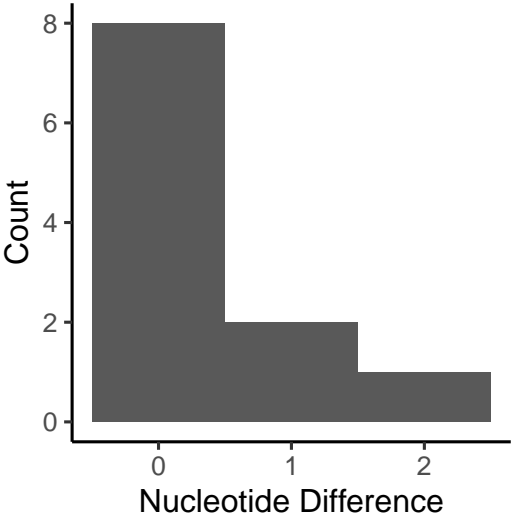


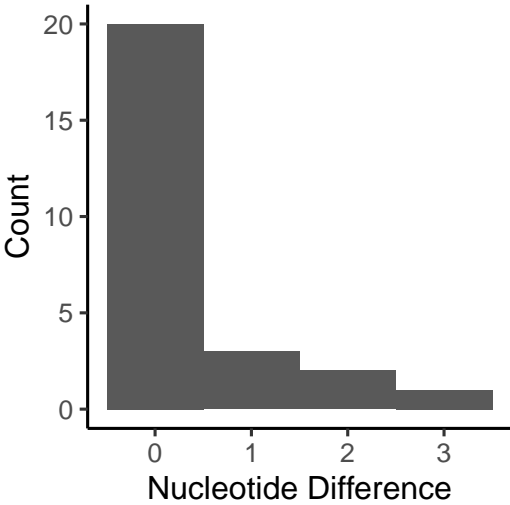
TRBV10–1*bp01

11 sequences assigned
8 (72.7%) exact matches, in which:
8 unique CDR3
4 unique J



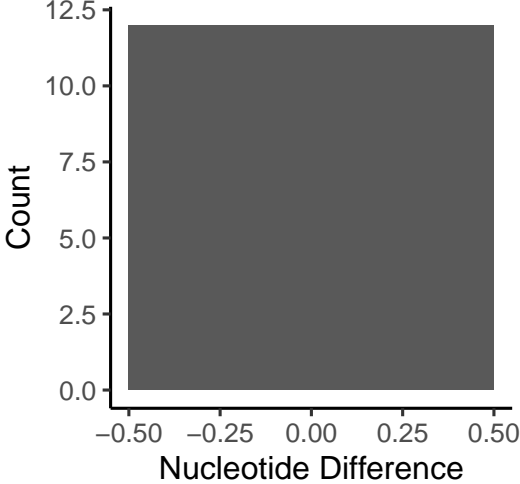
TRBV10–3*bp01

26 sequences assigned
20 (76.9%) exact matches, in which:
19 unique CDR3
7 unique J



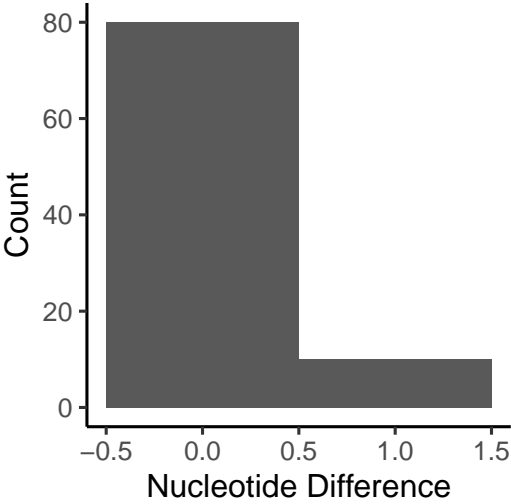
TRBV15*bp02

12 sequences assigned
12 (100%) exact matches, in which:
11 unique CDR3
6 unique J



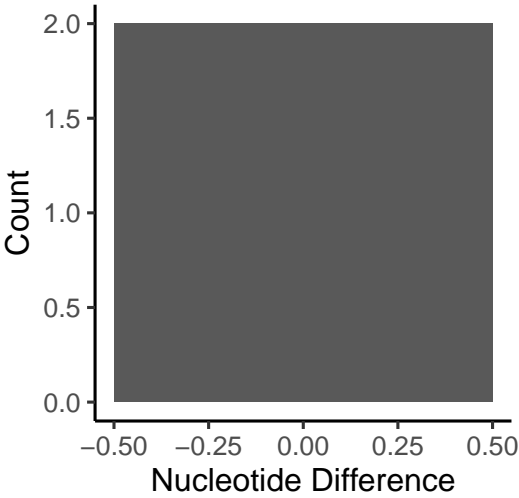
TRBV20–1*bp01

90 sequences assigned
80 (88.9%) exact matches, in which:
72 unique CDR3
12 unique J



