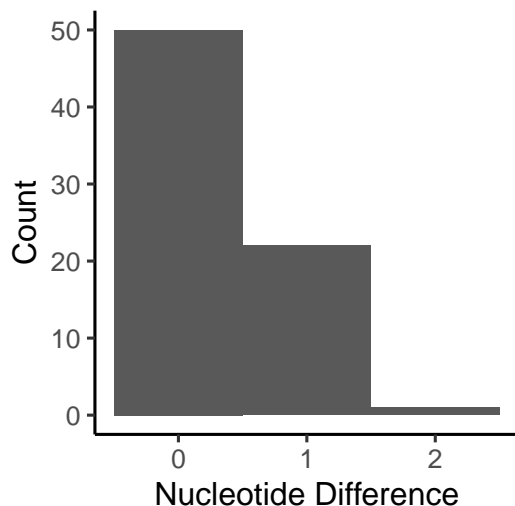


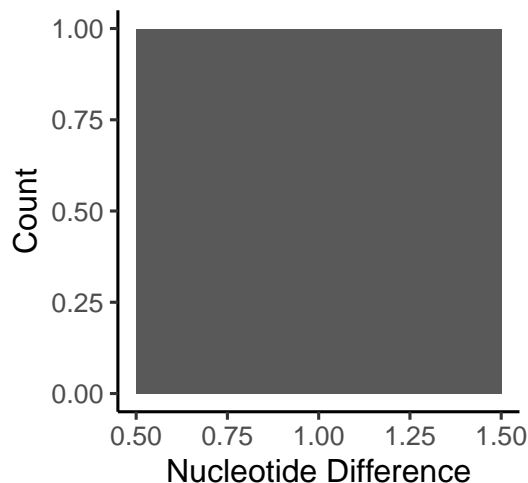
TRBV20–1*bp01

73 sequences assigned
50 (68.5%) exact matches, in which:
27 unique CDR3
10 unique J



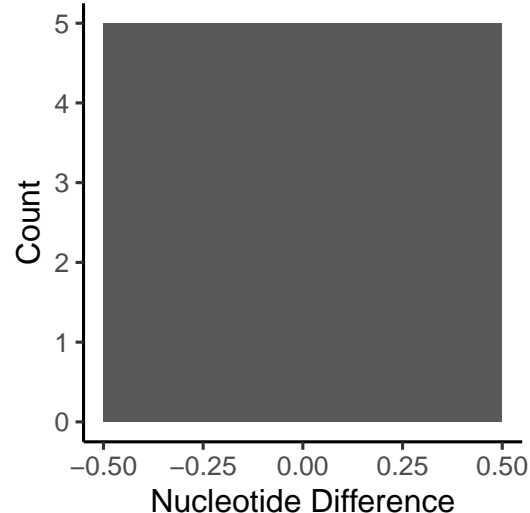
TRBV10–2*bp01

1 sequences assigned
No exact matches.



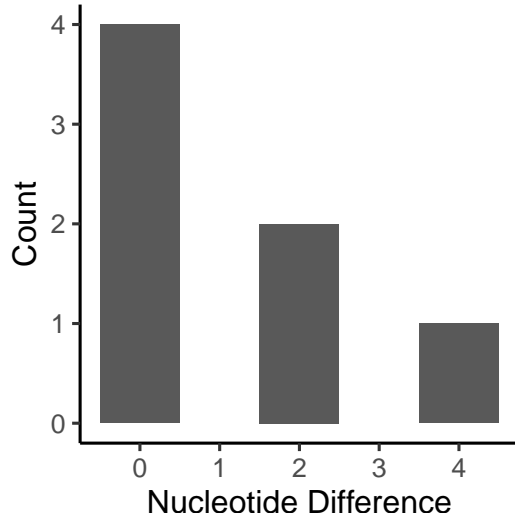
TRBV14*bp01

5 sequences assigned
5 (100%) exact matches, in which:
5 unique CDR3
3 unique J



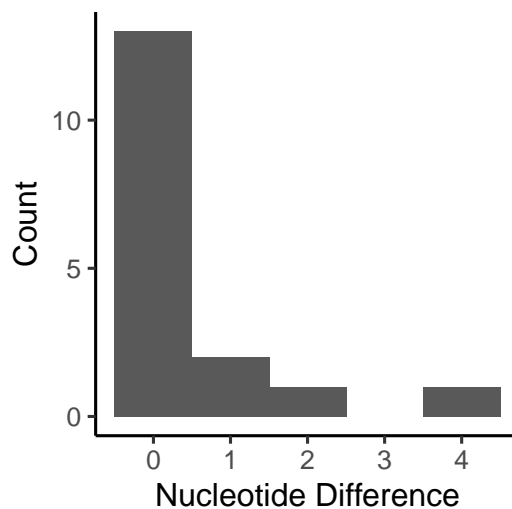
TRBV10–1*bp01

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



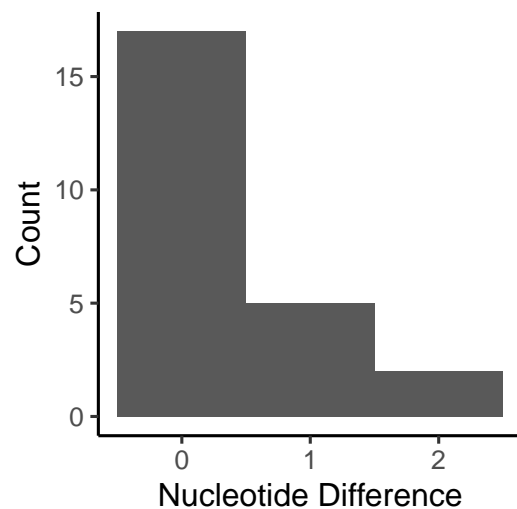
TRBV10–3*bp01

17 sequences assigned
13 (76.5%) exact matches, in which:
11 unique CDR3
7 unique J



