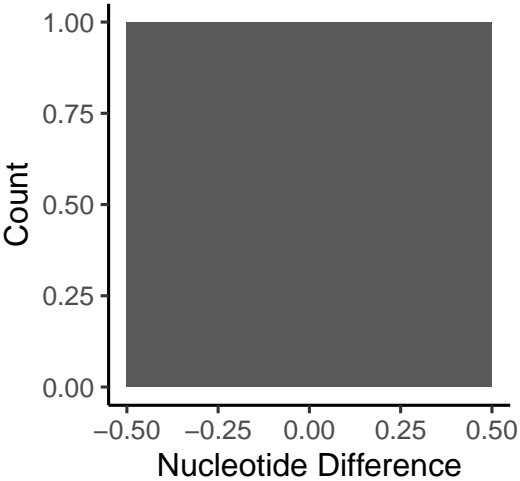
TRBV11-1*bp01

4 sequences assigned
4 (100%) exact matches, in which:
1 unique CDR3
1 unique J



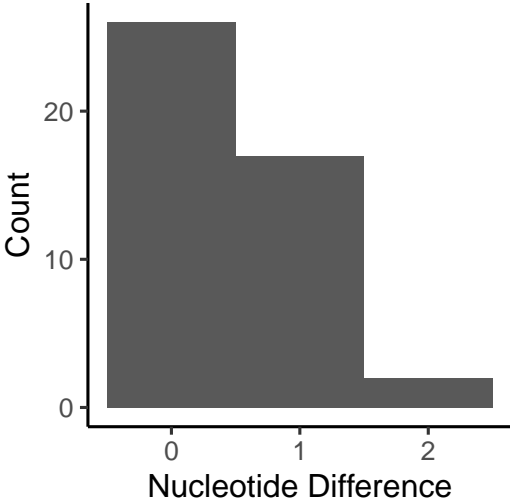
TRBV12-5*bp01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



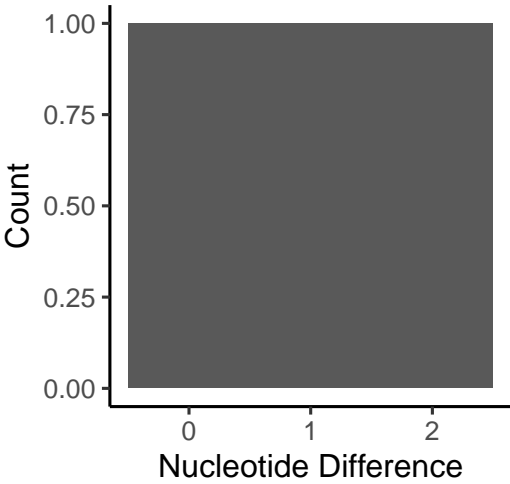
TRBV30*bp01

45 sequences assigned
26 (57.8%) exact matches, in which:
13 unique CDR3
4 unique J



