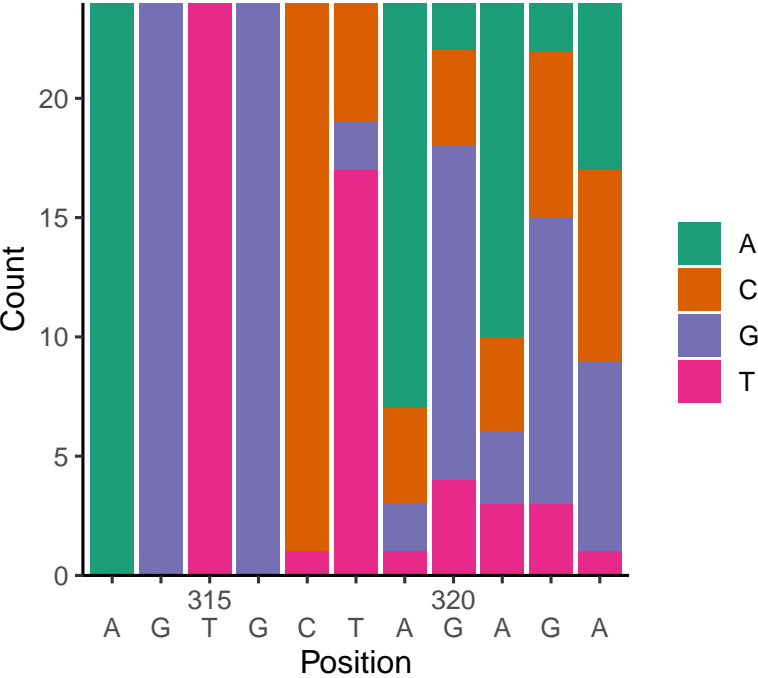
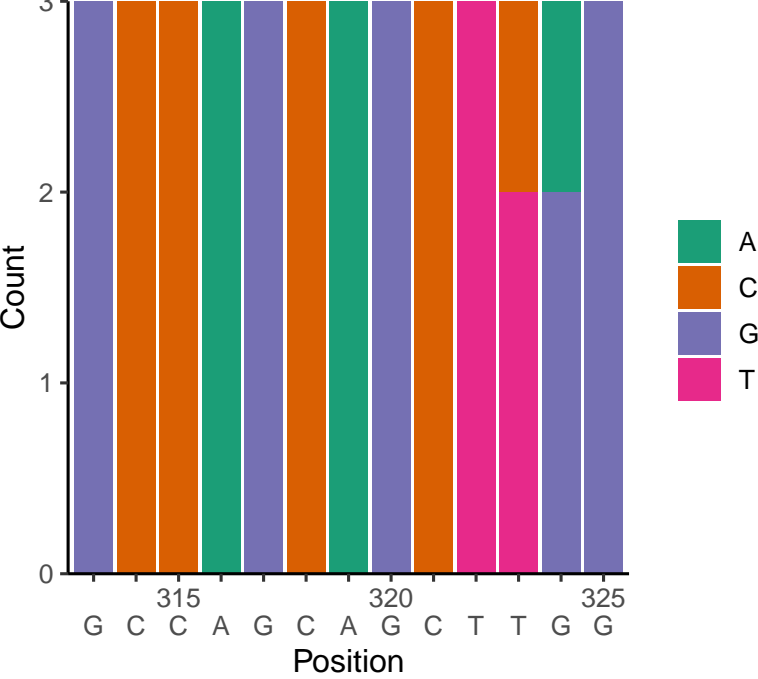


