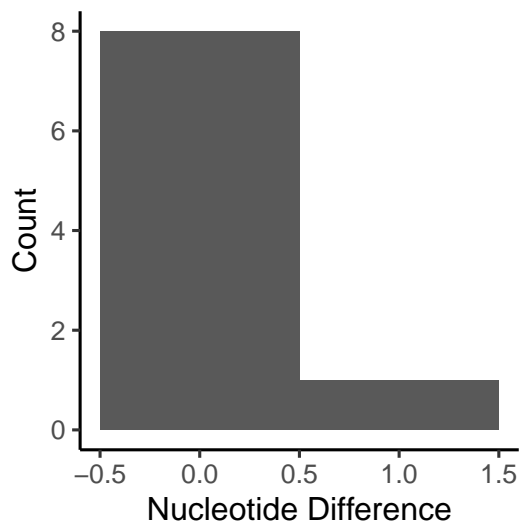


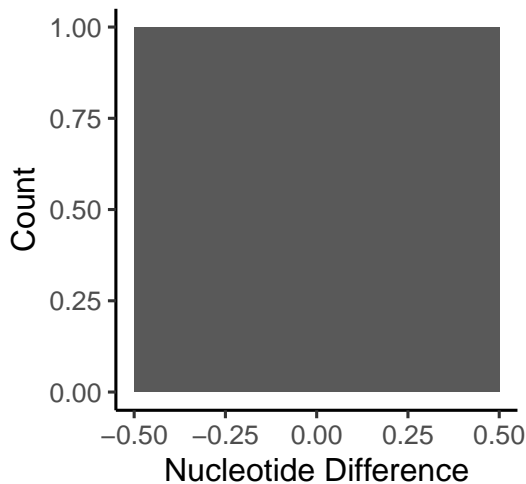
TRBV20–1*bp01

9 sequences assigned
8 (88.9%) exact matches, in which:
8 unique CDR3
7 unique J



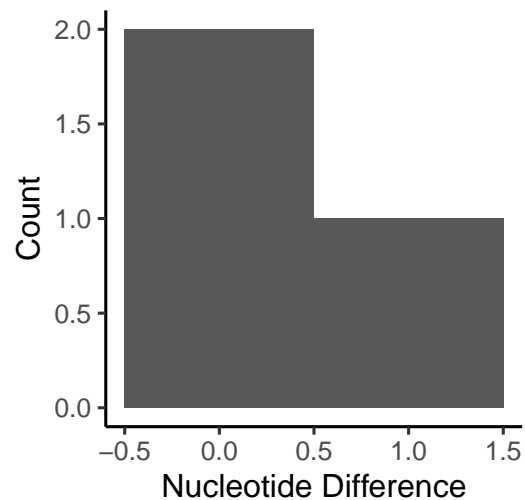
TRBV10–2*bp01

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



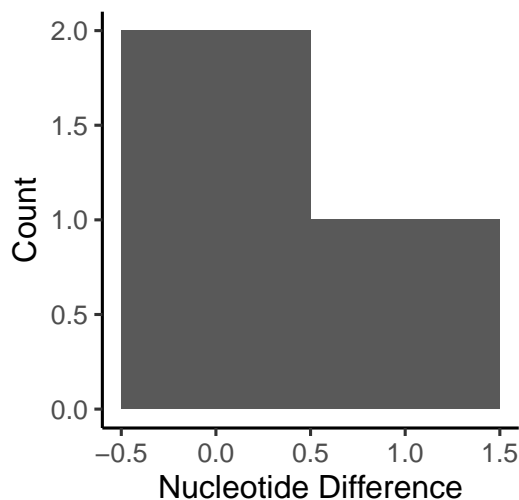
TRBV18*bp01

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



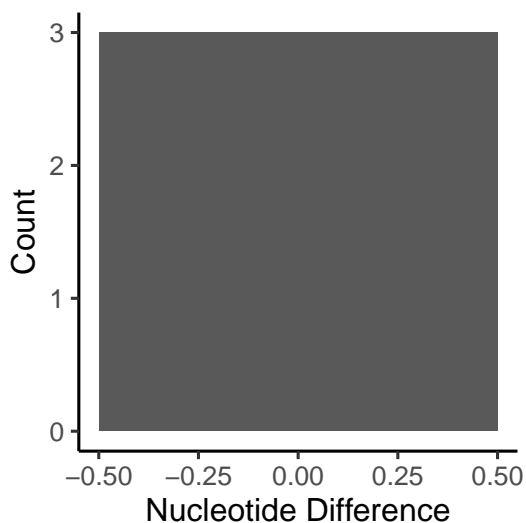
TRBV10–1*bp01

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



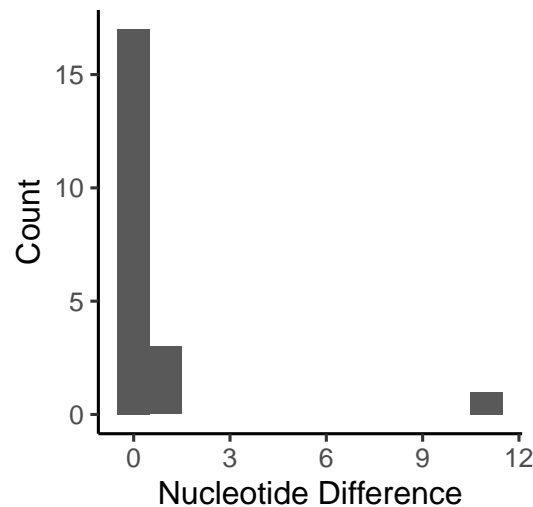
TRBV10–3*bp01

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



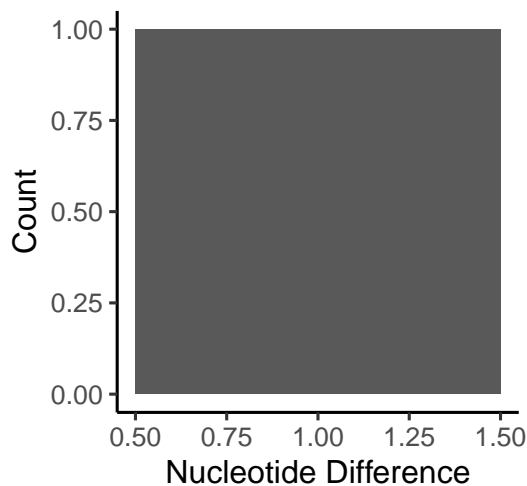
TRBV19*bp01

21 sequences assigned
17 (81%) exact matches, in which:
16 unique CDR3
9 unique J



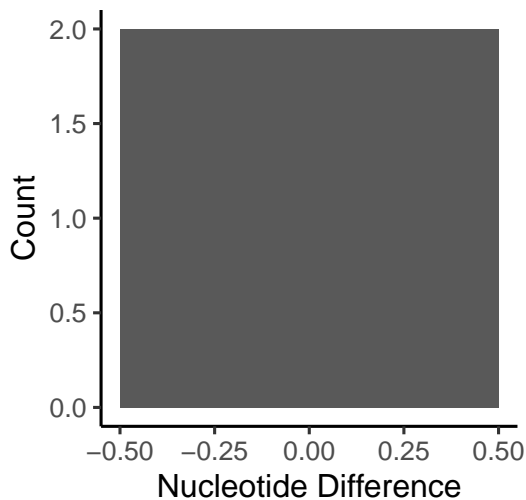
TRBV10–1*bp02

1 sequences assigned
No exact matches.



