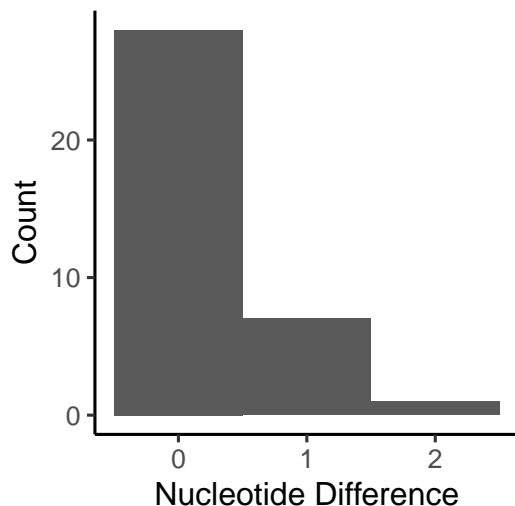


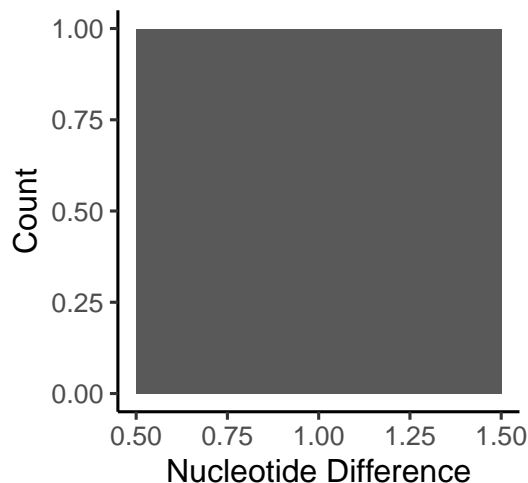
TRBV20–1*bp01

36 sequences assigned
28 (77.8%) exact matches, in which:
25 unique CDR3
12 unique J



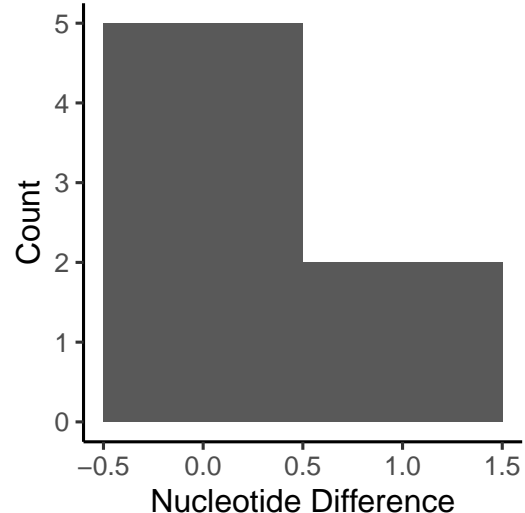
TRBV10–3*bp01

1 sequences assigned
No exact matches.



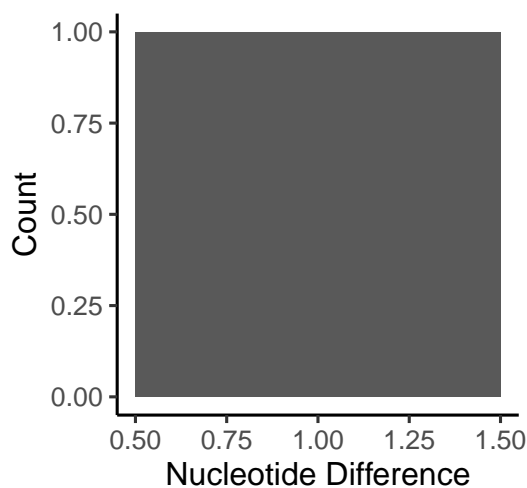
TRBV18*bp01

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



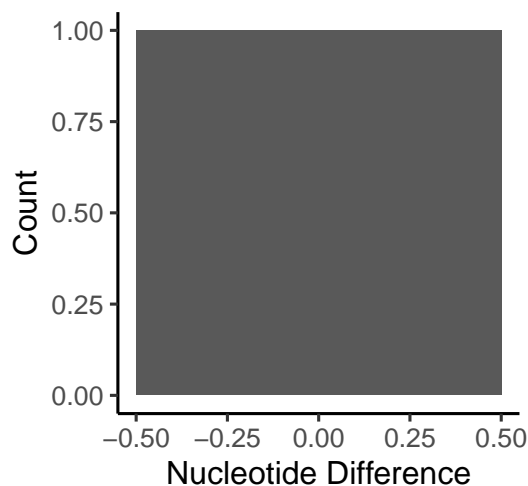
TRBV10–1*bp02

1 sequences assigned
No exact matches.