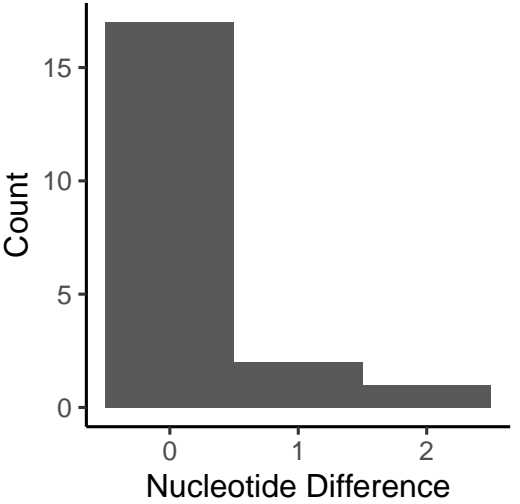
TRBV11-2*bp01

3 sequences assigned
1 (33.3%) exact matches, in which:
1 unique CDR3
1 unique J



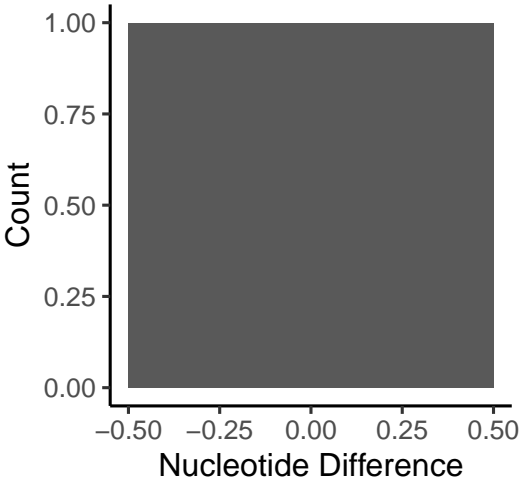
TRBV12-34*bp01

20 sequences assigned
17 (85%) exact matches, in which:
13 unique CDR3
7 unique J



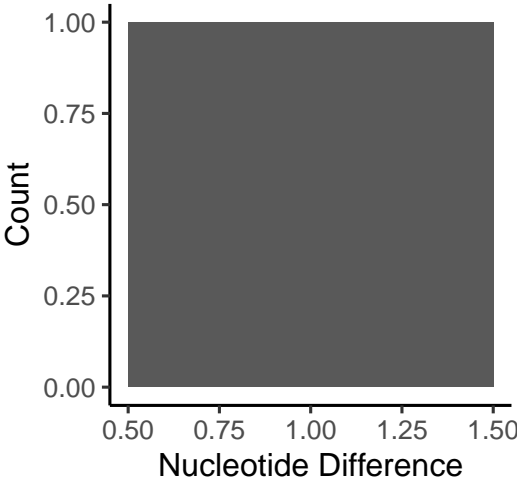
TRBV9*bp01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



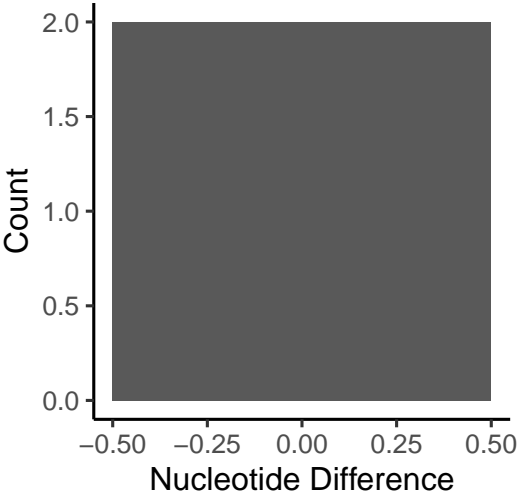
TRBV11-3*bp01

1 sequences assigned
No exact matches.



