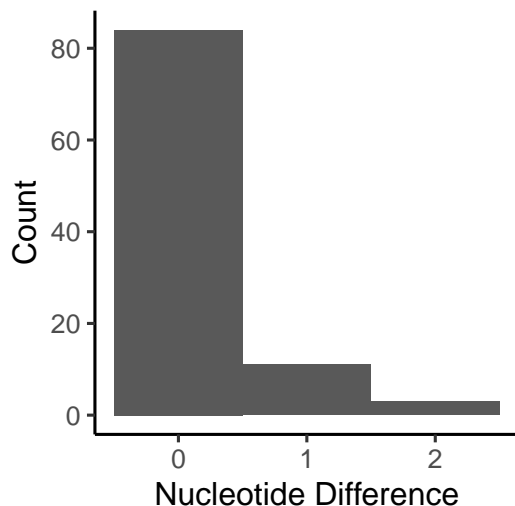


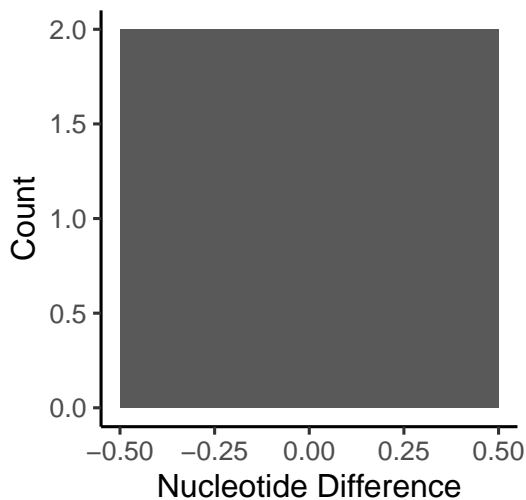
TRBV20–1*bp01

99 sequences assigned
84 (84.8%) exact matches, in which:
57 unique CDR3
12 unique J



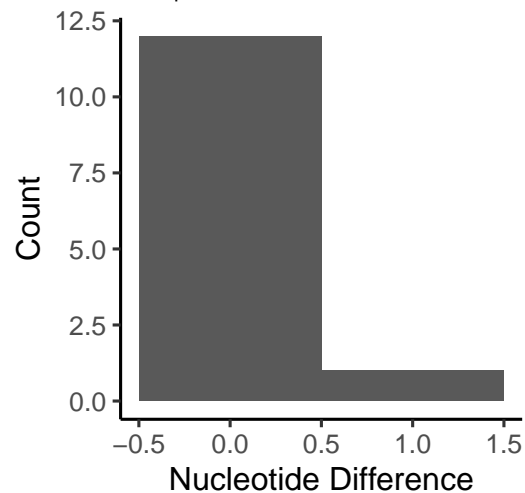
TRBV10–2*bp01

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



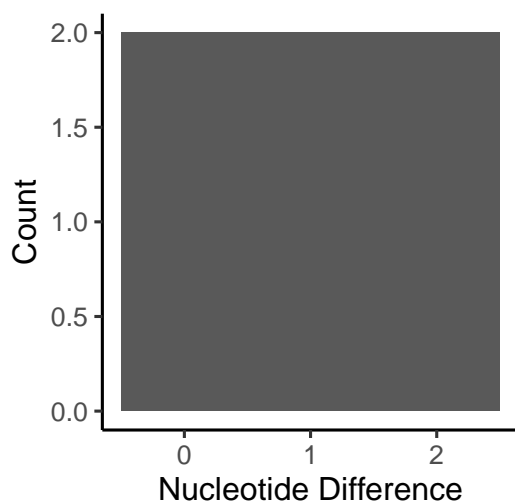
TRBV15*bp02

13 sequences assigned
12 (92.3%) exact matches, in which:
11 unique CDR3
6 unique J