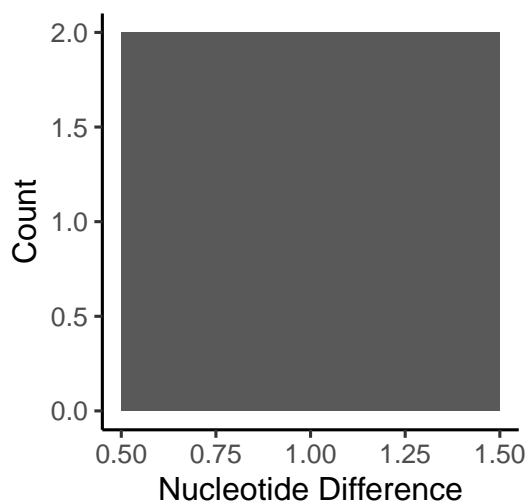
TRBV15*bp02

24 sequences assigned
17 (70.8%) exact matches, in which:
11 unique CDR3
7 unique J



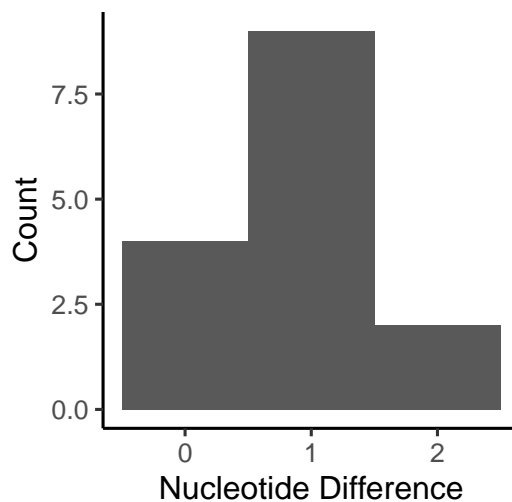
TRBV10–1*bp02

2 sequences assigned
No exact matches.



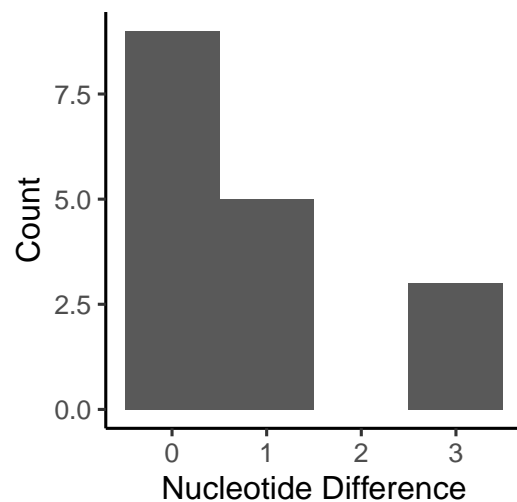
TRBV13*bp01

15 sequences assigned
4 (26.7%) exact matches, in which:
4 unique CDR3
4 unique J



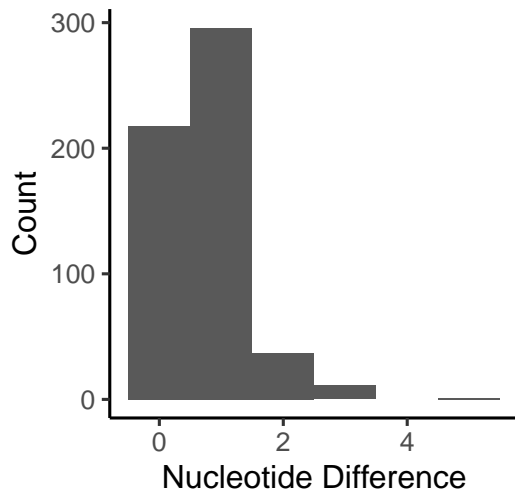
TRBV18*bp01

17 sequences assigned
9 (52.9%) exact matches, in which:
6 unique CDR3
5 unique J



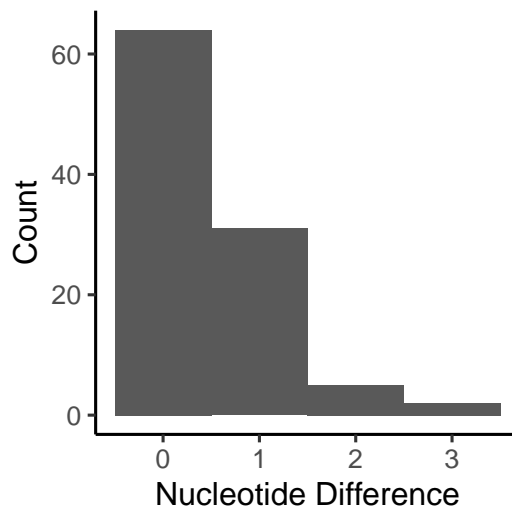
TRBV19*bp01

563 sequences assigned
218 (38.7%) exact matches, in which:
132 unique CDR3
12 unique J



