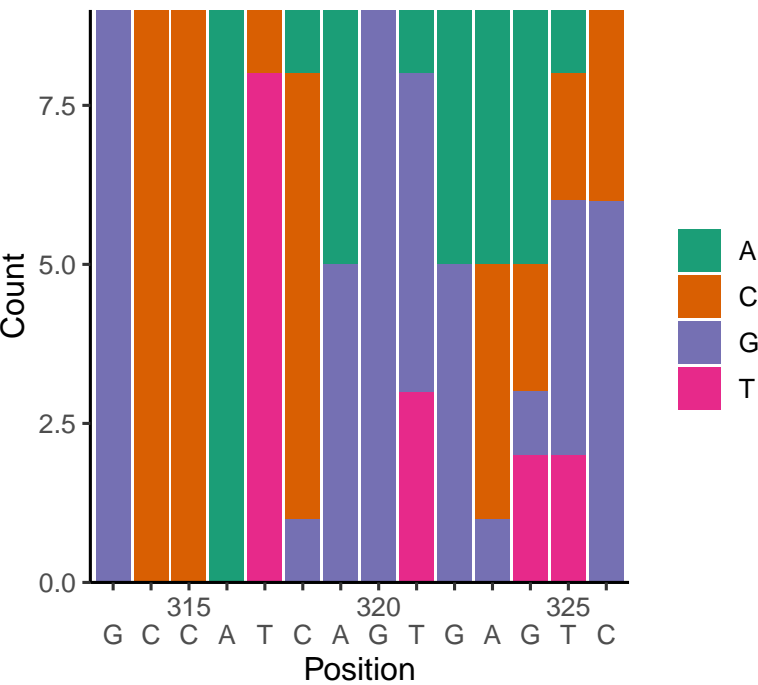
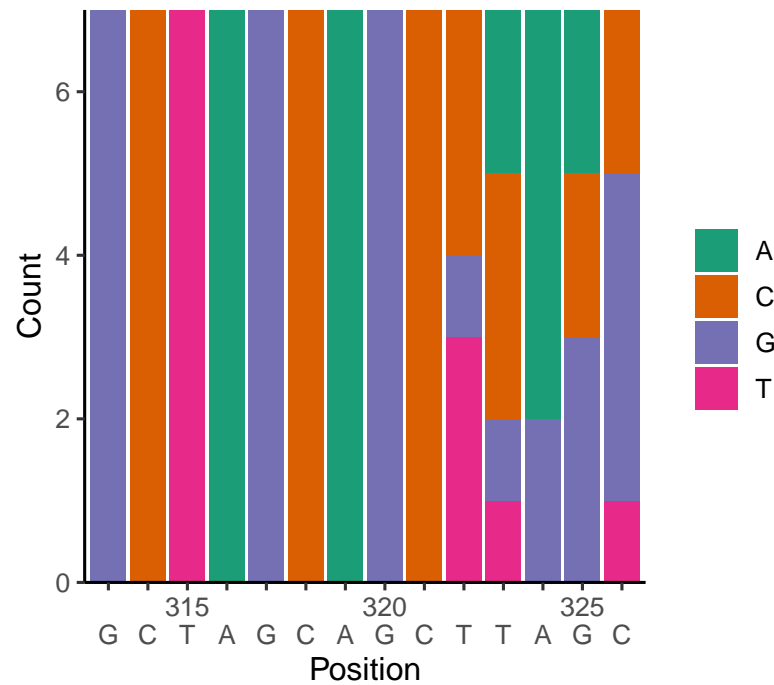