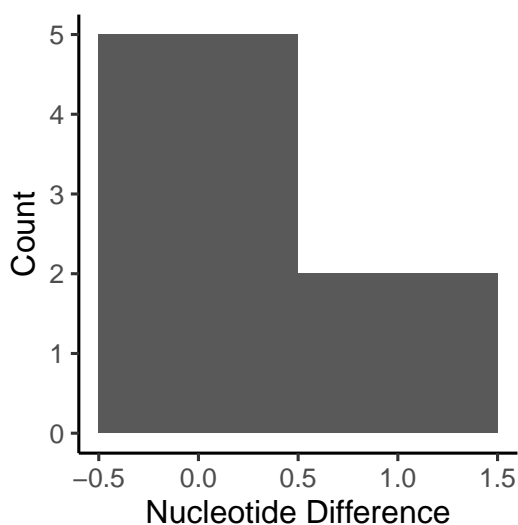
TRBV10–1*bp01

6 sequences assigned
2 (33.3%) exact matches, in which:
1 unique CDR3
1 unique J



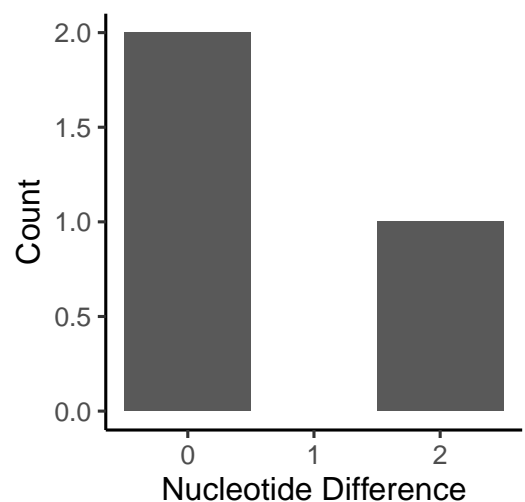
TRBV10–3*bp01

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



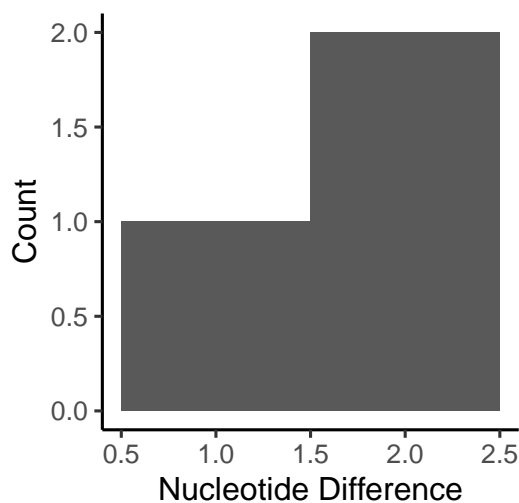
TRBV18*bp01

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



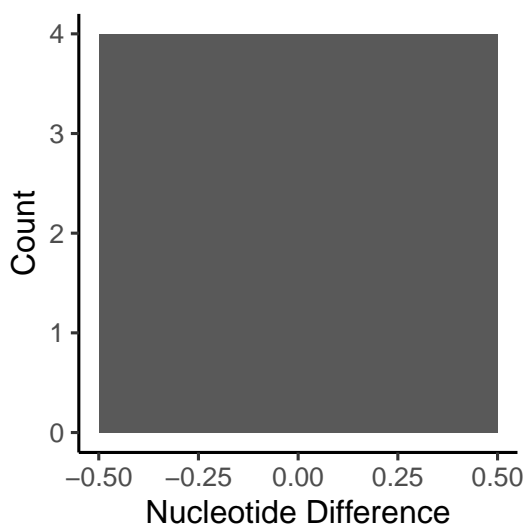
TRBV10–1*bp02

3 sequences assigned
No exact matches.



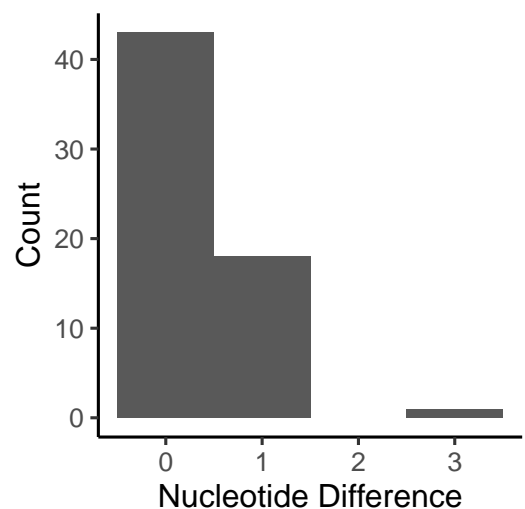
TRBV14*bp01

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



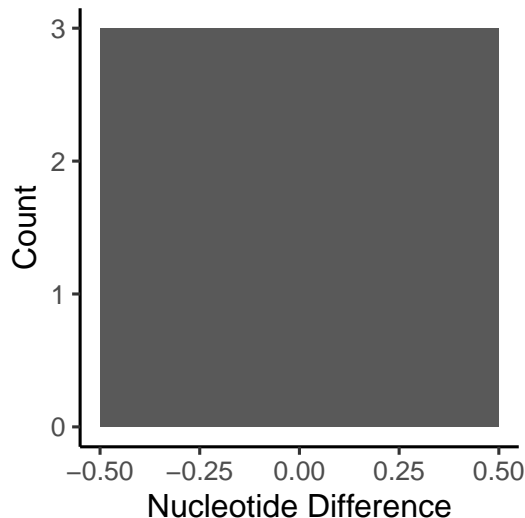
TRBV19*bp01

67 sequences assigned
43 (64.2%) exact matches, in which:
33 unique CDR3
12 unique J



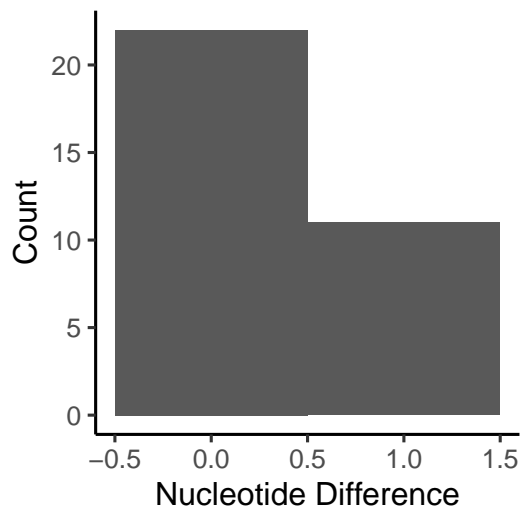
TRBV2*bp01

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



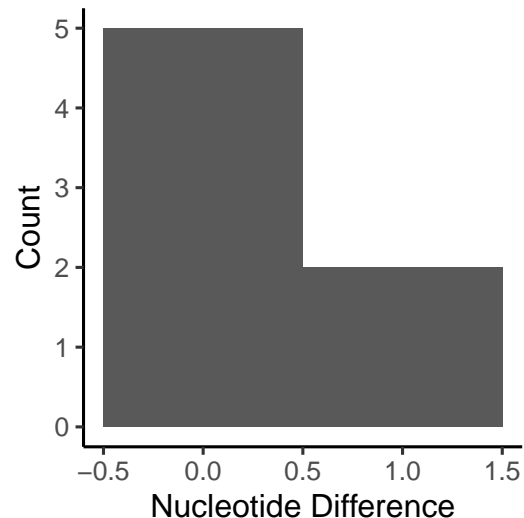
TRBV30*bp01

33 sequences assigned
22 (66.7%) exact matches, in which:
14 unique CDR3
9 unique J