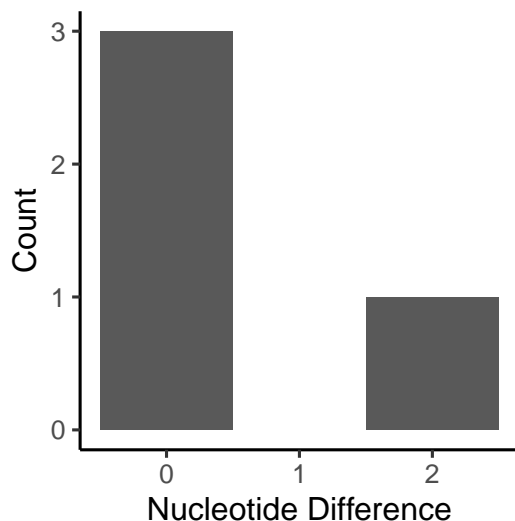
TRBV4-2*bp01

2 sequences assigned
2 (100%) exact matches, in which:
2 unique CDR3
2 unique J



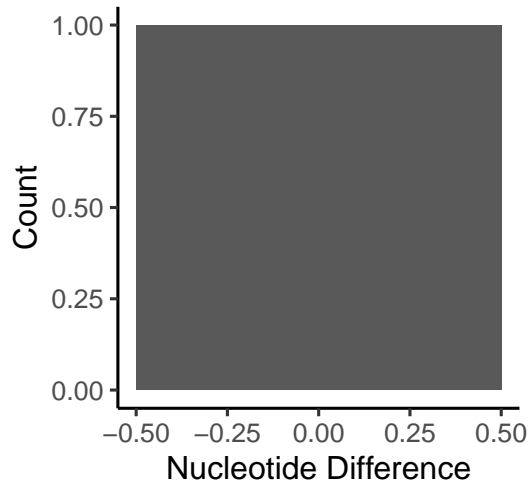
TRBV5-1*bp01

4 sequences assigned
3 (75%) exact matches, in which:
3 unique CDR3
2 unique J



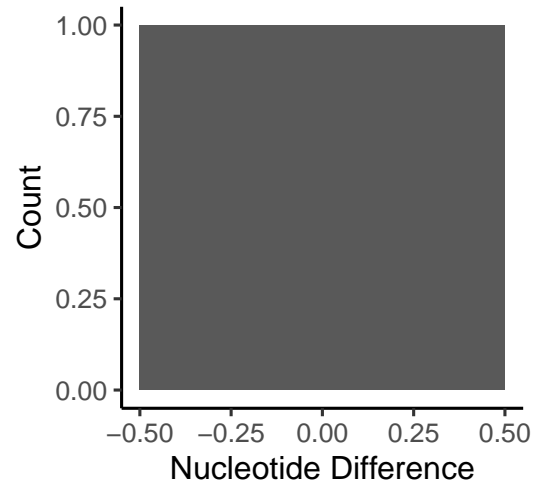
TRBV5-5*bp01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



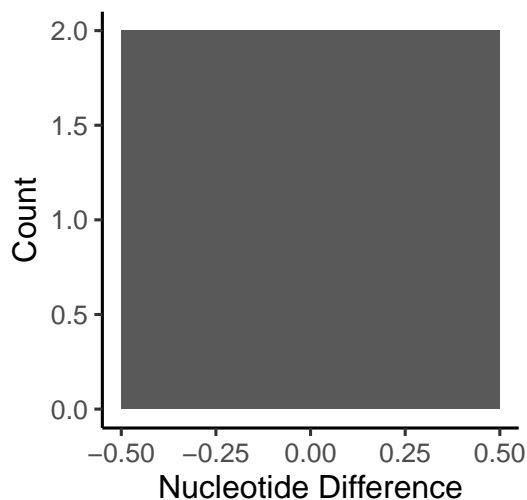
TRBV6-4*bp01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



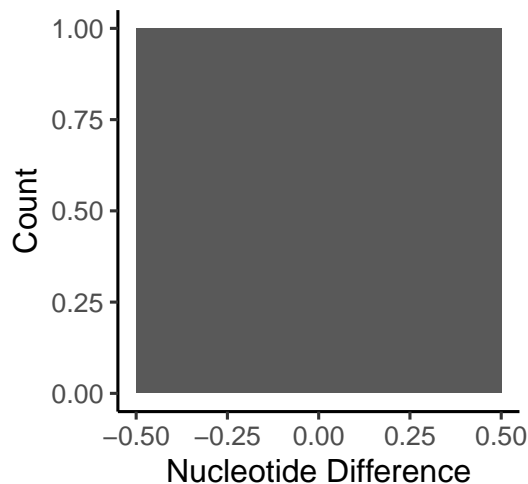
TRBV5-3*bp01

2 sequences assigned
2 (100%) exact matches, in which:
2 unique CDR3
2 unique J



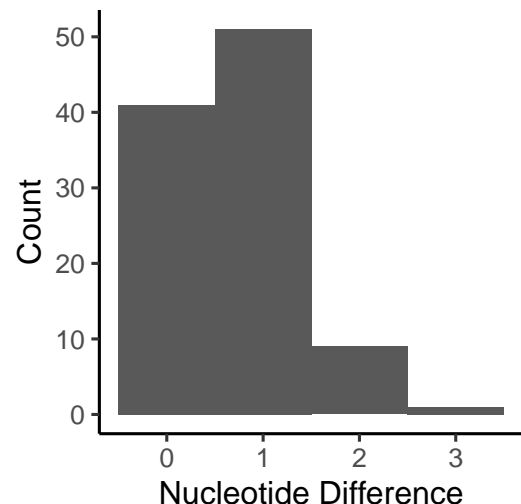
TRBV5-6*bp01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