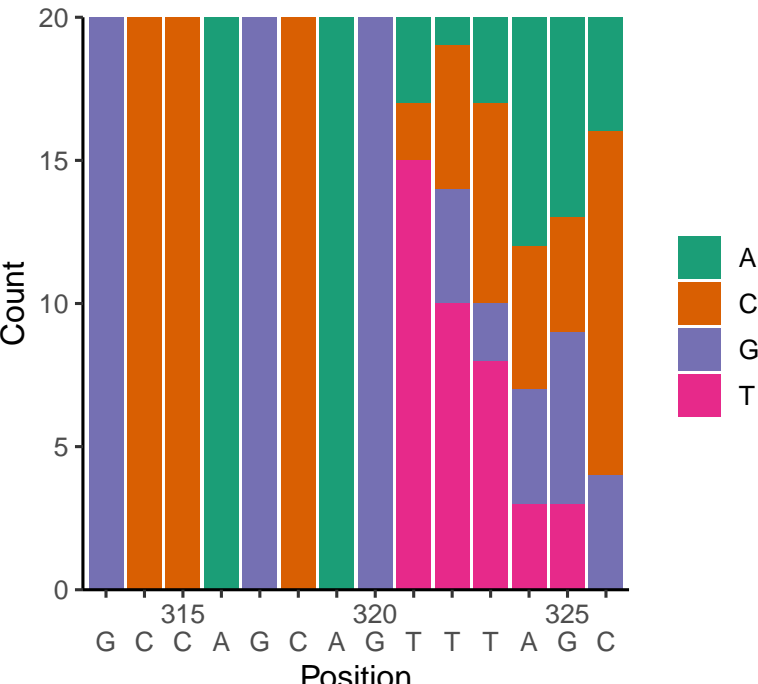
Gene TRBV10-3*03_313GCCATCAGTGA



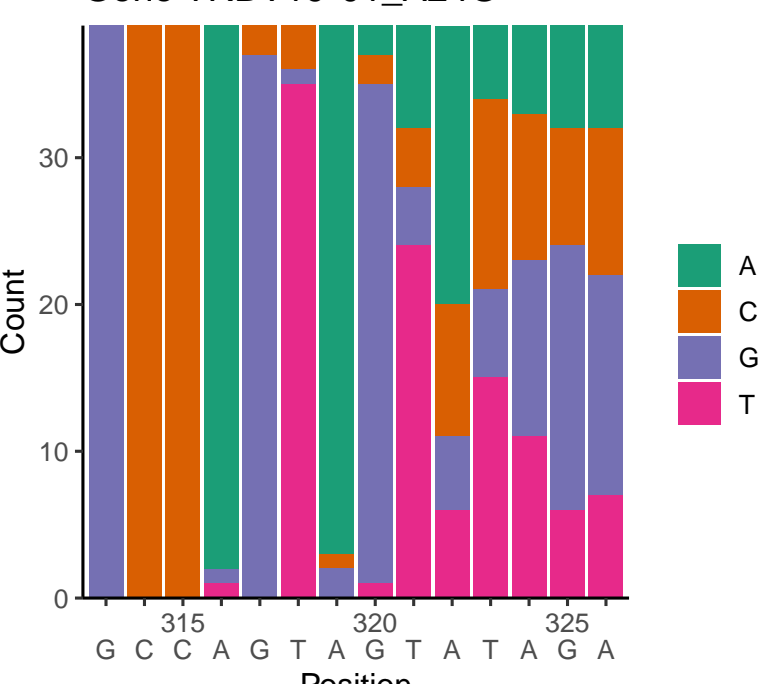
Gene TRBV7-7*01_C315T



Gene TRBV12-4*01_C87T



Gene TRBV19*01_A24G



Gene TRBV10-3*03_313GCCATCAGTGAGTC326



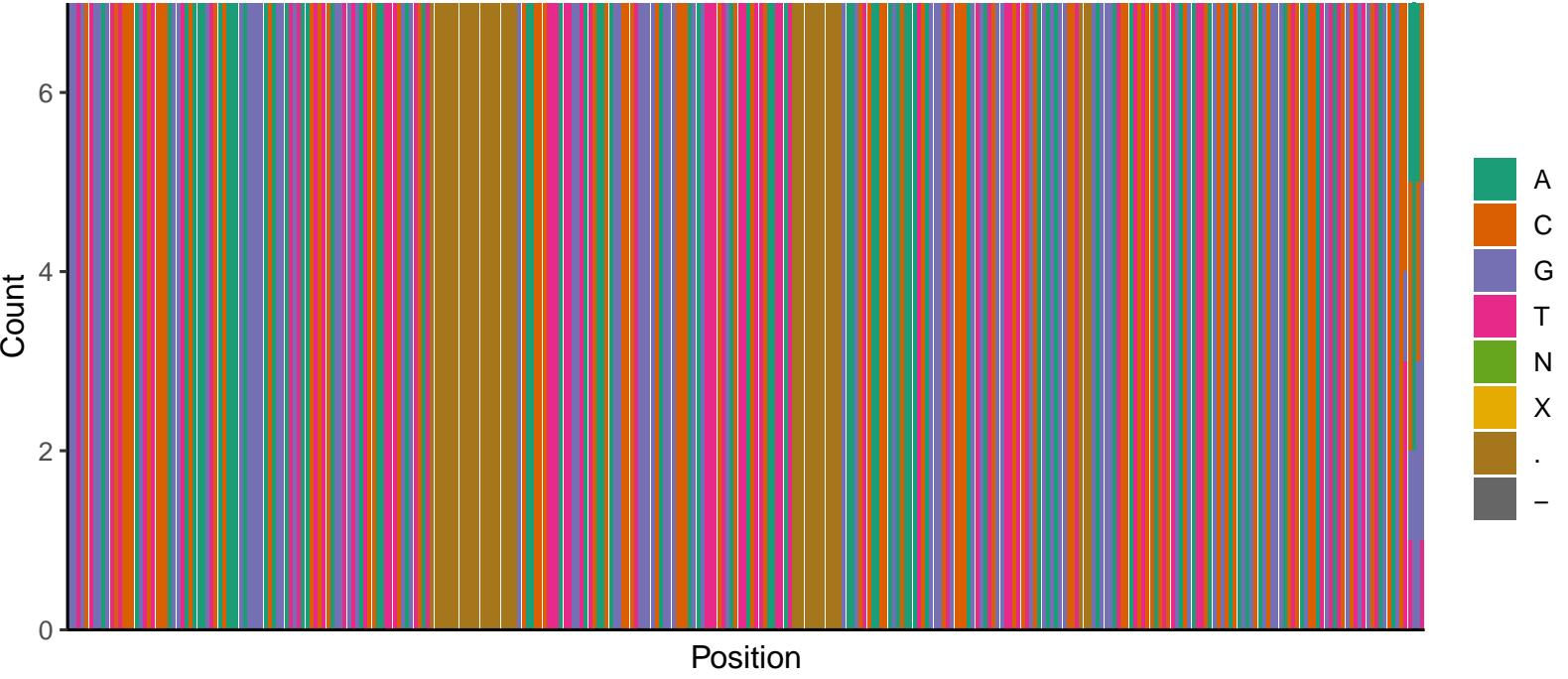
Gene TRBV12-4*01_C87T



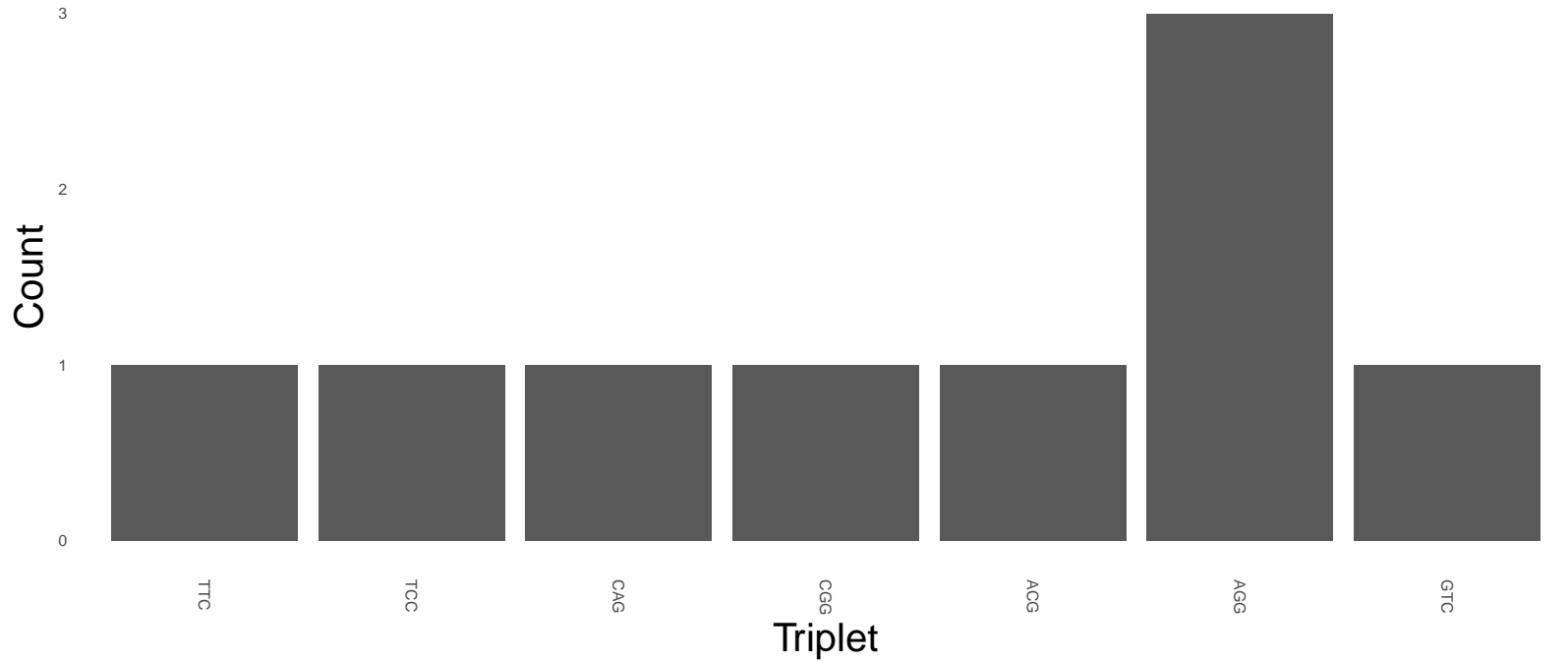
Gene TRBV19*01_A24G



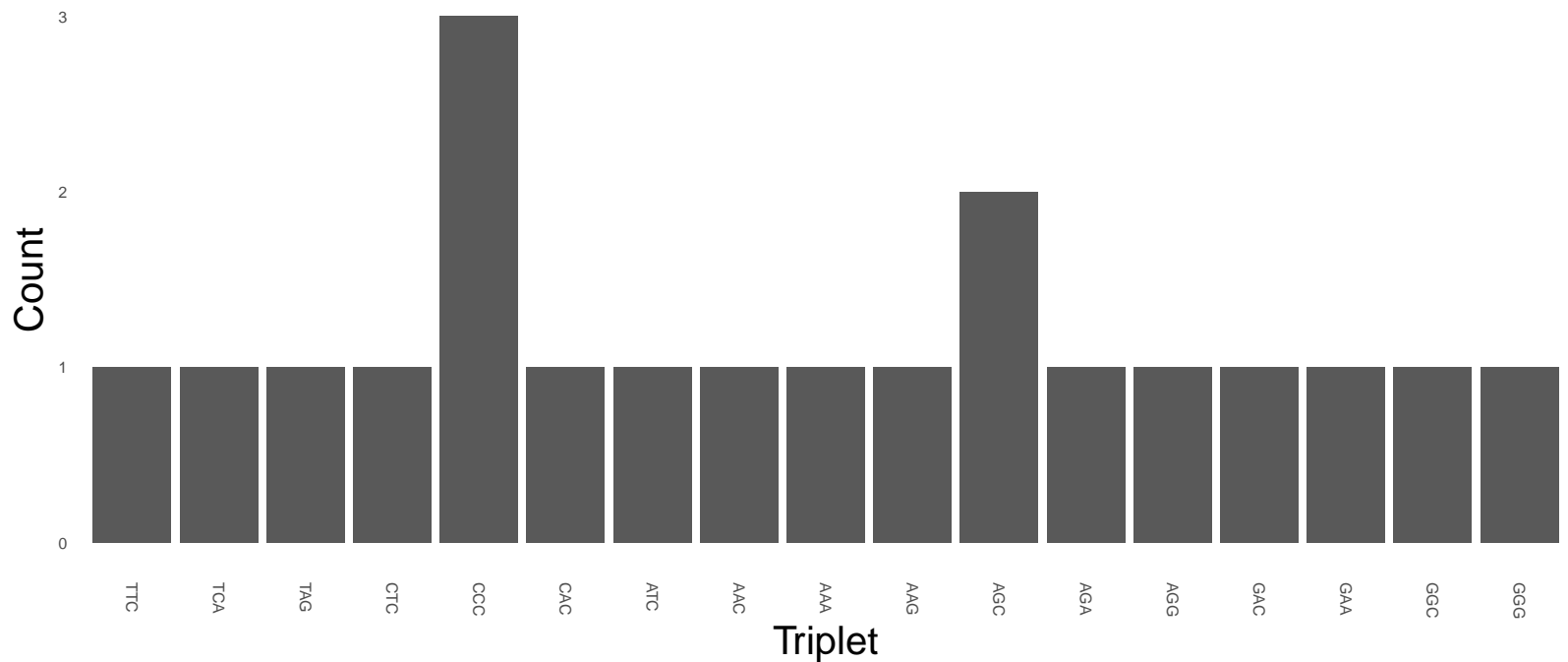
Gene TRBV7-7*01_C315T



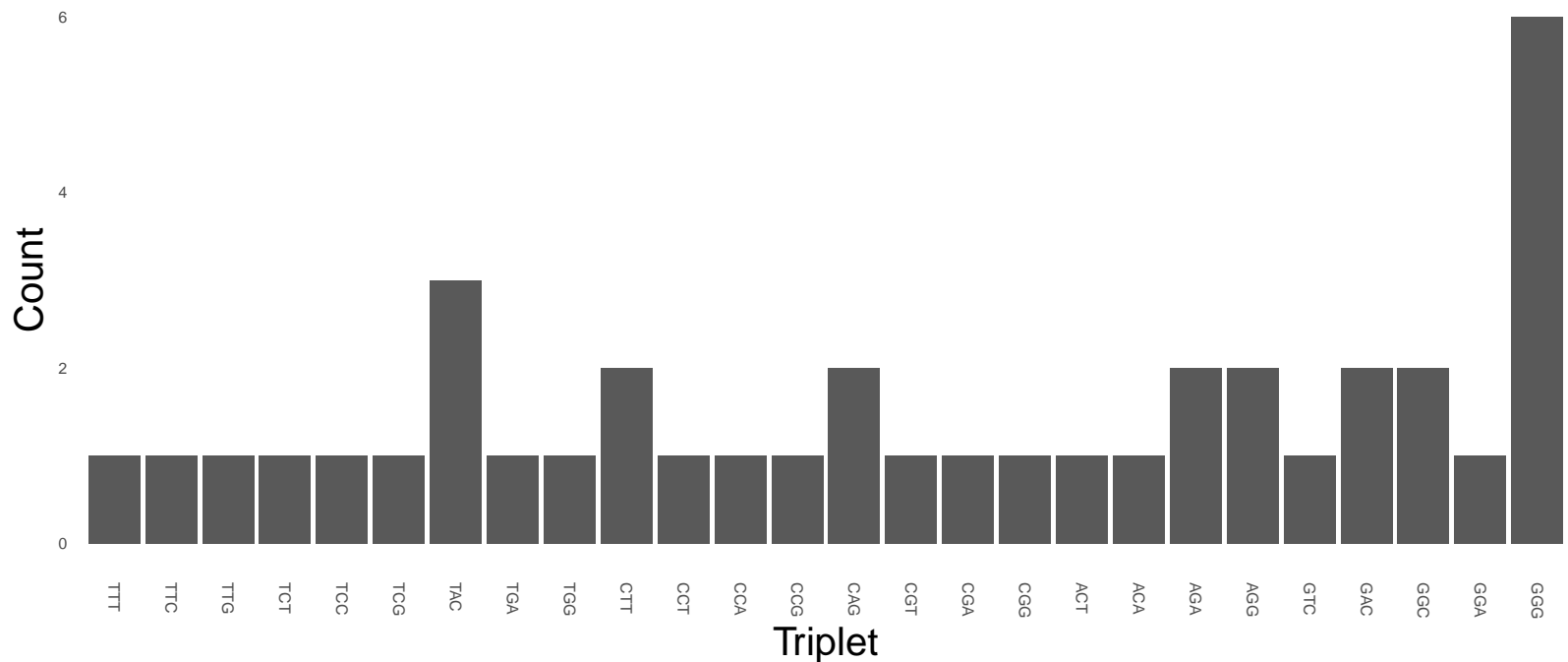
TRBV10–3*03_313GCCATCAGTGAGTC326– Final 3 nucleotides as a triplet