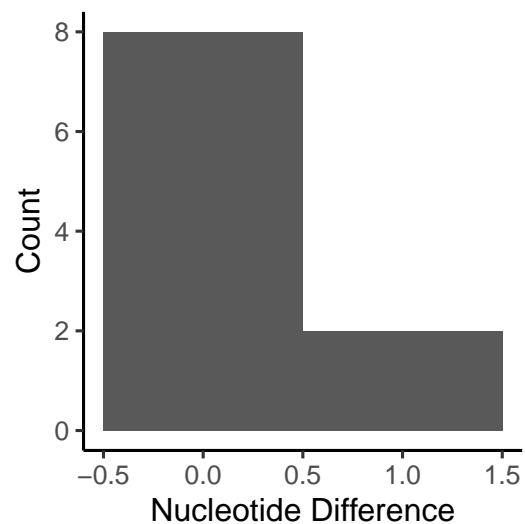
TRBV15*bp02

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



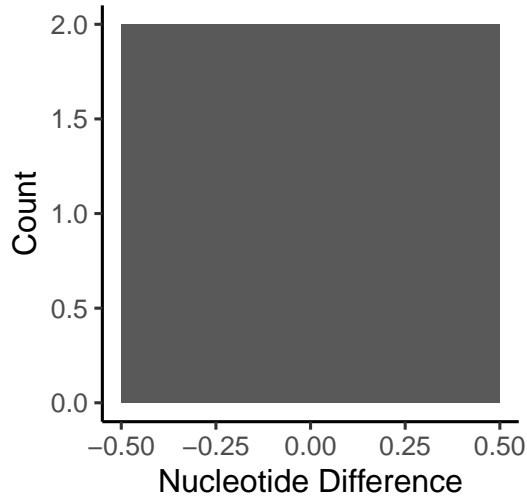
TRBV27*bp01

10 sequences assigned
8 (80%) exact matches, in which:
6 unique CDR3
4 unique J



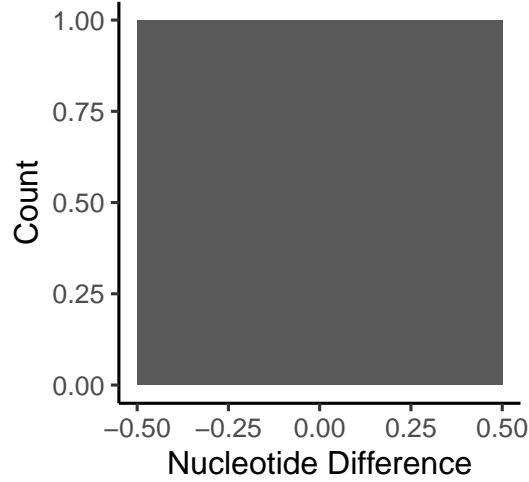
TRBV28*bp01

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



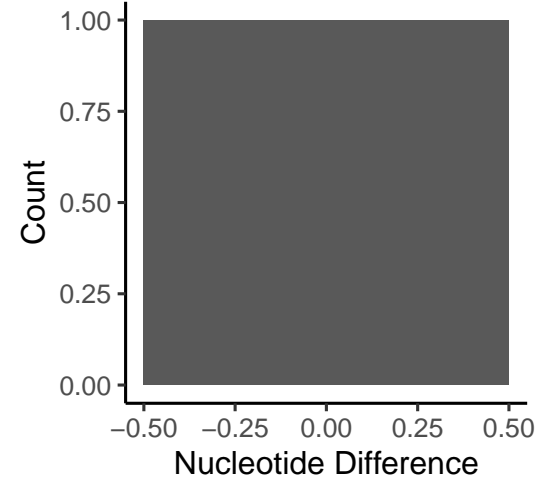
TRBV11-3*bp01

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