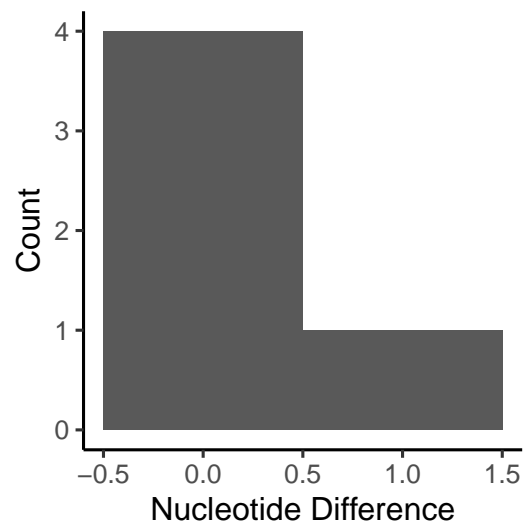
TRBV30*bp01

102 sequences assigned
64 (62.7%) exact matches, in which:
27 unique CDR3
12 unique J



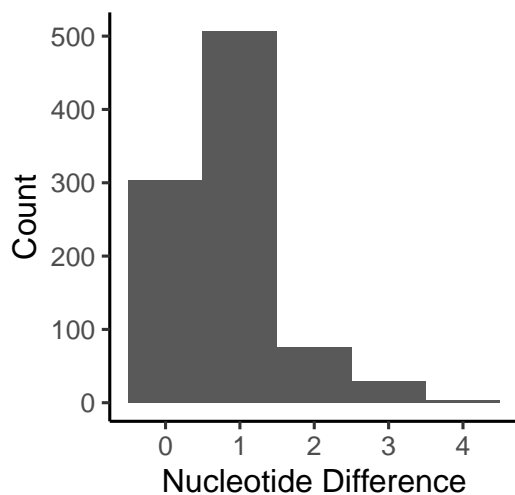
TRBV11-2*bp01

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



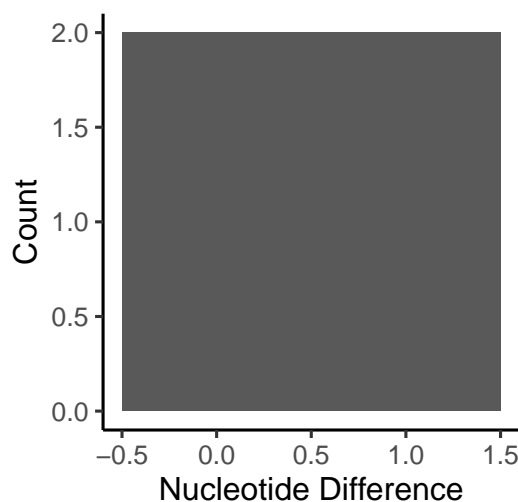
TRBV27*bp01

918 sequences assigned
303 (33%) exact matches, in which:
186 unique CDR3
12 unique J