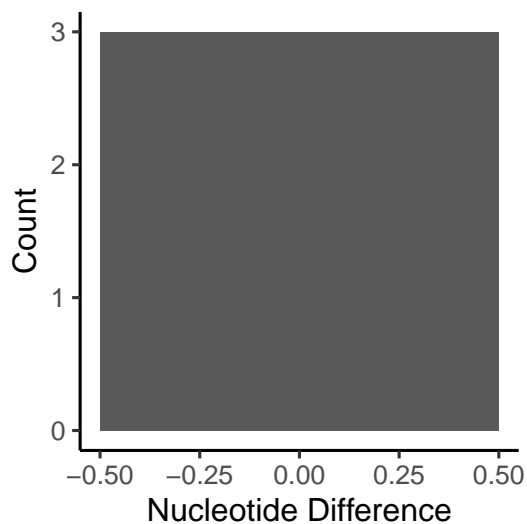
TRBV6-5*bp01

102 sequences assigned
41 (40.2%) exact matches, in which:
30 unique CDR3
8 unique J



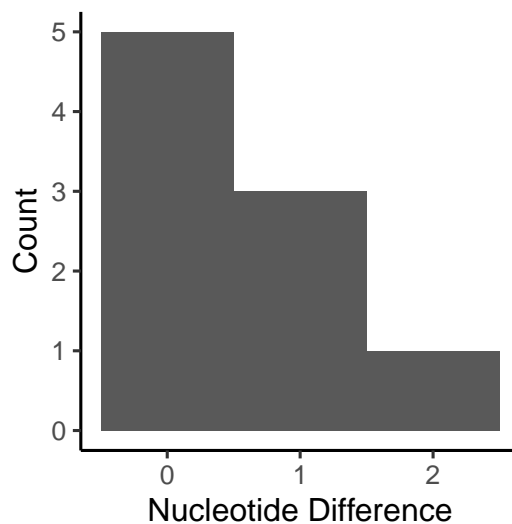
TRBV5-4*bp01

3 sequences assigned
3 (100%) exact matches, in which:
3 unique CDR3
3 unique J



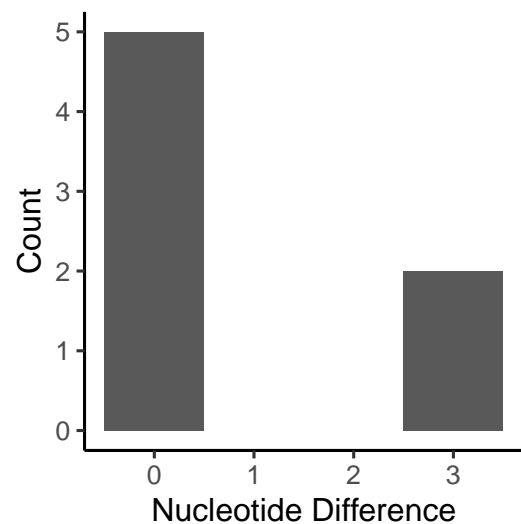
TRBV6-1*bp01

9 sequences assigned
5 (55.6%) exact matches, in which:
4 unique CDR3
3 unique J



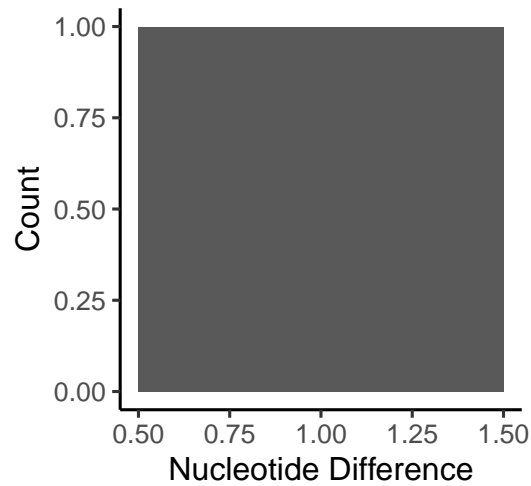
TRBV6-6*bp01

7 sequences assigned
5 (71.4%) exact matches, in which:
4 unique CDR3
3 unique J



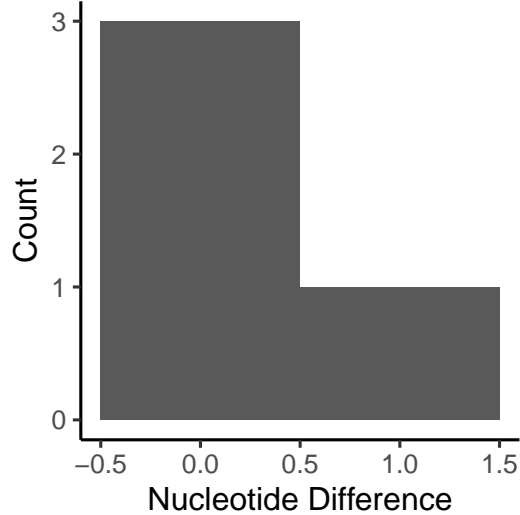