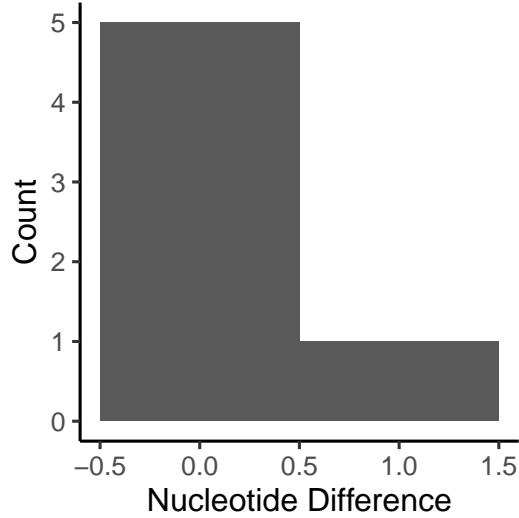
TRBV3-1*bp01

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



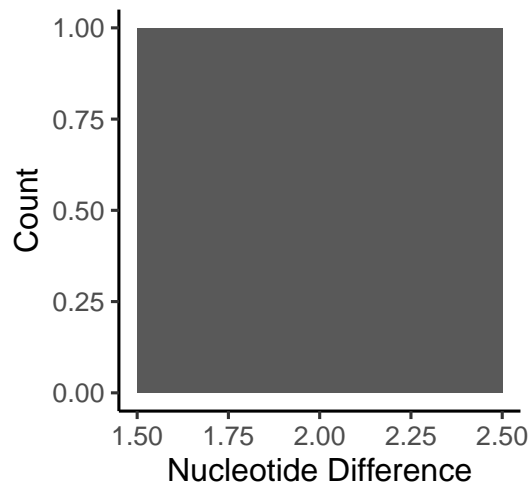
TRBV30*bp01

6 sequences assigned
5 (83.3%) exact matches, in which:
5 unique CDR3
5 unique J



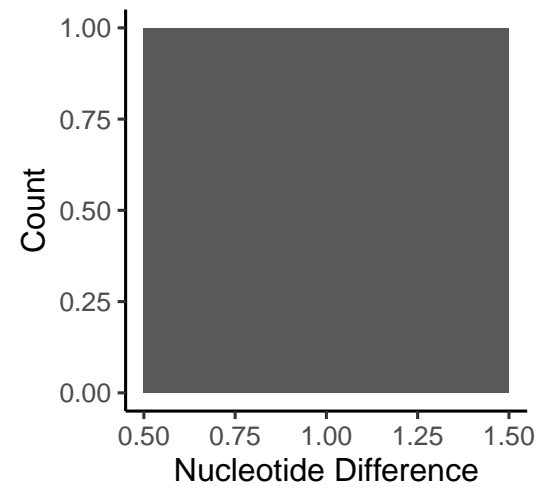
TRBV12-5*bp01

1 sequences assigned
No exact matches.



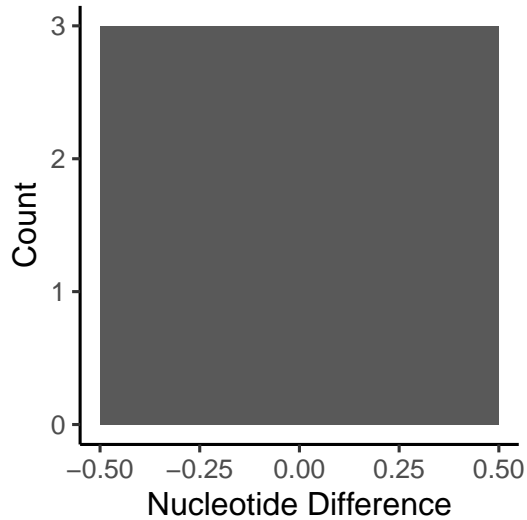
TRBV23-1*bp01

1 sequences assigned
No exact matches.



