

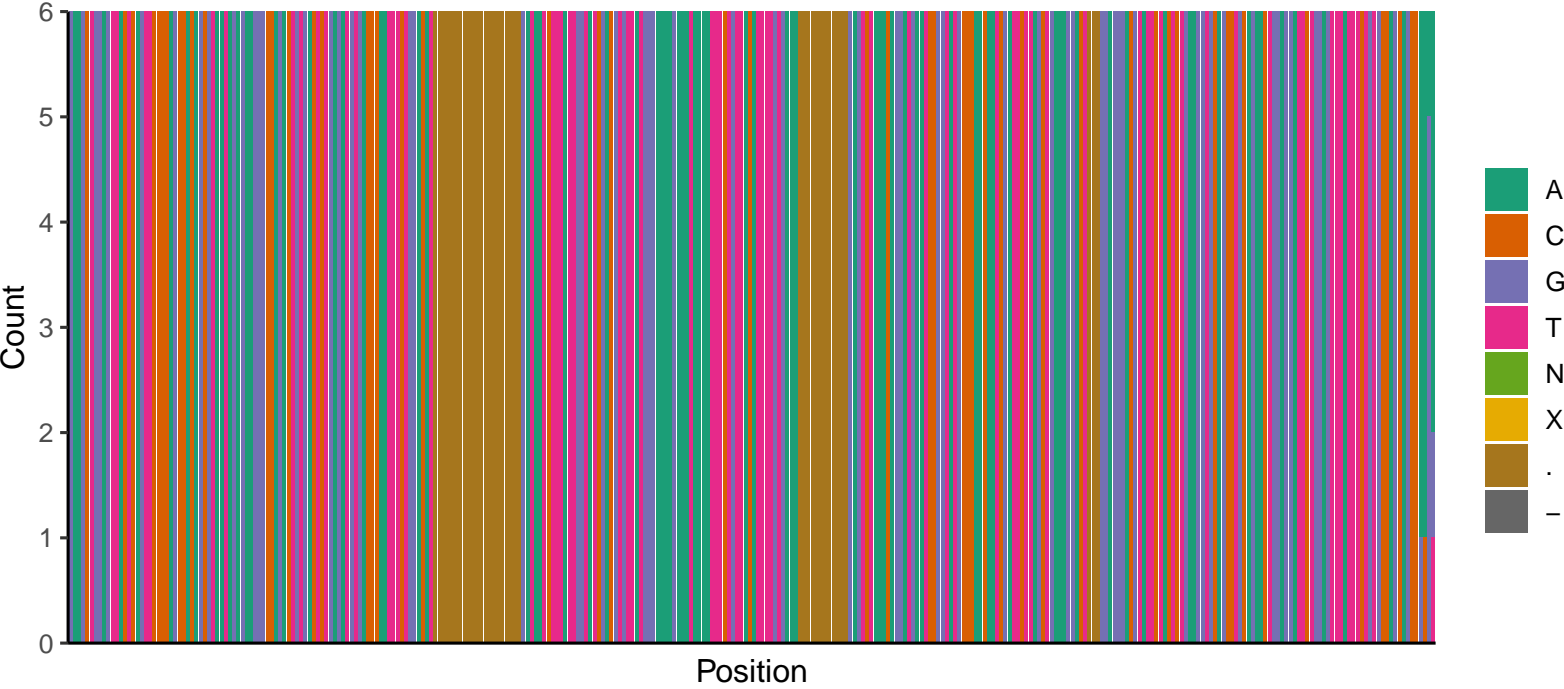
Gene TRBV10-3*03_313GCCATCAGTGAGTC326