TRBV6–6*bp03

1 sequences assigned
No exact matches.



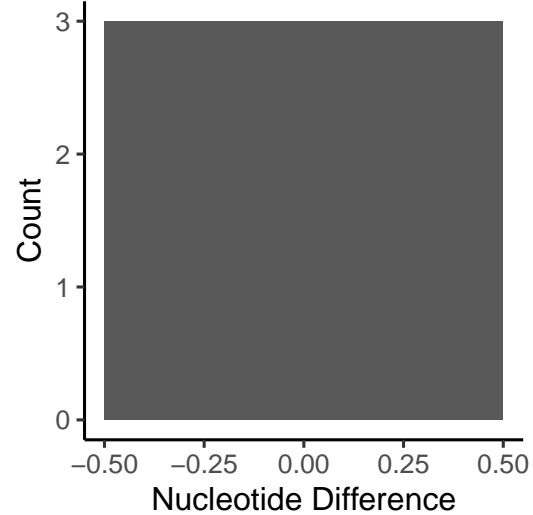
TRBV7–6*bp01

4 sequences assigned
3 (75%) exact matches, in which:
3 unique CDR3
1 unique J



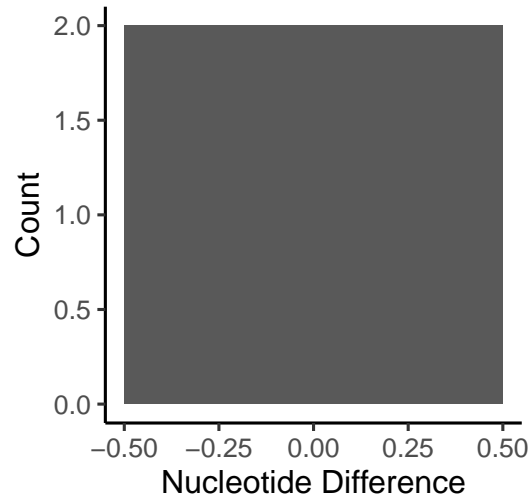
TRBV29–1*bp01

3 sequences assigned
3 (100%) exact matches, in which:
2 unique CDR3
2 unique J



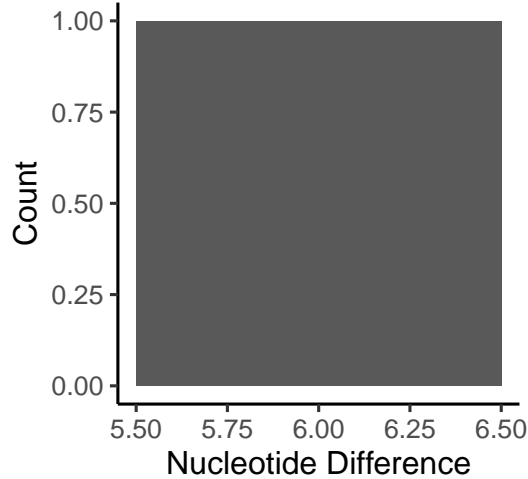
TRBV7–2*bp02

2 sequences assigned
2 (100%) exact matches, in which:
2 unique CDR3
1 unique J



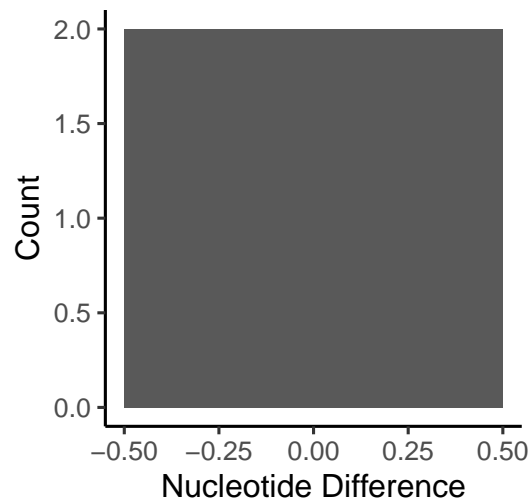
TRBV7–7*bp01

1 sequences assigned
No exact matches.