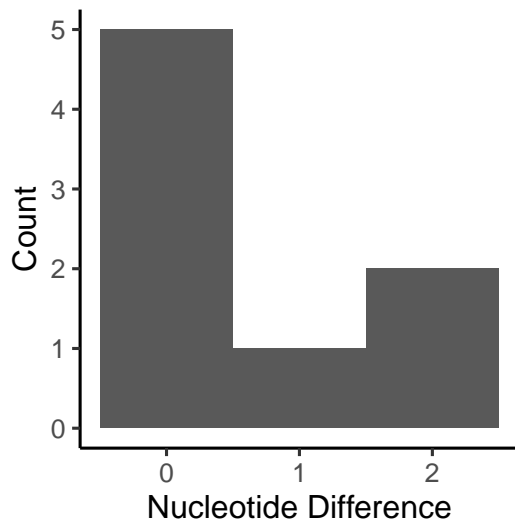
TRBV21-1*bp01

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



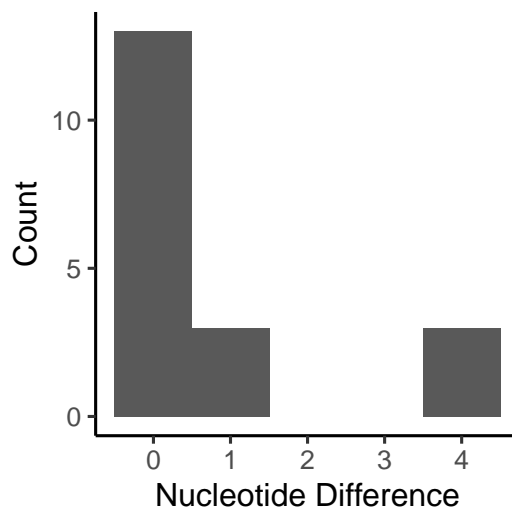
TRBV27*bp01

8 sequences assigned
5 (62.5%) exact matches, in which:
5 unique CDR3
5 unique J



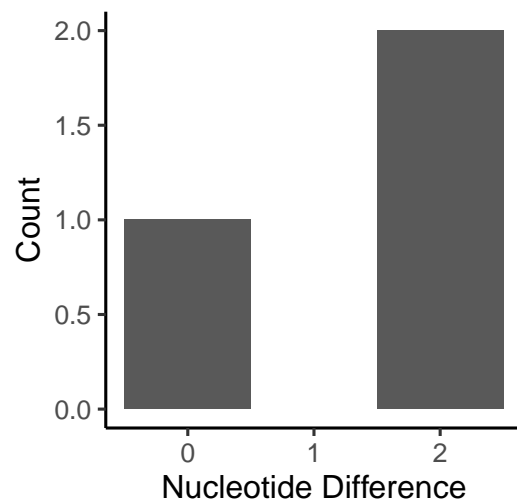
TRBV30*bp02

19 sequences assigned
13 (68.4%) exact matches, in which:
8 unique CDR3
5 unique J