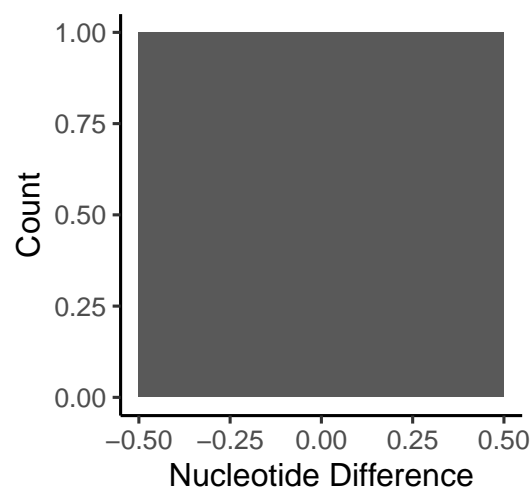
TRBV11-1*bp01

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



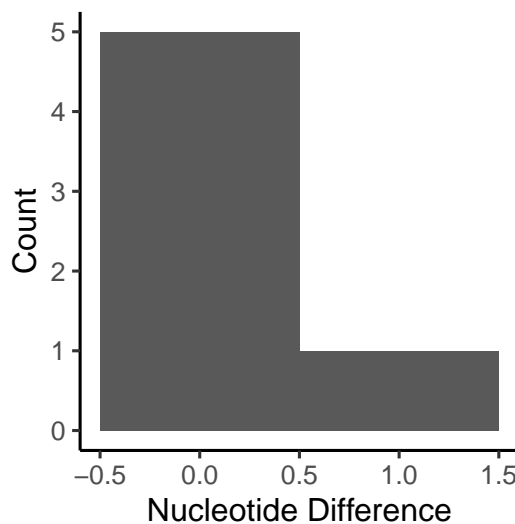
TRBV11-3*bp01

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



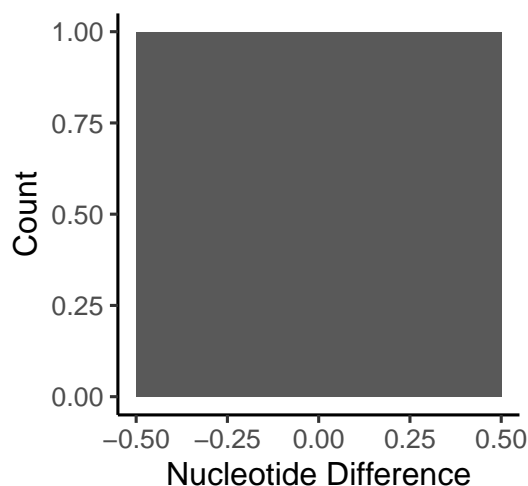
TRBV28*bp01

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



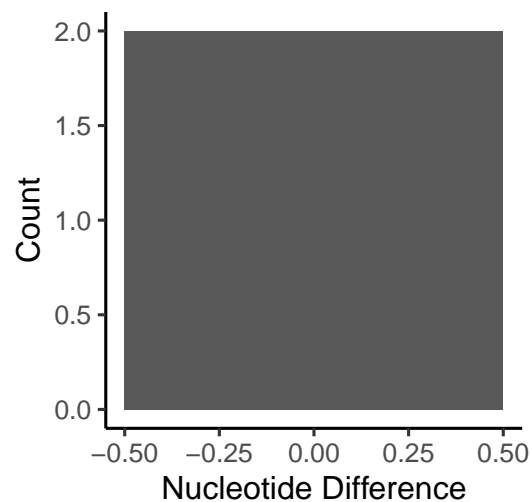
TRBV21-1*bp02

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



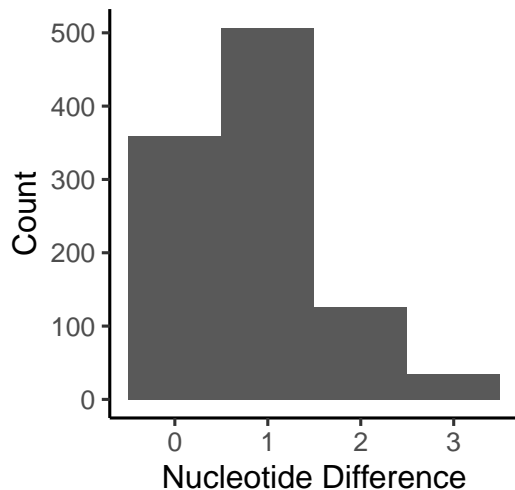
TRBV12-5*bp01

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



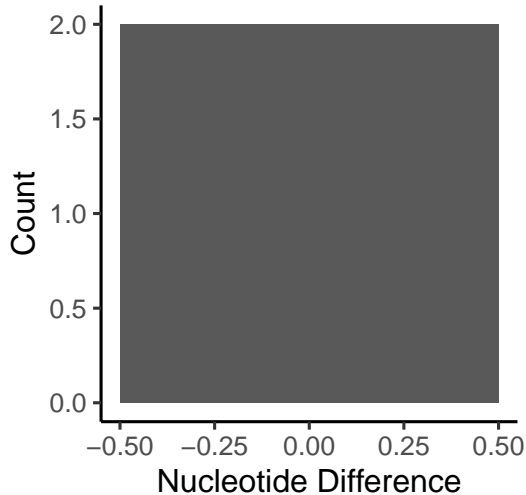
TRBV12–34*bp01

1027 sequences assigned
359 (35%) exact matches, in which:
168 unique CDR3
12 unique J