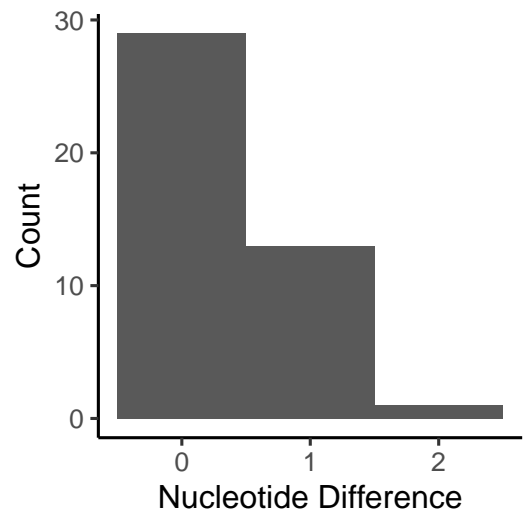
TRBV13*bp01

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



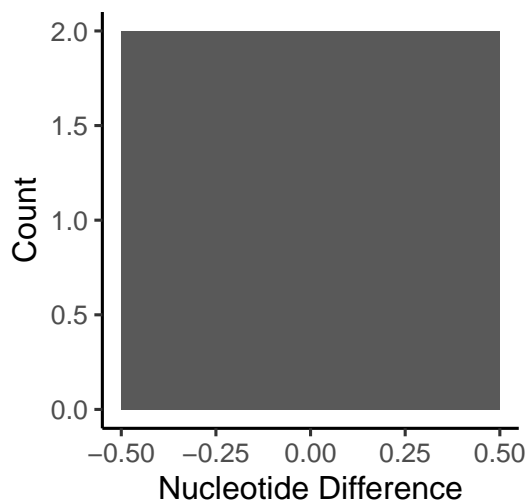
TRBV19*bp01

43 sequences assigned
29 (67.4%) exact matches, in which:
27 unique CDR3
9 unique J



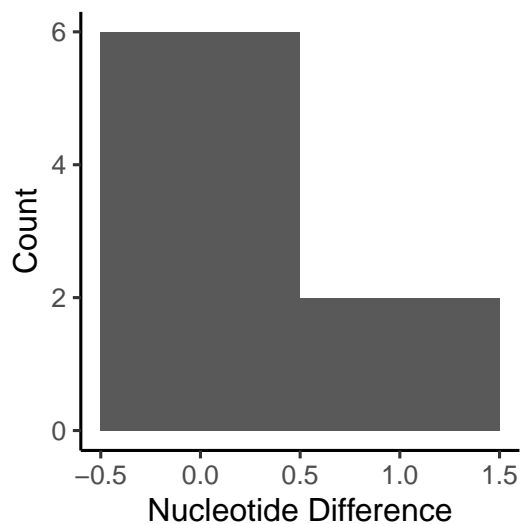
TRBV10–2*bp01

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



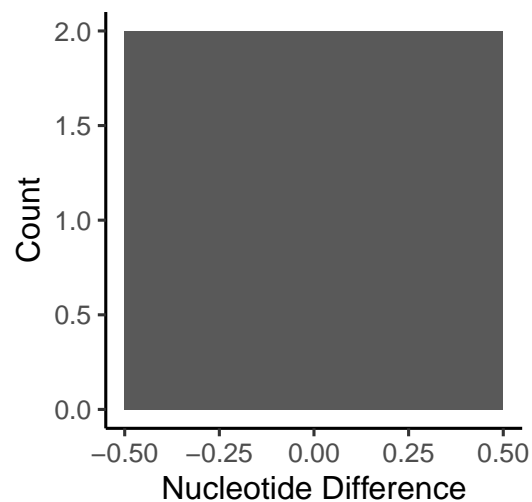
TRBV15*bp02

8 sequences assigned
6 (75%) exact matches, in which:
6 unique CDR3
5 unique J



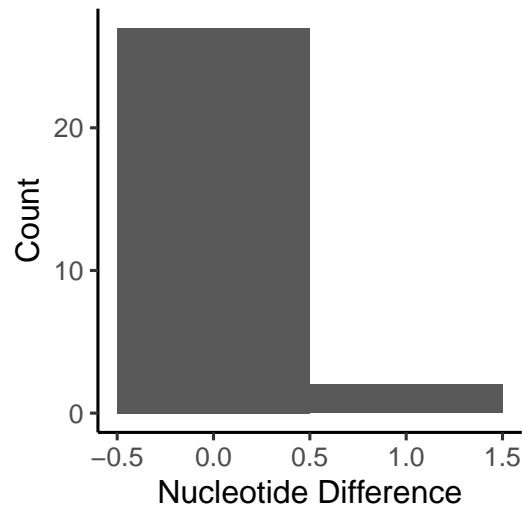
TRBV2*bp01

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



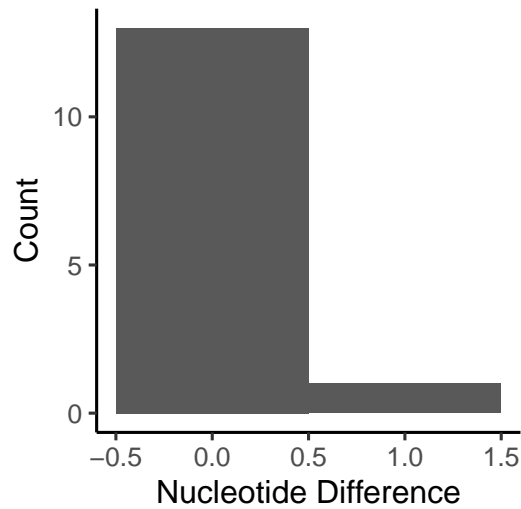
TRBV27*bp01

29 sequences assigned
27 (93.1%) exact matches, in which:
25 unique CDR3
9 unique J