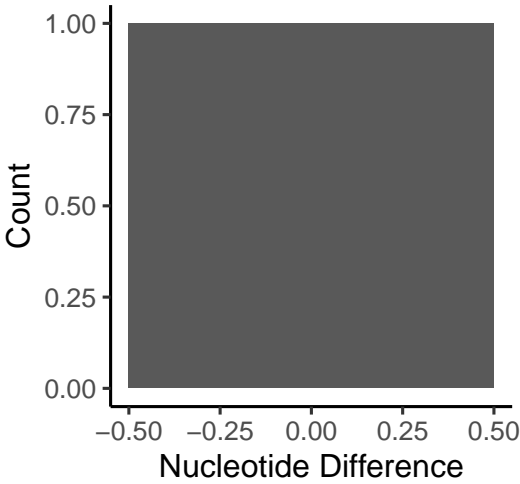
TRBV13*bp01

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



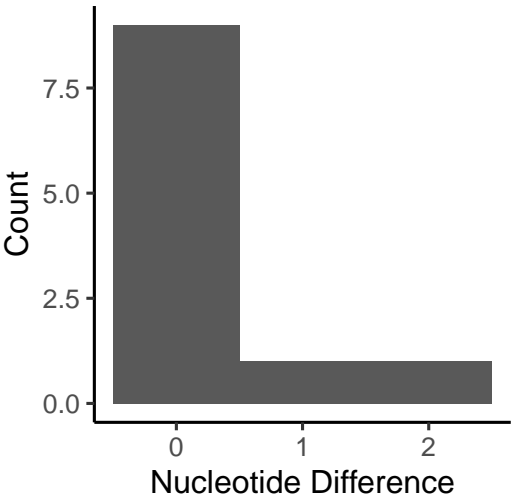
TRBV18*bp01

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



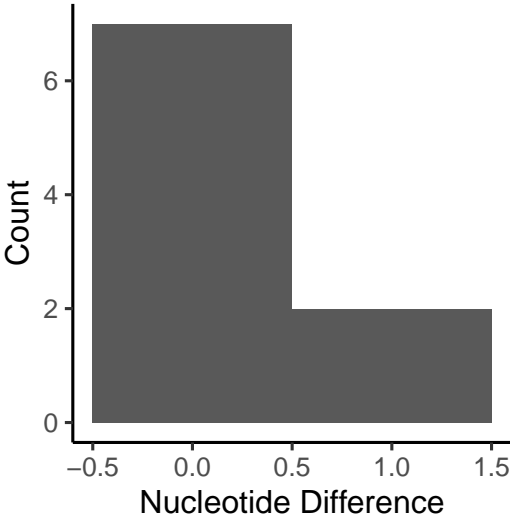
TRBV10–2*bp01

11 sequences assigned
9 (81.8%) exact matches, in which:
9 unique CDR3
3 unique J



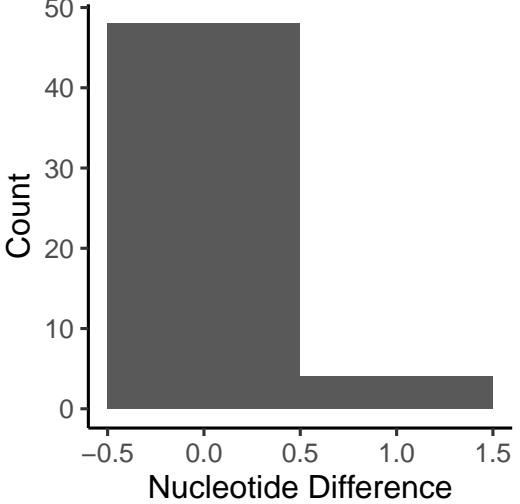
TRBV14*bp01

9 sequences assigned
7 (77.8%) exact matches, in which:
7 unique CDR3
5 unique J



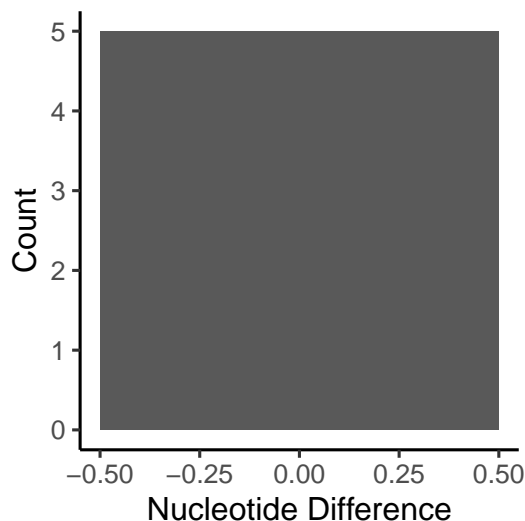
TRBV19*bp01

53 sequences assigned
48 (90.6%) exact matches, in which:
46 unique CDR3
12 unique J



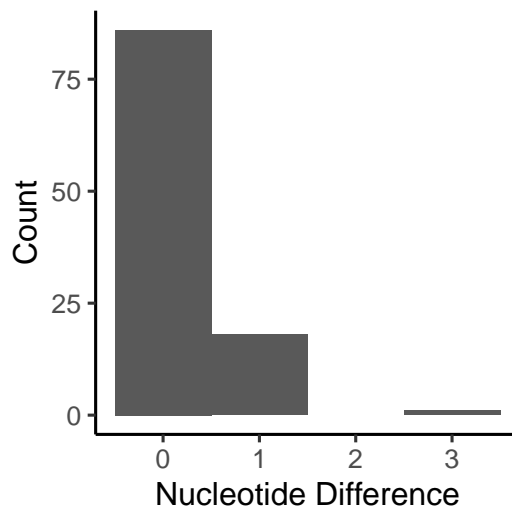
TRBV2*bp01

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



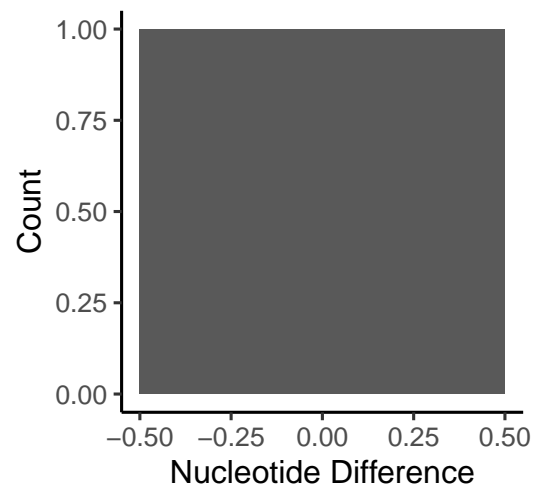
TRBV30*bp01

105 sequences assigned
86 (81.9%) exact matches, in which:
65 unique CDR3
12 unique J



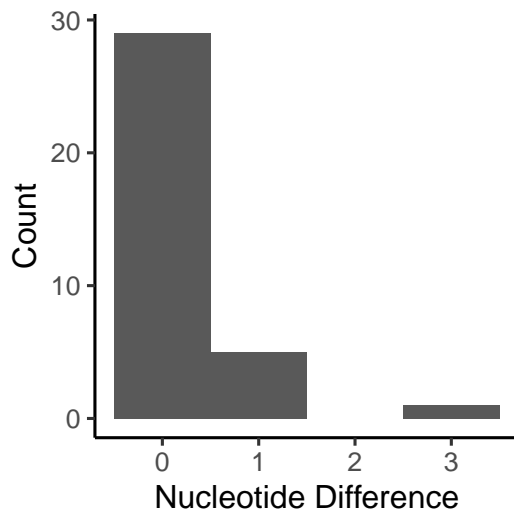
TRBV21-1*bp01

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



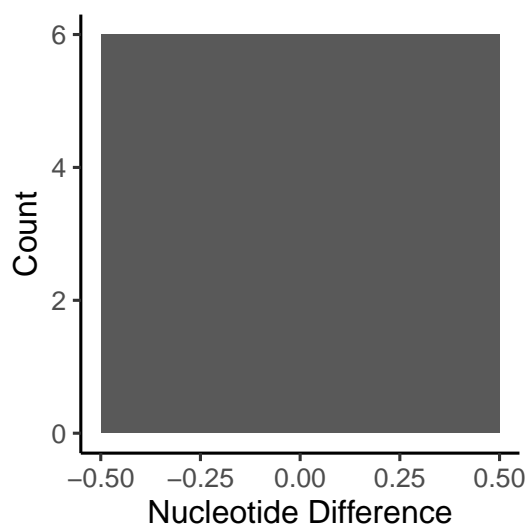
TRBV27*bp01

35 sequences assigned
29 (82.9%) exact matches, in which:
28 unique CDR3
10 unique J