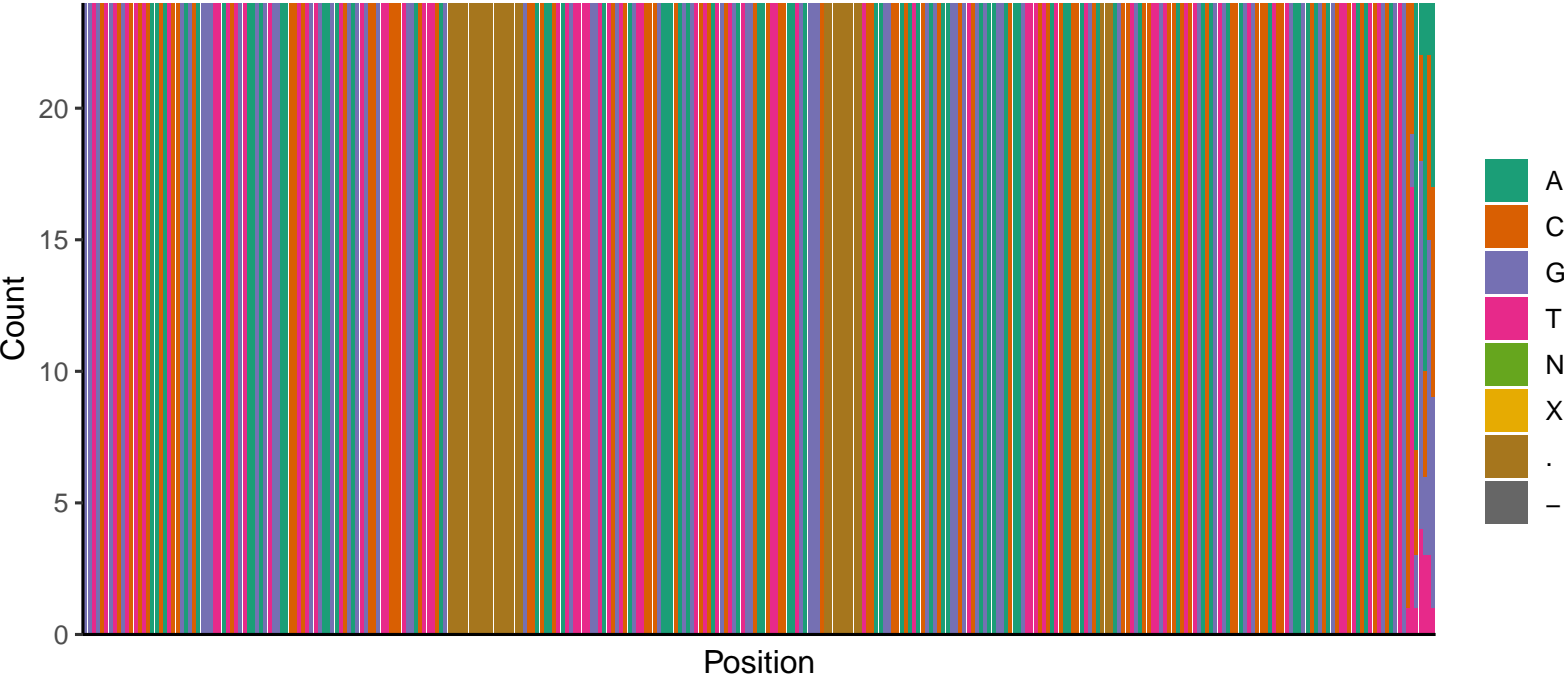
Gene TRBV20-1*02_319AGAGA323



Gene TRBV5-5*02_322TTGG325



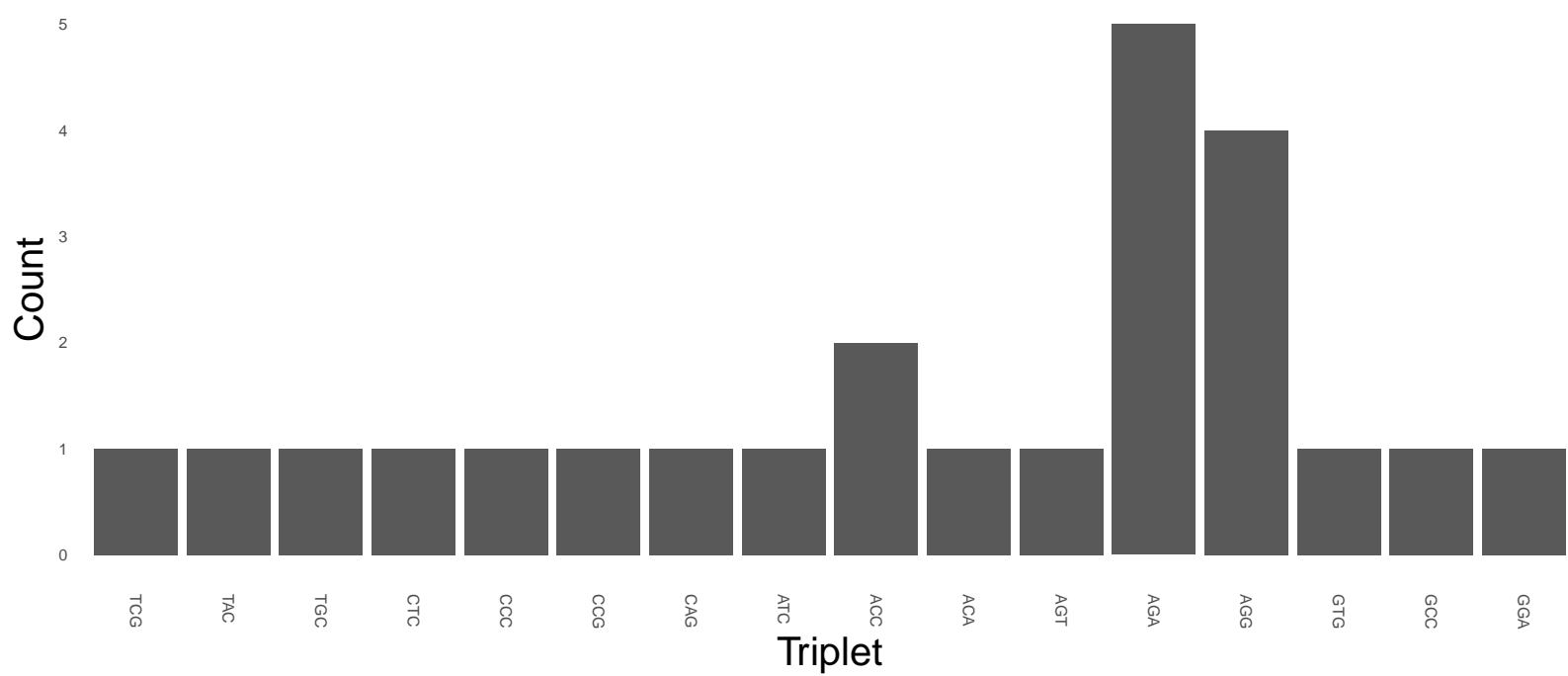
Gene TRBV20-1*02_319AGAGA323



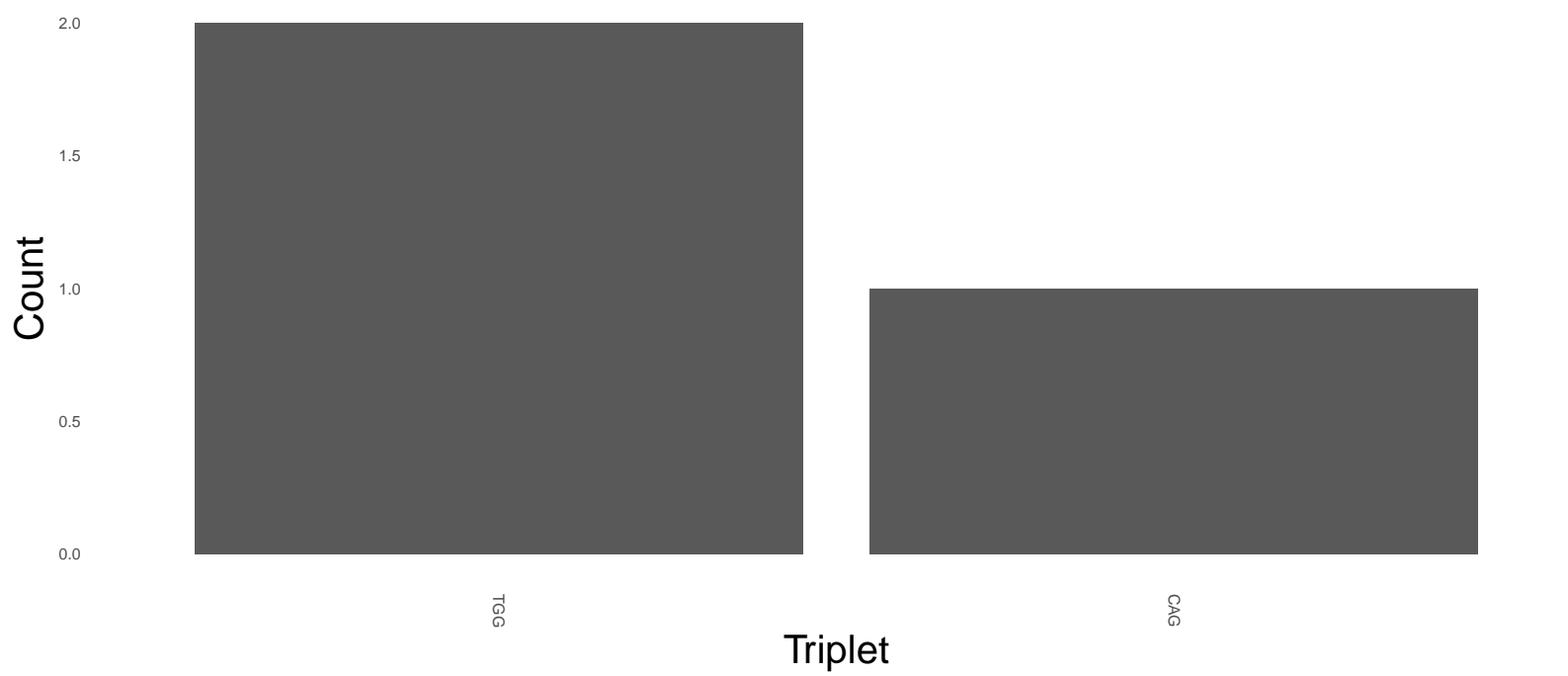
Gene TRBV5-5*02_322TTGG325



TRBV20–1*02_319AGAGA323– Final 3 nucleotides as a triplet

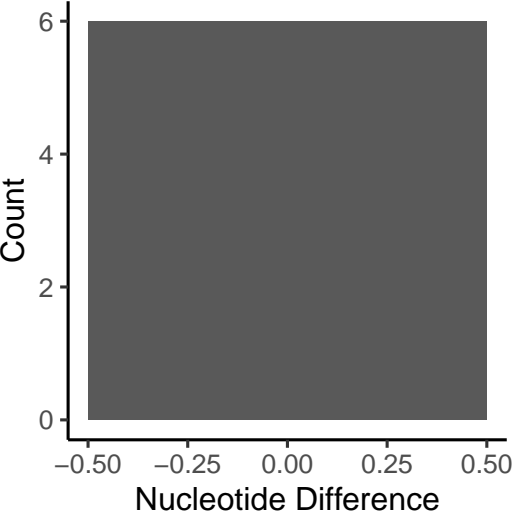


TRBV5–5*02_322TTGG325– Final 3 nucleotides as a triplet



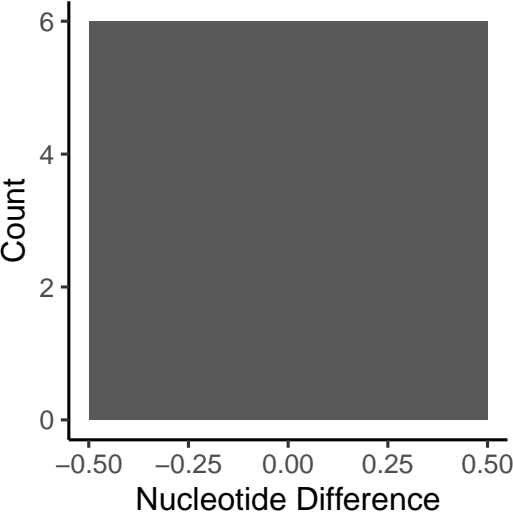
TRBV10–1*01

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



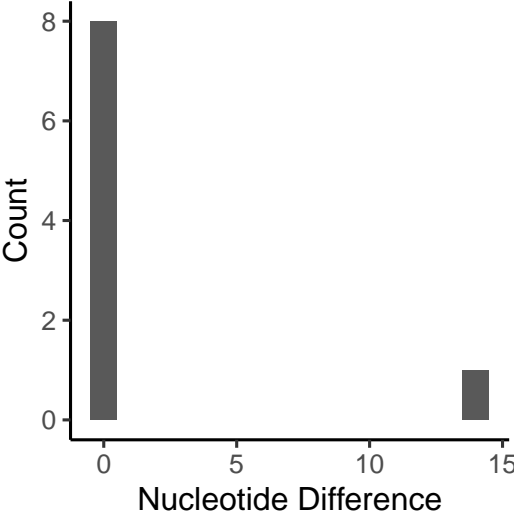
TRBV10–2*01

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



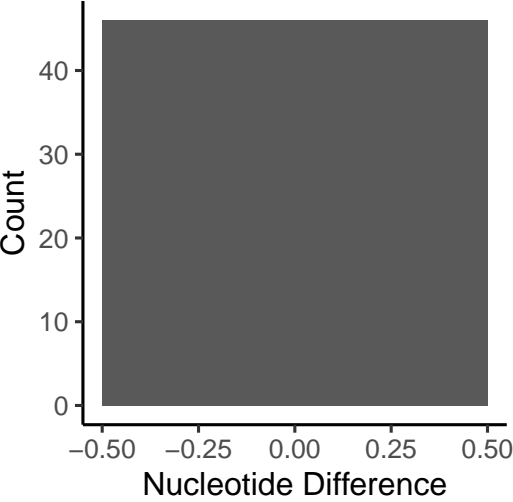
TRBV13*01

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



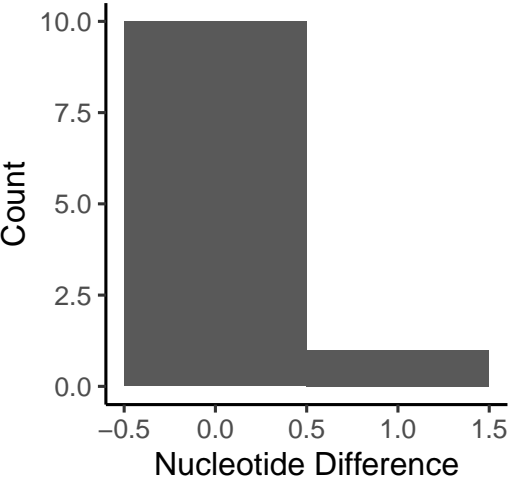
TRBV20–1*01

46 sequences assigned
46 (100%) exact matches, in which:
46 unique CDR3
10 unique J



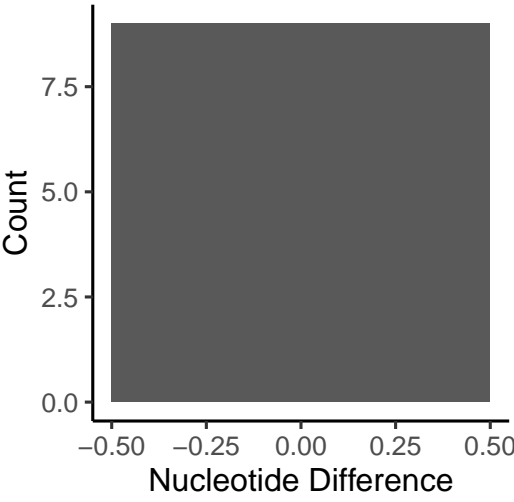
TRBV10–3*02

11 sequences assigned
10 (90.9%) exact matches, in which:
10 unique CDR3
6 unique J