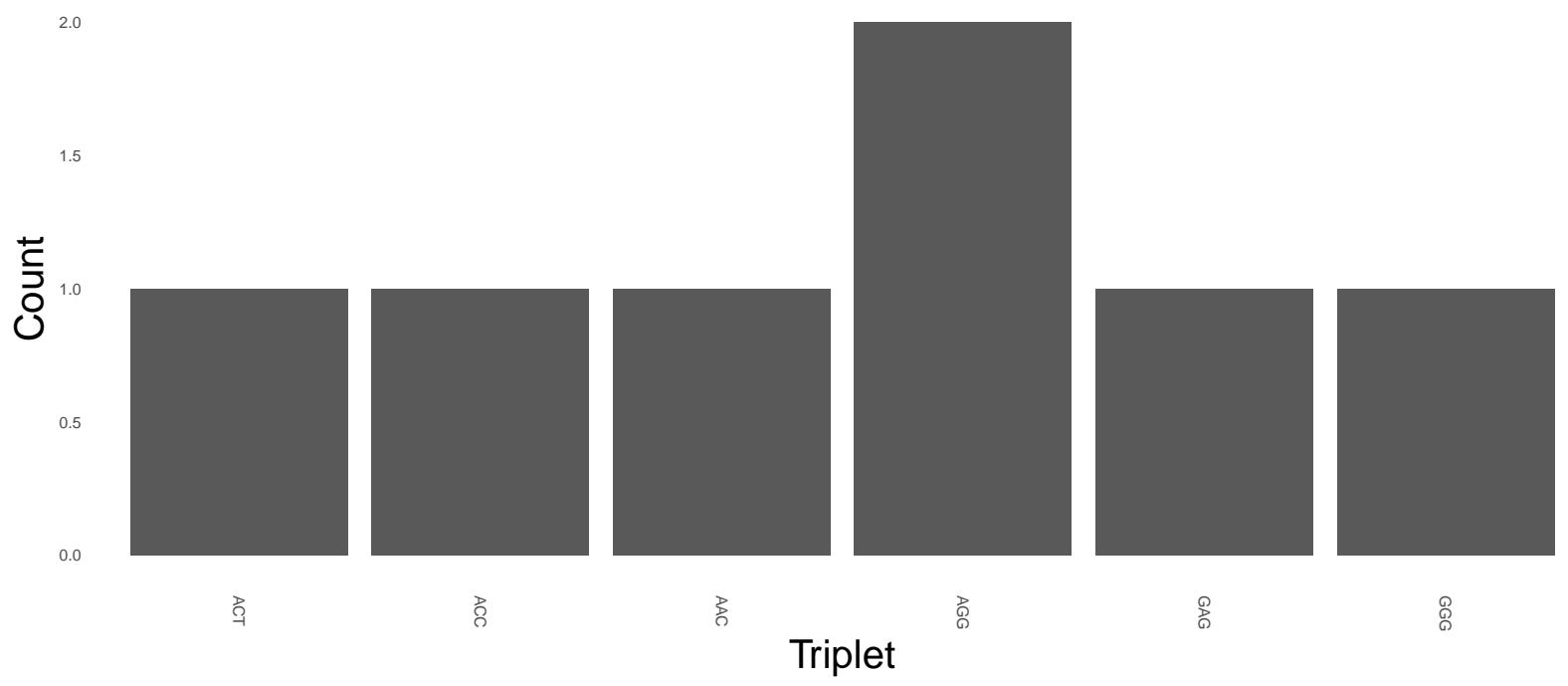
TRBV12–4*01_C87T– Final 3 nucleotides as a triplet



TRBV19*01_A24G– Final 3 nucleotides as a triplet

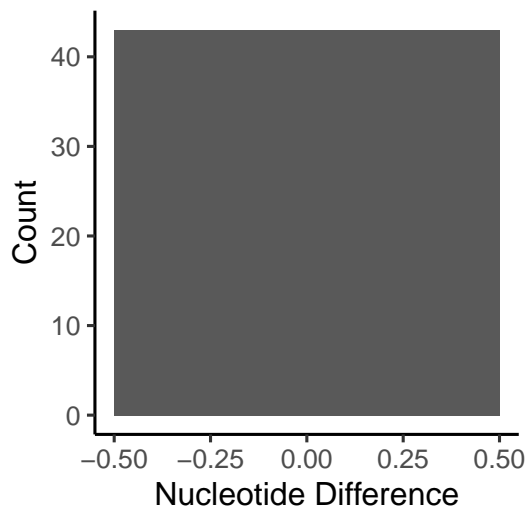


TRBV7-7*01_C315T- Final 3 nucleotides as a triplet



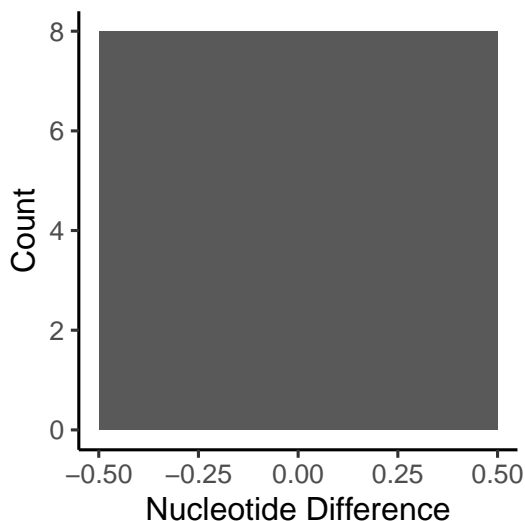
TRBV20–1*01

43 sequences assigned
43 (100%) exact matches, in which:
43 unique CDR3
10 unique J