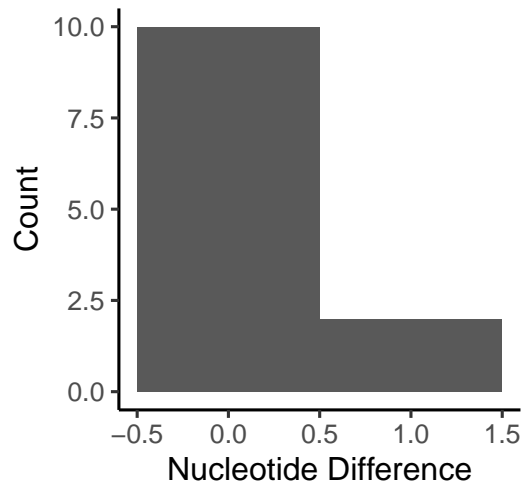
TRBV11-2*bp01

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



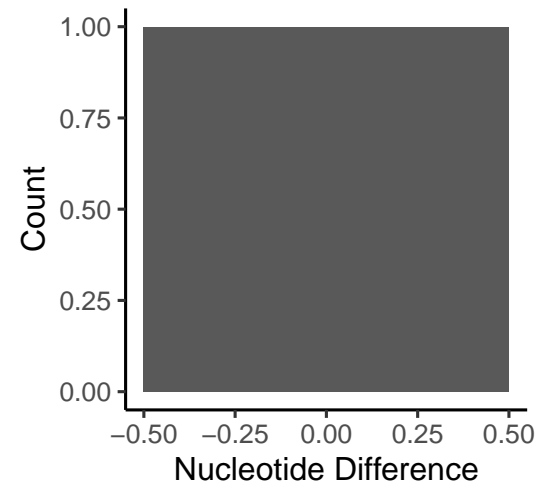
TRBV12-34*bp01

12 sequences assigned
10 (83.3%) exact matches, in which:
9 unique CDR3
5 unique J