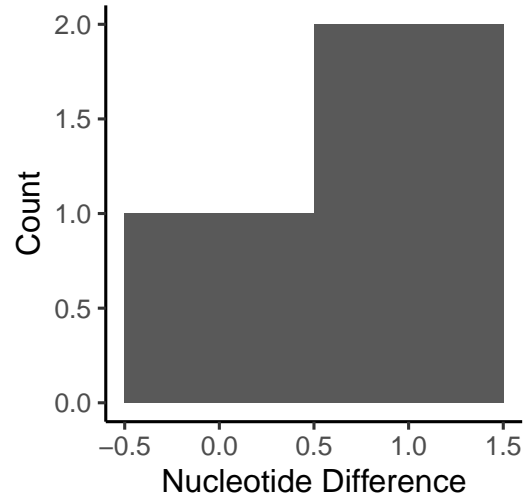
TRBV4–2*bp01

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



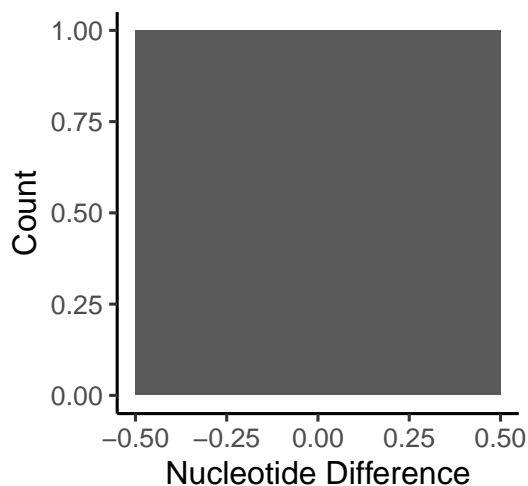
TRBV5–3*bp01

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



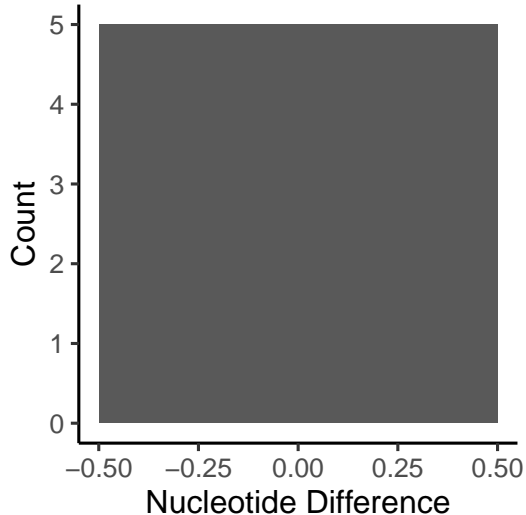
TRBV4–1*bp01

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



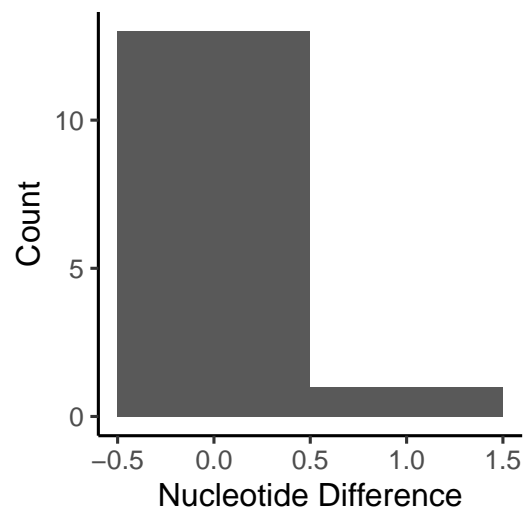
TRBV4–3*bp01

5 sequences assigned
5 (100%) exact matches, in which:
3 unique CDR3
3 unique J



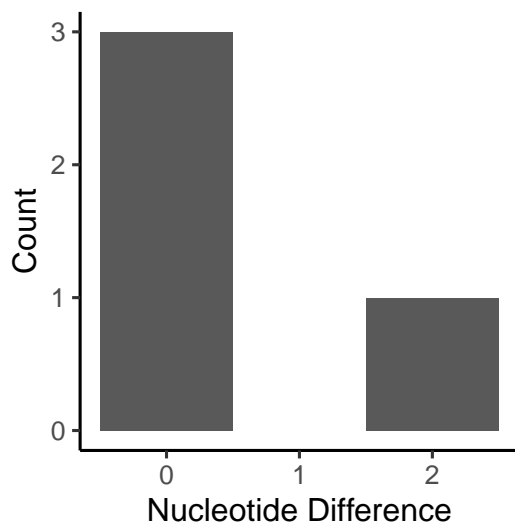
TRBV5–4*bp01

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



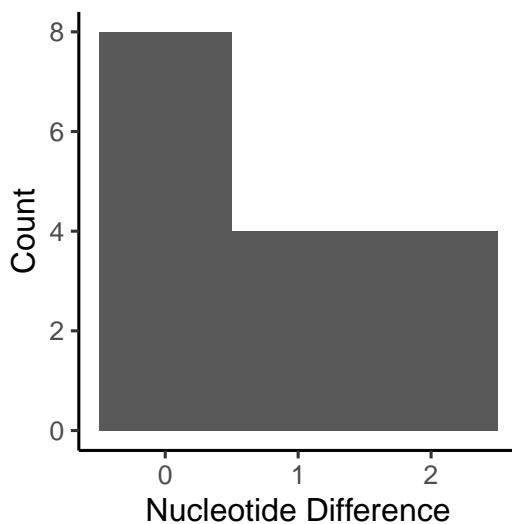
TRBV24–1*bp02

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



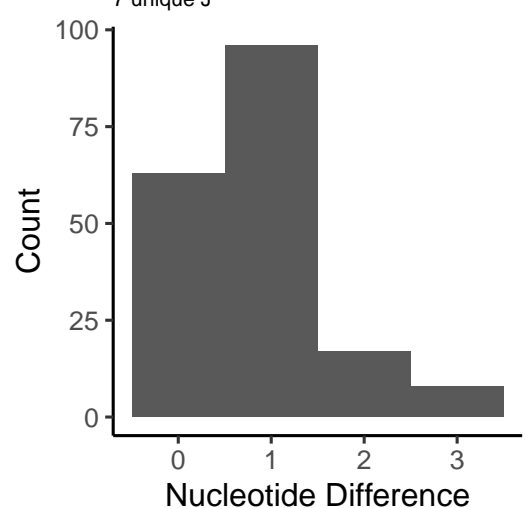
TRBV5–1*bp01

16 sequences assigned
8 (50%) exact matches, in which:
7 unique CDR3
4 unique J



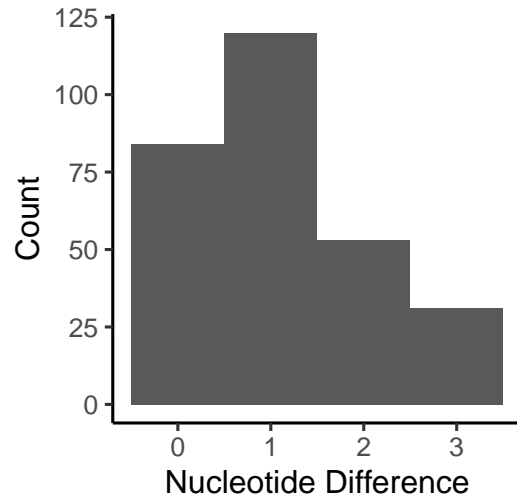
TRBV5–5*bp01

184 sequences assigned
63 (34.2%) exact matches, in which:
35 unique CDR3
7 unique J



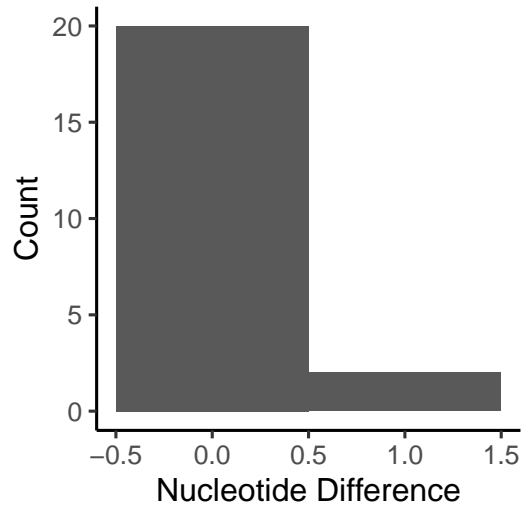
TRBV5-6*bp01

288 sequences assigned
84 (29.2%) exact matches, in which:
42 unique CDR3
7 unique J