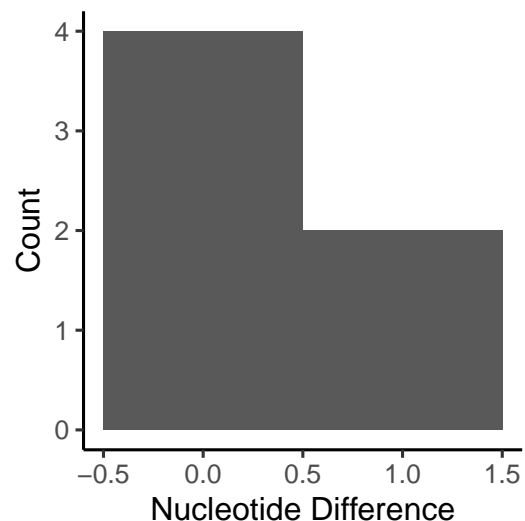
TRBV10–2*01

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



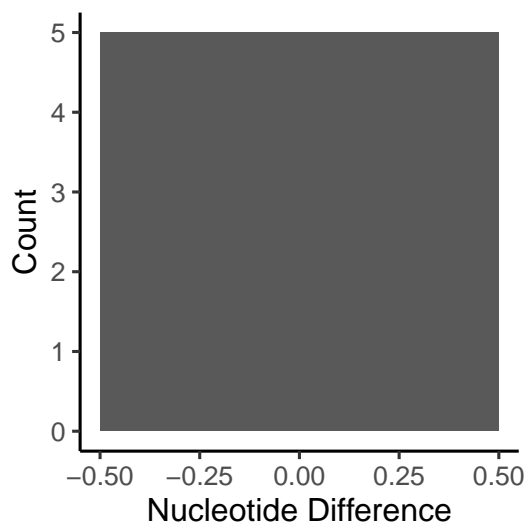
TRBV13*01

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



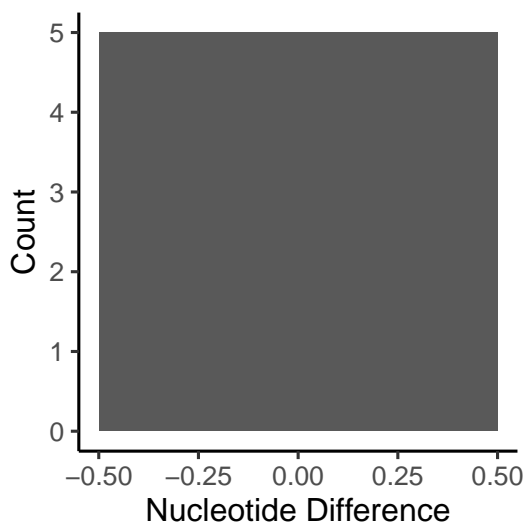
TRBV10–1*01

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



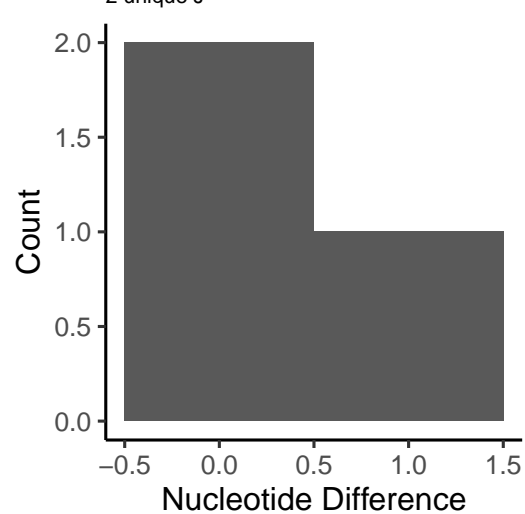
TRBV10–3*02

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



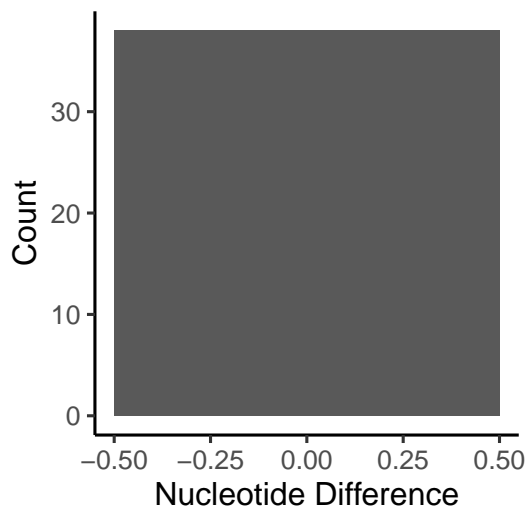
TRBV14*01

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



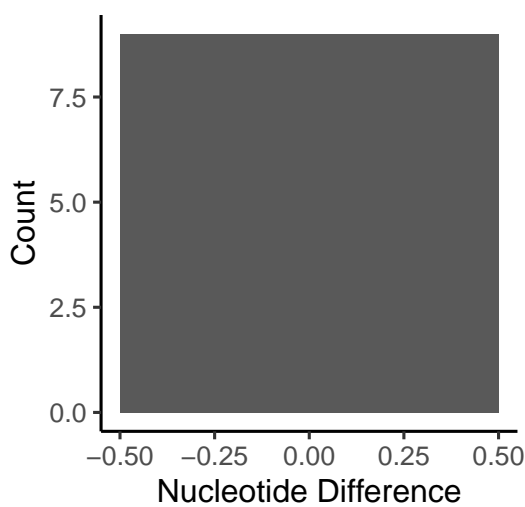
TRBV20–1*05

38 sequences assigned
38 (100%) exact matches, in which:
38 unique CDR3
13 unique J



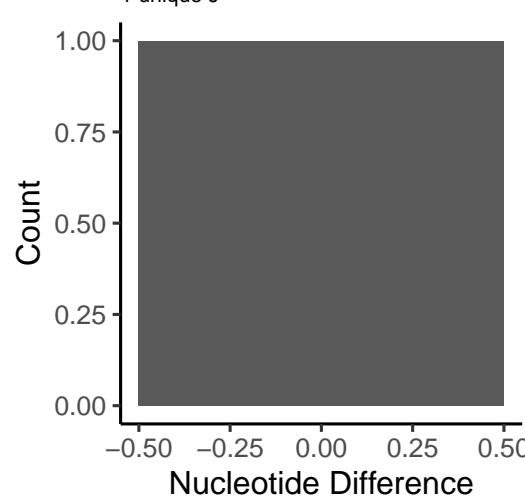
TRBV10–3*03_313GCCA

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



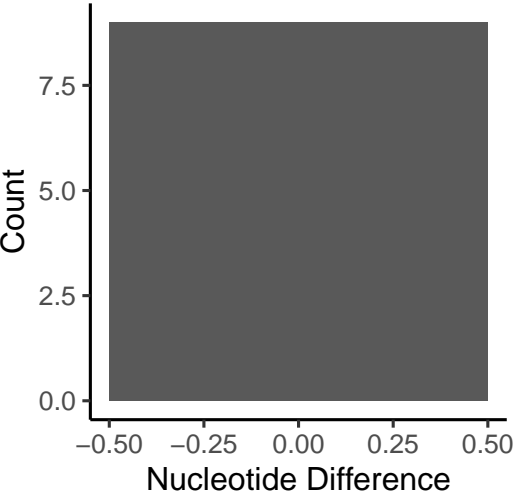
TRBV14*02

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



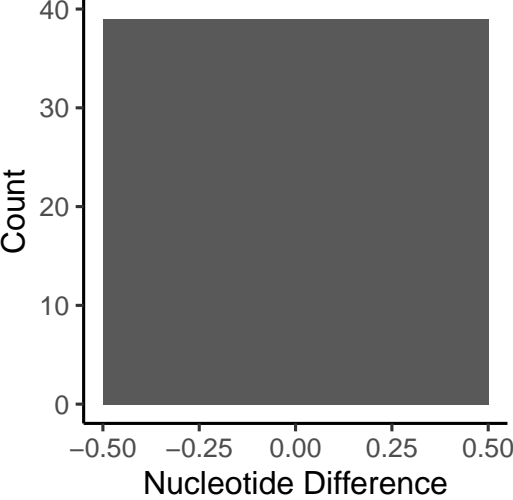
TRBV15*02

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



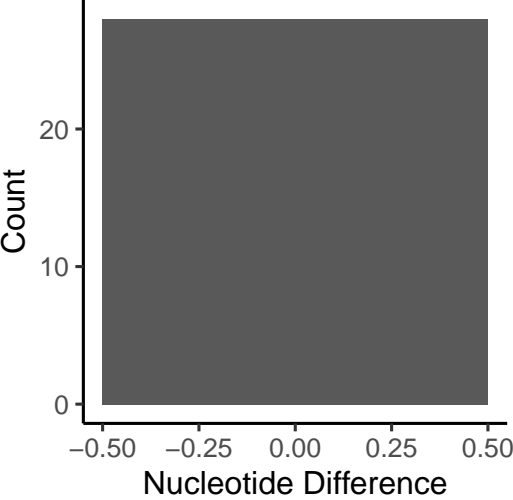
TRBV19*01_A24G

39 sequences assigned
39 (100%) exact matches, in which:
39 unique CDR3
12 unique J



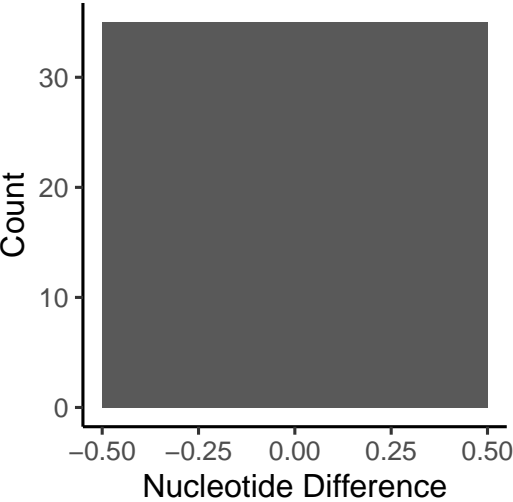
TRBV28*01

28 sequences assigned
28 (100%) exact matches, in which:
28 unique CDR3
11 unique J



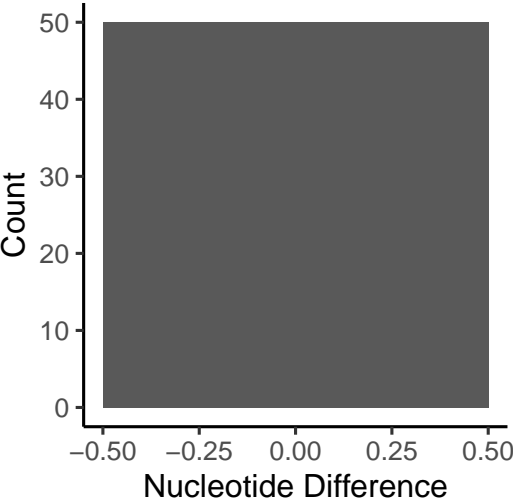
TRBV18*01

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



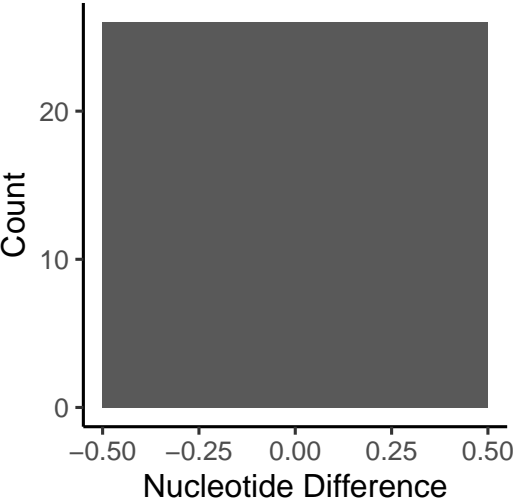
TRBV2*01

50 sequences assigned
50 (100%) exact matches, in which:
50 unique CDR3
11 unique J