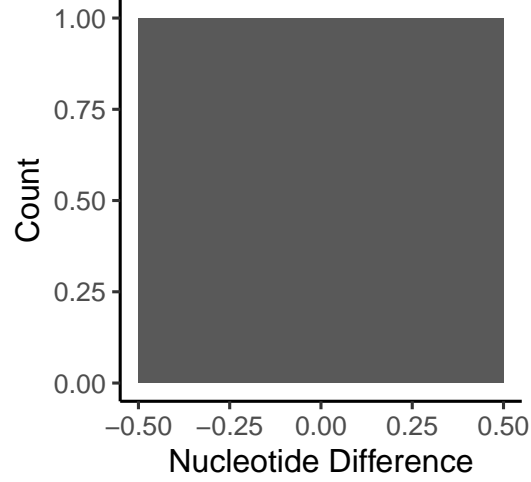
TRBV24-1*bp01

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



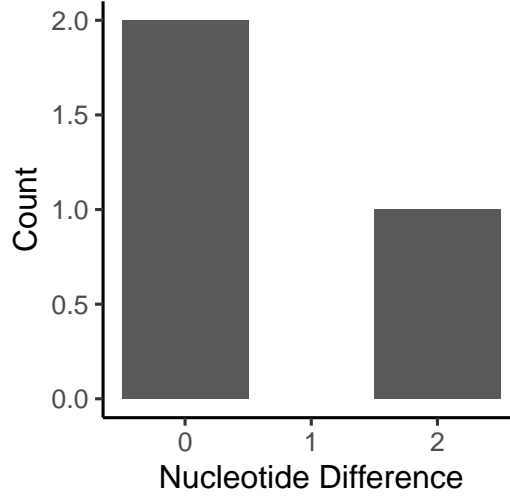
TRBV4-3*bp01

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



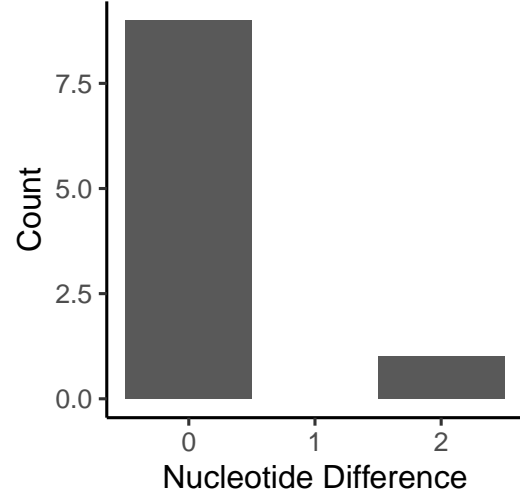
TRBV5-4*bp01

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



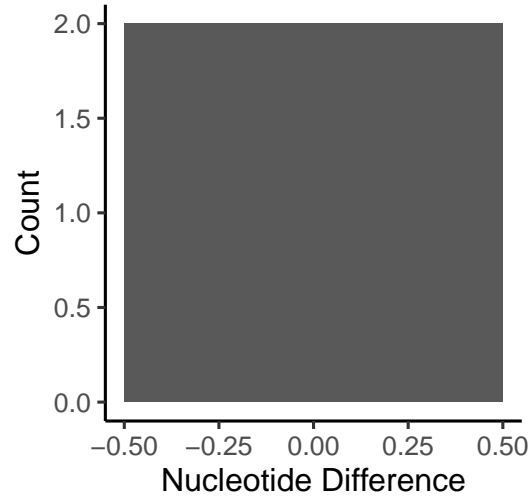
TRBV6-1*bp01

10 sequences assigned
9 (90%) exact matches, in which:
9 unique CDR3
7 unique J



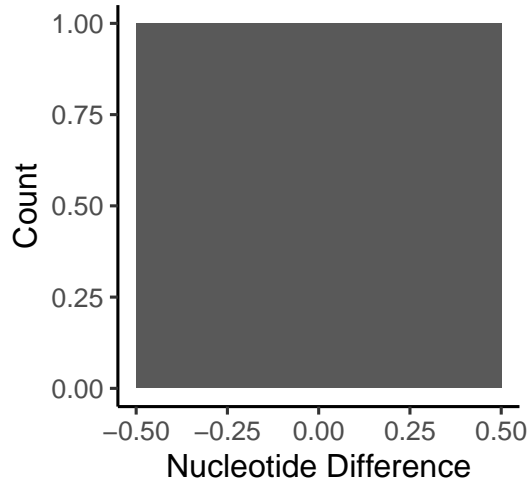
TRBV5-1*bp01

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



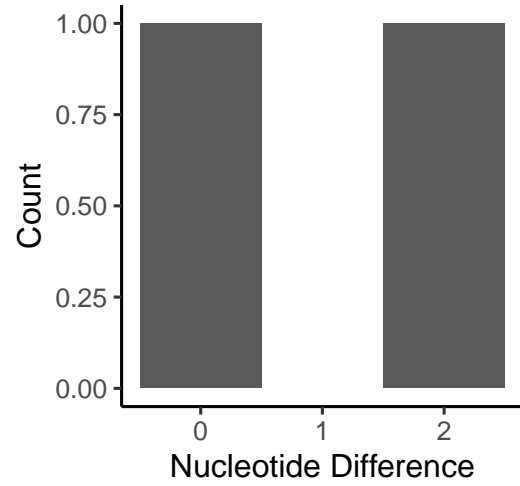
TRBV5-5*bp01

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



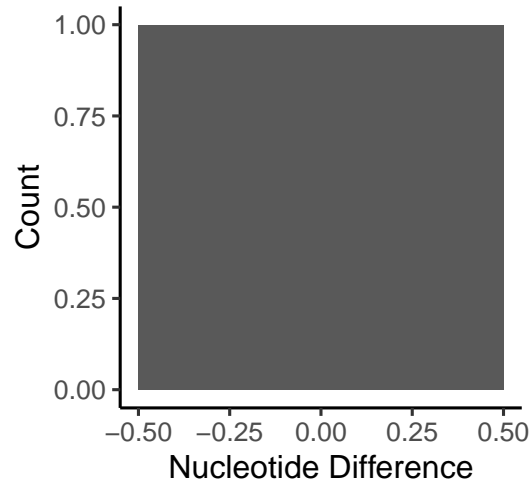
TRBV6-4*bp01