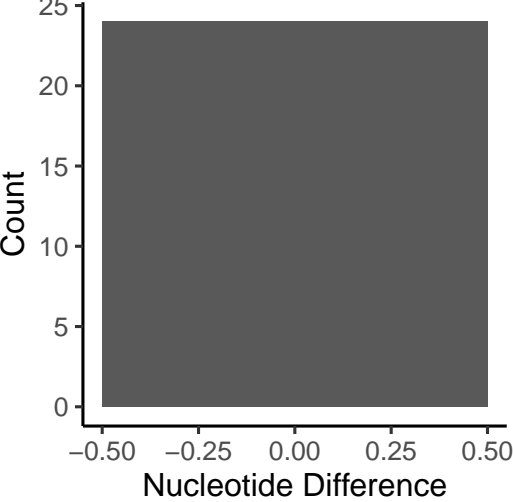
TRBV14*01

9 sequences assigned
9 (100%) exact matches, in which:
9 unique CDR3
7 unique J



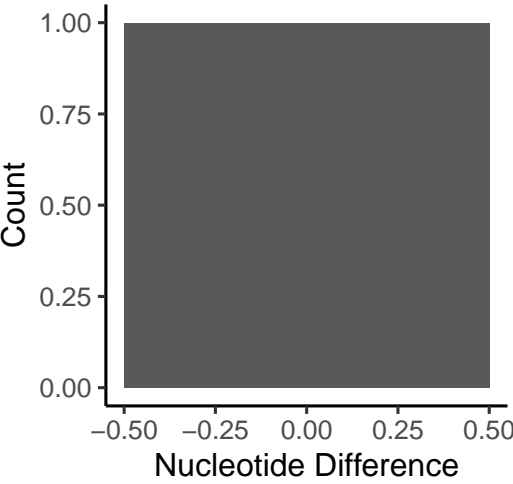
TRBV20–1*02_319AGAG/

24 sequences assigned
24 (100%) exact matches, in which:
24 unique CDR3
12 unique J



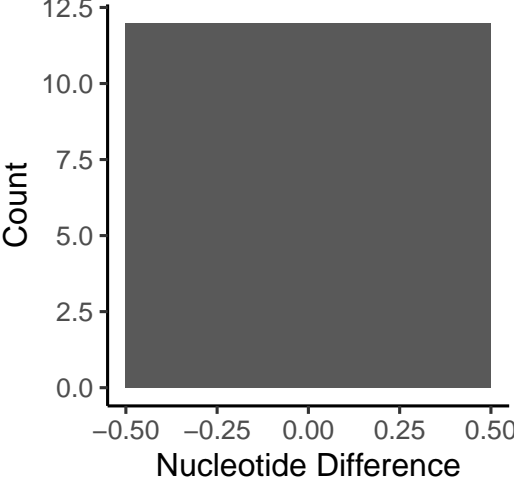
TRBV1*01

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



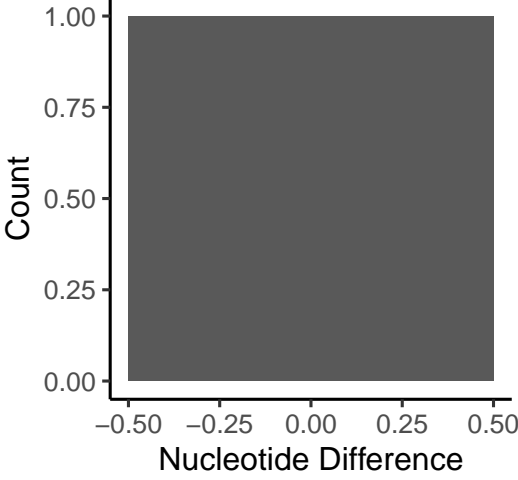
TRBV15*02

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



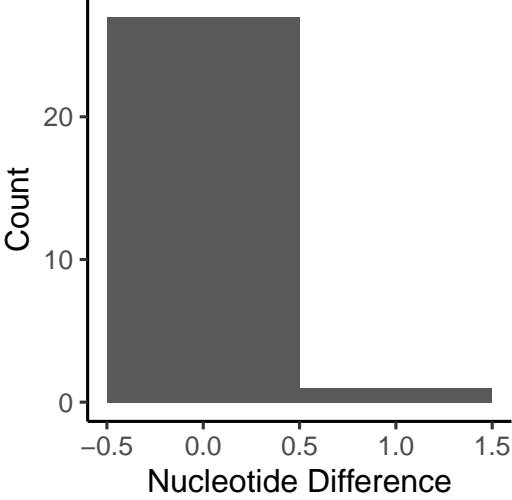
TRBV16*01

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



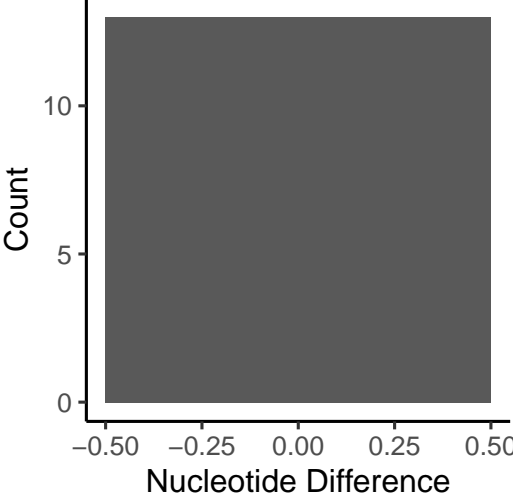
TRBV2*01

28 sequences assigned
27 (96.4%) exact matches, in which:
27 unique CDR3
9 unique J



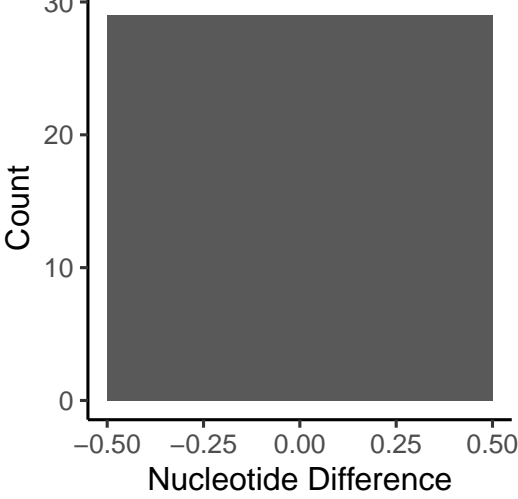
TRBV30*01

13 sequences assigned
13 (100%) exact matches, in which:
13 unique CDR3
9 unique J



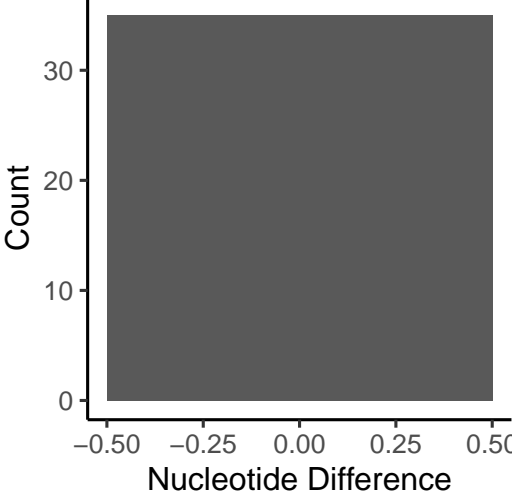
TRBV18*01

29 sequences assigned
29 (100%) exact matches, in which:
29 unique CDR3
11 unique J



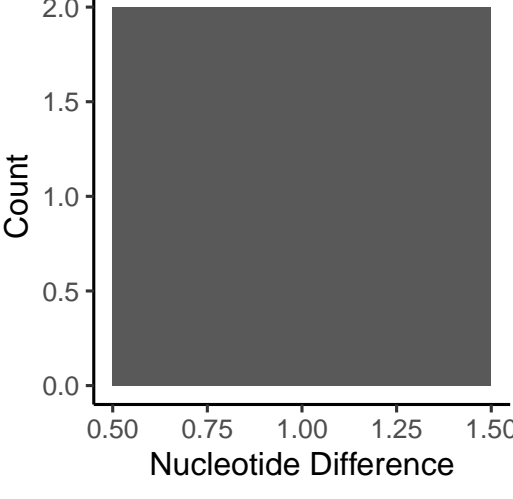
TRBV27*01

35 sequences assigned
35 (100%) exact matches, in which:
35 unique CDR3
11 unique J



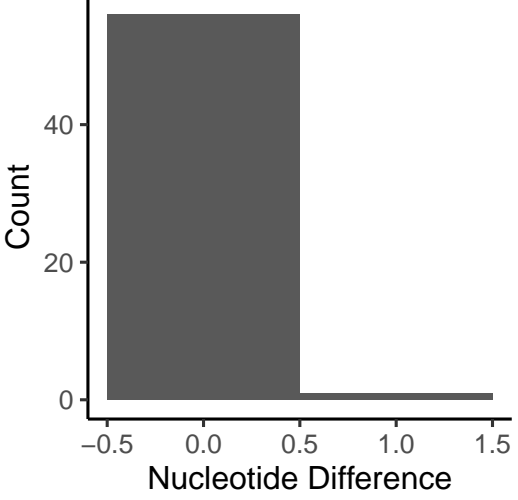
TRBV30*03

2 sequences assigned
No exact matches.