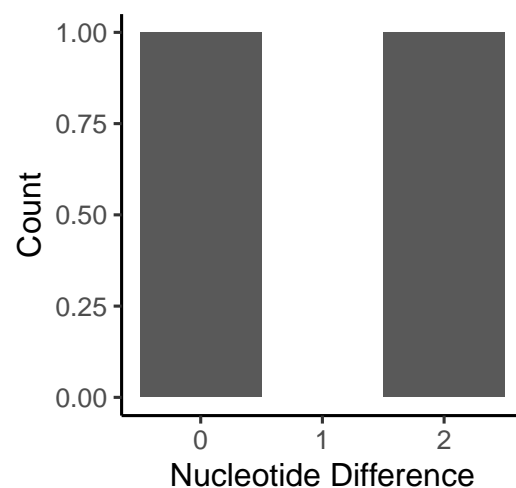
TRBV9*bp01

6 sequences assigned
6 (100%) exact matches, in which:
6 unique CDR3
5 unique J



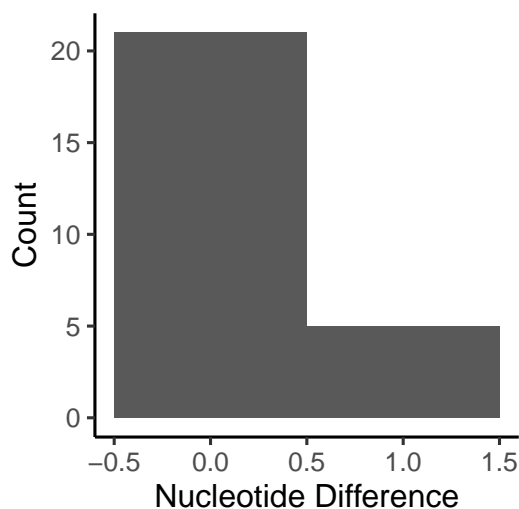
TRBV21-1*bp02

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



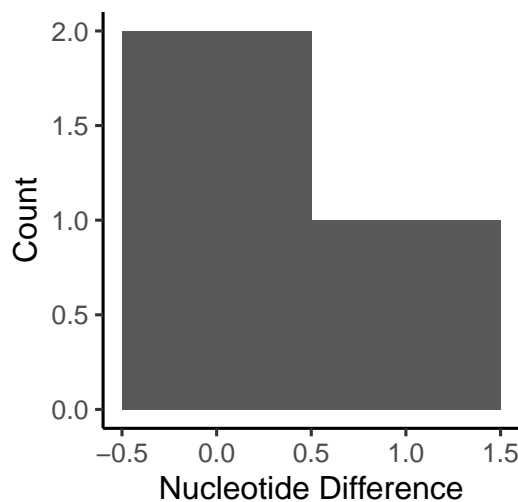
TRBV28*bp01

26 sequences assigned
21 (80.8%) exact matches, in which:
20 unique CDR3
9 unique J



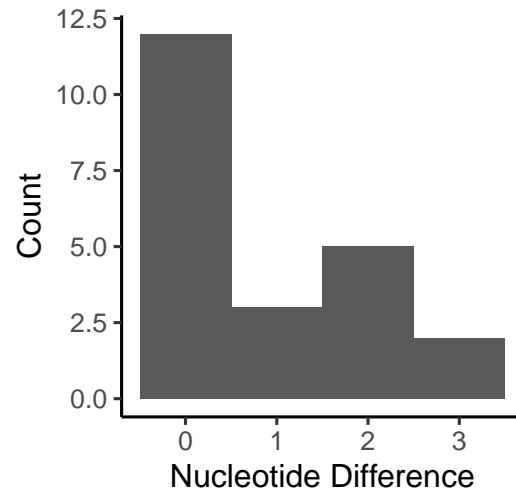
TRBV11-1*bp01

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



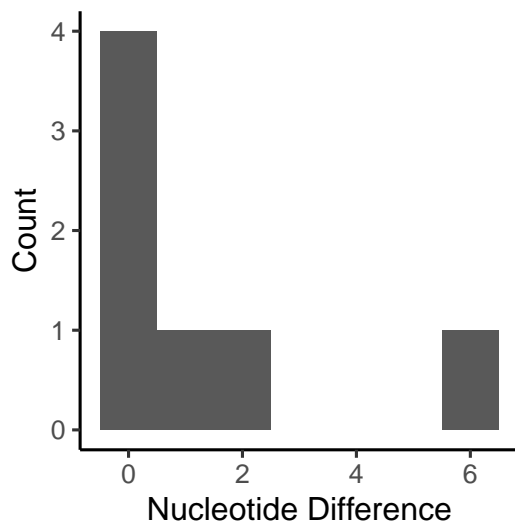
TRBV11-2*bp01

22 sequences assigned
12 (54.5%) exact matches, in which:
11 unique CDR3
6 unique J