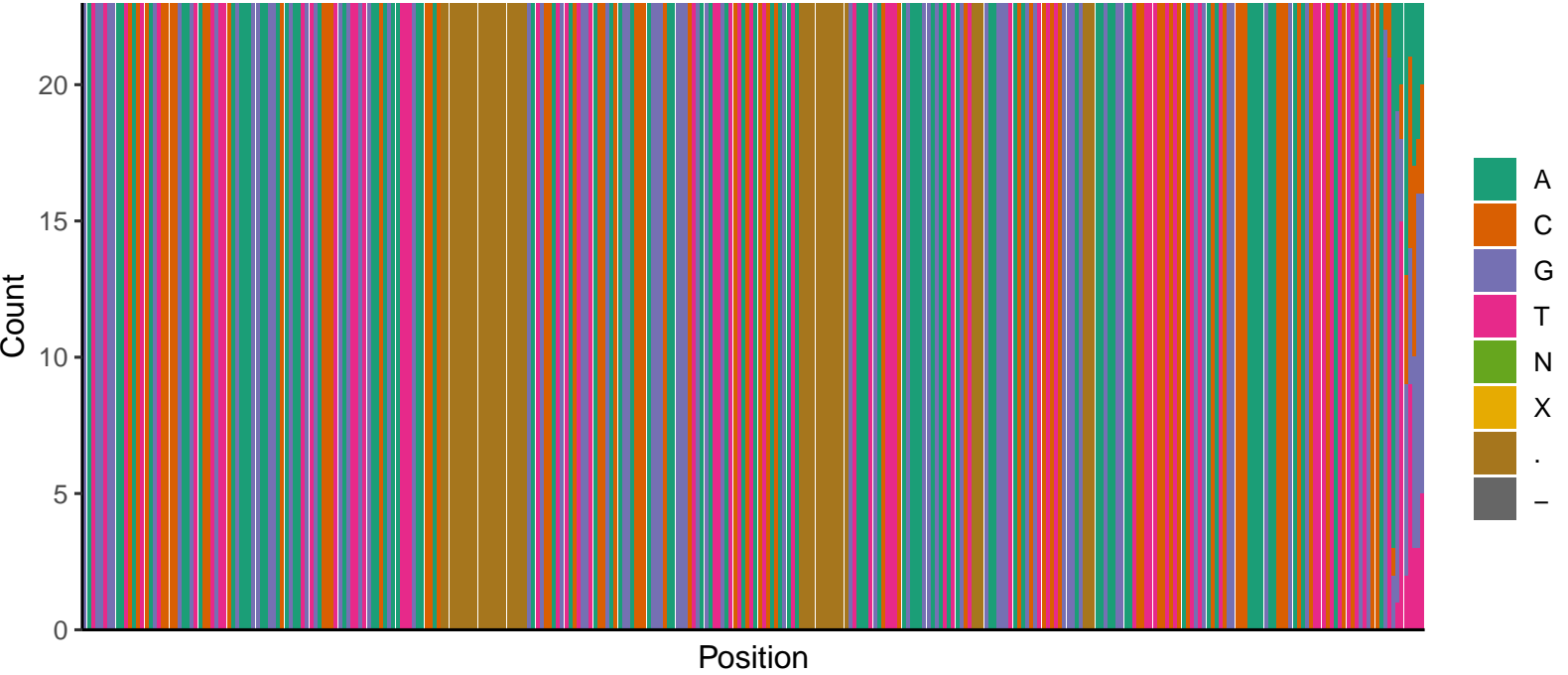
Gene TRBV12-4*01_C87T



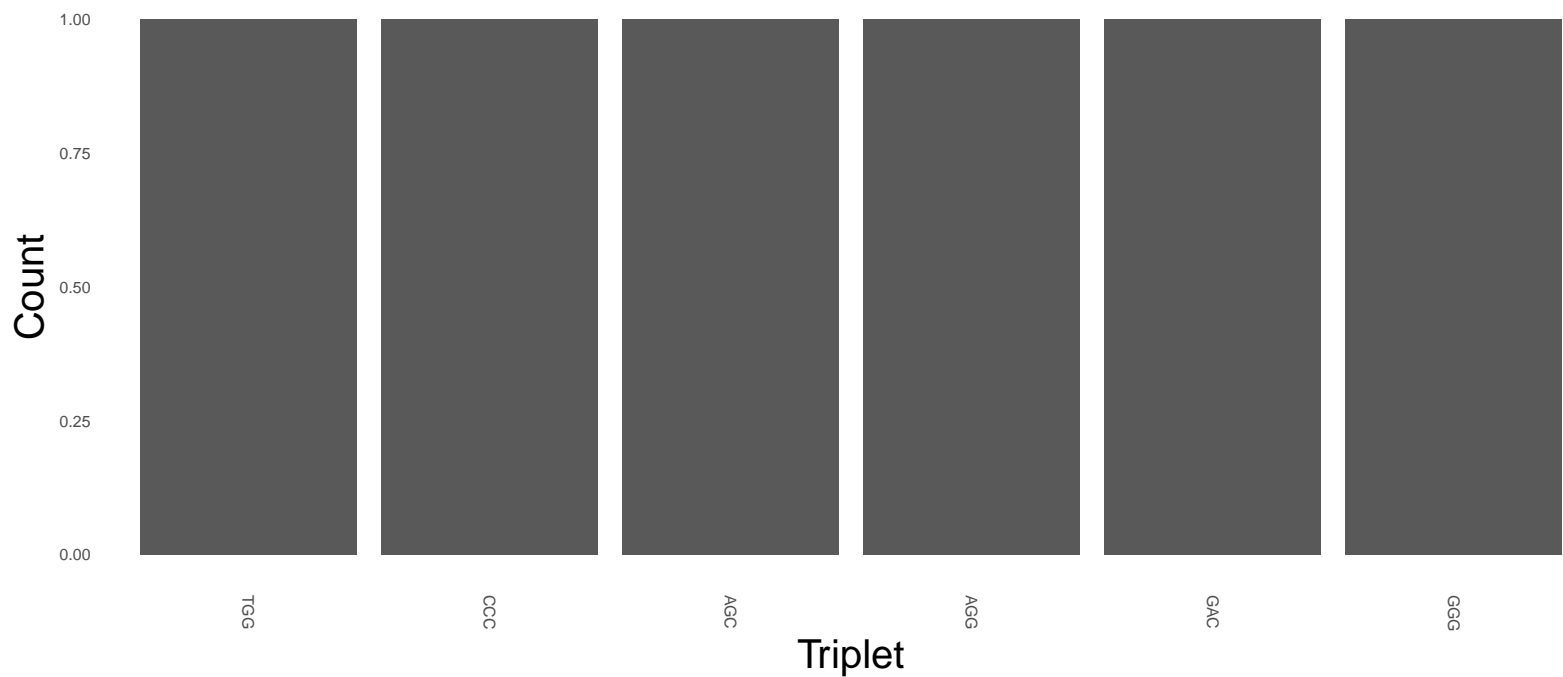