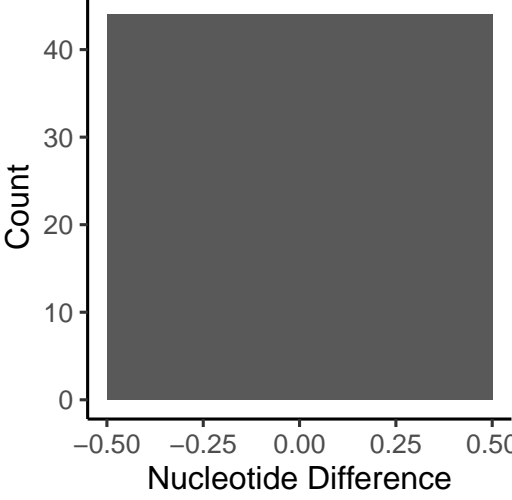
TRBV19*01

57 sequences assigned
56 (98.2%) exact matches, in which:
56 unique CDR3
13 unique J



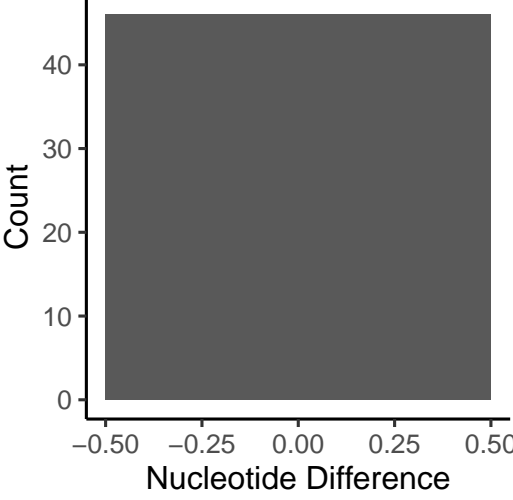
TRBV28*01

44 sequences assigned
44 (100%) exact matches, in which:
44 unique CDR3
10 unique J



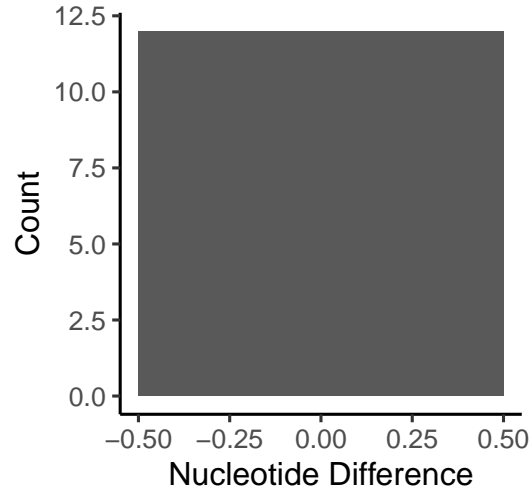
TRBV9*01

46 sequences assigned
46 (100%) exact matches, in which:
46 unique CDR3
11 unique J



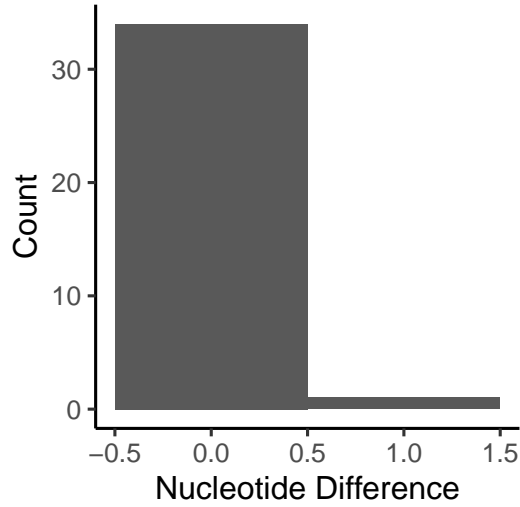
TRBV21-1*01

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



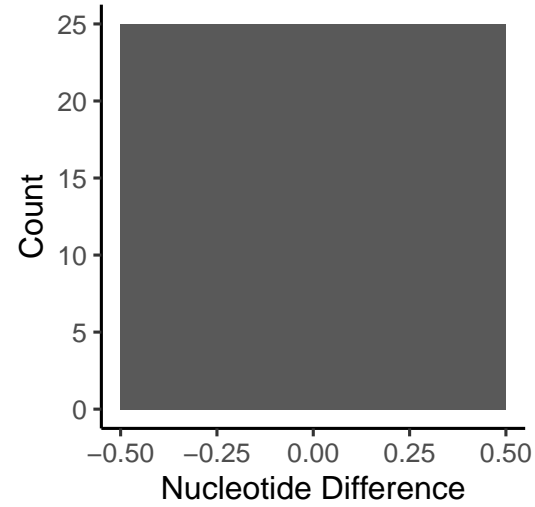
TRBV11-2*01

35 sequences assigned
34 (97.1%) exact matches, in which:
33 unique CDR3
13 unique J



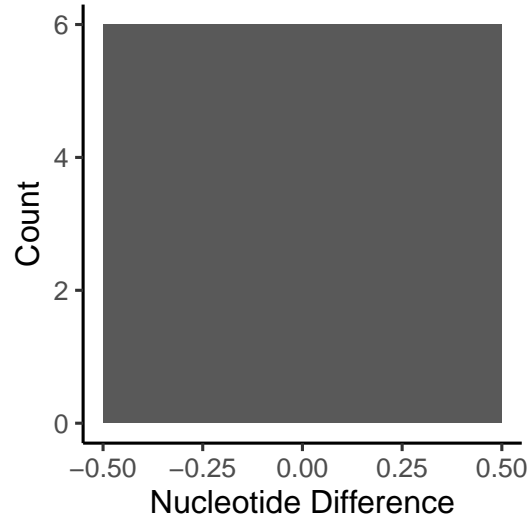
TRBV12-3*01

25 sequences assigned
25 (100%) exact matches, in which:
25 unique CDR3
11 unique J



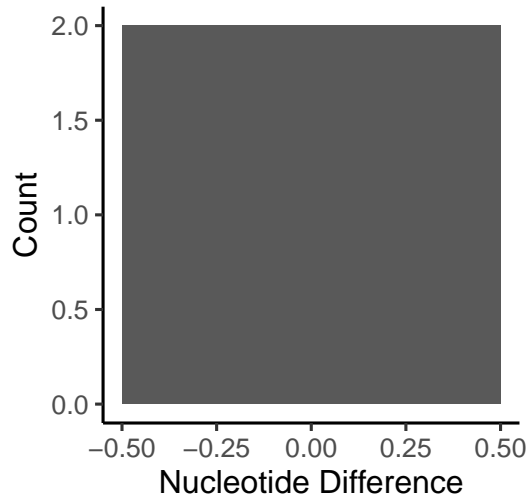
TRBV11-1*01

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



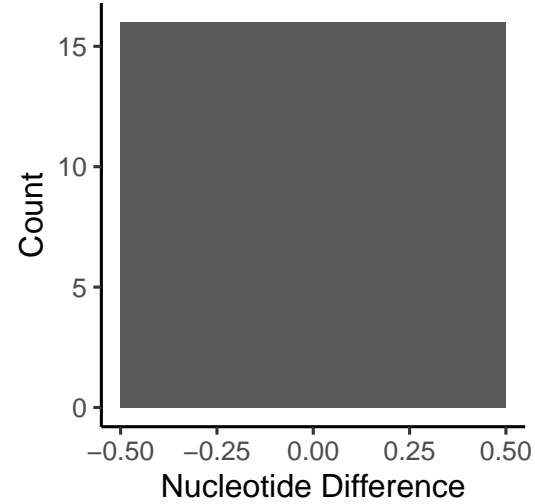
TRBV11-3*01

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



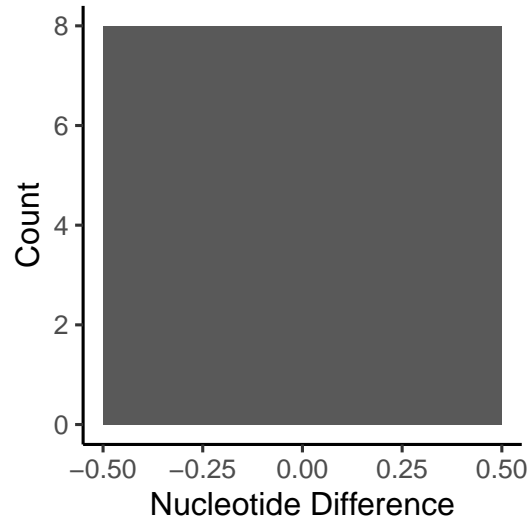
TRBV12-4*01

16 sequences assigned
16 (100%) exact matches, in which:
16 unique CDR3
10 unique J