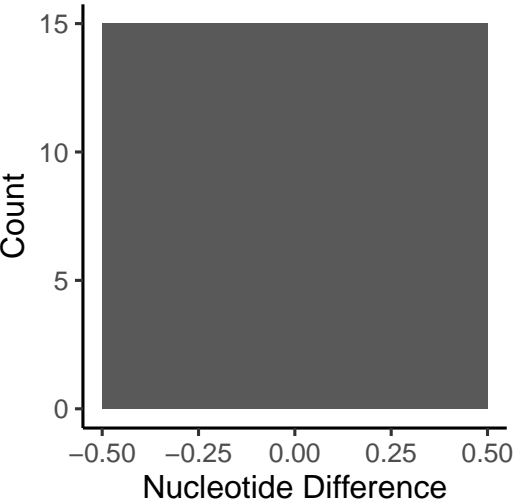
TRBV30*01

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



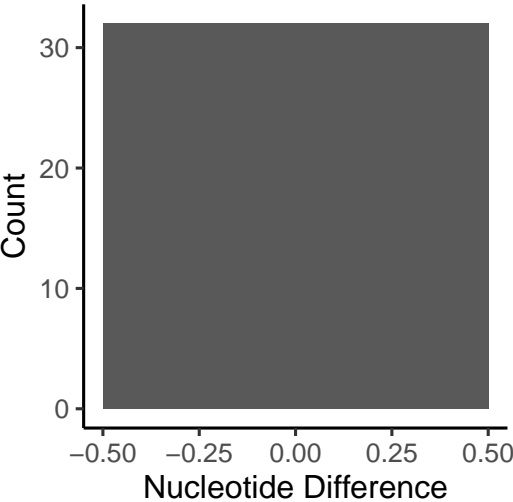
TRBV19*01

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



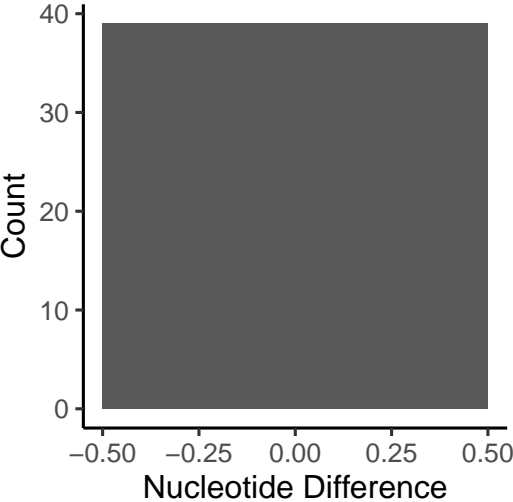
TRBV27*01

32 sequences assigned
32 (100%) exact matches, in which:
32 unique CDR3
9 unique J



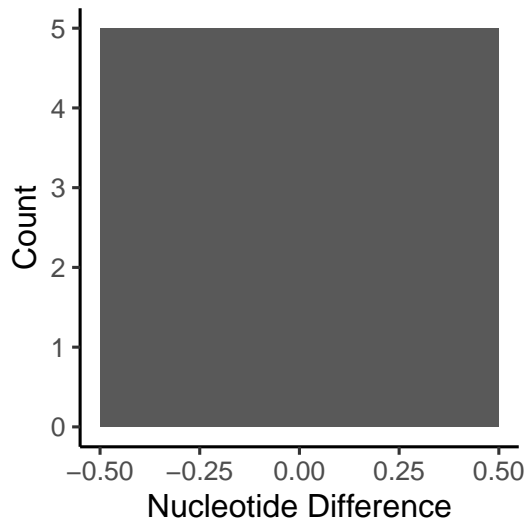
TRBV9*01

39 sequences assigned
39 (100%) exact matches, in which:
39 unique CDR3
11 unique J



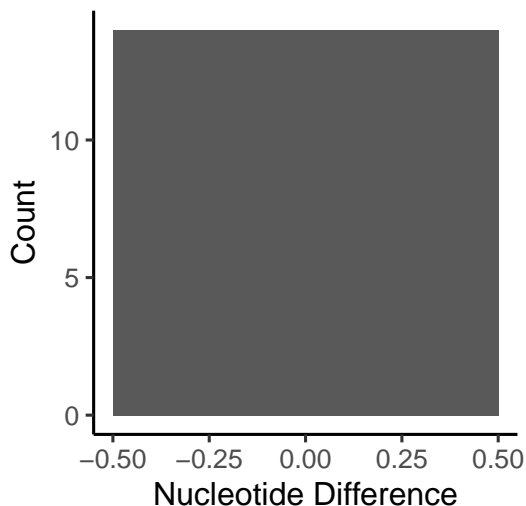
TRBV11–1*01

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



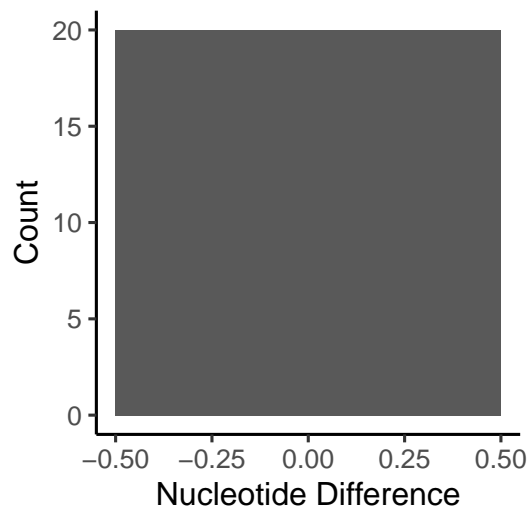
TRBV11–3*04

14 sequences assigned
14 (100%) exact matches, in which:
14 unique CDR3
5 unique J



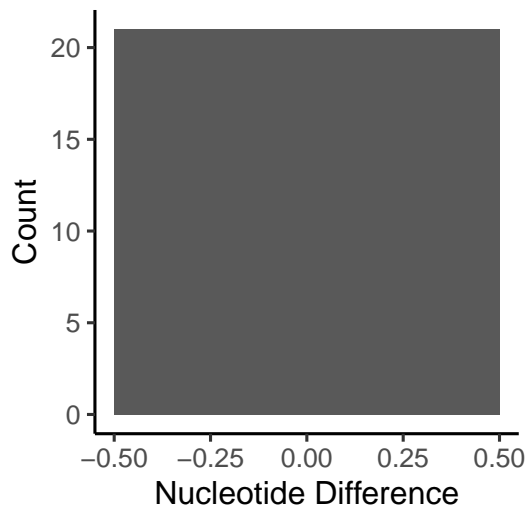
TRBV12–4*01_C87T

20 sequences assigned
20 (100%) exact matches, in which:
20 unique CDR3
9 unique J



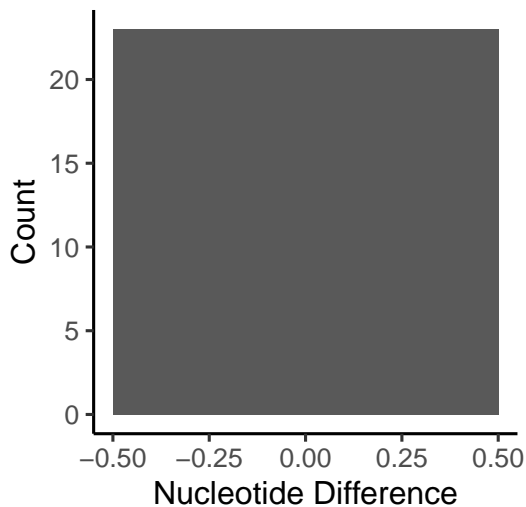
TRBV11–2*01

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



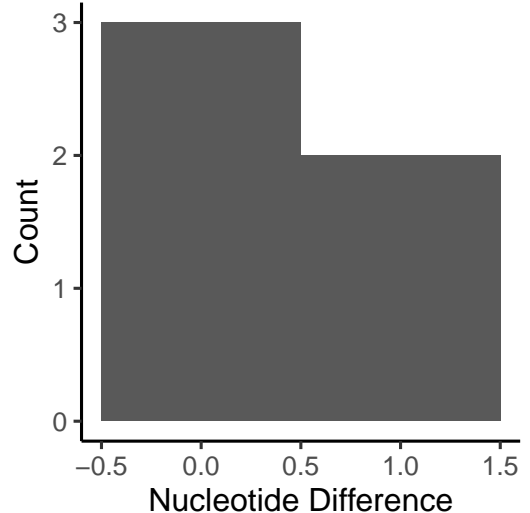
TRBV12–3*01

23 sequences assigned
23 (100%) exact matches, in which:
23 unique CDR3
10 unique J