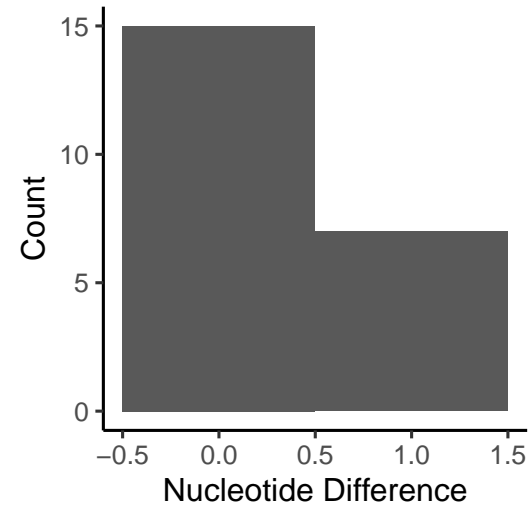
TRBV6-4*bp01

22 sequences assigned
20 (90.9%) exact matches, in which:
11 unique CDR3
9 unique J



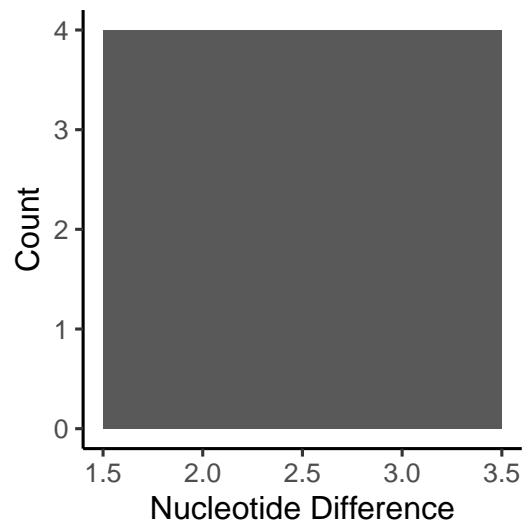
TRBV6-23*bp01

22 sequences assigned
15 (68.2%) exact matches, in which:
13 unique CDR3
5 unique J



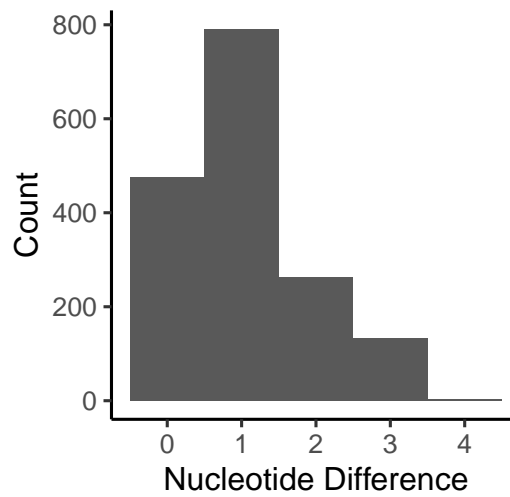
TRBV5-7*bp01

8 sequences assigned
No exact matches.



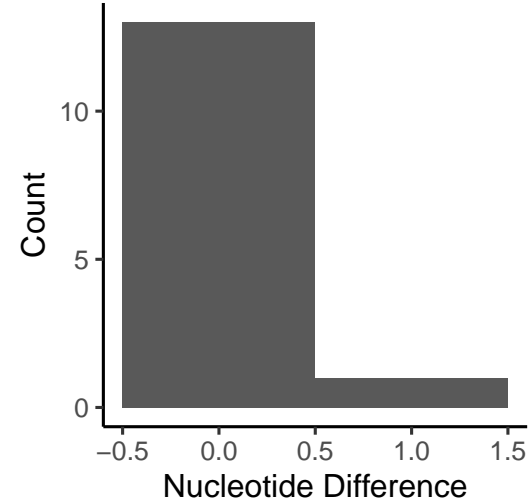
TRBV6-5*bp01

1666 sequences assigned
476 (28.6%) exact matches, in which:
265 unique CDR3
13 unique J



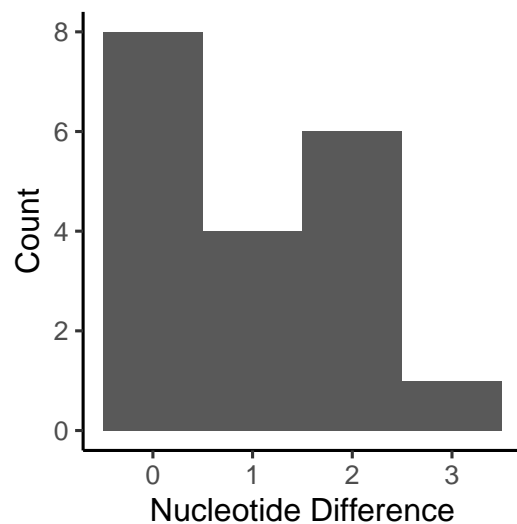
TRBV7-2*bp01

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



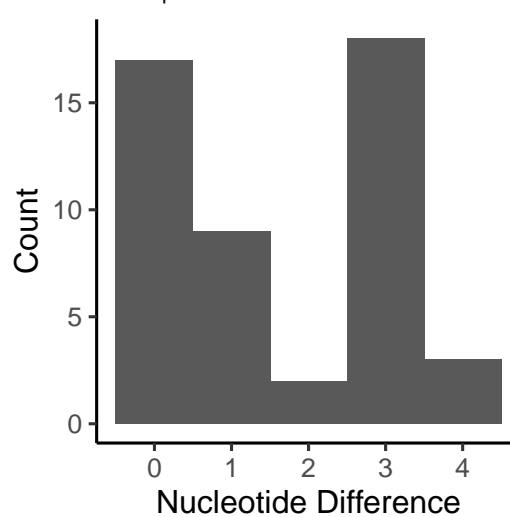
TRBV6-1*bp01

19 sequences assigned
8 (42.1%) exact matches, in which:
8 unique CDR3
5 unique J