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



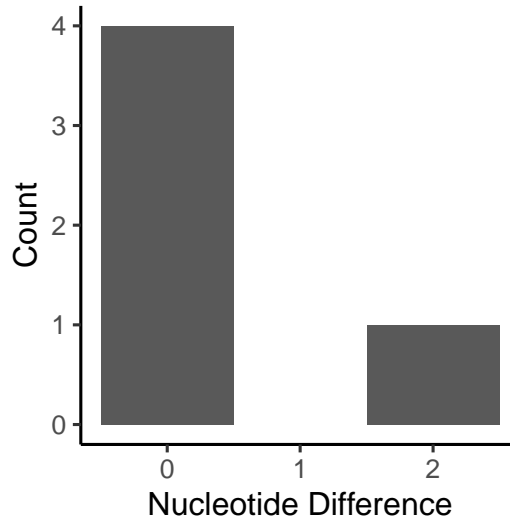
TRBV25-1*bp01

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



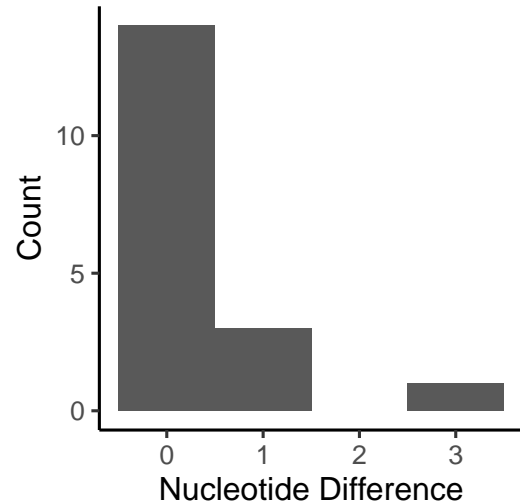
TRBV5-6*bp01

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



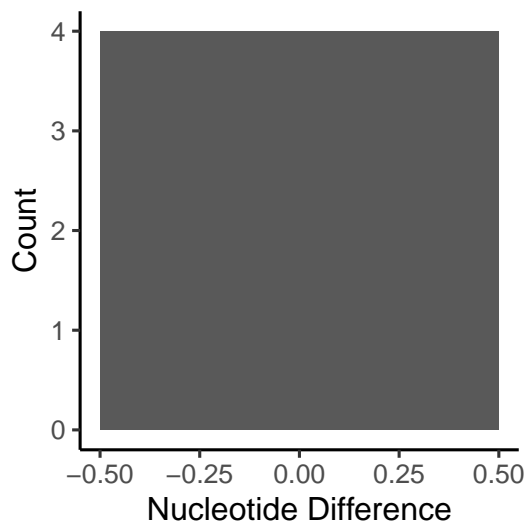
TRBV6-5*bp01

18 sequences assigned
14 (77.8%) exact matches, in which:
14 unique CDR3
8 unique J



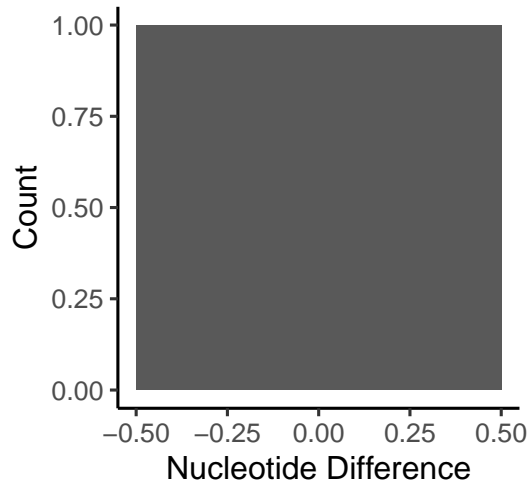
TRBV6–6*bp01

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



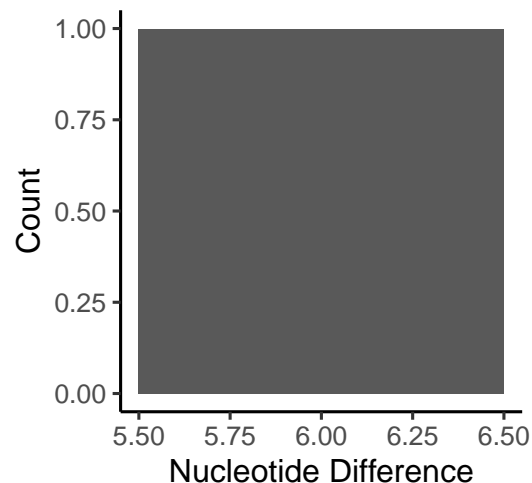
TRBV7–3*bp01

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



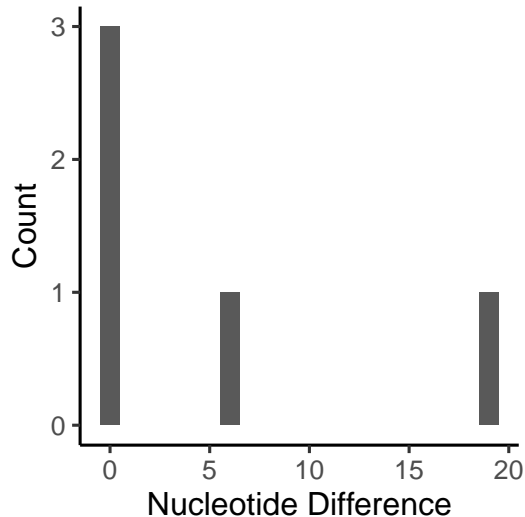
TRBV7–7*bp01

1 sequences assigned
No exact matches.