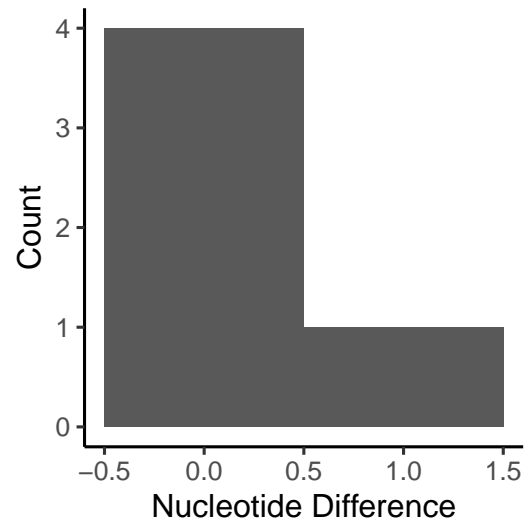
TRBV30*bp02

14 sequences assigned
13 (92.9%) exact matches, in which:
13 unique CDR3
6 unique J



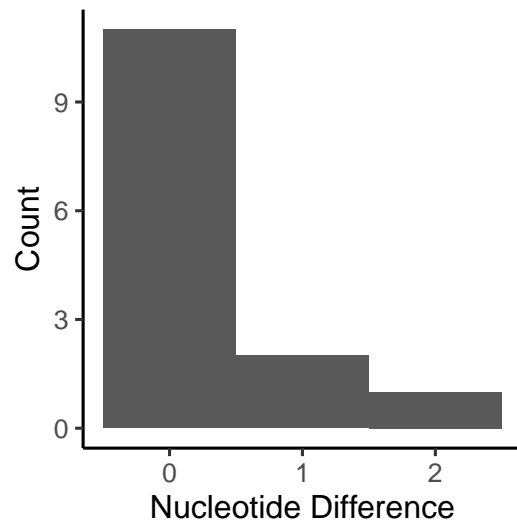
TRBV21-1*bp01

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



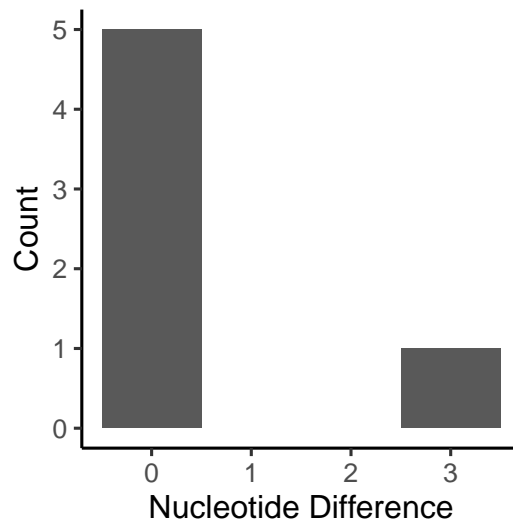
TRBV28*bp01

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



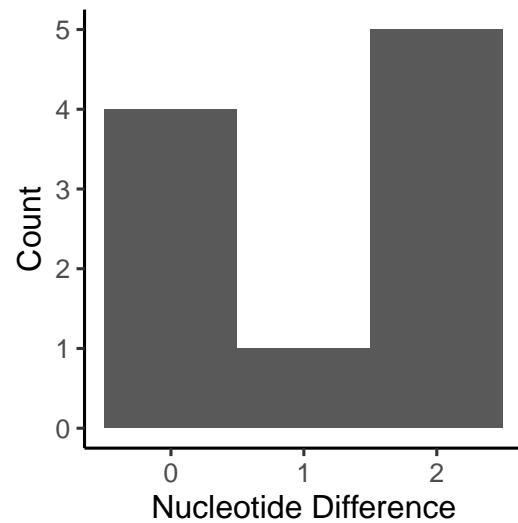
TRBV9*bp01

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



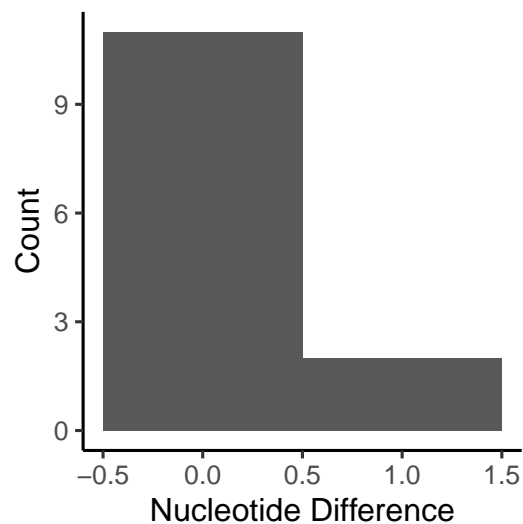
TRBV11-2*bp01

10 sequences assigned
4 (40%) exact matches, in which:
4 unique CDR3
4 unique J



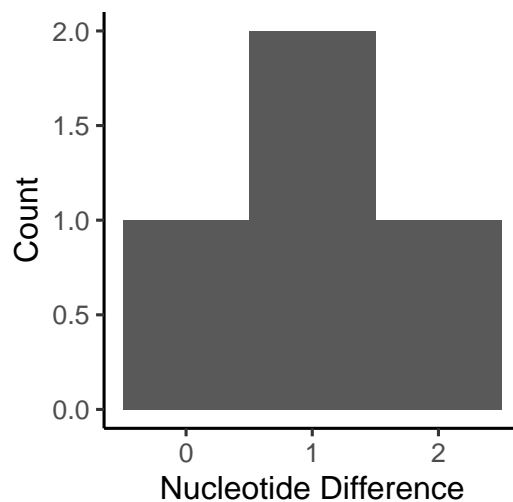
TRBV30*bp01

13 sequences assigned
11 (84.6%) exact matches, in which:
11 unique CDR3
6 unique J



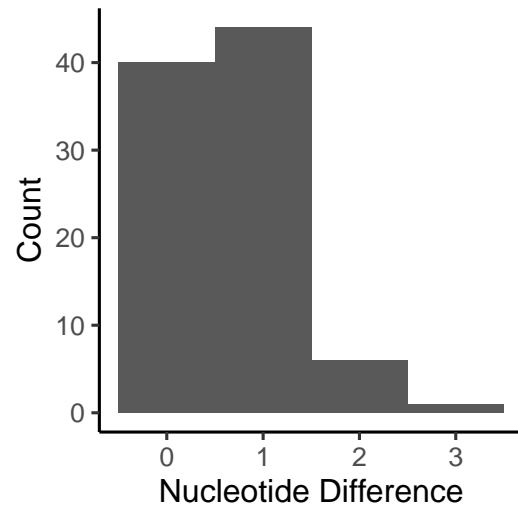
TRBV11-1*bp01

4 sequences assigned
1 (25%) exact matches, in which:
1 unique CDR3
1 unique J



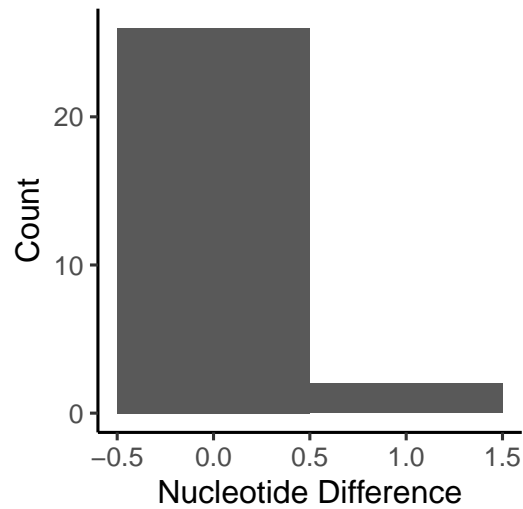
TRBV11-3*bp01

91 sequences assigned
40 (44%) exact matches, in which:
32 unique CDR3
8 unique J



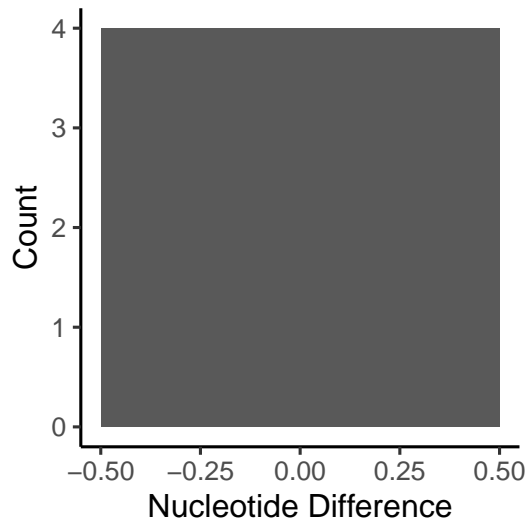
TRBV12–34*bp01

28 sequences assigned
26 (92.9%) exact matches, in which:
24 unique CDR3
8 unique J