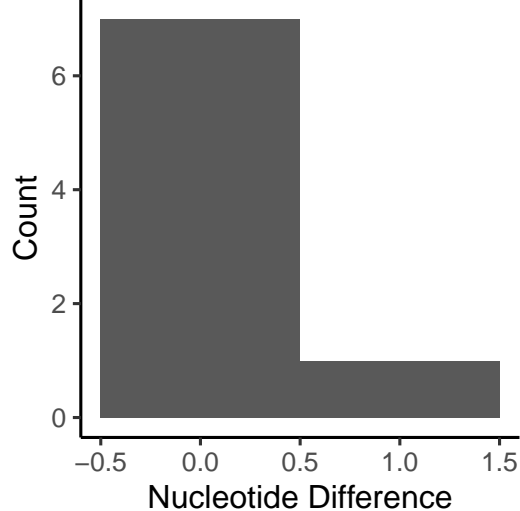
TRBV7–3*bp01

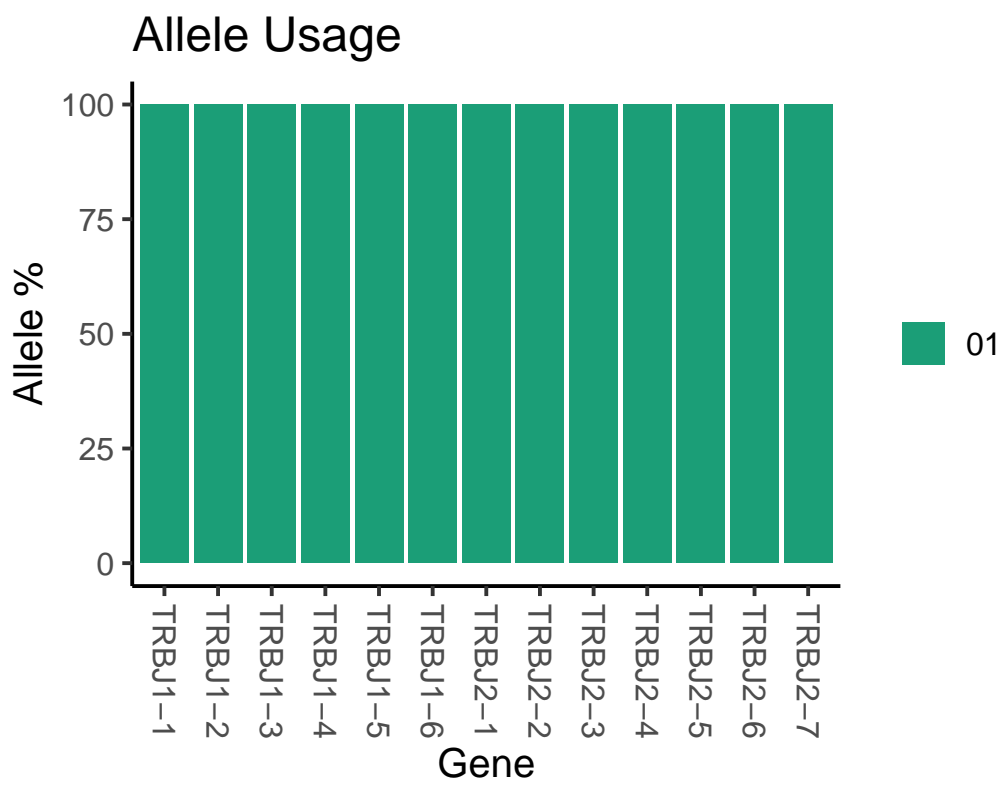
2 sequences assigned
2 (100%) exact matches, in which:
2 unique CDR3
2 unique J



TRBV7–9*bp01

8 sequences assigned
7 (87.5%) exact matches, in which:
7 unique CDR3
5 unique J





[illegible]