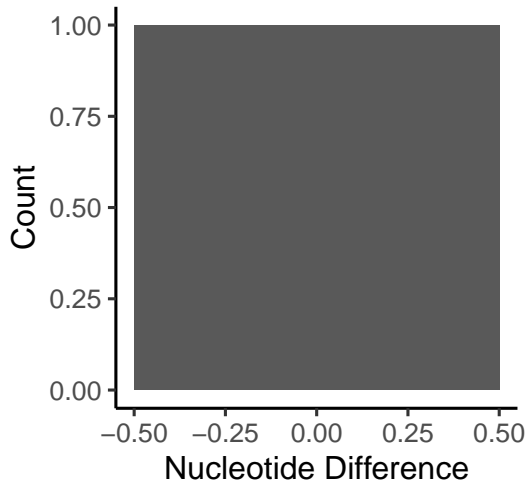
TRBV11–3*bp01

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



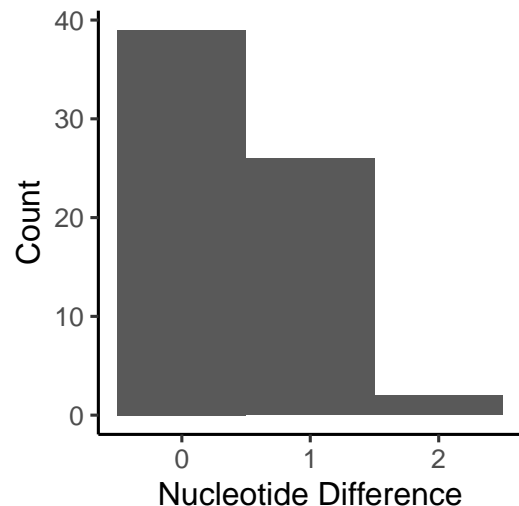
TRBV23–1*bp01

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



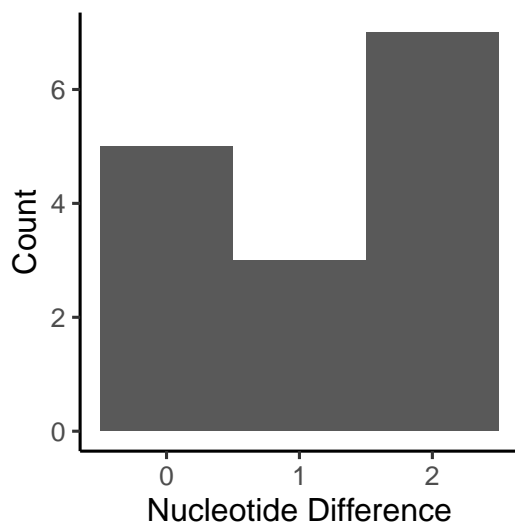
TRBV24–1*bp01

67 sequences assigned
39 (58.2%) exact matches, in which:
31 unique CDR3
8 unique J



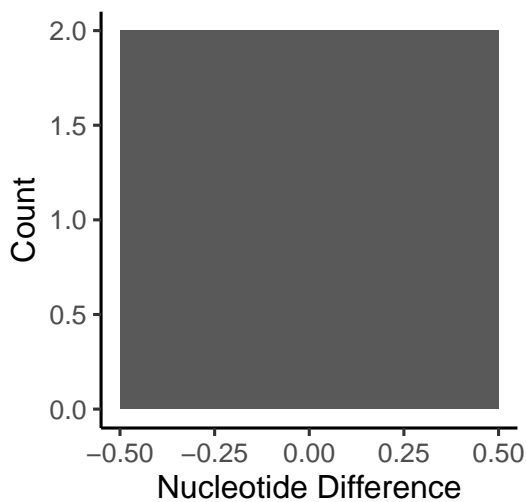
TRBV12–5*bp01

15 sequences assigned
5 (33.3%) exact matches, in which:
4 unique CDR3
3 unique J



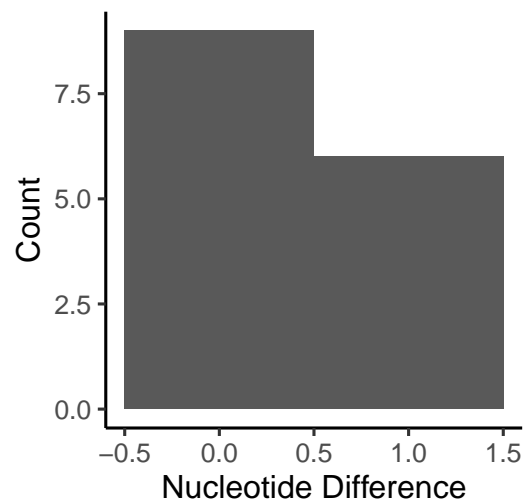
TRBV3–2*bp01

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



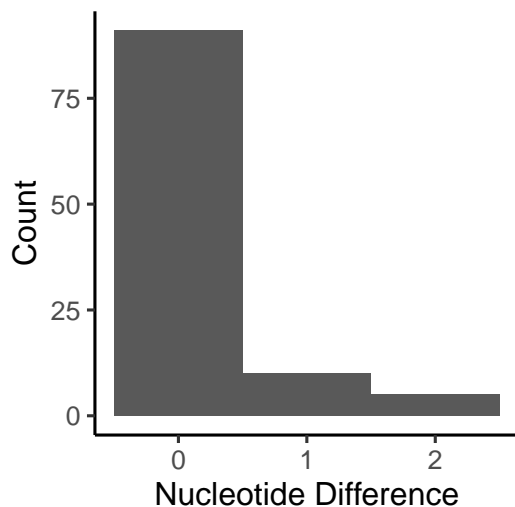
TRBV24–1*bp02

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



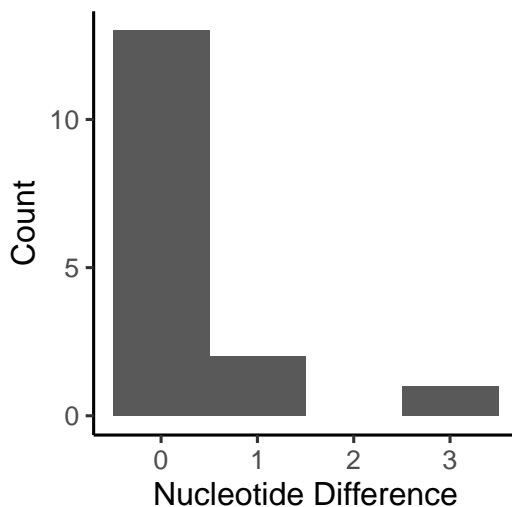
TRBV12–34*bp01

106 sequences assigned
91 (85.8%) exact matches, in which:
69 unique CDR3
13 unique J