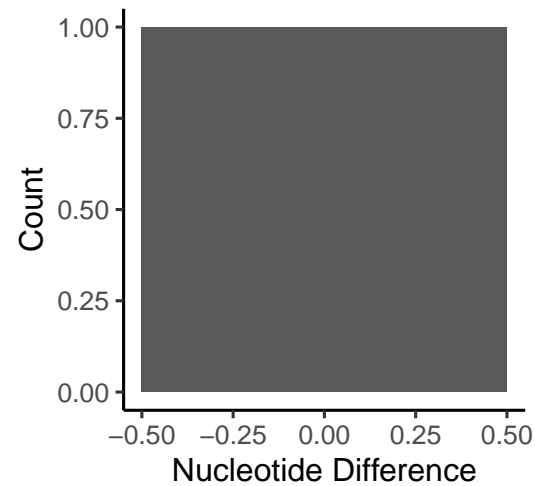
TRBV4–3*bp01

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



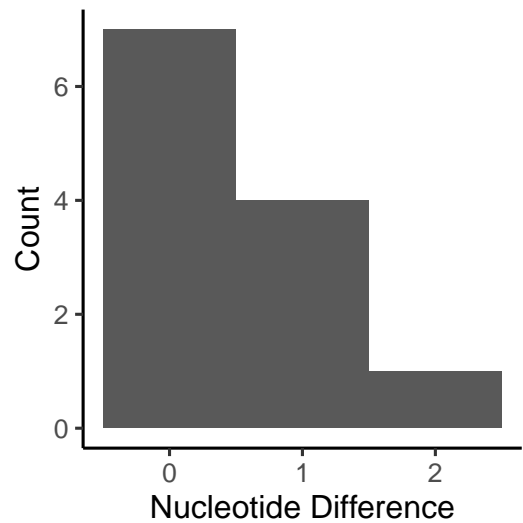
TRBV5–4*bp01

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



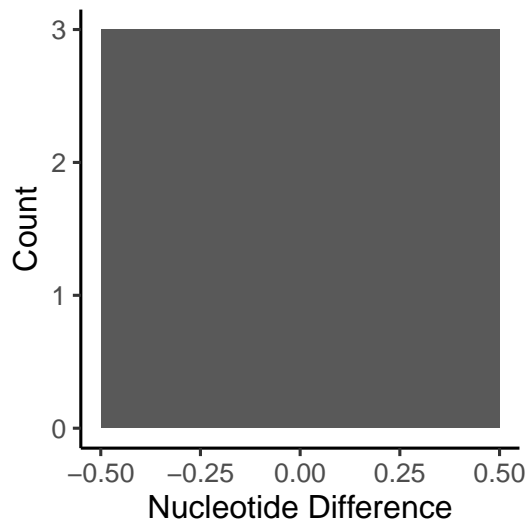
TRBV24–1*bp01

12 sequences assigned
7 (58.3%) exact matches, in which:
7 unique CDR3
5 unique J