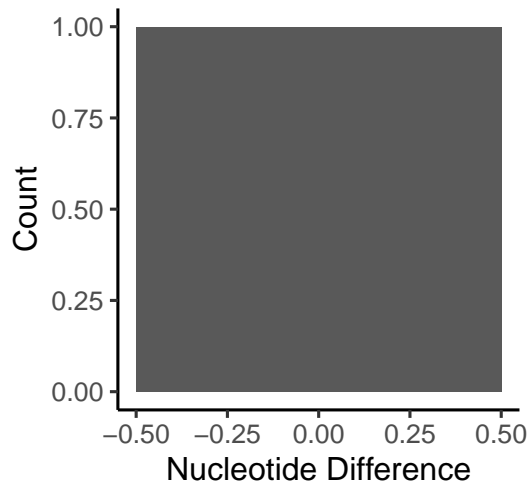
TRBV21-1*02

8 sequences assigned
8 (100%) exact matches, in which:
8 unique CDR3
7 unique J



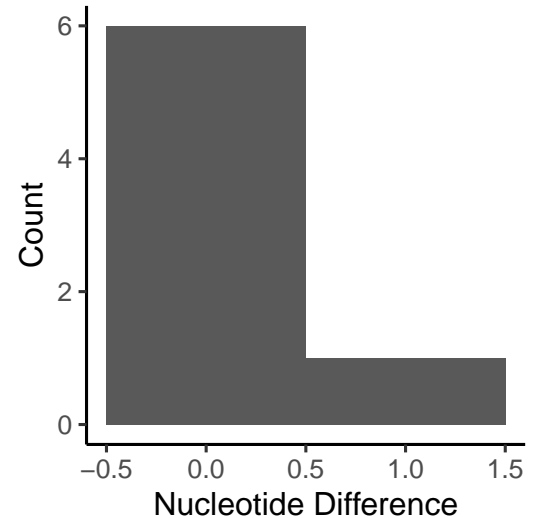
TRBV12-1*01

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



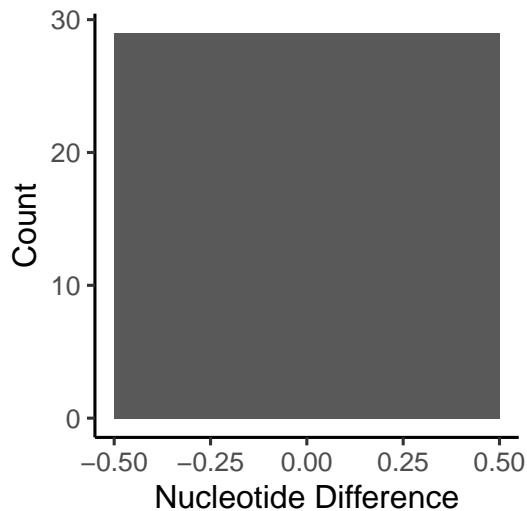
TRBV12-5*01

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



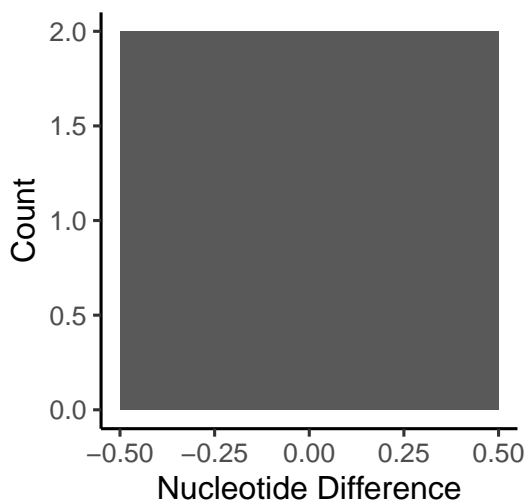
TRBV3–1*01

29 sequences assigned
29 (100%) exact matches, in which:
29 unique CDR3
9 unique J



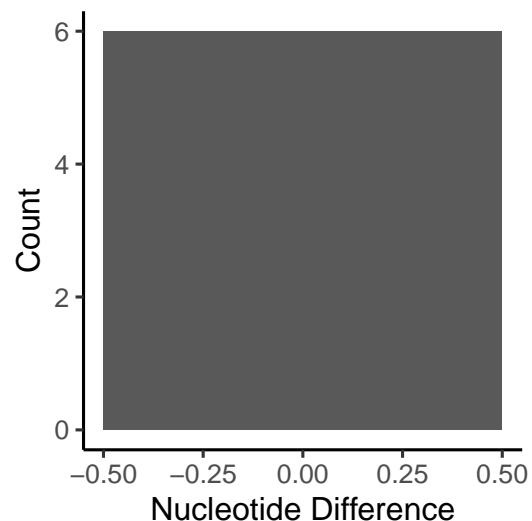
TRBV3–2*02

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



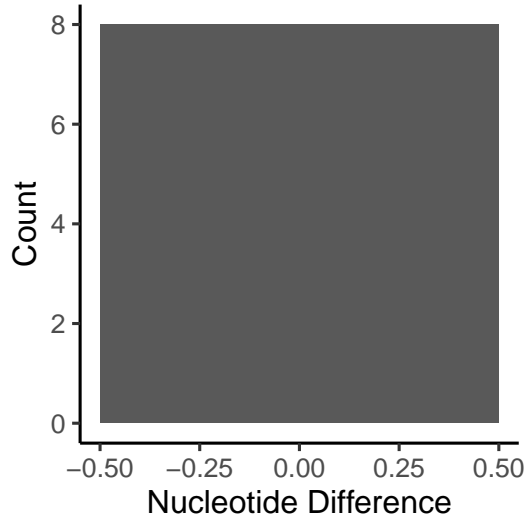
TRBV24–1*02

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



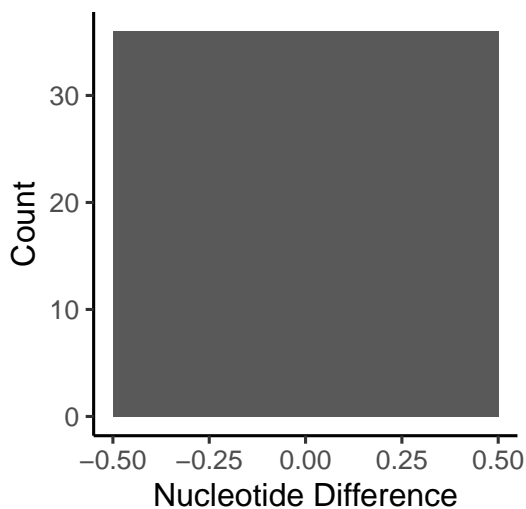
TRBV23–1*01

8 sequences assigned
8 (100%) exact matches, in which:
8 unique CDR3
7 unique J



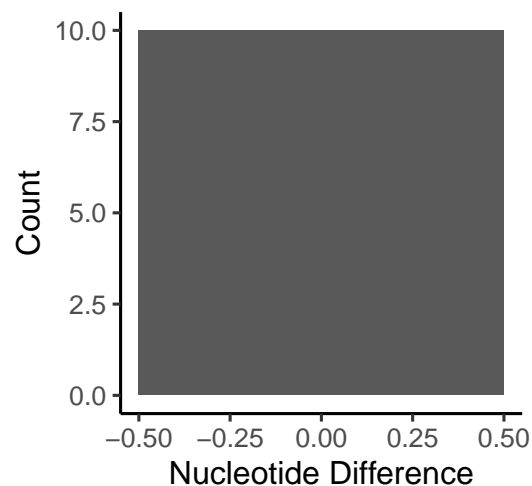
TRBV4–1*01

36 sequences assigned
36 (100%) exact matches, in which:
35 unique CDR3
10 unique J



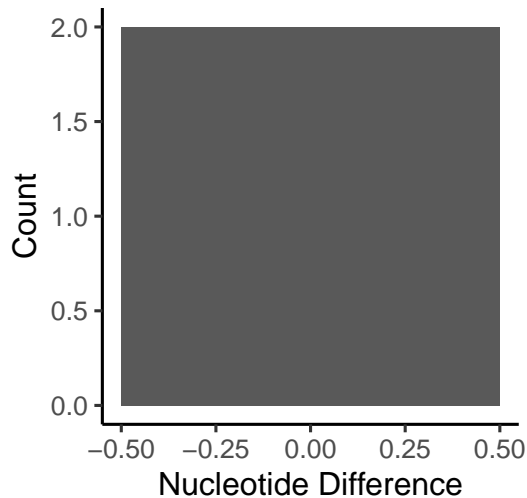
TRBV4–2*01

10 sequences assigned
10 (100%) exact matches, in which:
10 unique CDR3
6 unique J



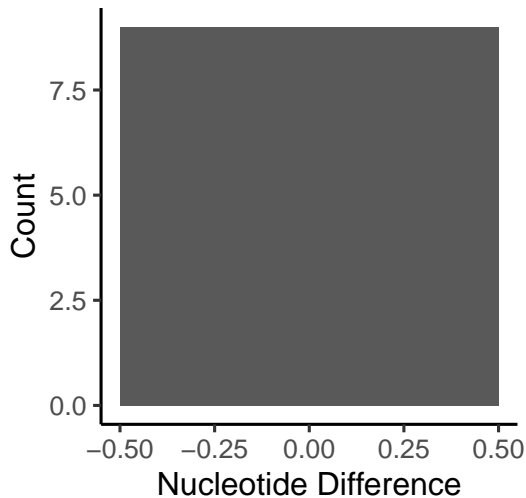
TRBV3–2*01

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



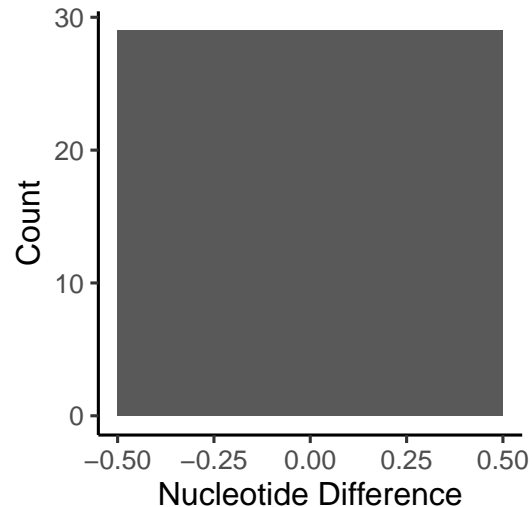
TRBV24–1*01

9 sequences assigned
9 (100%) exact matches, in which:
9 unique CDR3
6 unique J



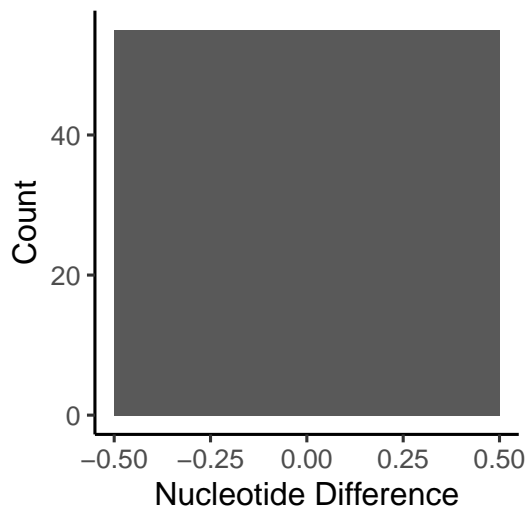
TRBV4–3*01

29 sequences assigned
29 (100%) exact matches, in which:
29 unique CDR3
10 unique J