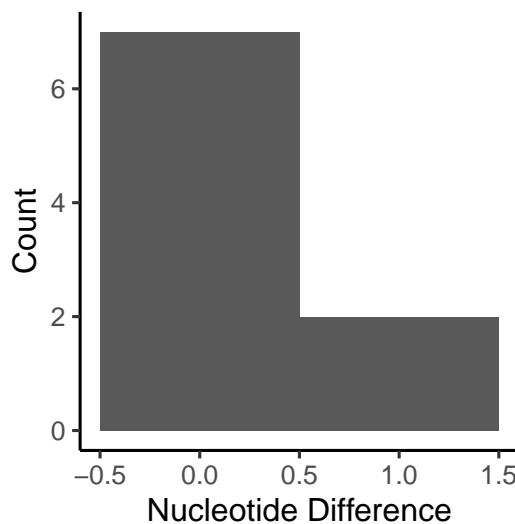
TRBV11-1*bp01

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



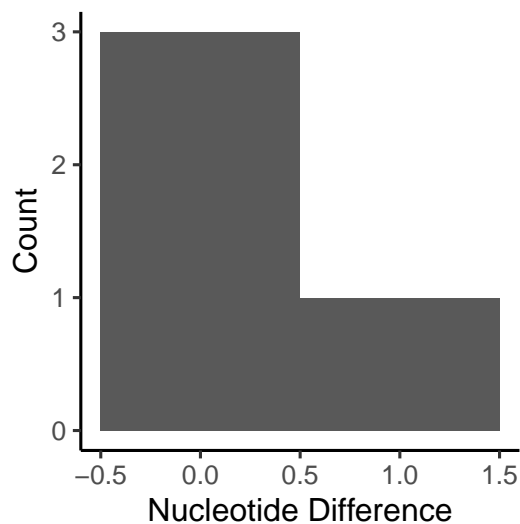
TRBV28*bp01

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



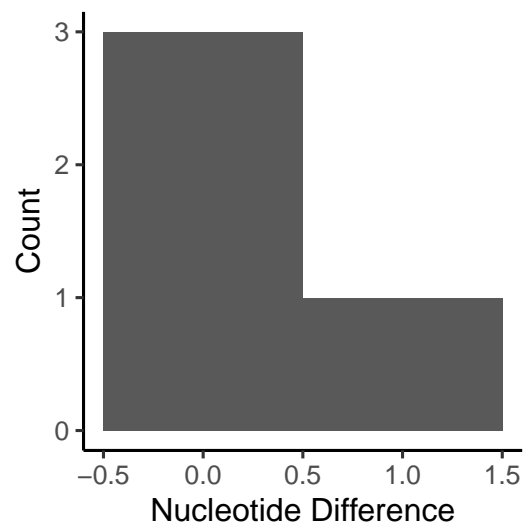
TRBV9*bp01

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



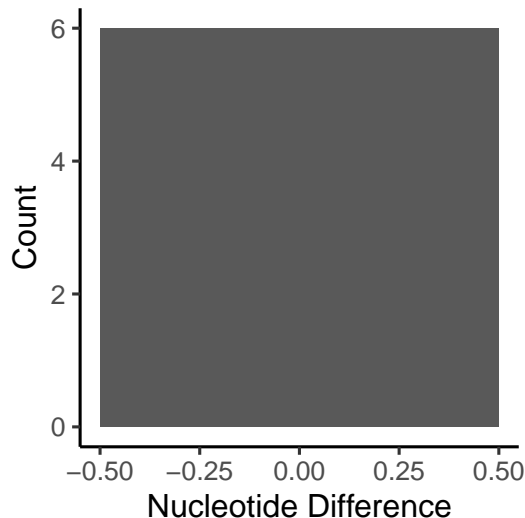
TRBV21-1*bp02

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



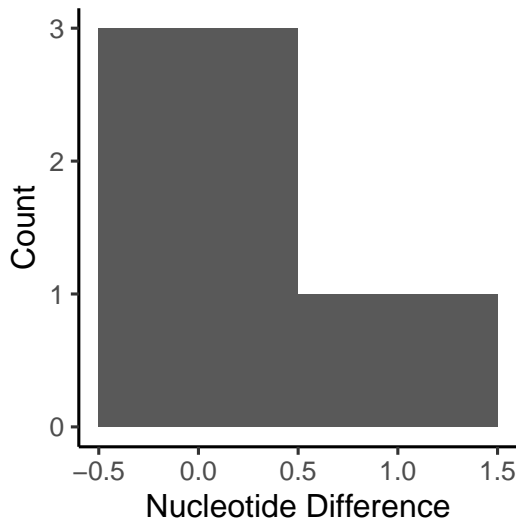
TRBV11–2*bp01

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



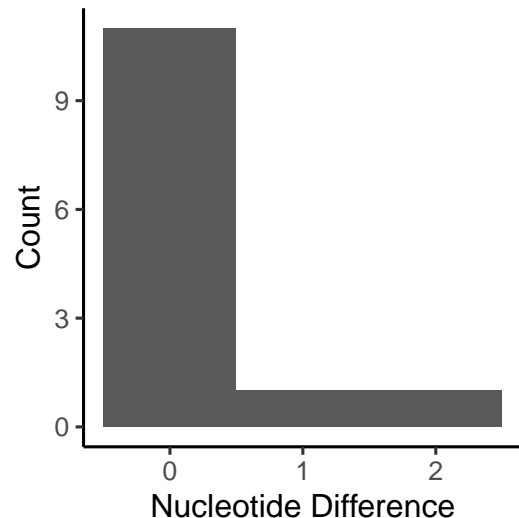
TRBV12–5*bp01

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



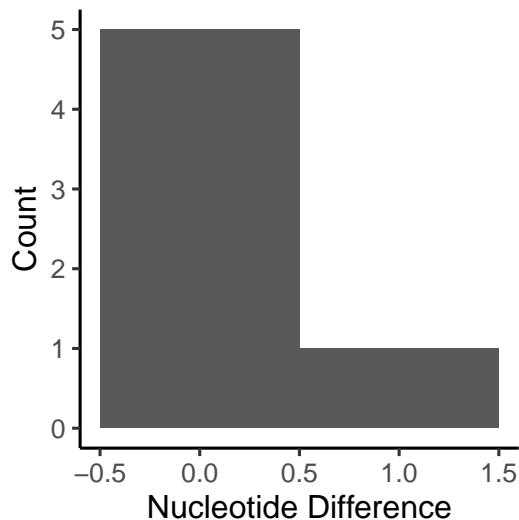
TRBV4–1*bp01

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



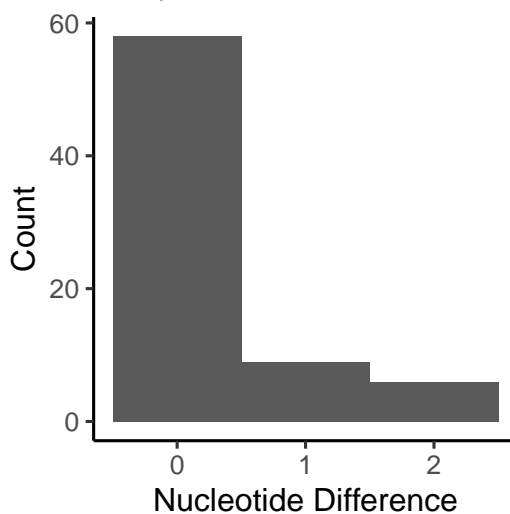
TRBV11–2*bp03

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



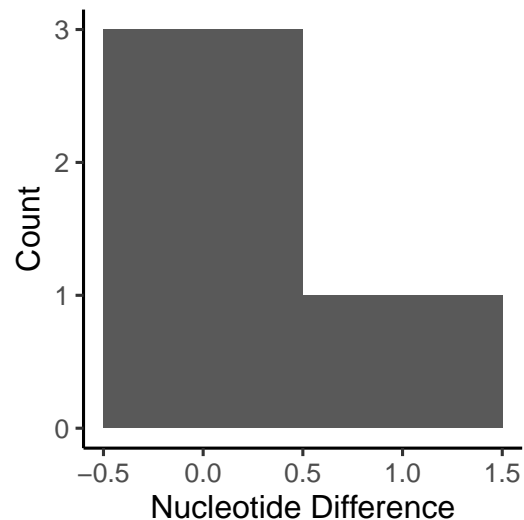
TRBV12–34*bp01

73 sequences assigned
58 (79.5%) exact matches, in which:
34 unique CDR3
12 unique J