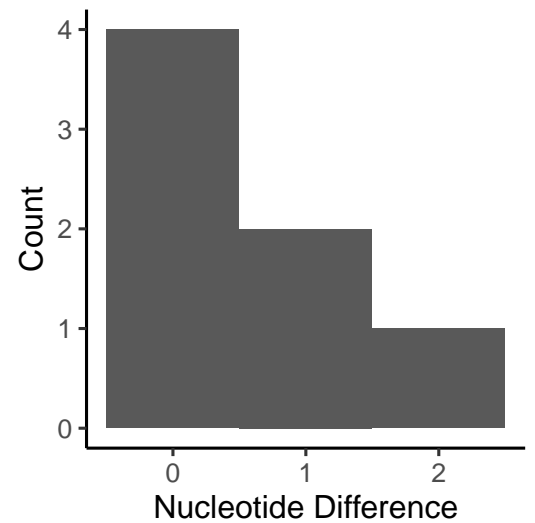
TRBV5–1*bp01

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



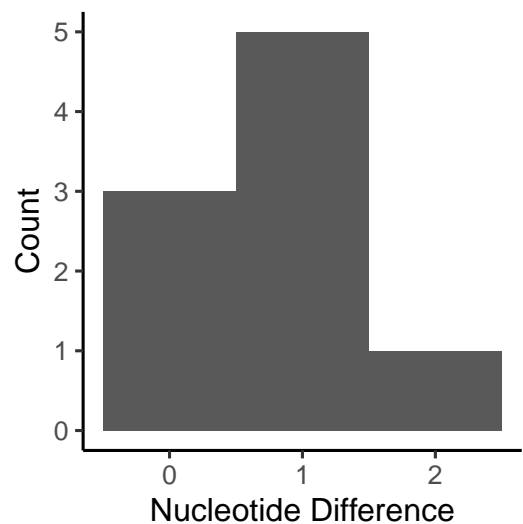
TRBV5–5*bp01

7 sequences assigned
4 (57.1%) exact matches, in which:
4 unique CDR3
4 unique J



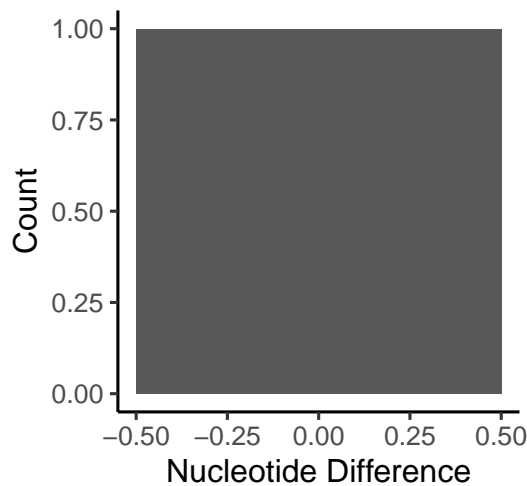
TRBV4–2*bp01

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



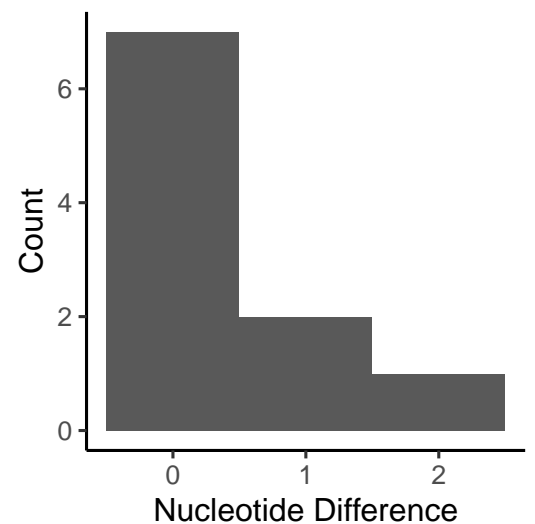
TRBV25–1*bp01

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



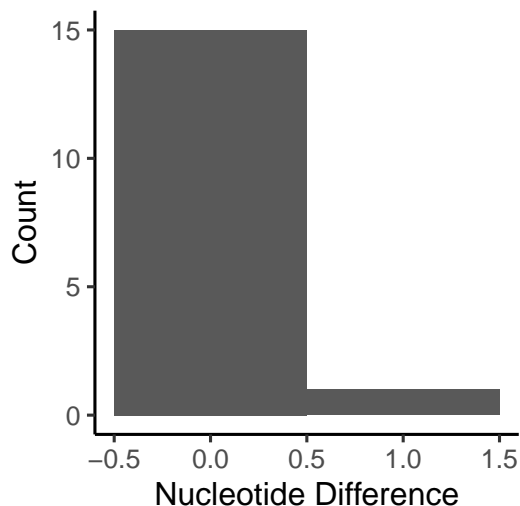
TRBV5–6*bp01

10 sequences assigned
7 (70%) exact matches, in which:
7 unique CDR3
5 unique J



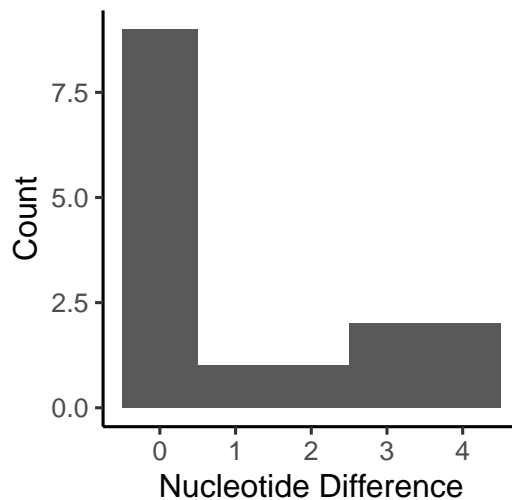
TRBV6-1*bp01

16 sequences assigned
15 (93.8%) exact matches, in which:
14 unique CDR3
8 unique J



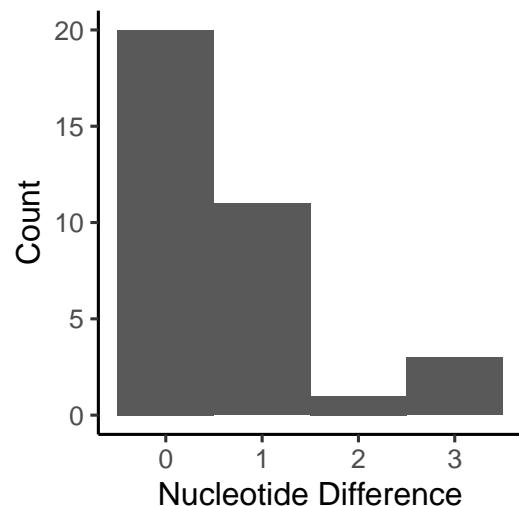
TRBV6-6*bp01

15 sequences assigned
9 (60%) exact matches, in which:
7 unique CDR3
3 unique J



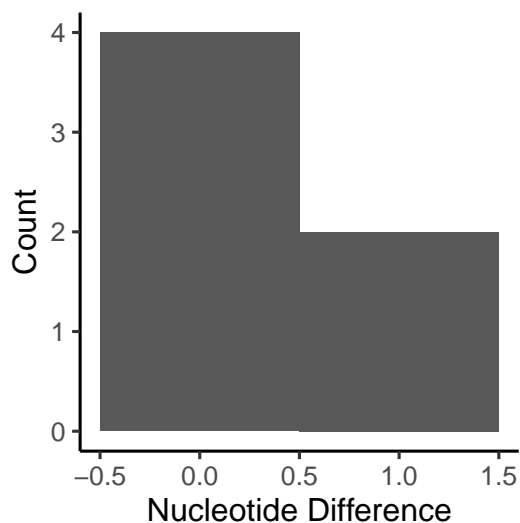
TRBV7-3*bp01

35 sequences assigned
20 (57.1%) exact matches, in which:
16 unique CDR3
7 unique J