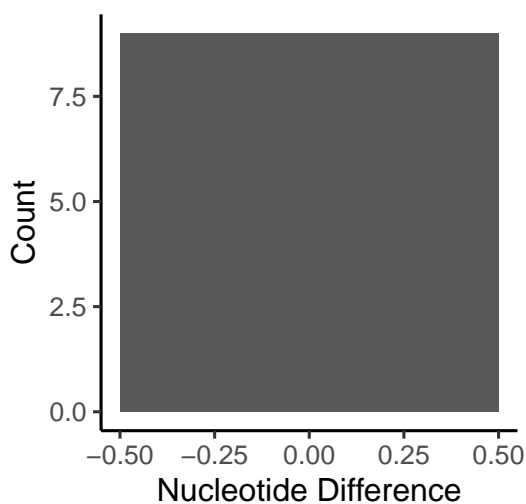
TRBV5-1*01

55 sequences assigned
55 (100%) exact matches, in which:
55 unique CDR3
12 unique J



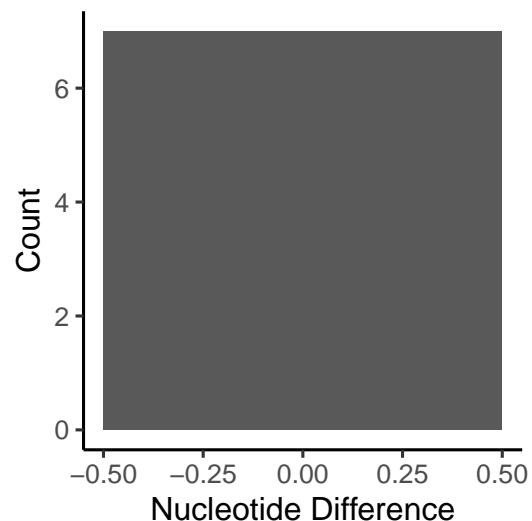
TRBV5-5*01

9 sequences assigned
9 (100%) exact matches, in which:
9 unique CDR3
7 unique J



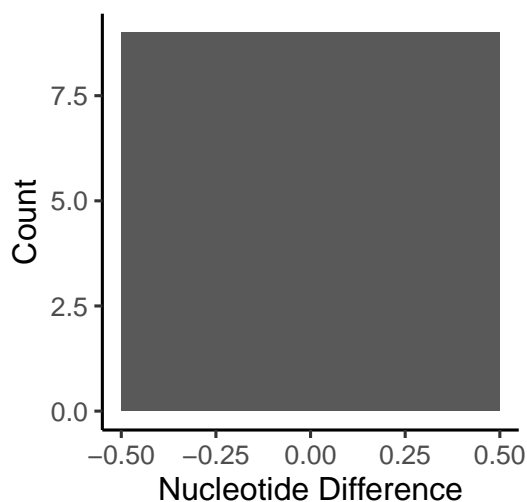
TRBV5-8*01

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



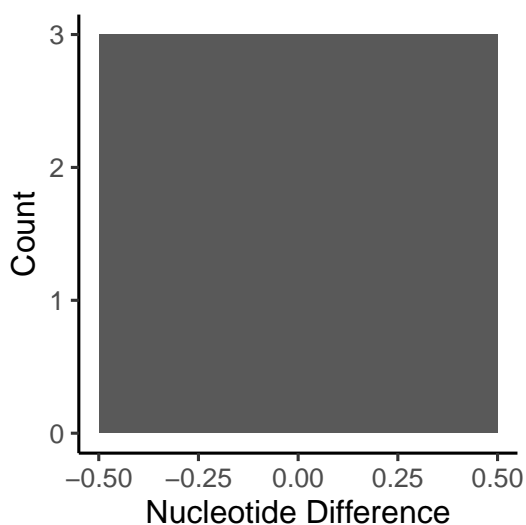
TRBV25-1*01

9 sequences assigned
9 (100%) exact matches, in which:
9 unique CDR3
7 unique J



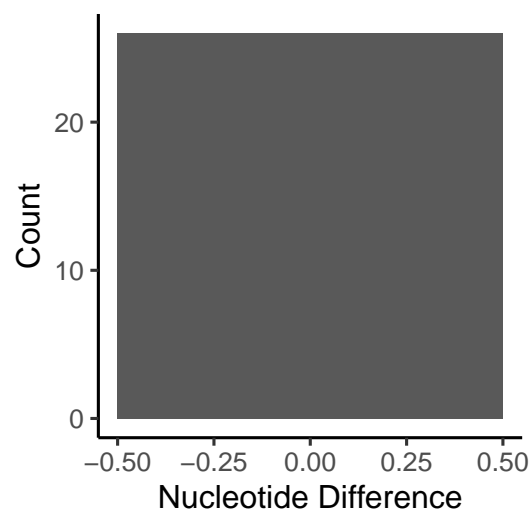
TRBV5-5*02_322TTGG32!

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



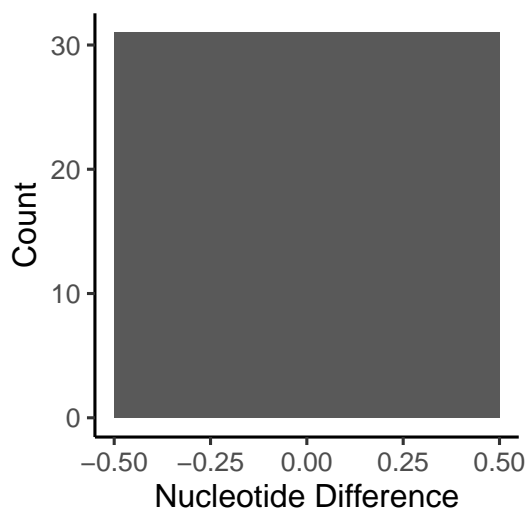
TRBV6-1*01

26 sequences assigned
26 (100%) exact matches, in which:
26 unique CDR3
10 unique J



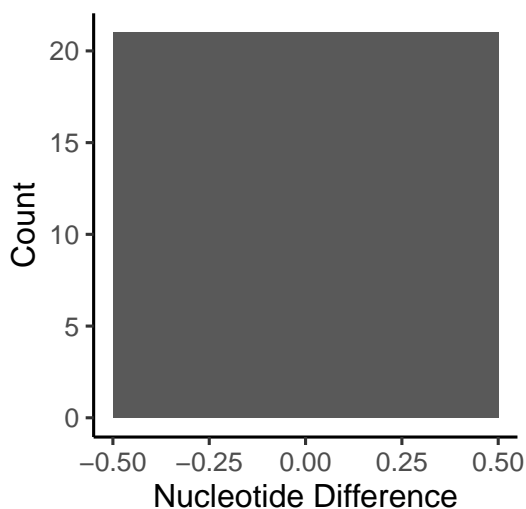
TRBV5-4*01

31 sequences assigned
31 (100%) exact matches, in which:
31 unique CDR3
12 unique J



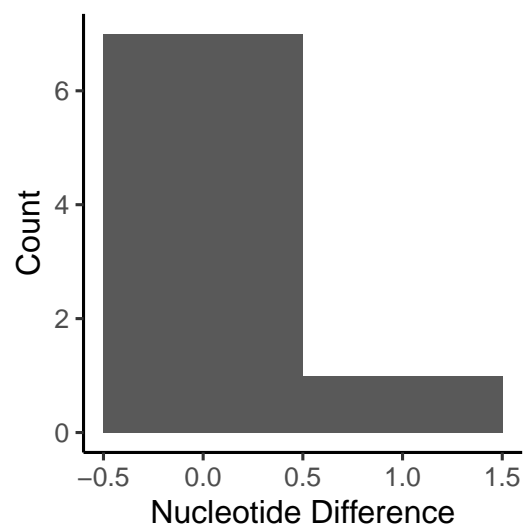
TRBV5-6*01

21 sequences assigned
21 (100%) exact matches, in which:
21 unique CDR3
9 unique J



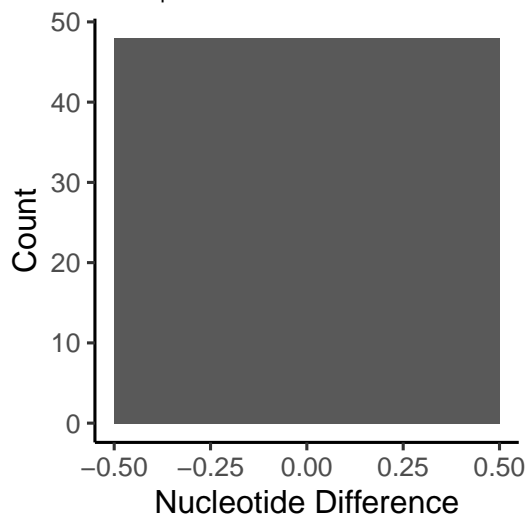
TRBV6-4*01

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



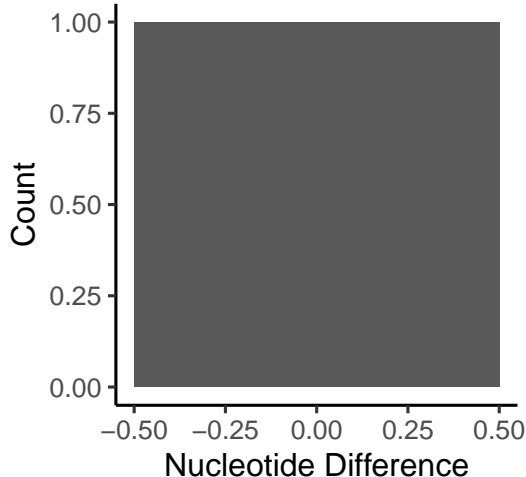
TRBV6-5*01

48 sequences assigned
48 (100%) exact matches, in which:
48 unique CDR3
13 unique J