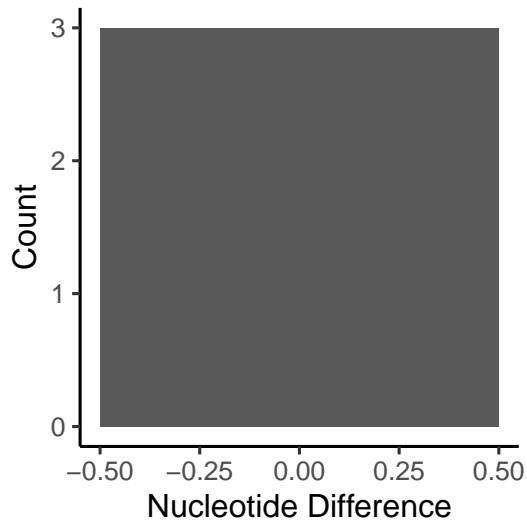
TRBV12–5*01

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



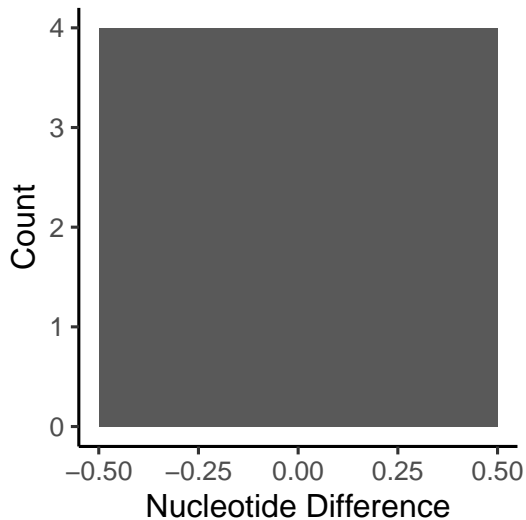
TRBV11–3*01

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



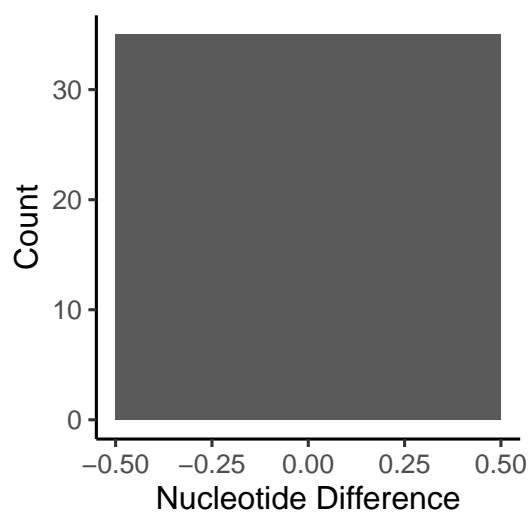
TRBV12–4*01

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



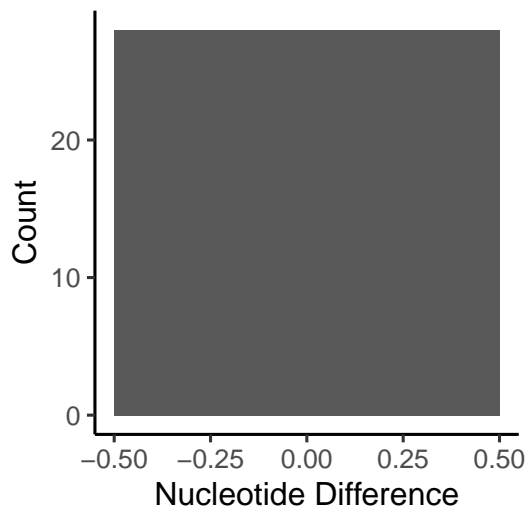
TRBV3–1*01

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



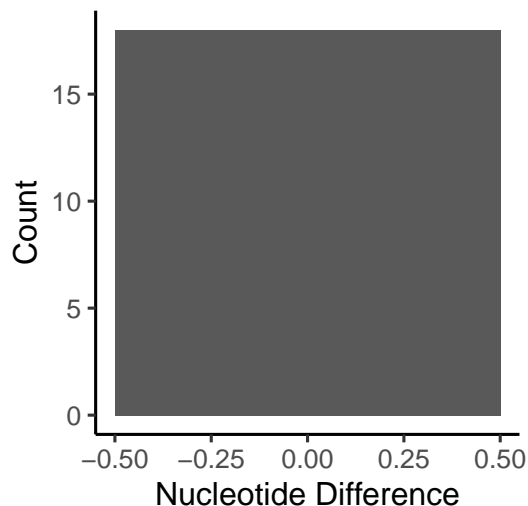
TRBV4-1*01

28 sequences assigned
28 (100%) exact matches, in which:
28 unique CDR3
10 unique J



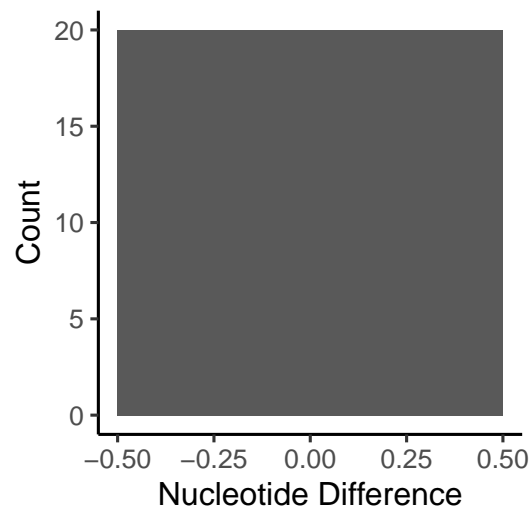
TRBV4-3*01

18 sequences assigned
18 (100%) exact matches, in which:
18 unique CDR3
9 unique J



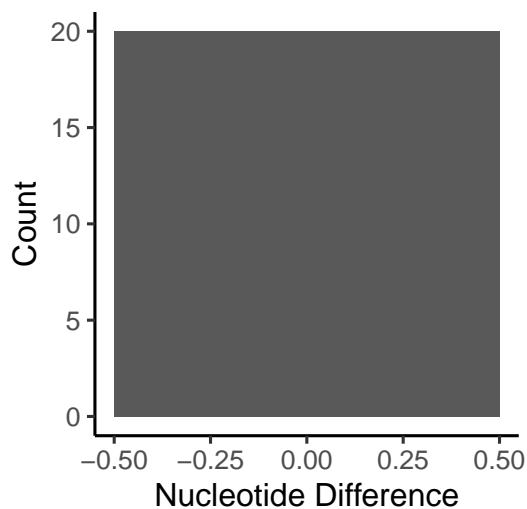
TRBV5-4*01

20 sequences assigned
20 (100%) exact matches, in which:
20 unique CDR3
7 unique J



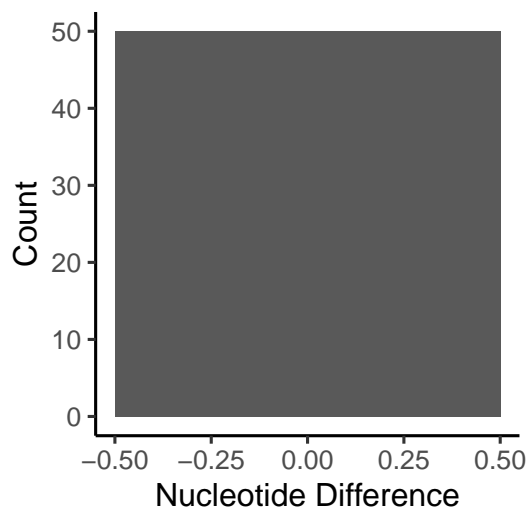
TRBV24-1*01

20 sequences assigned
20 (100%) exact matches, in which:
20 unique CDR3
8 unique J



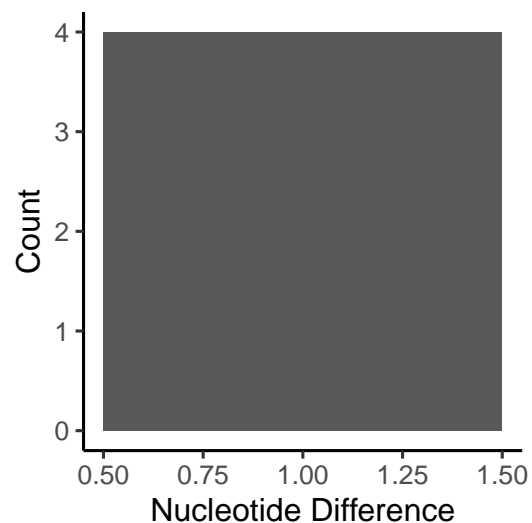
TRBV5-1*01

50 sequences assigned
50 (100%) exact matches, in which:
50 unique CDR3
12 unique J



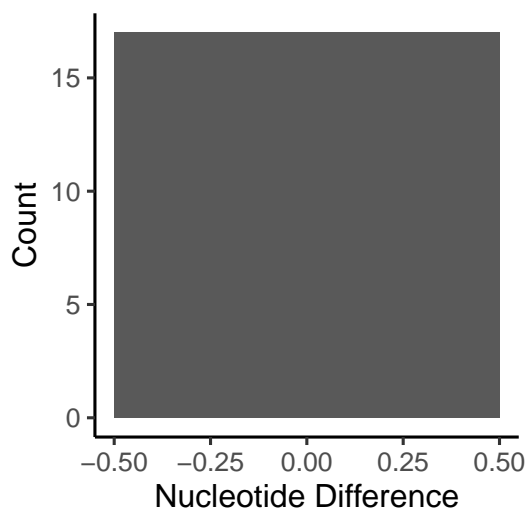
TRBV5-5*01

4 sequences assigned
No exact matches.