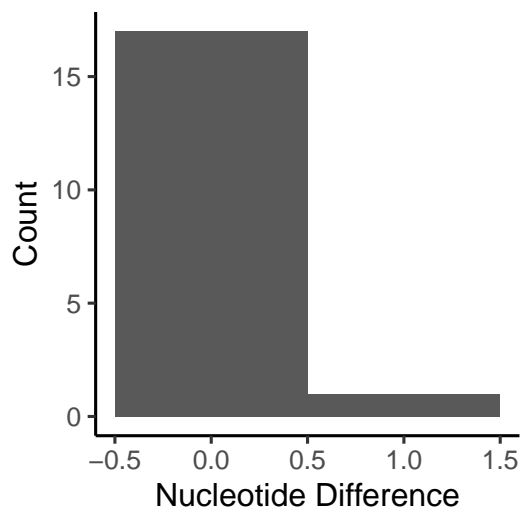
TRBV6-4*bp01

6 sequences assigned
4 (66.7%) exact matches, in which:
4 unique CDR3
4 unique J



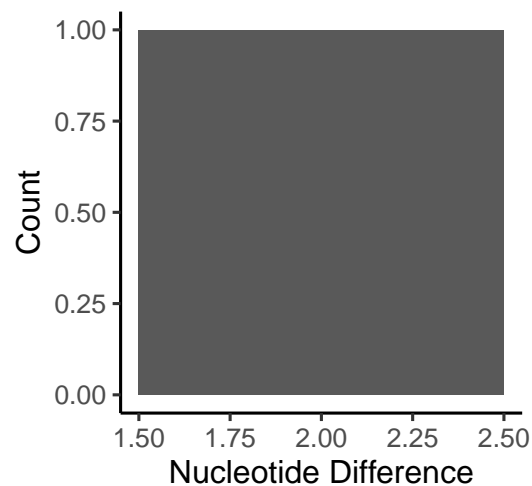
TRBV6-23*bp01

18 sequences assigned
17 (94.4%) exact matches, in which:
17 unique CDR3
8 unique J



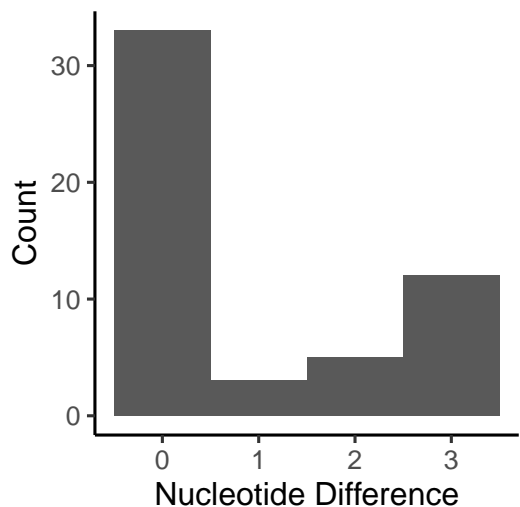
TRBV7-4*bp01

1 sequences assigned
No exact matches.



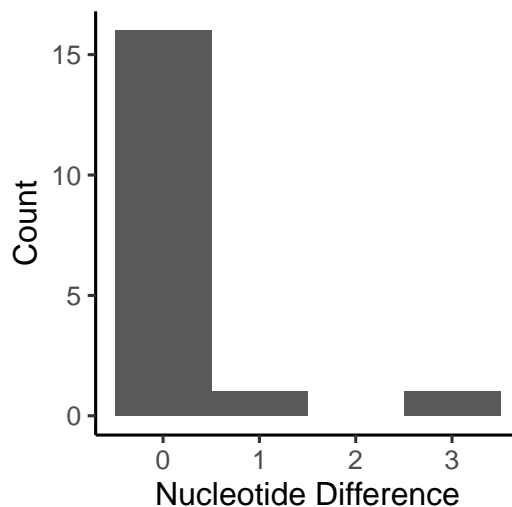
TRBV6-5*bp01

53 sequences assigned
33 (62.3%) exact matches, in which:
33 unique CDR3
11 unique J



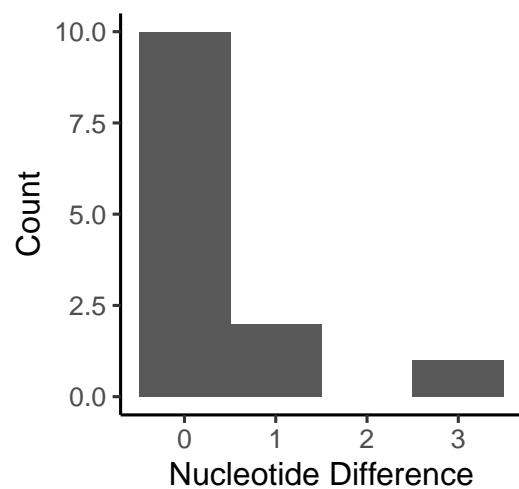
TRBV7-2*bp01

18 sequences assigned
16 (88.9%) exact matches, in which:
16 unique CDR3
8 unique J