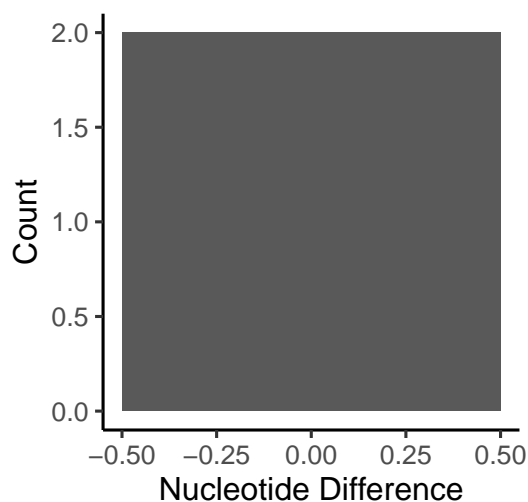
TRBV6–23*bp01

5 sequences assigned
3 (60%) exact matches, in which:
3 unique CDR3
3 unique J



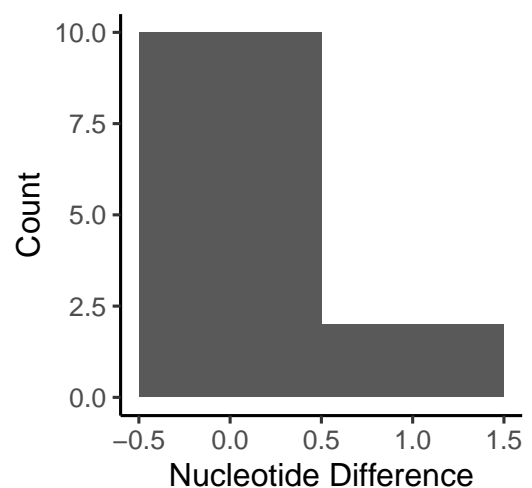
TRBV7–4*bp01

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



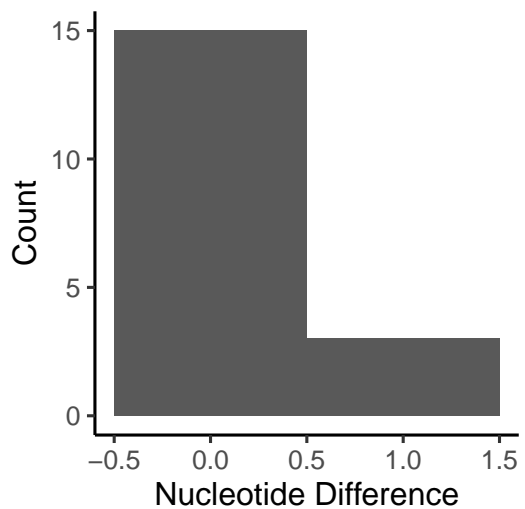
TRBV7–9*bp01

12 sequences assigned
10 (83.3%) exact matches, in which:
9 unique CDR3
7 unique J



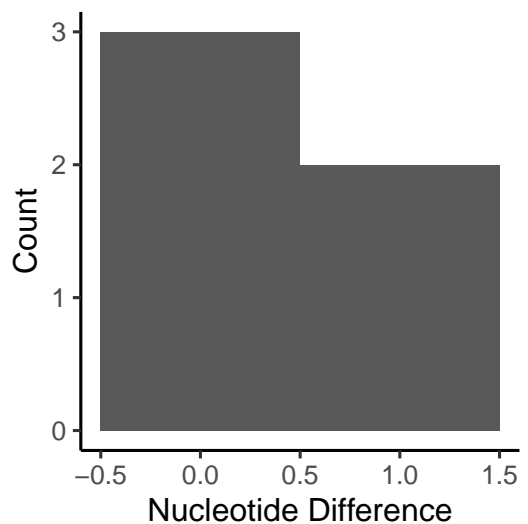
TRBV7–2*bp01

18 sequences assigned
15 (83.3%) exact matches, in which:
14 unique CDR3
9 unique J