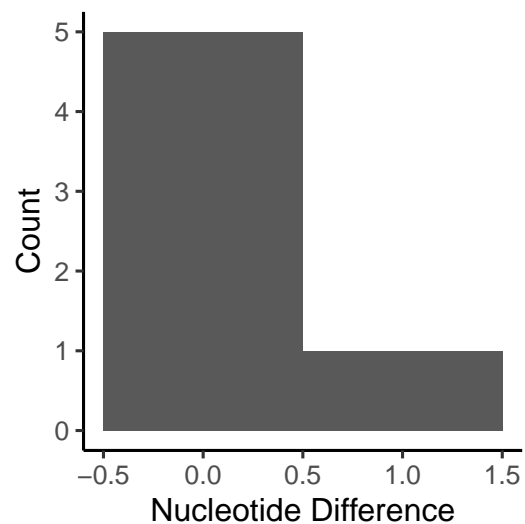
TRBV4–1*bp01

16 sequences assigned
13 (81.2%) exact matches, in which:
11 unique CDR3
5 unique J



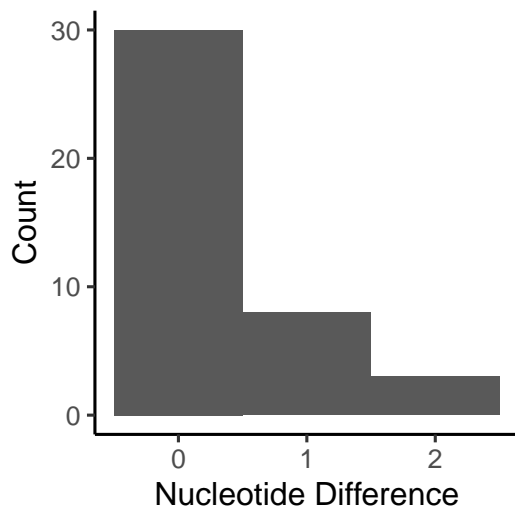
TRBV4–2*bp01

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



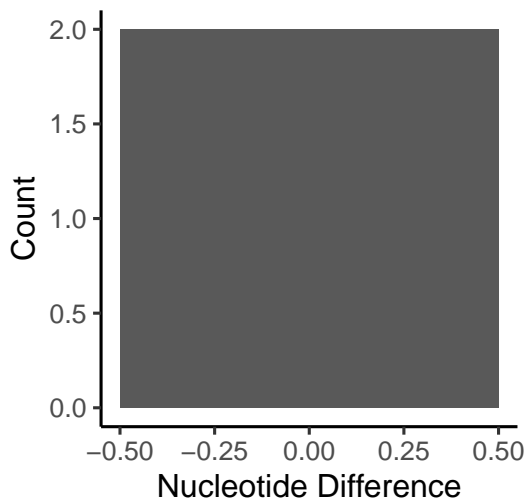
TRBV4-3*bp01

41 sequences assigned
30 (73.2%) exact matches, in which:
26 unique CDR3
12 unique J



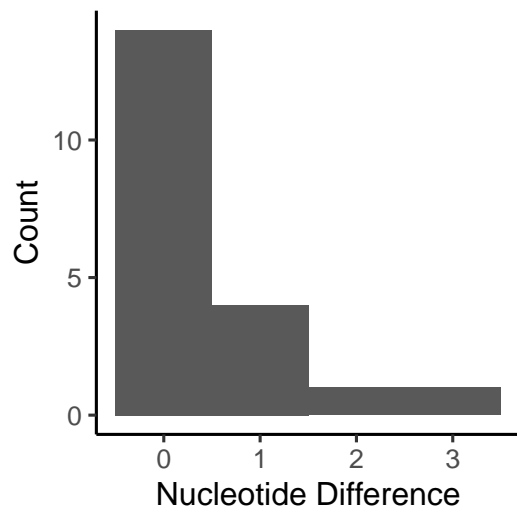
TRBV5-3*bp01

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



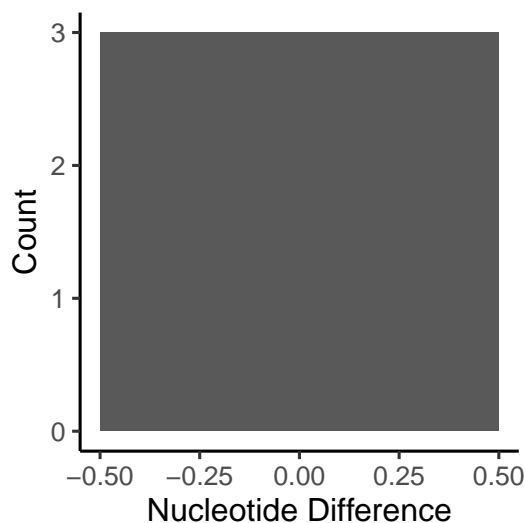
TRBV5-6*bp01

20 sequences assigned
14 (70%) exact matches, in which:
13 unique CDR3
9 unique J