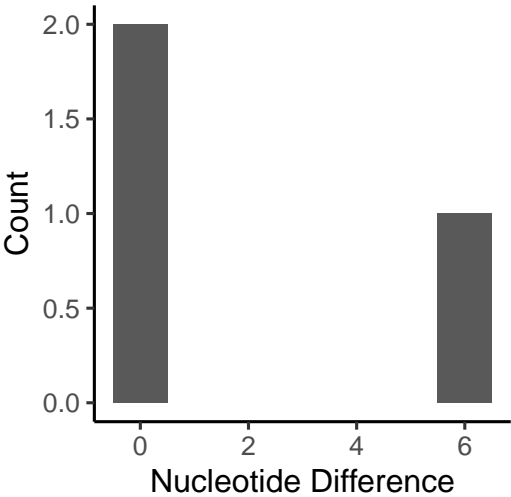
TRBV7-6*bp01

13 sequences assigned
10 (76.9%) exact matches, in which:
7 unique CDR3
4 unique J



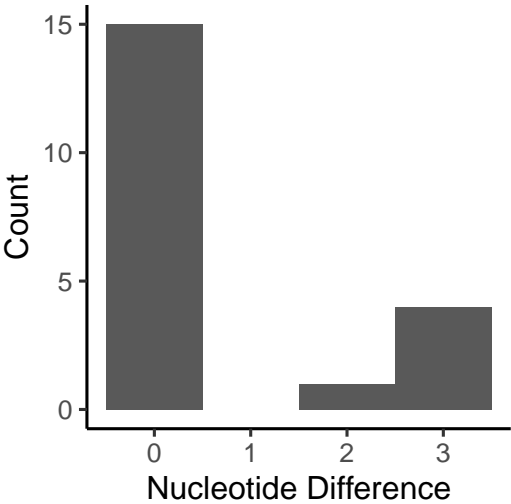
TRBV7-7*bp01

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



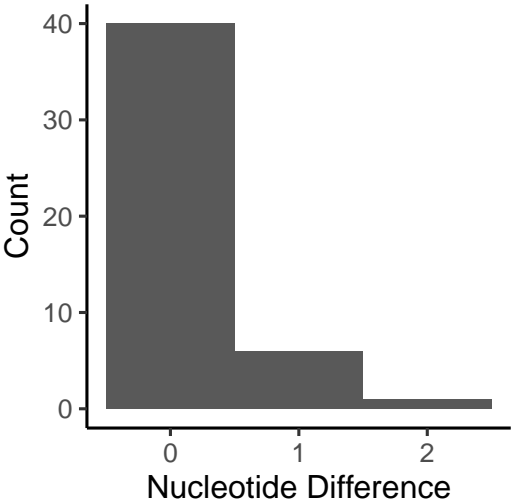
TRBV7-9*bp01

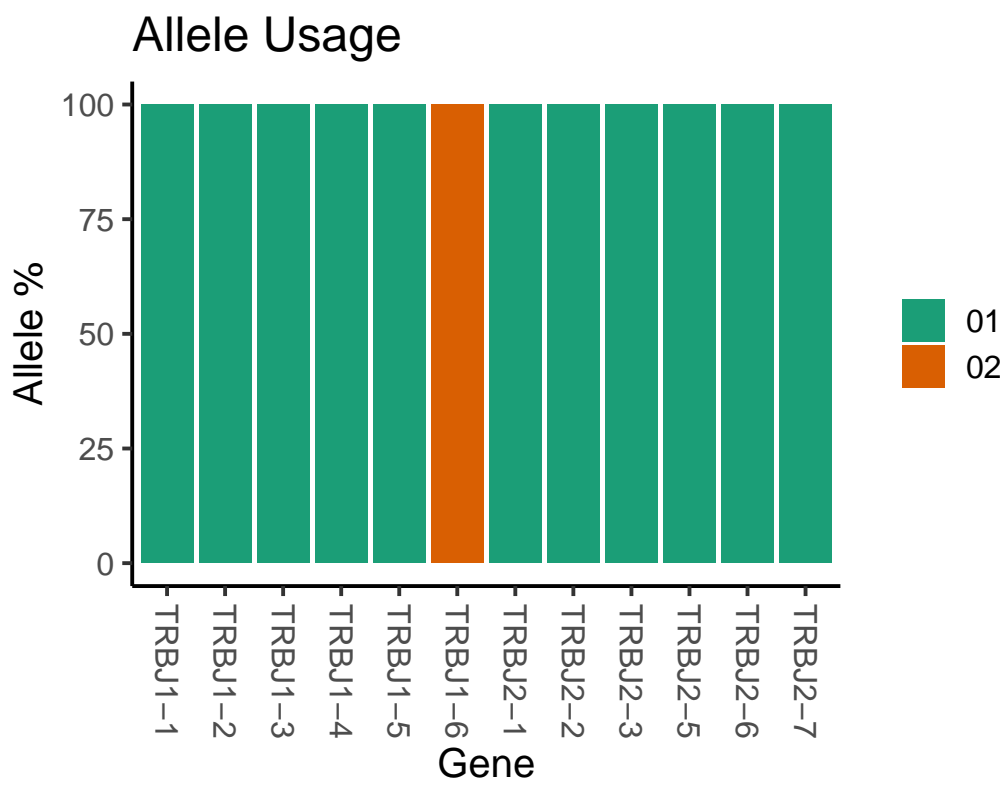
20 sequences assigned
15 (75%) exact matches, in which:
14 unique CDR3
6 unique J



TRBV29-1*bp01

47 sequences assigned
40 (85.1%) exact matches, in which:
40 unique CDR3
9 unique J





[illegible]