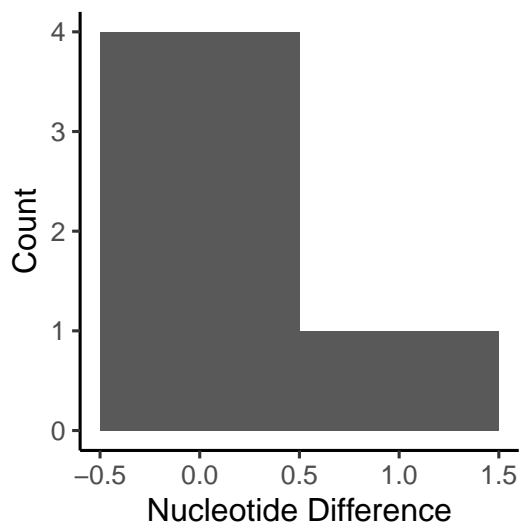
TRBV24–1*bp02

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



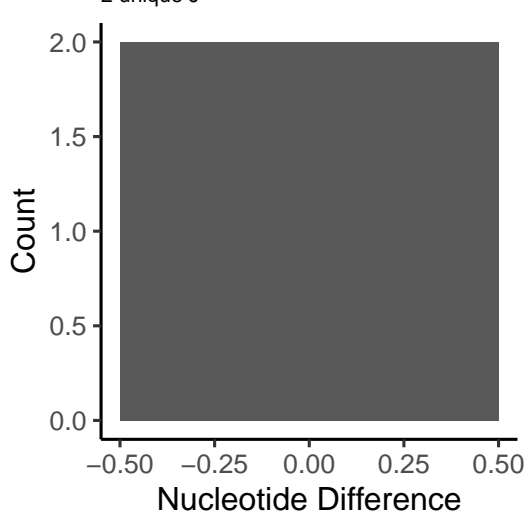
TRBV11–3*bp01

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



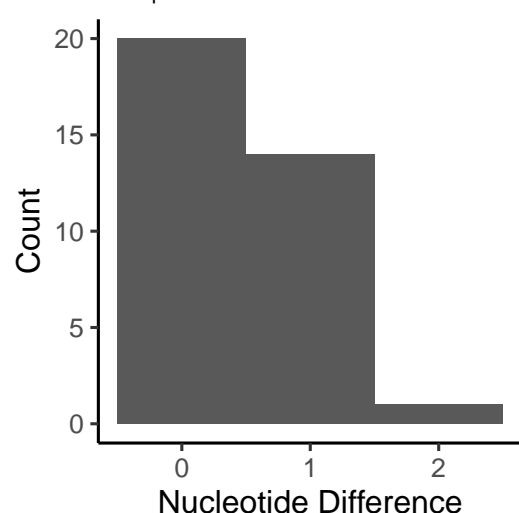
TRBV24–1*bp01

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



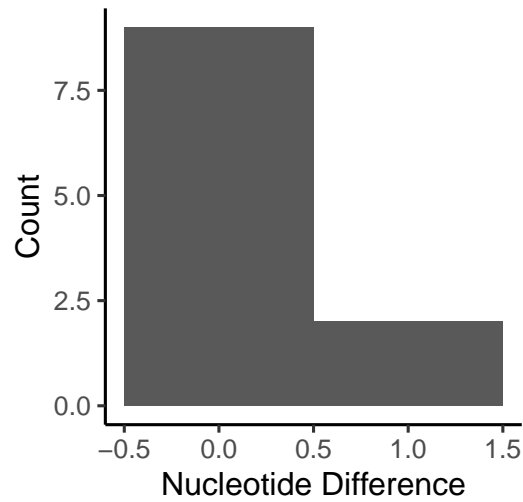
TRBV4–2*bp01

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



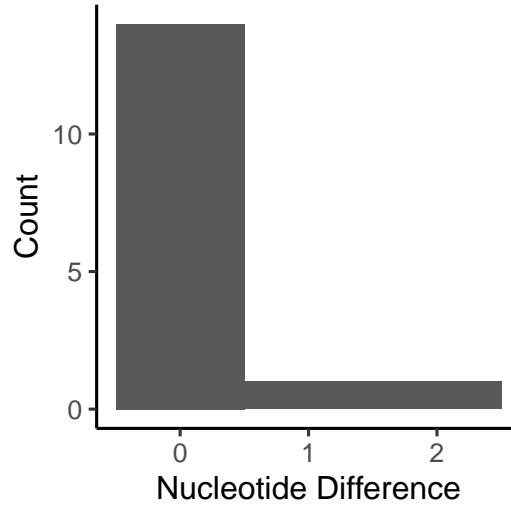
TRBV25–1*bp01

12 sequences assigned
9 (75%) exact matches, in which:
4 unique CDR3
3 unique J



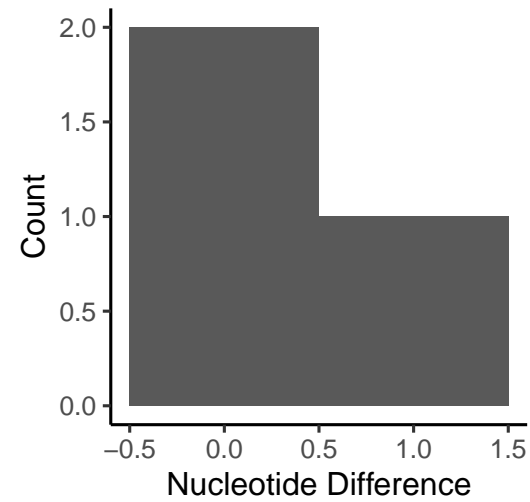
TRBV5–4*bp01

16 sequences assigned
14 (87.5%) exact matches, in which:
10 unique CDR3
7 unique J