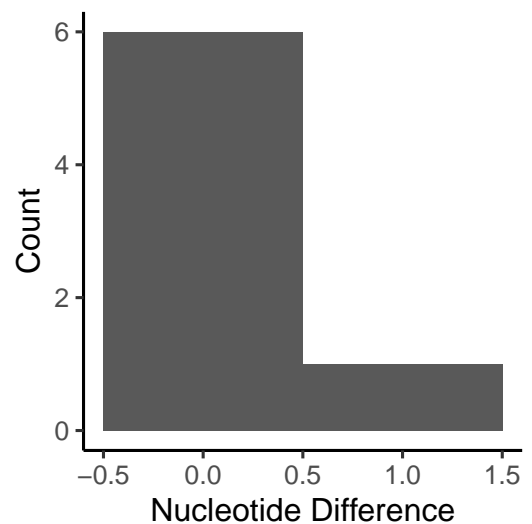
TRBV7–6*bp01

5 sequences assigned
3 (60%) exact matches, in which:
3 unique CDR3
3 unique J



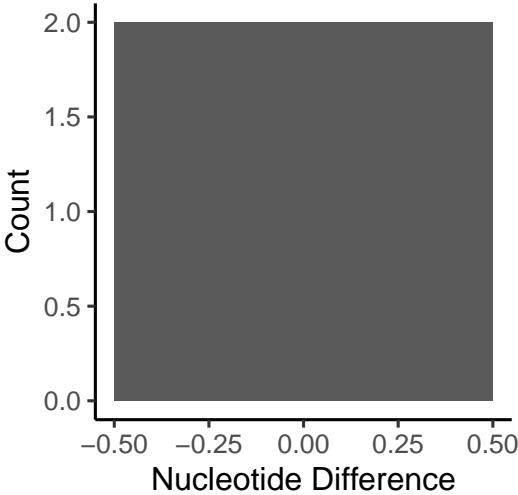
TRBV29–1*bp01

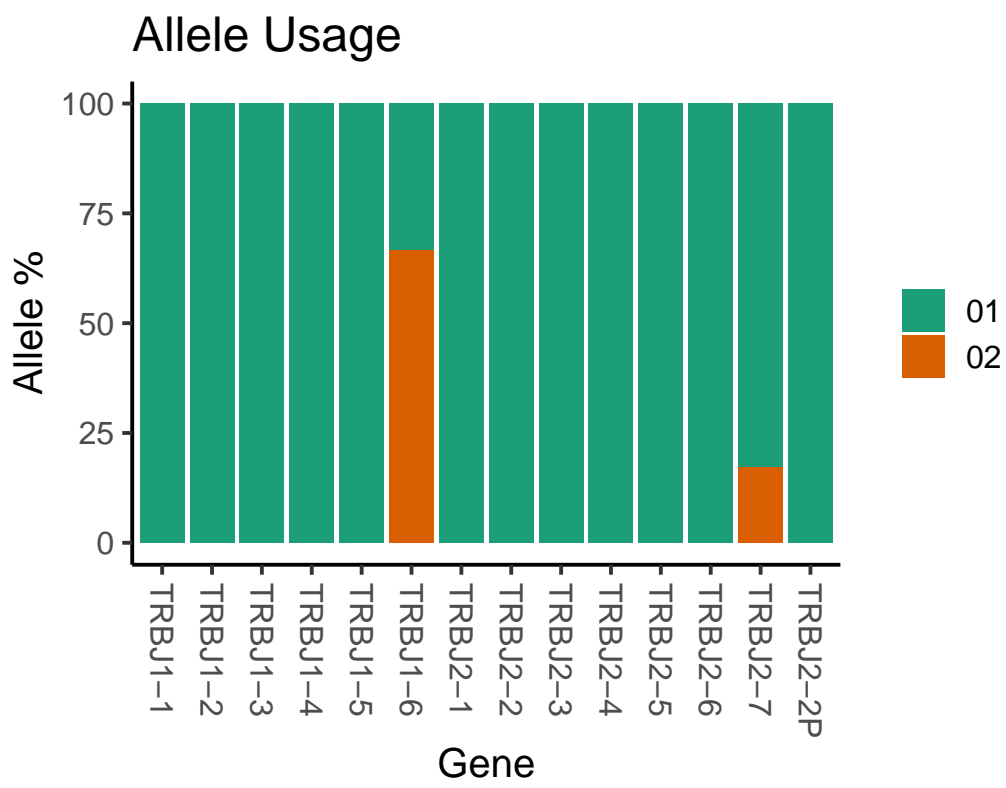
7 sequences assigned
6 (85.7%) exact matches, in which:
5 unique CDR3
5 unique J



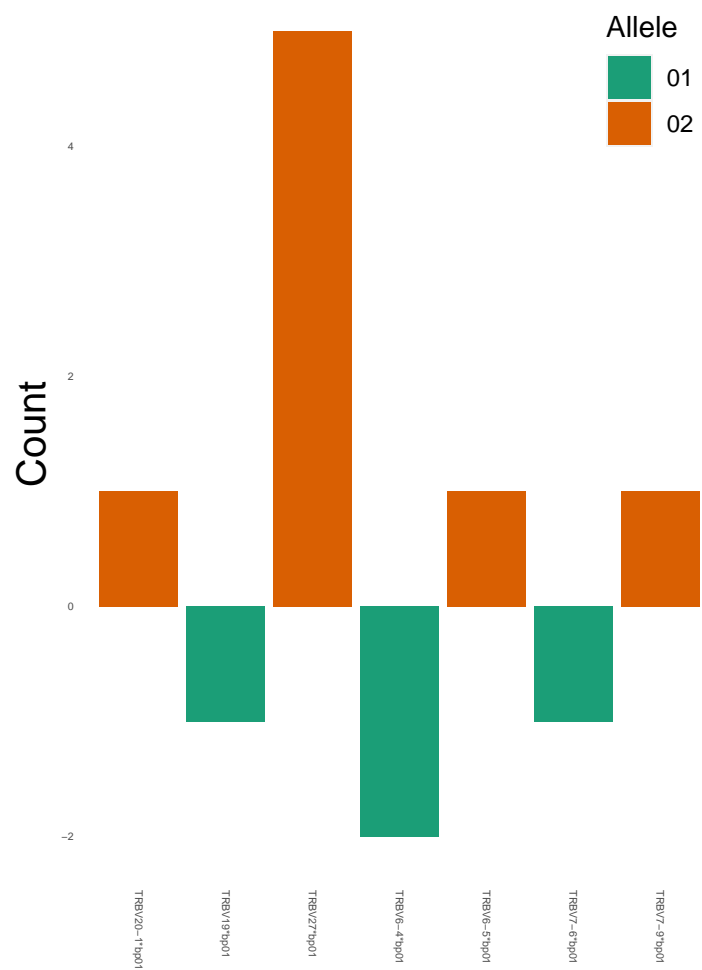
TRBV29-1*bp03

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





Sequence Count by TRBJ1–6 allele usage



[illegible]