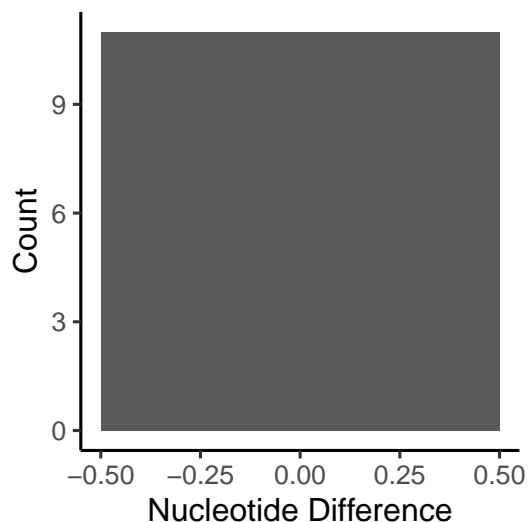
TRBV4-2*01

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



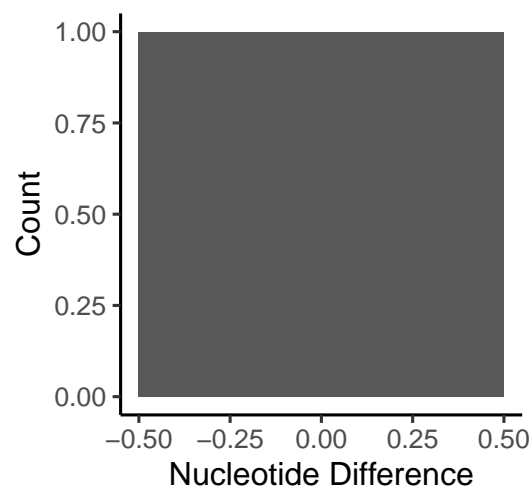
TRBV25-1*01

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



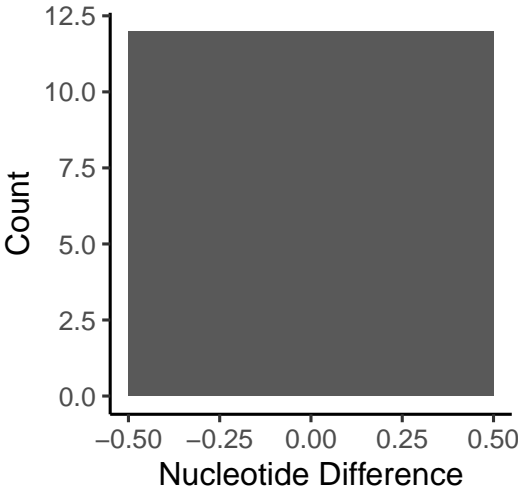
TRBV5-5*02

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



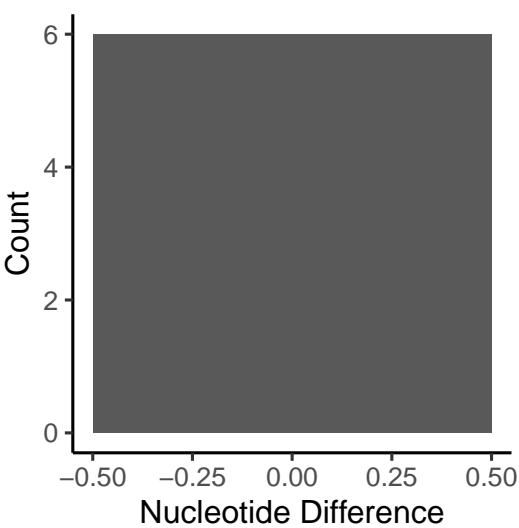
TRBV5–6*01

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



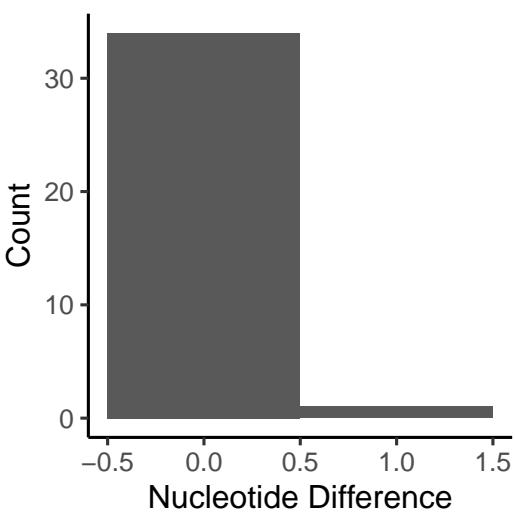
TRBV6–4*01

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



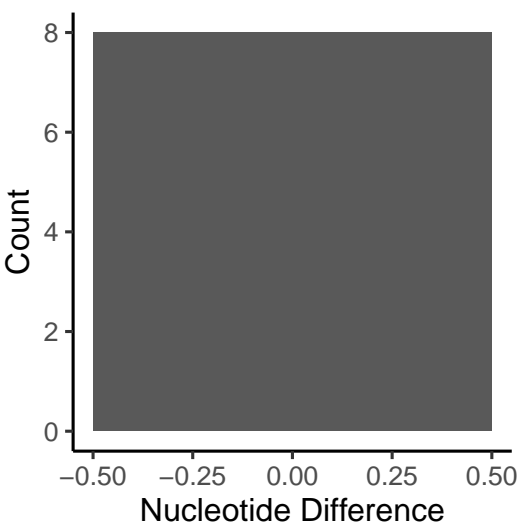
TRBV6–23*01

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



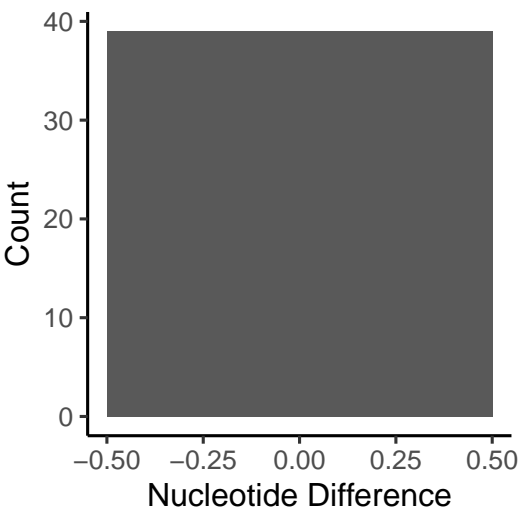
TRBV5–8*01

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



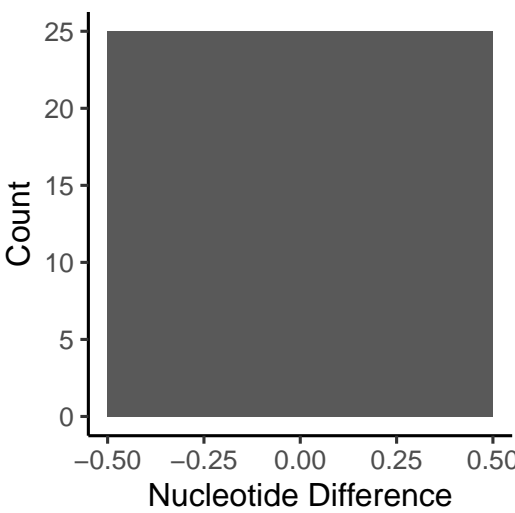
TRBV6–5*01

39 sequences assigned
39 (100%) exact matches, in which:
39 unique CDR3
10 unique J



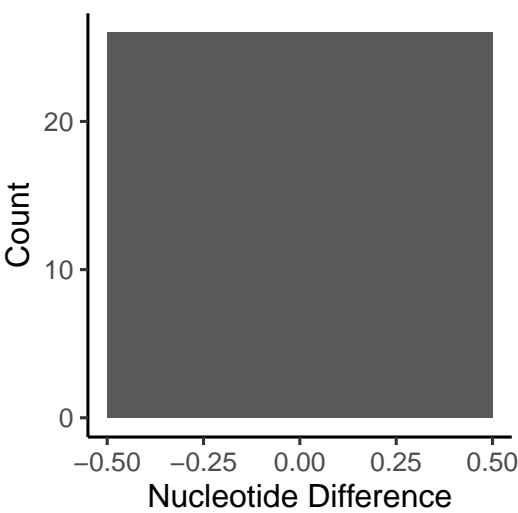
TRBV7–2*01

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



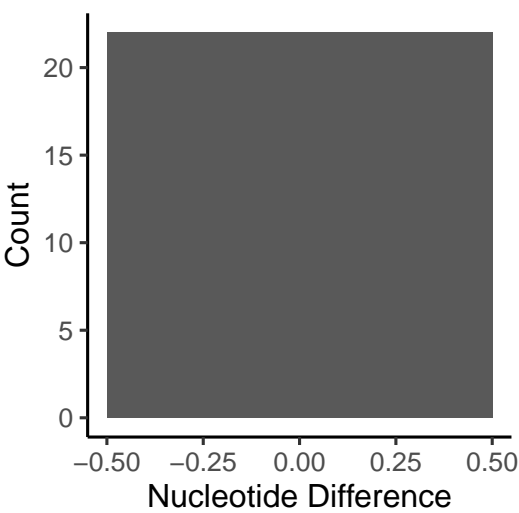
TRBV6–1*01

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



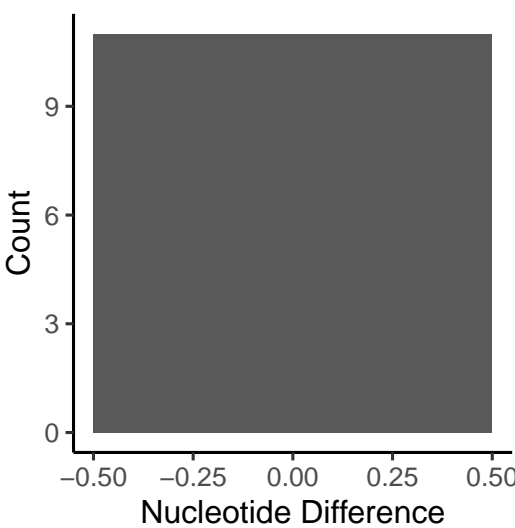
TRBV6–6*01

22 sequences assigned
22 (100%) exact matches, in which:
22 unique CDR3
9 unique J