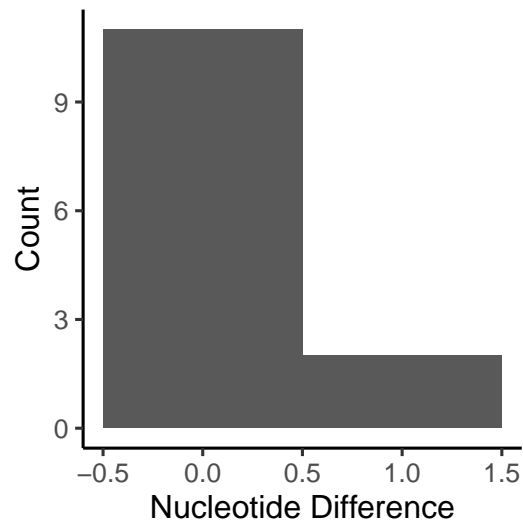
TRBV5–8*bp01

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



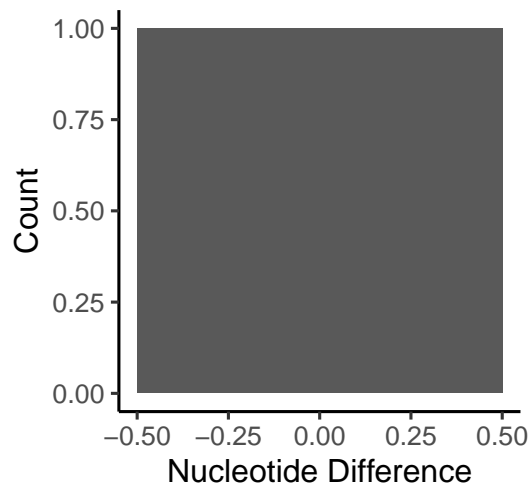
TRBV5–1*bp01

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



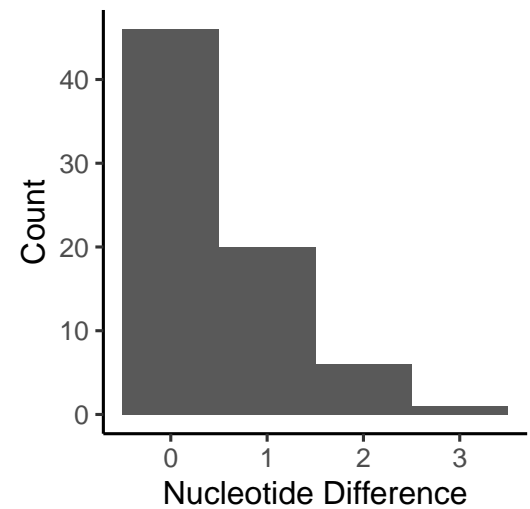
TRBV5–5*bp01

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



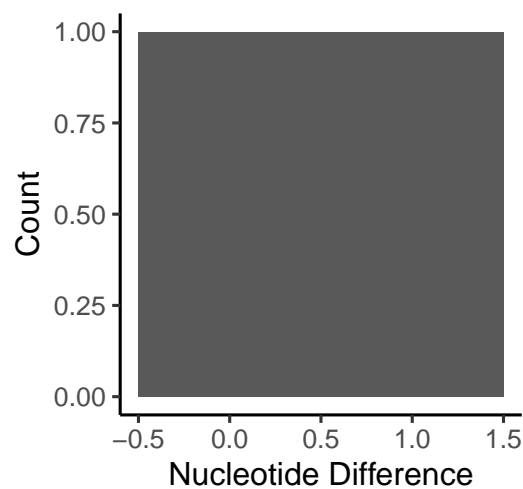
TRBV6–1*bp01

73 sequences assigned
46 (63%) exact matches, in which:
40 unique CDR3
12 unique J



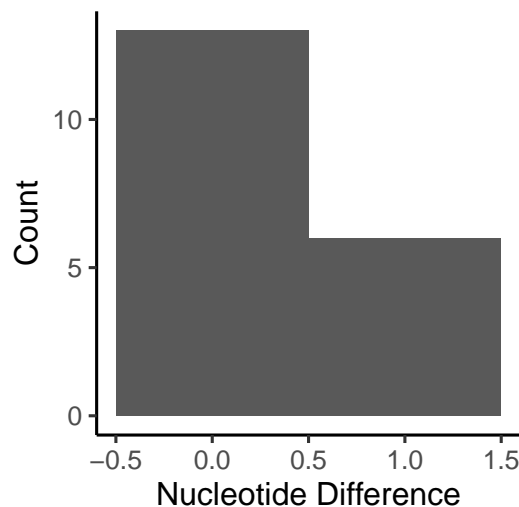
TRBV5–3*bp01

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



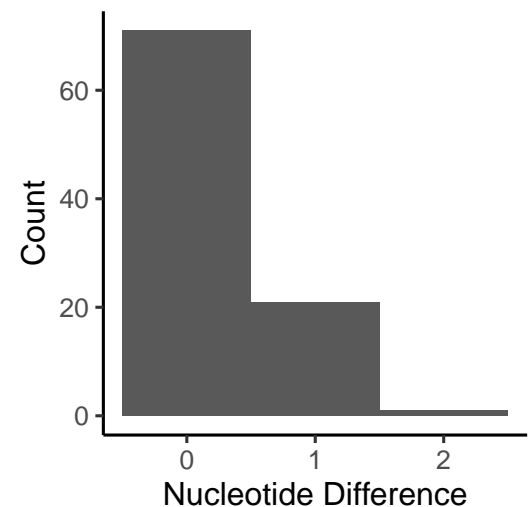
TRBV5–6*bp01

19 sequences assigned
13 (68.4%) exact matches, in which:
12 unique CDR3
8 unique J



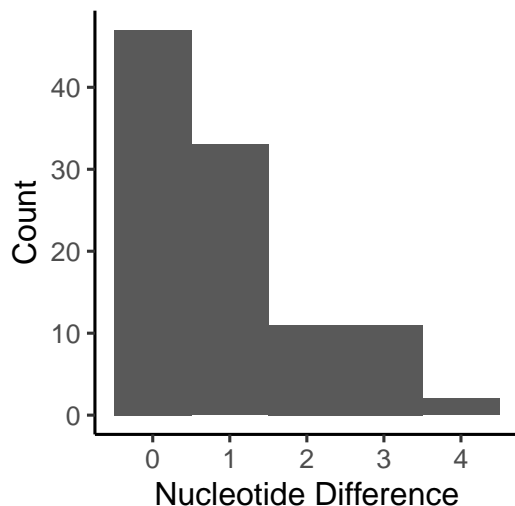
TRBV6–4*bp01

93 sequences assigned
71 (76.3%) exact matches, in which:
53 unique CDR3
10 unique J



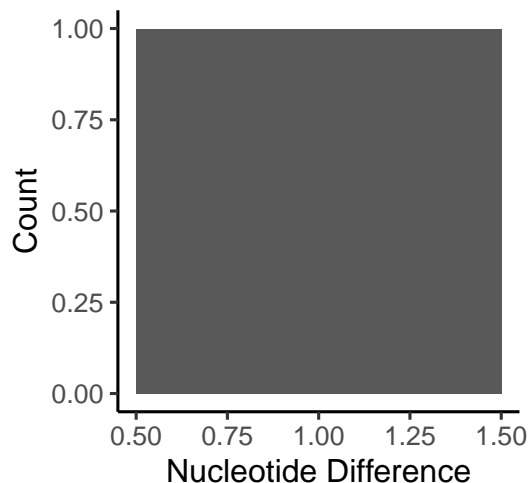
TRBV6-5*bp01

105 sequences assigned
47 (44.8%) exact matches, in which:
38 unique CDR3
11 unique J



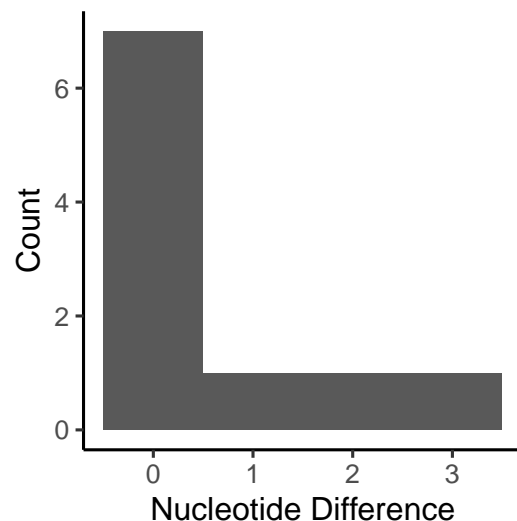
TRBV6-8*bp01

1 sequences assigned
No exact matches.