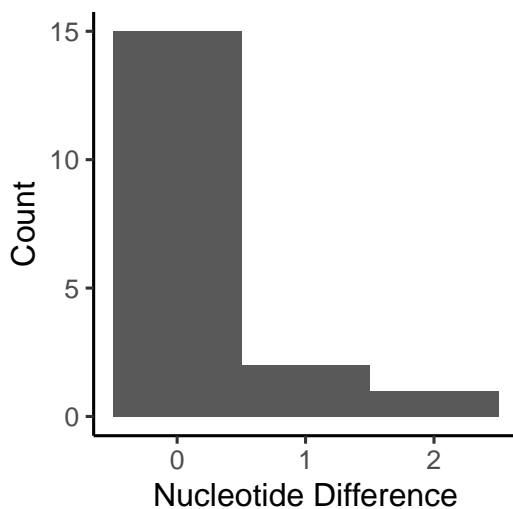
TRBV25-1*bp01

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



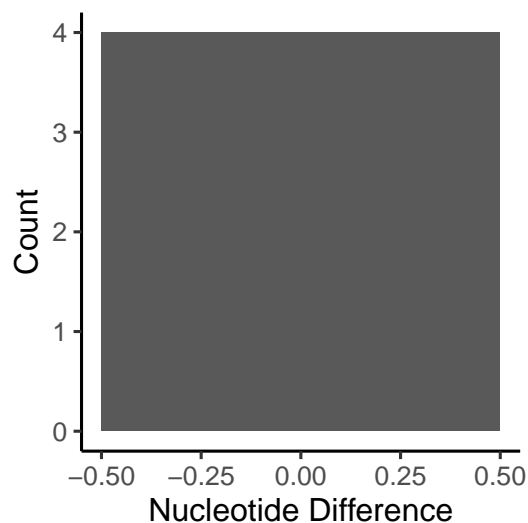
TRBV5-4*bp01

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



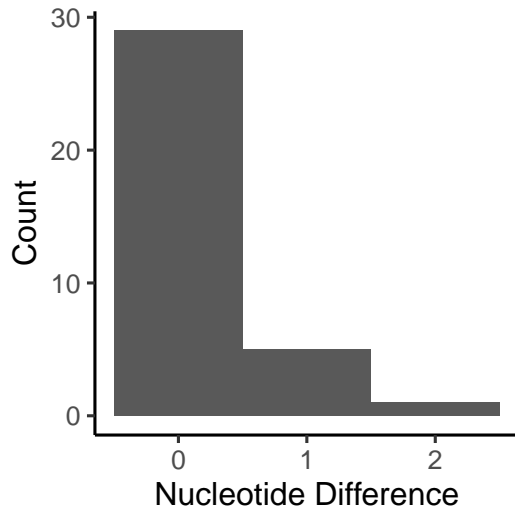
TRBV5-8*bp01

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



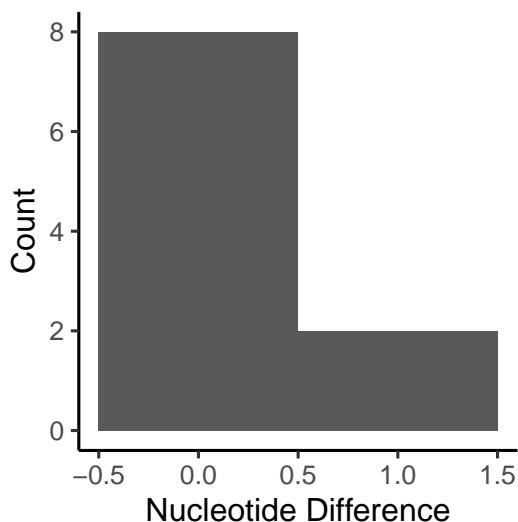
TRBV5-1*bp01

35 sequences assigned
29 (82.9%) exact matches, in which:
27 unique CDR3
9 unique J



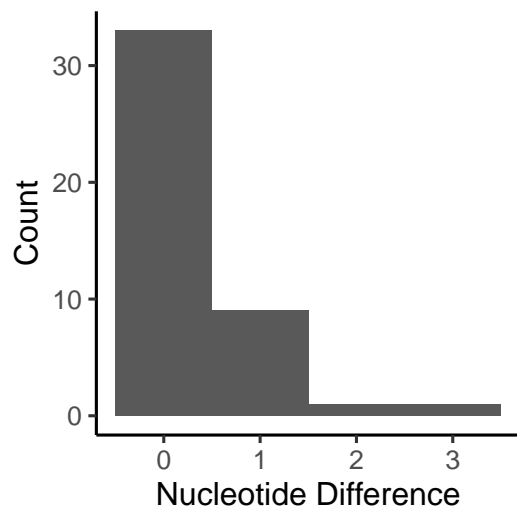
TRBV5-5*bp01

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



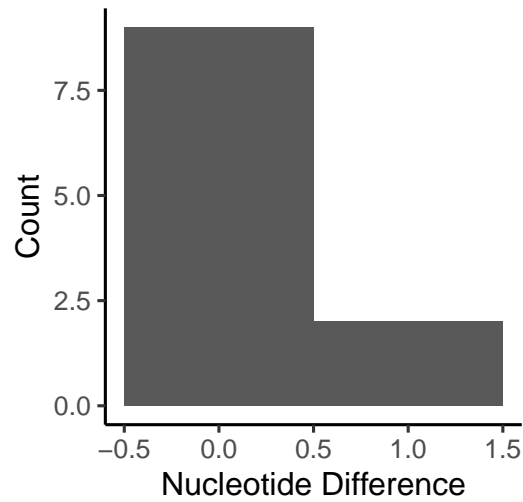
TRBV6-1*bp01

44 sequences assigned
33 (75%) exact matches, in which:
32 unique CDR3
11 unique J



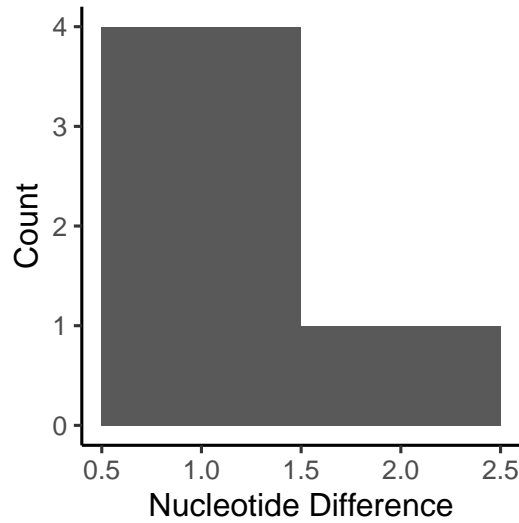
TRBV6-4*bp01

11 sequences assigned
9 (81.8%) exact matches, in which:
7 unique CDR3
5 unique J



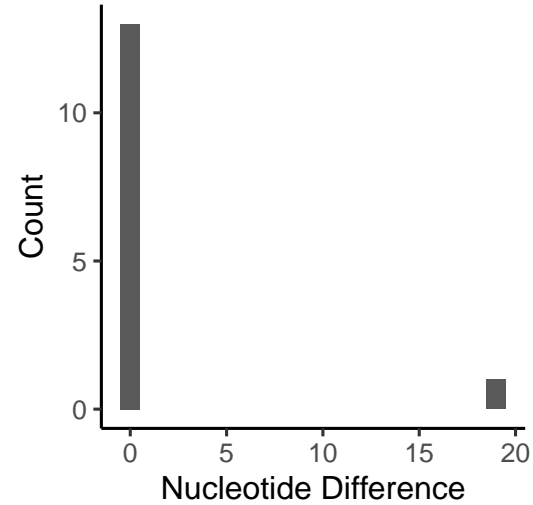
TRBV6-6*bp03

5 sequences assigned
No exact matches.