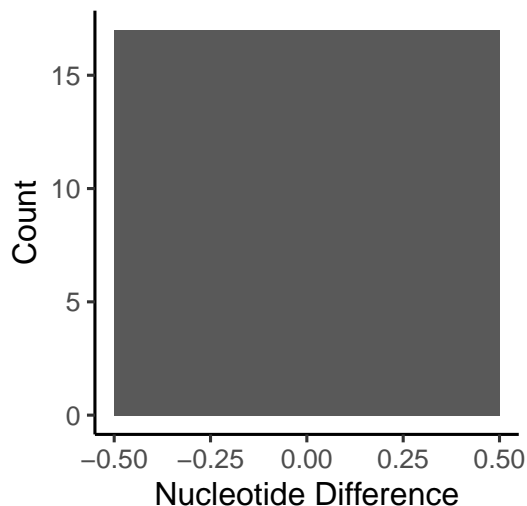
TRBV7–2*02

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



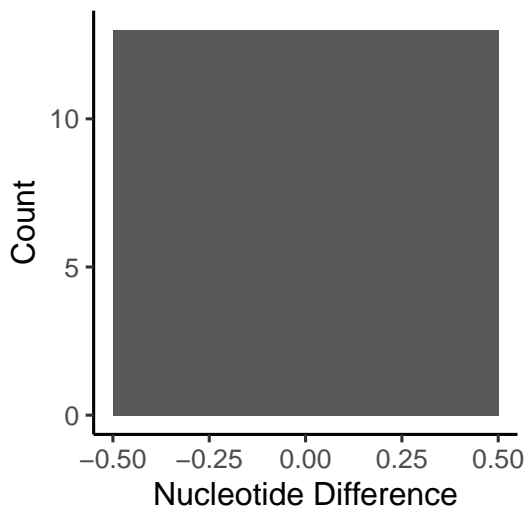
TRBV7-3*01

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



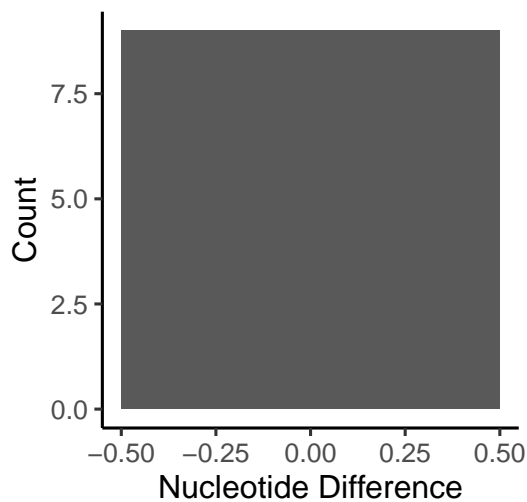
TRBV7-8*01

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



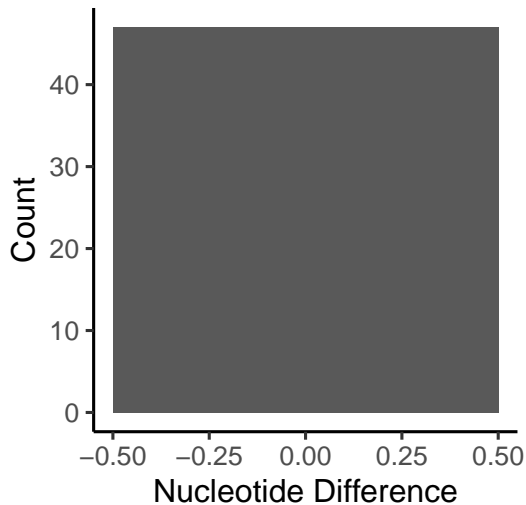
TRBV7-6*01

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



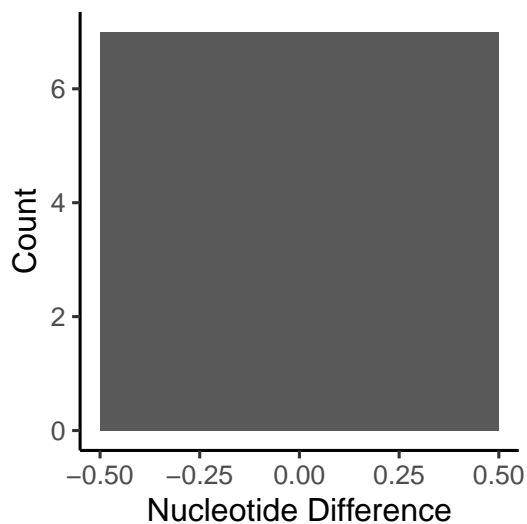
TRBV7-9*03

47 sequences assigned
47 (100%) exact matches, in which:
47 unique CDR3
11 unique J



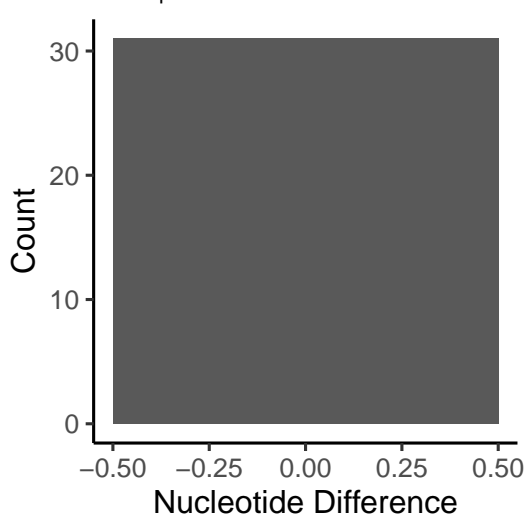
TRBV7-7*01_C315T

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

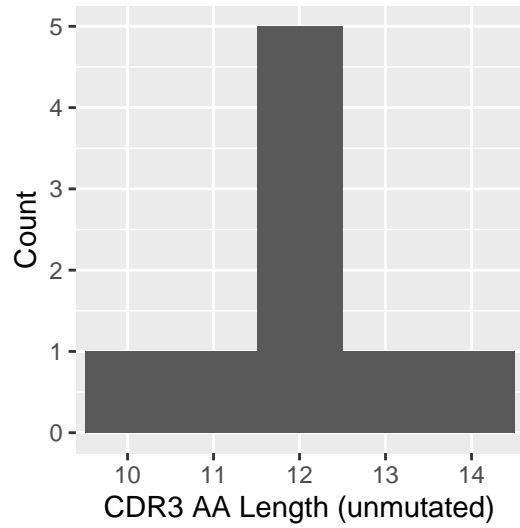


TRBV29-1*01

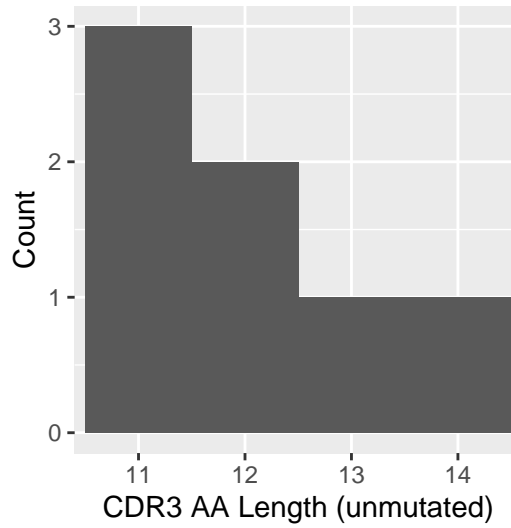
31 sequences assigned
31 (100%) exact matches, in which:
31 unique CDR3
10 unique J



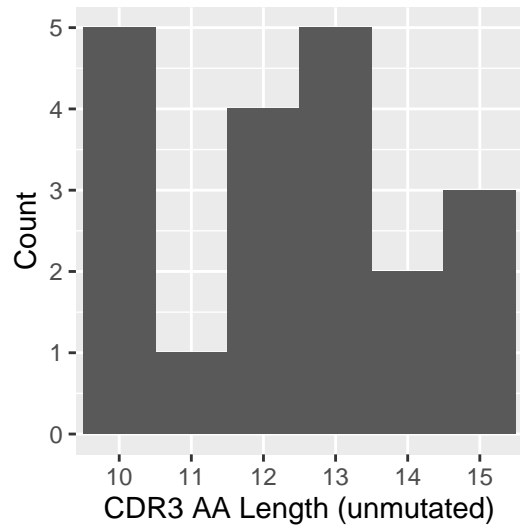
TRBV10-3*03_313GCCATCA



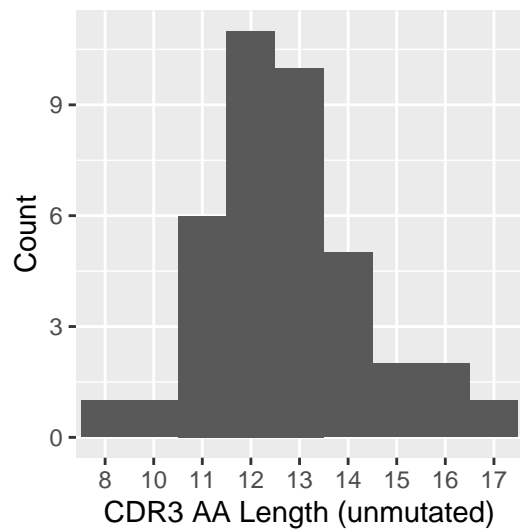
TRBV7-7*01_C315T

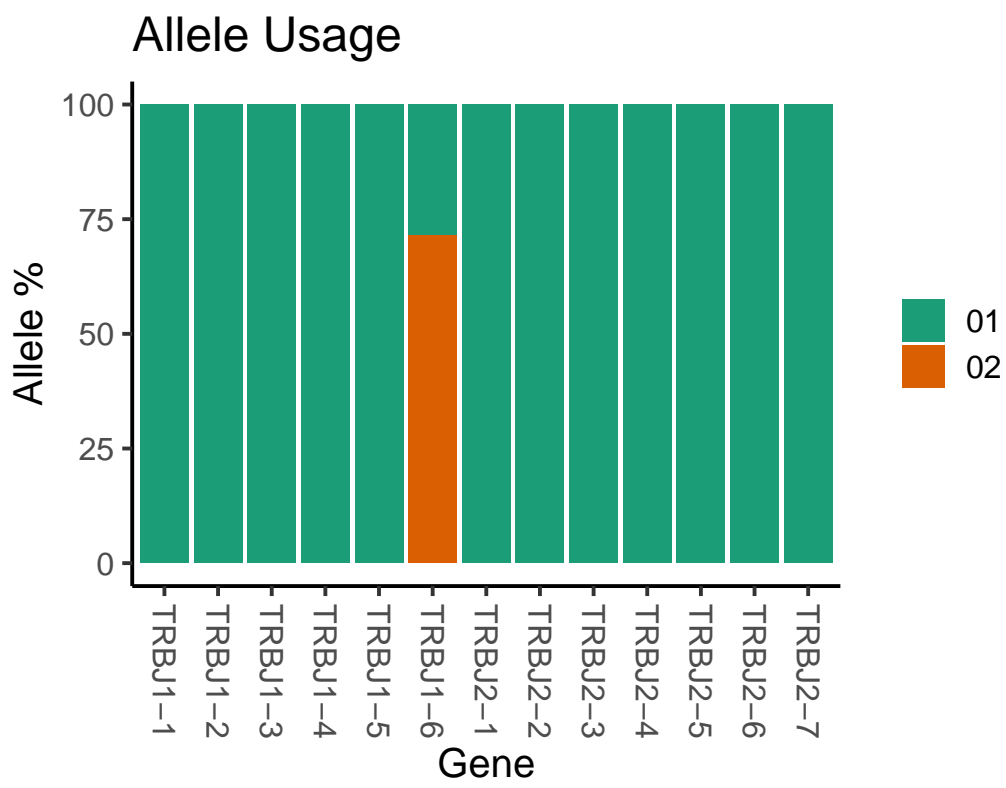


TRBV12-4*01_C87T



TRBV19*01_A24G





Sequence Count by TRBJ1–6 allele usage



Warning – no inferred sequences found.

Warning – no inferred sequences found.