Gene TRBV14*02_322CAAGA326



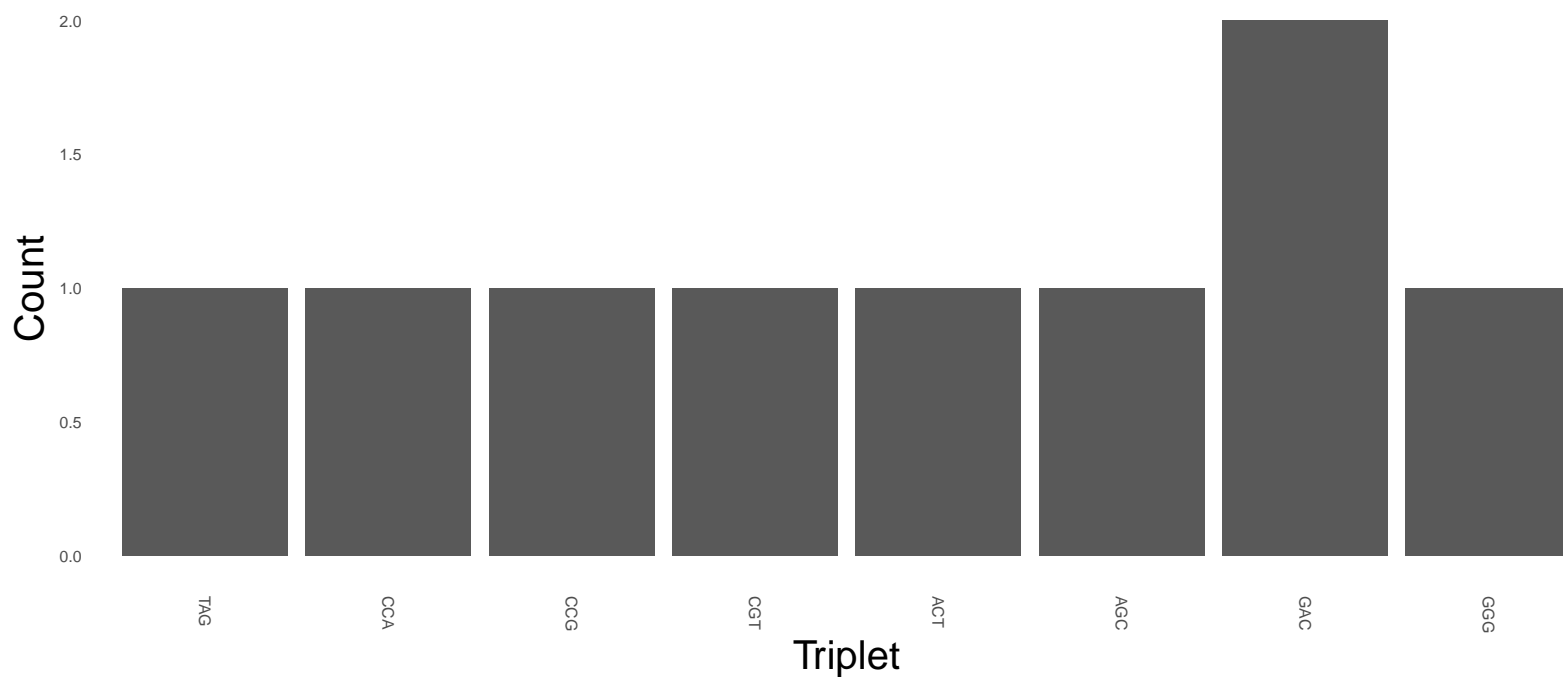
Gene TRBV19*01_A24G