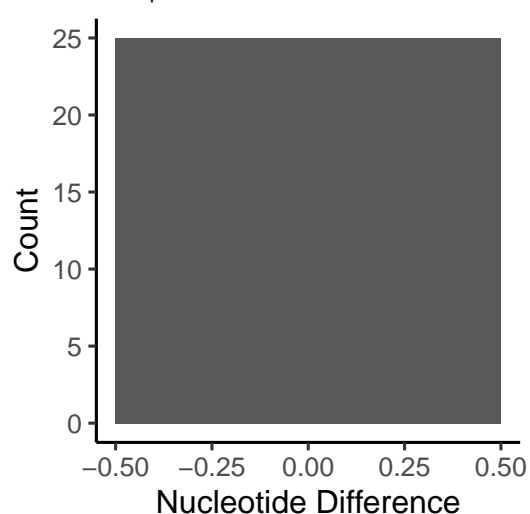
TRBV6-8*01

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



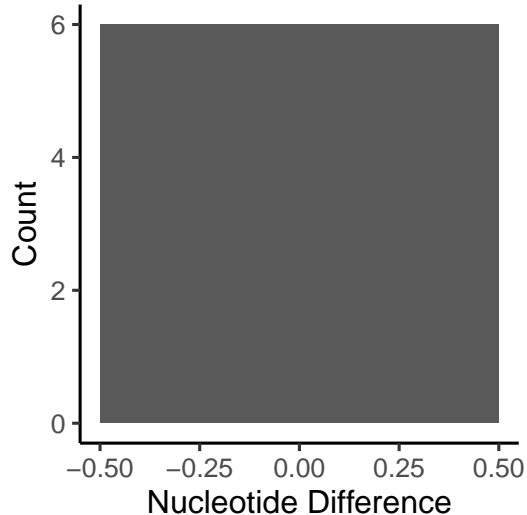
TRBV7-3*01

25 sequences assigned
25 (100%) exact matches, in which:
25 unique CDR3
8 unique J



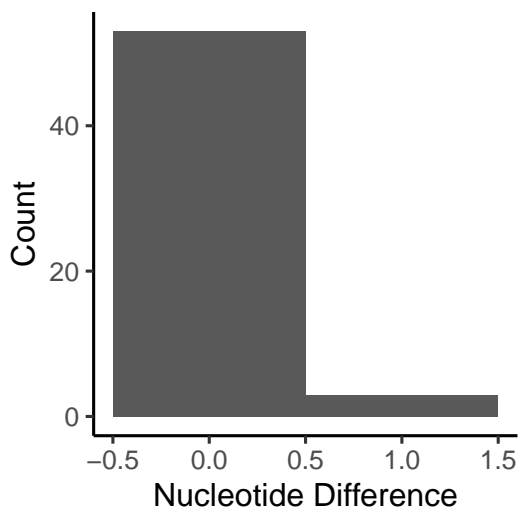
TRBV6-6*01

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



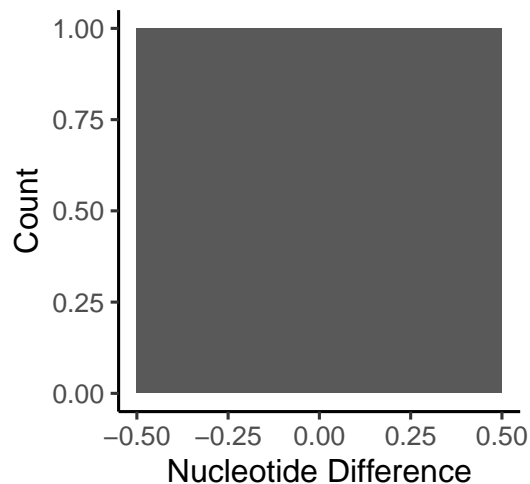
TRBV6-23*01

56 sequences assigned
53 (94.6%) exact matches, in which:
53 unique CDR3
11 unique J



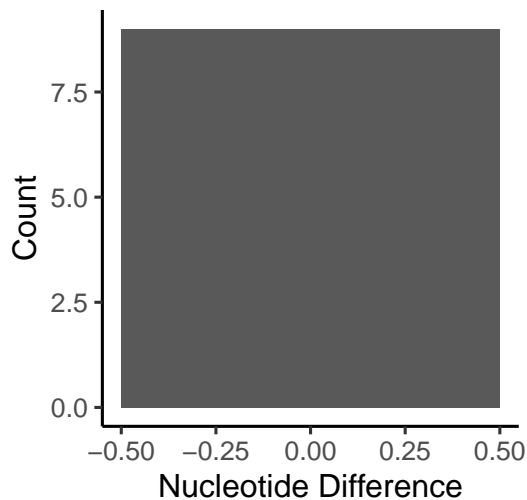
TRBV7-4*01

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



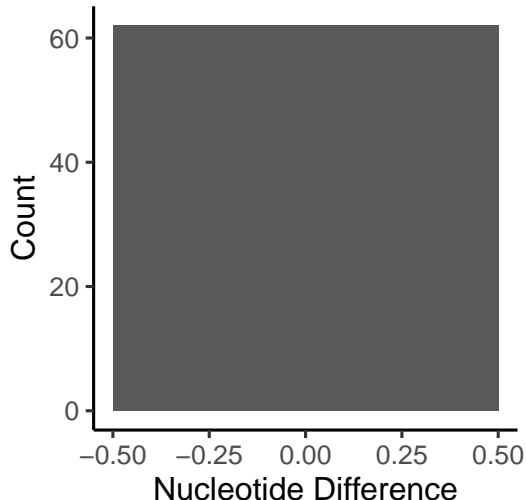
TRBV6-6*02

9 sequences assigned
9 (100%) exact matches, in which:
9 unique CDR3
6 unique J



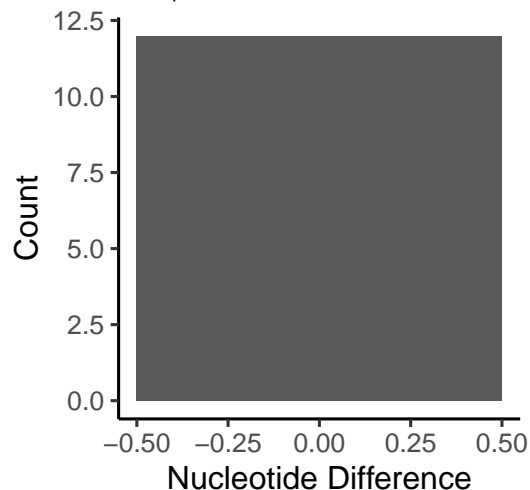
TRBV7-2*01

62 sequences assigned
62 (100%) exact matches, in which:
62 unique CDR3
12 unique J



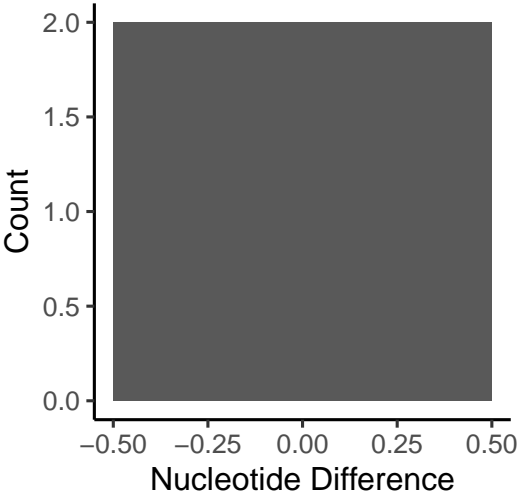
TRBV7-6*01

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



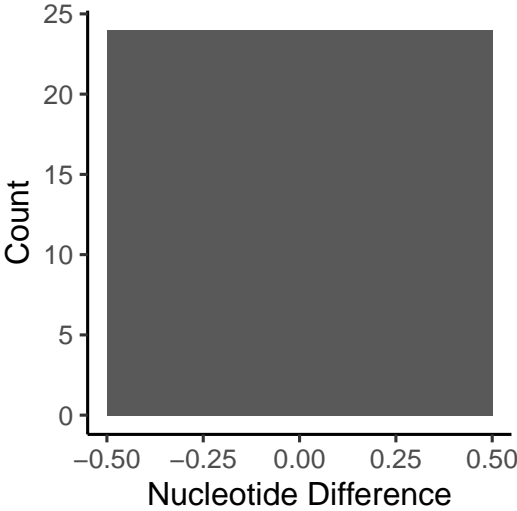
TRBV7-7*01

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



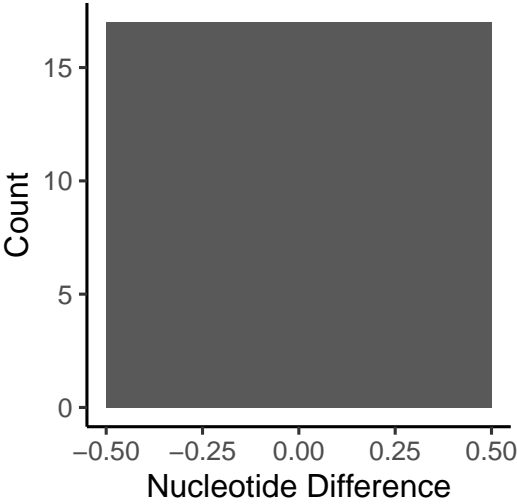
TRBV7-9*03

24 sequences assigned
24 (100%) exact matches, in which:
24 unique CDR3
8 unique J



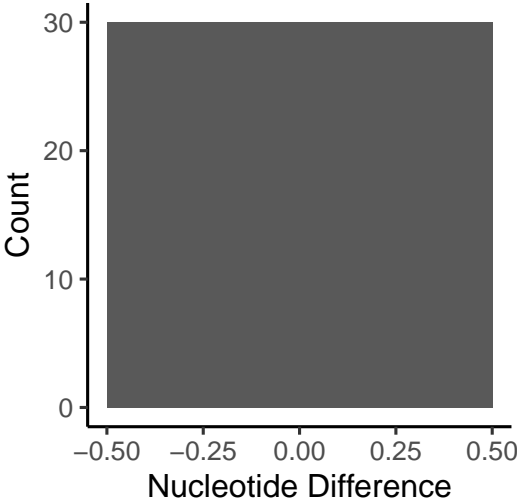
TRBV7-8*01

17 sequences assigned
17 (100%) exact matches, in which:
17 unique CDR3
9 unique J