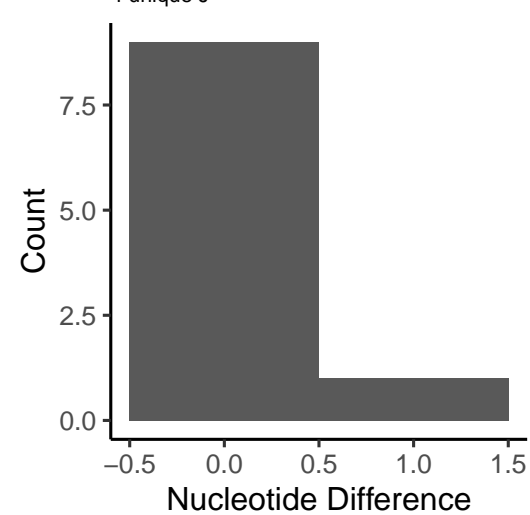
TRBV6-6*bp01

49 sequences assigned
17 (34.7%) exact matches, in which:
13 unique CDR3
11 unique J



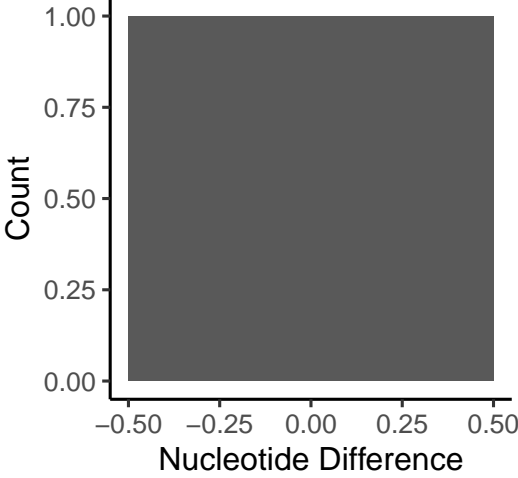
TRBV7-2*bp02

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



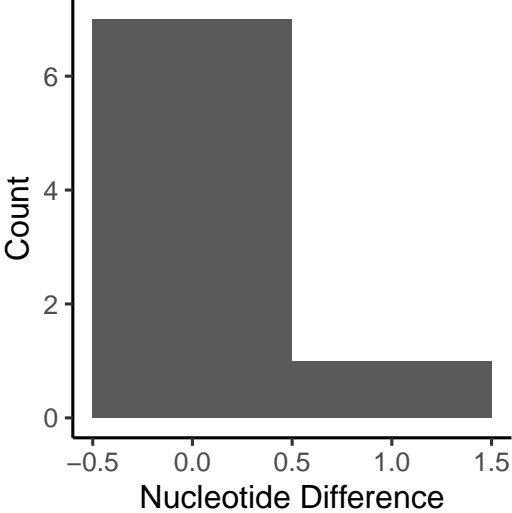
TRBV7-3*bp01

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



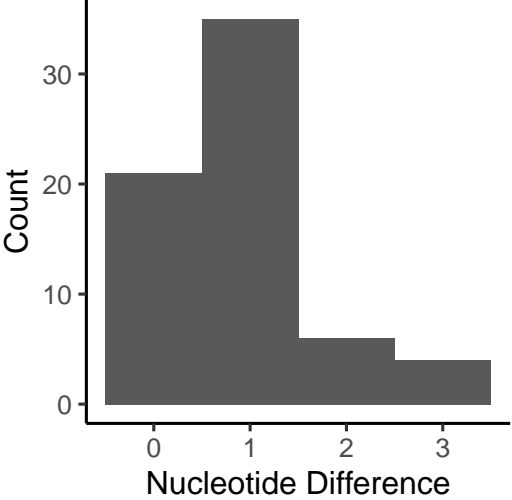
TRBV7-9*bp01

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



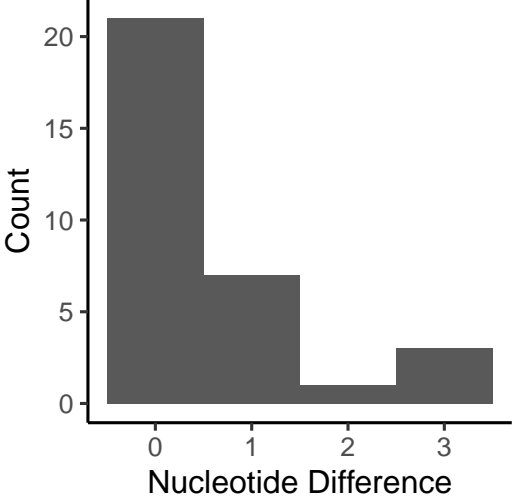
TRBV29-1*bp03

66 sequences assigned
21 (31.8%) exact matches, in which:
11 unique CDR3
2 unique J