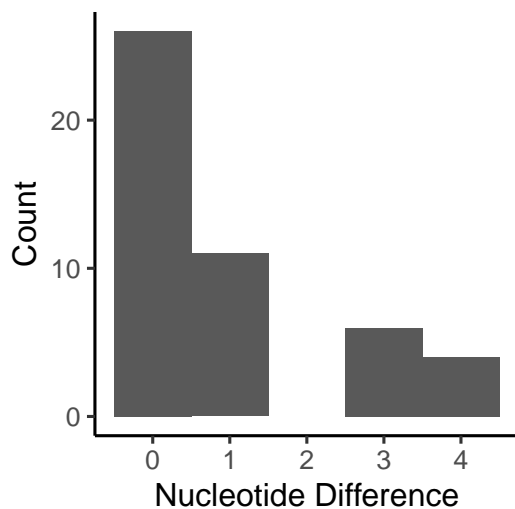
TRBV7-2*bp01

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



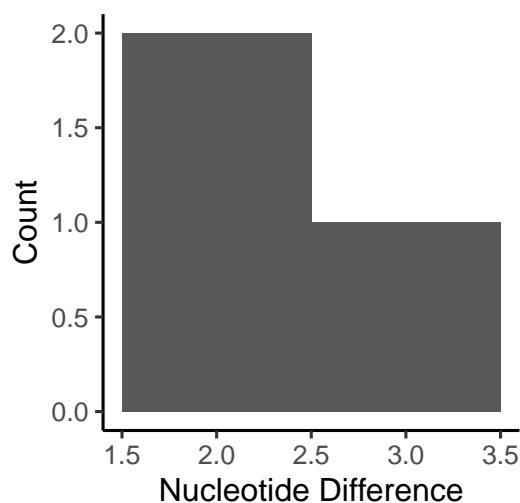
TRBV6-6*bp01

47 sequences assigned
26 (55.3%) exact matches, in which:
23 unique CDR3
10 unique J



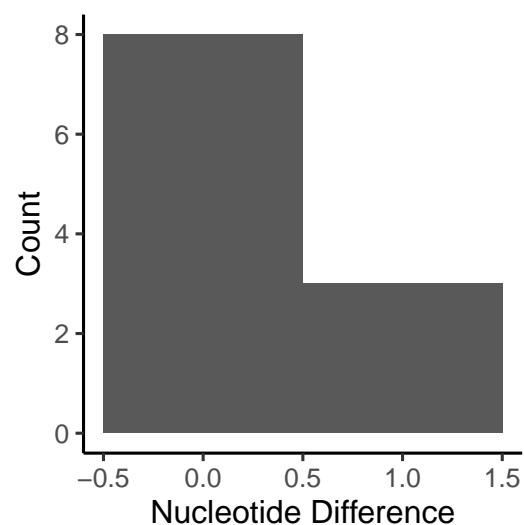
TRBV6-9*bp01

4 sequences assigned
No exact matches.



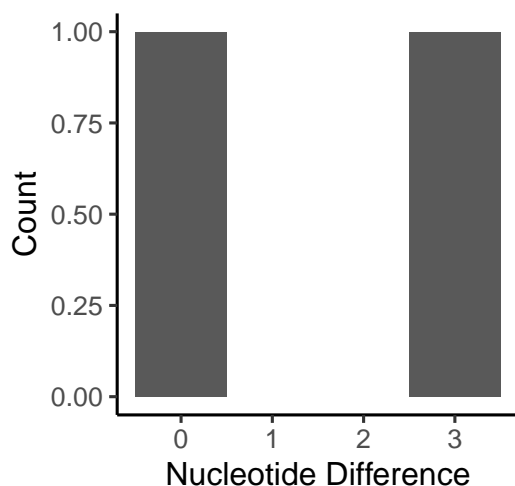
TRBV7-2*bp02

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



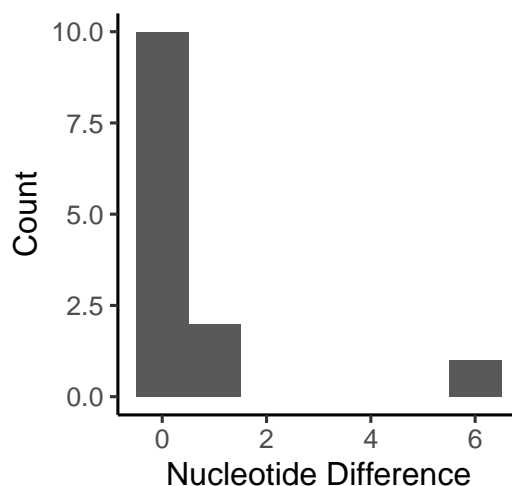
TRBV6-7*bp01

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



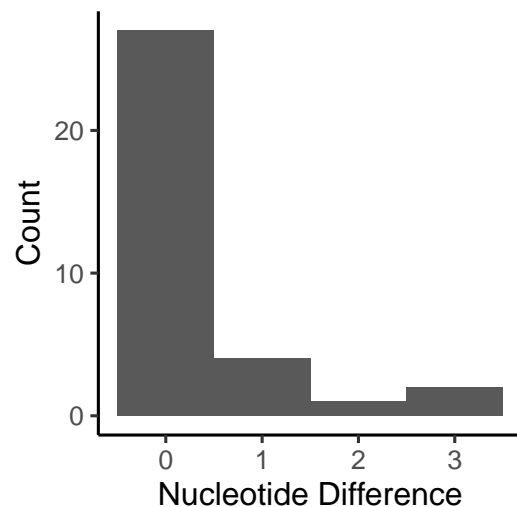
TRBV6-23*bp01

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



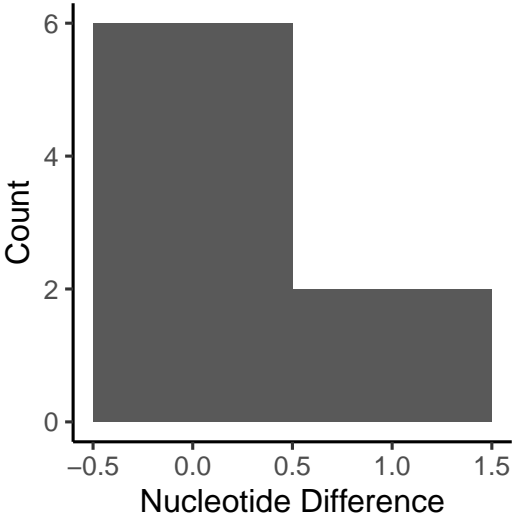
TRBV7-3*bp01

34 sequences assigned
27 (79.4%) exact matches, in which:
17 unique CDR3
9 unique J