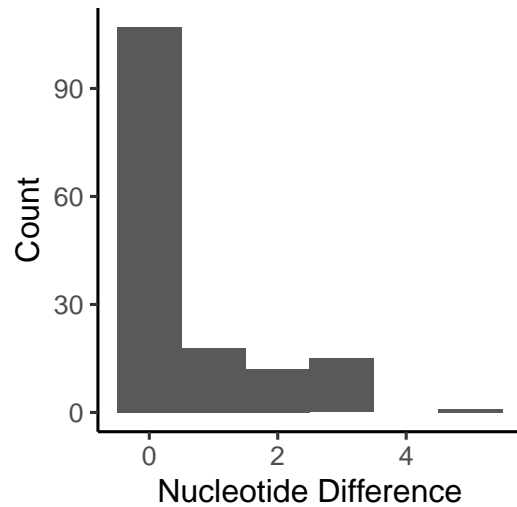
TRBV6-23*bp01

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



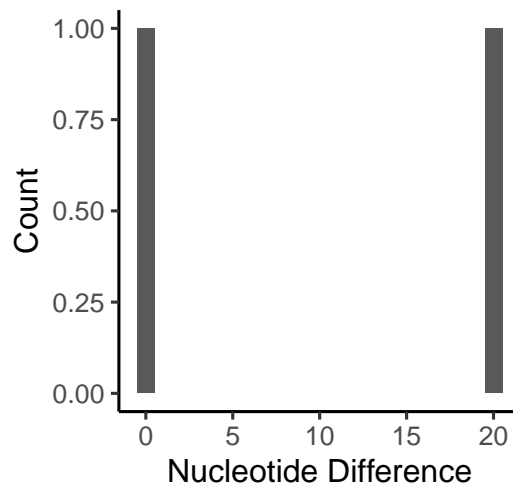
TRBV6-5*bp01

153 sequences assigned
107 (69.9%) exact matches, in which:
95 unique CDR3
11 unique J



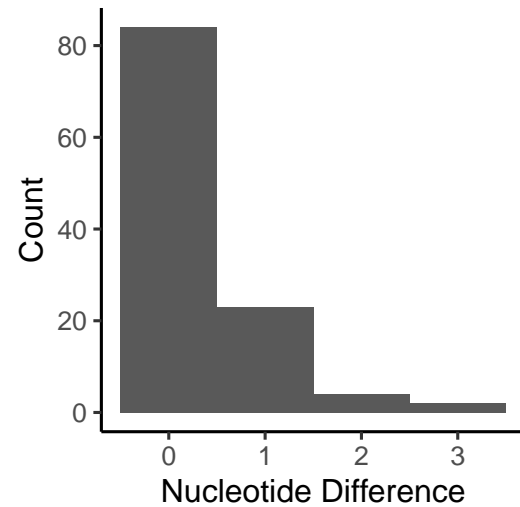
TRBV6-8*bp01

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



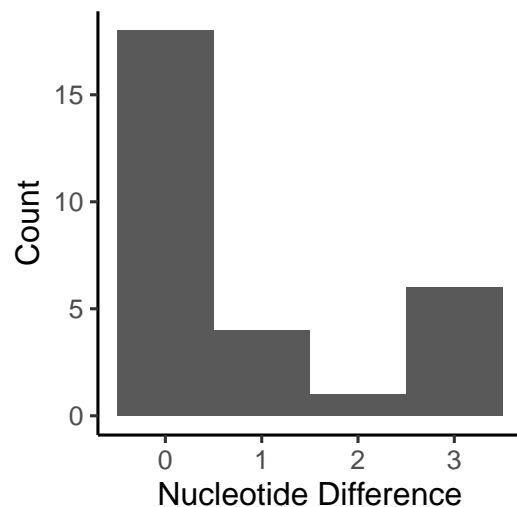
TRBV7-2*bp01

113 sequences assigned
84 (74.3%) exact matches, in which:
51 unique CDR3
11 unique J



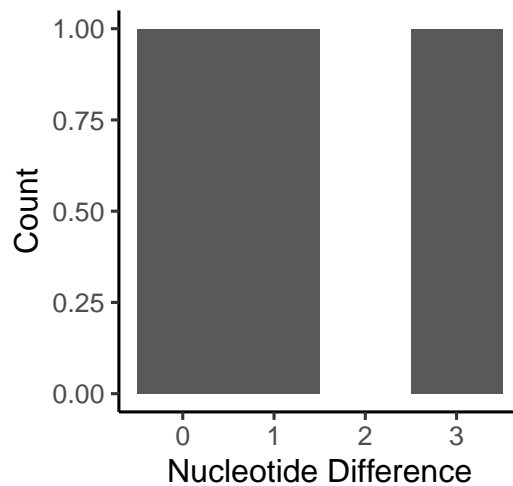
TRBV6-6*bp01

29 sequences assigned
18 (62.1%) exact matches, in which:
15 unique CDR3
6 unique J



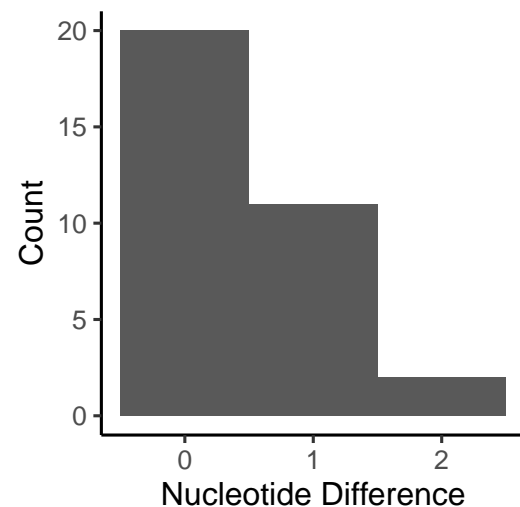
TRBV6-9*bp01

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



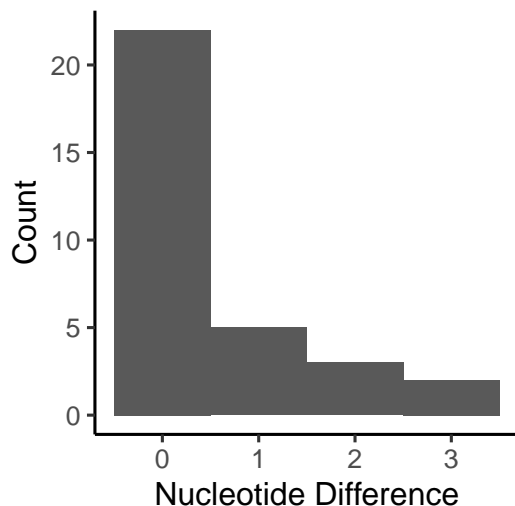
TRBV7-2*bp02

33 sequences assigned
20 (60.6%) exact matches, in which:
14 unique CDR3
7 unique J