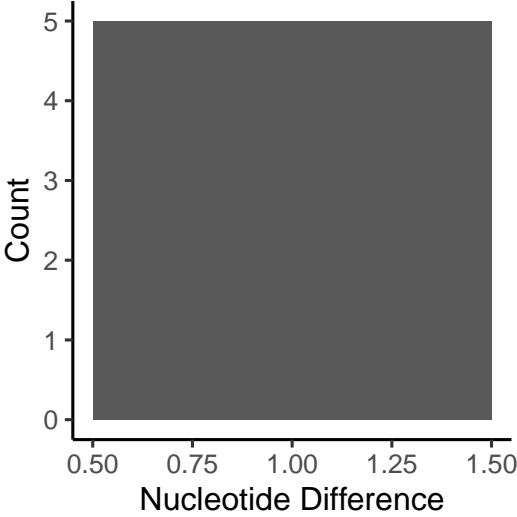
TRBV7-6*bp01

32 sequences assigned
21 (65.6%) exact matches, in which:
11 unique CDR3
6 unique J



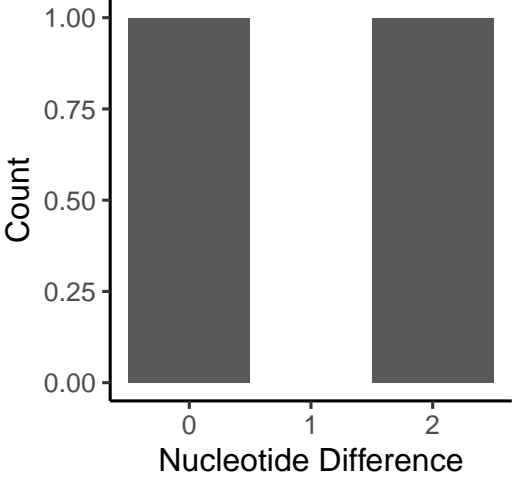
TRBV7-9*bp03

5 sequences assigned
No exact matches.



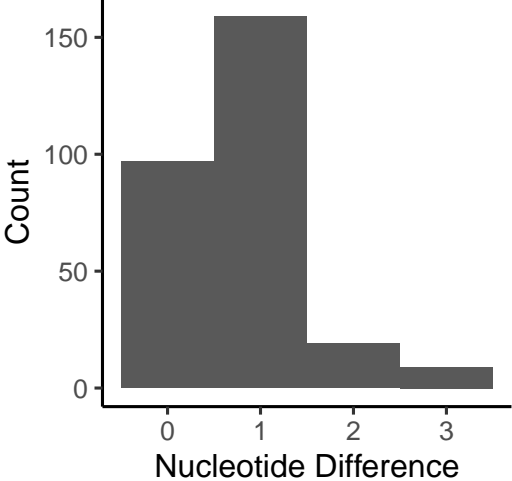
TRBV7-7*bp01

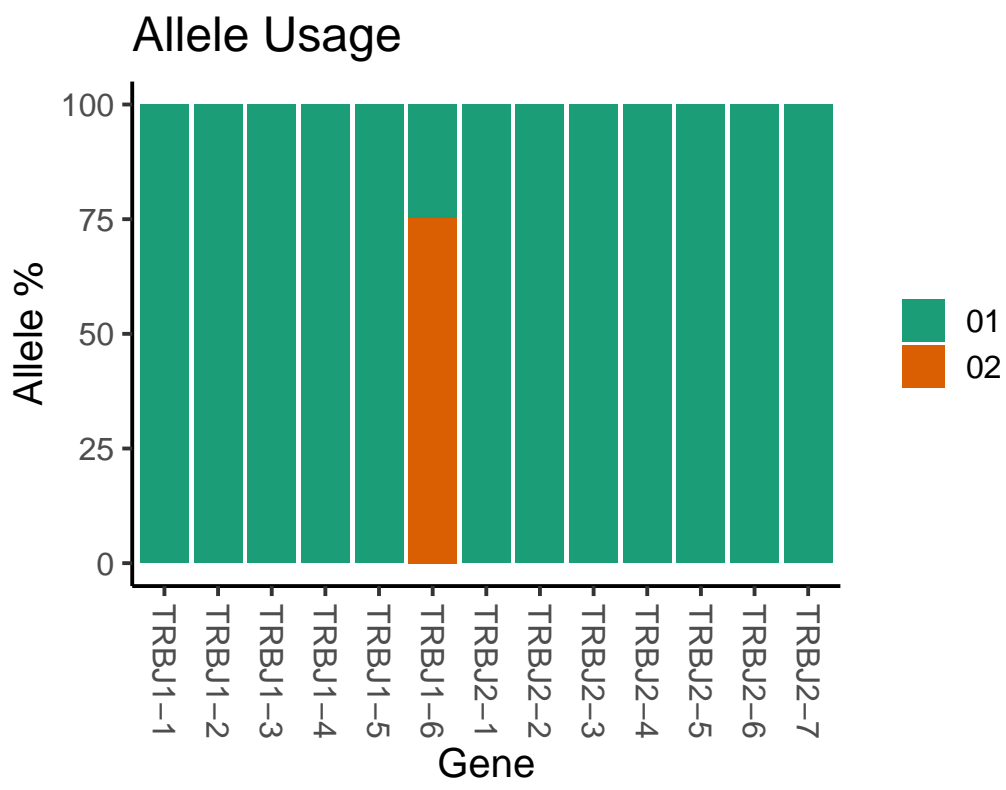
2 sequences assigned
1 (50%) exact matches, in which:
1 unique CDR3
1 unique J



TRBV29-1*bp01

284 sequences assigned
97 (34.2%) exact matches, in which:
63 unique CDR3
9 unique J





[illegible]