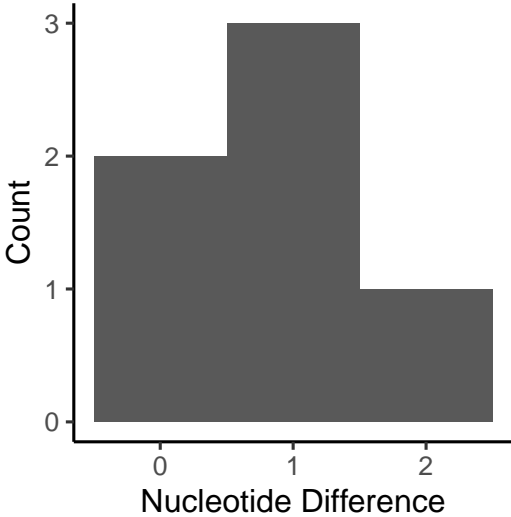
TRBV7-6*bp01

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



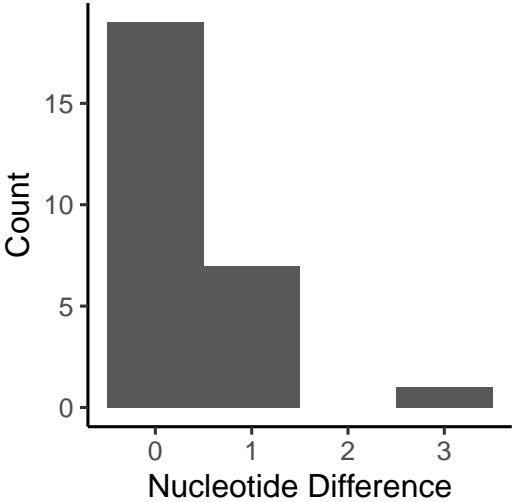
TRBV29-1*bp03

6 sequences assigned
2 (33.3%) exact matches, in which:
1 unique CDR3
1 unique J



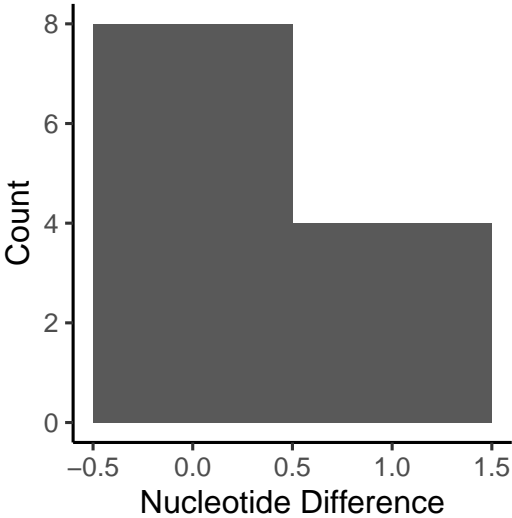
TRBV7-9*bp01

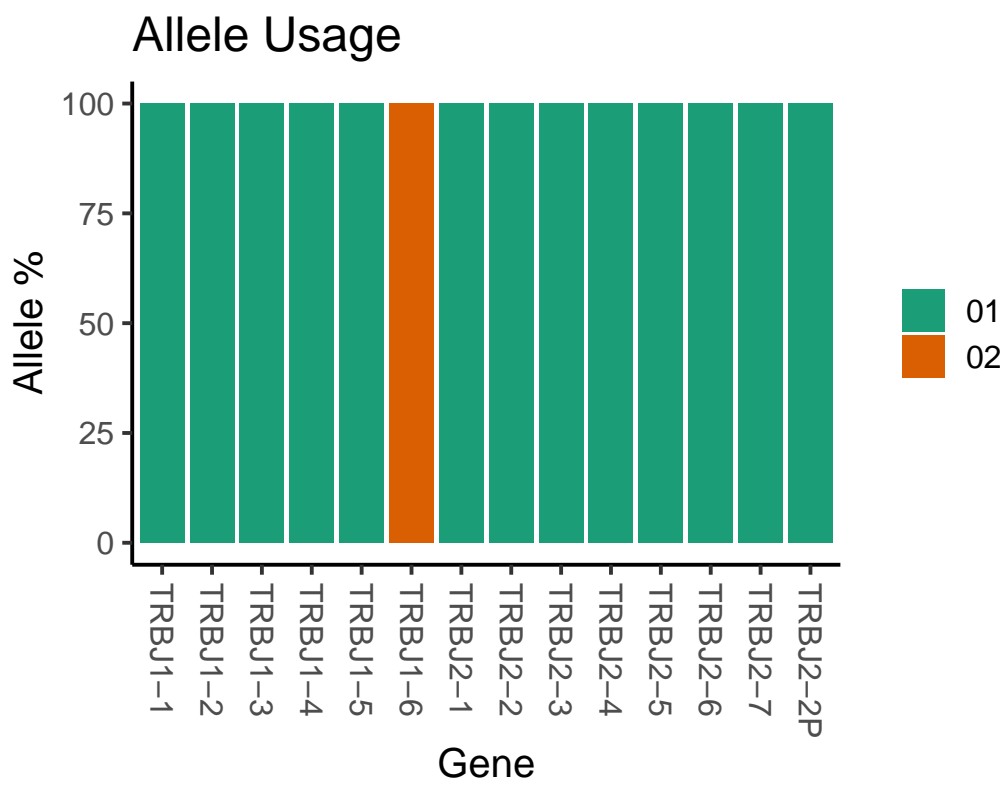
27 sequences assigned
19 (70.4%) exact matches, in which:
9 unique CDR3
6 unique J



TRBV29-1*bp01

12 sequences assigned
8 (66.7%) exact matches, in which:
6 unique CDR3
5 unique J





Warning – no inferred sequences found.

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