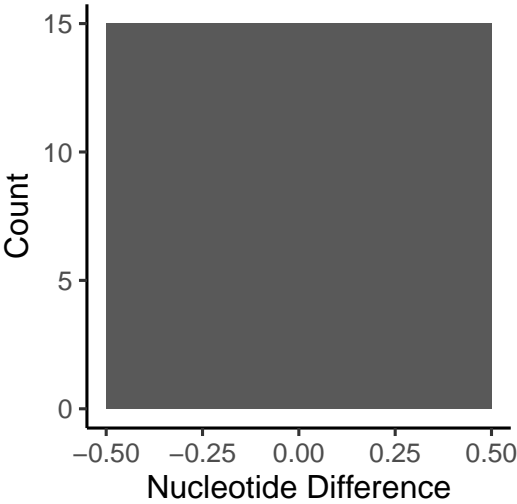
TRBV29-1*01

30 sequences assigned
30 (100%) exact matches, in which:
30 unique CDR3
10 unique J

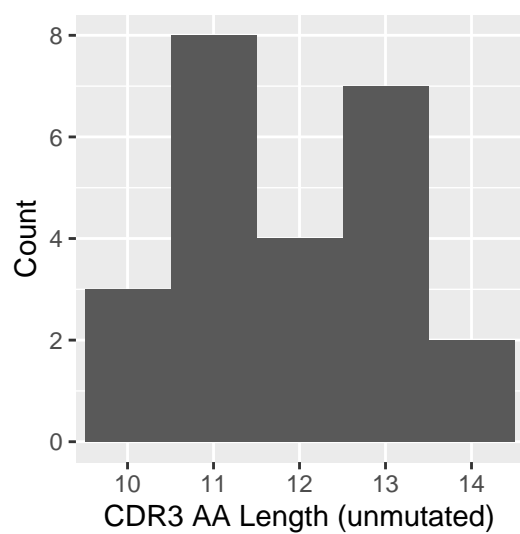


TRBV7-9*01

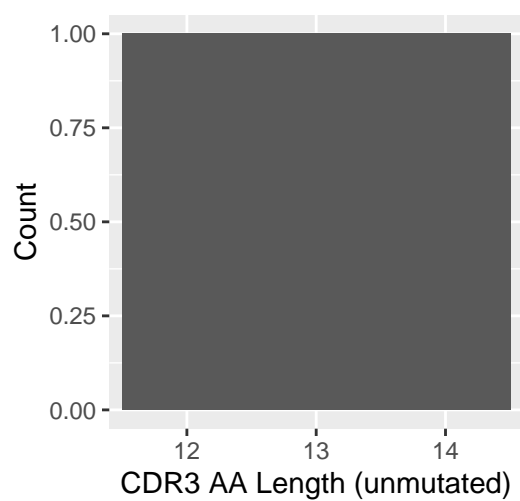
15 sequences assigned
15 (100%) exact matches, in which:
15 unique CDR3
7 unique J

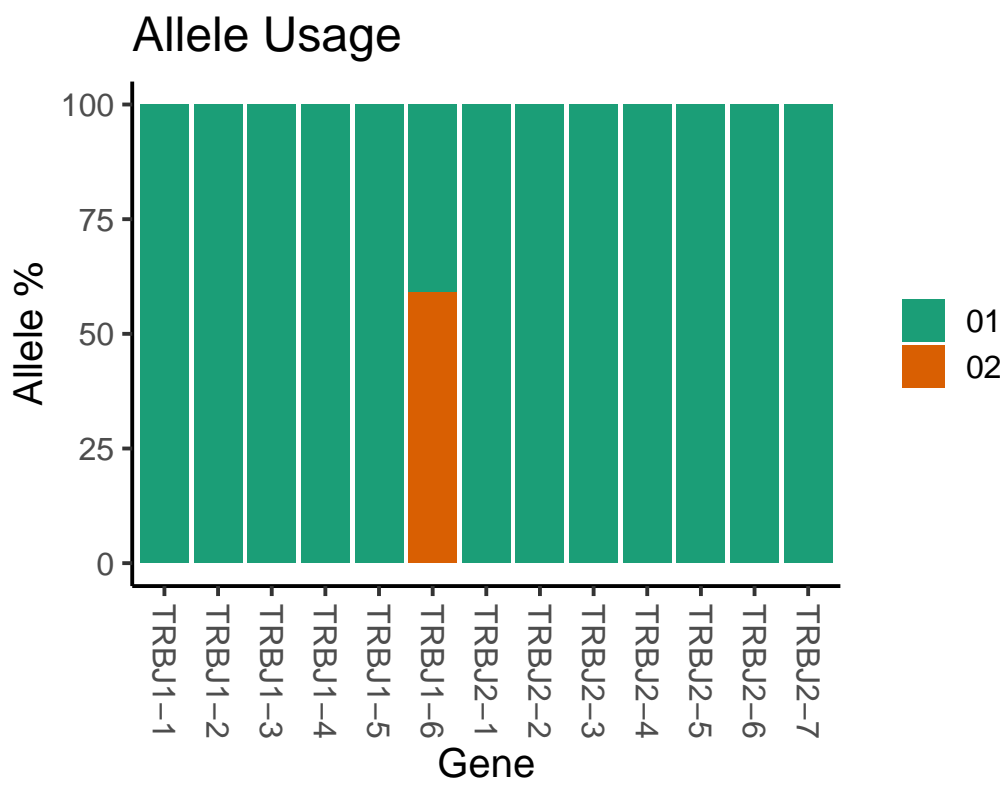


TRBV20-1*02_319AGAGA323

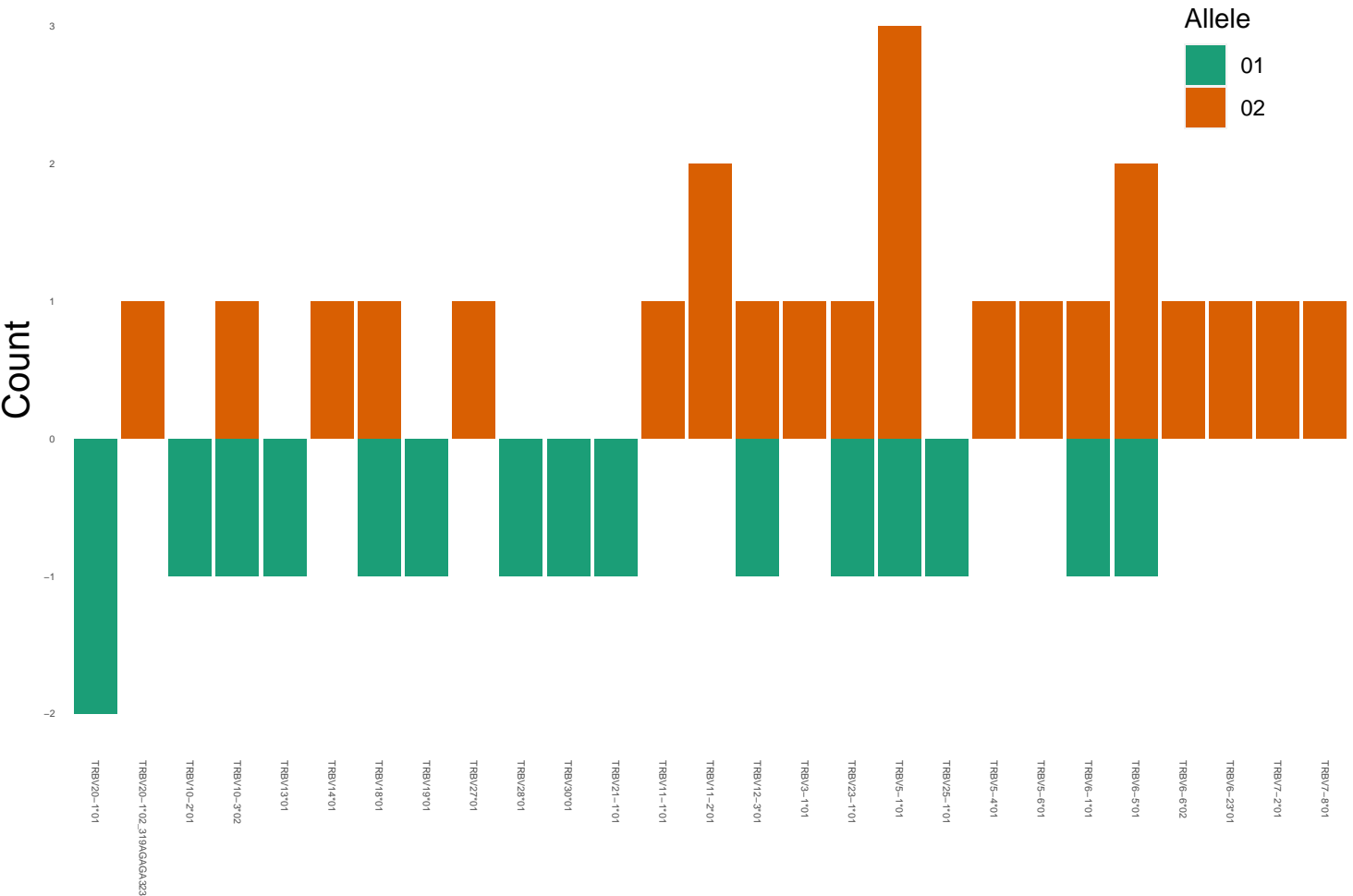


TRBV5-5*02_322TTGG325





Sequence Count by TRBJ1–6 allele usage



Warning – no inferred sequences found.

Warning – no inferred sequences found.