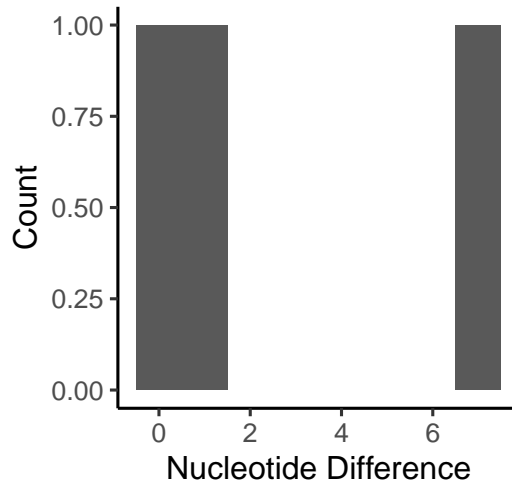
TRBV7-3*bp01

32 sequences assigned
22 (68.8%) exact matches, in which:
21 unique CDR3
7 unique J



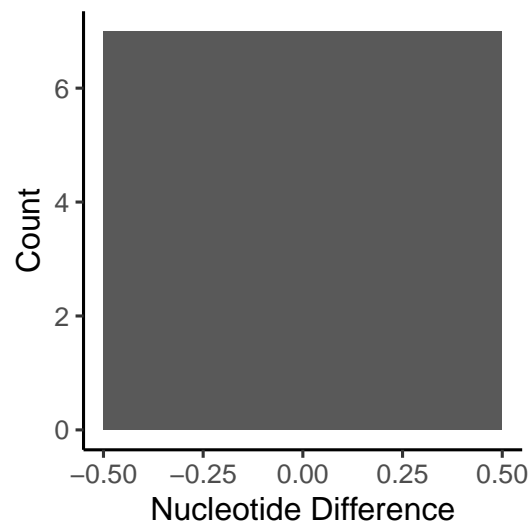
TRBV7-7*bp01

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



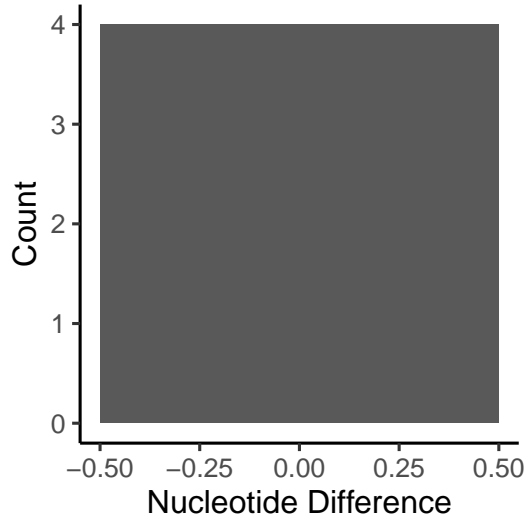
TRBV29-1*bp03

7 sequences assigned
7 (100%) exact matches, in which:
7 unique CDR3
6 unique J



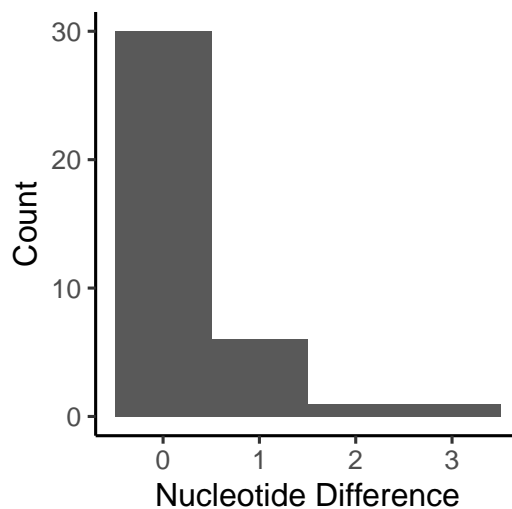
TRBV7-4*bp01

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



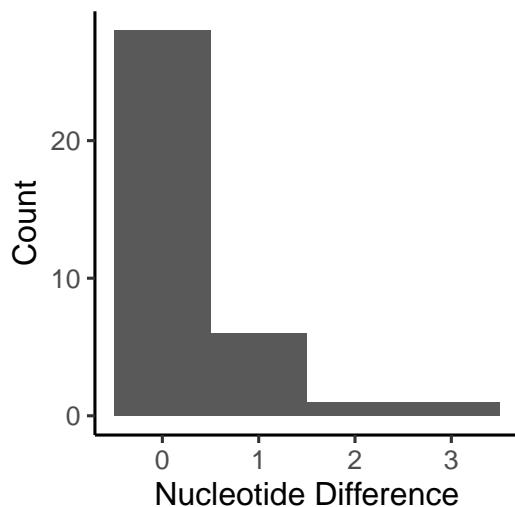
TRBV7-9*bp01

38 sequences assigned
30 (78.9%) exact matches, in which:
27 unique CDR3
8 unique J



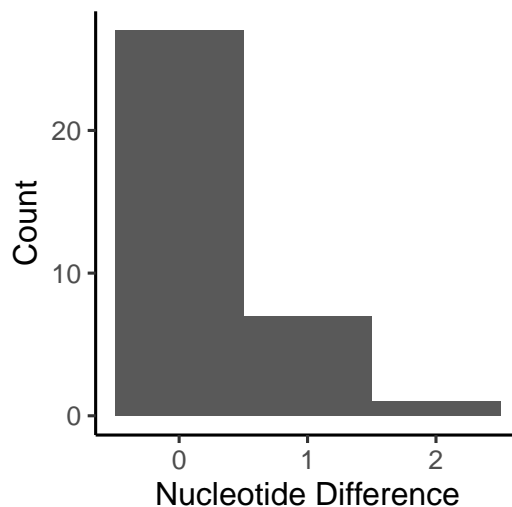
TRBV7-6*bp01

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



TRBV29-1*bp01

35 sequences assigned
27 (77.1%) exact matches, in which:
21 unique CDR3
10 unique J





[illegible]