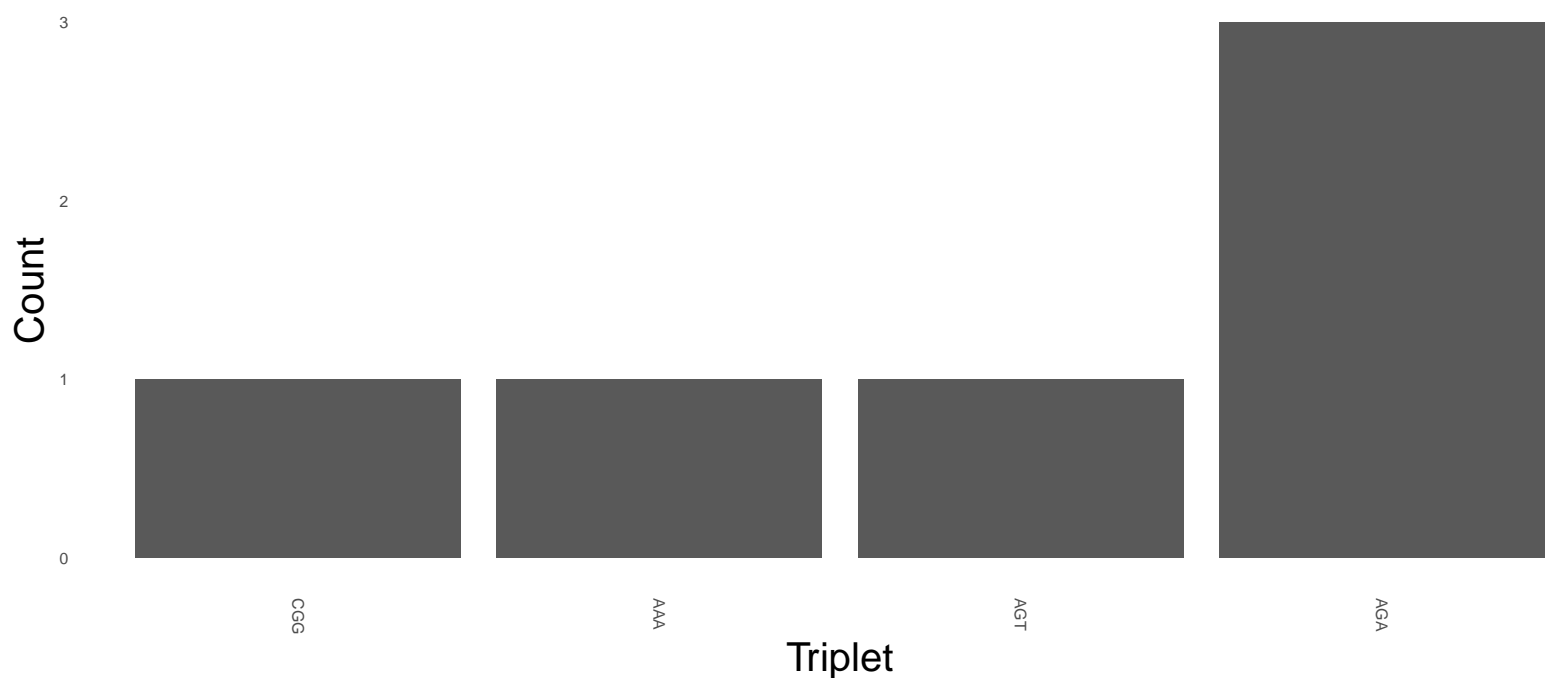
TRBV10–3*03_313GCCATCAGTGAGTC326– Final 3 nucleotides as a triplet



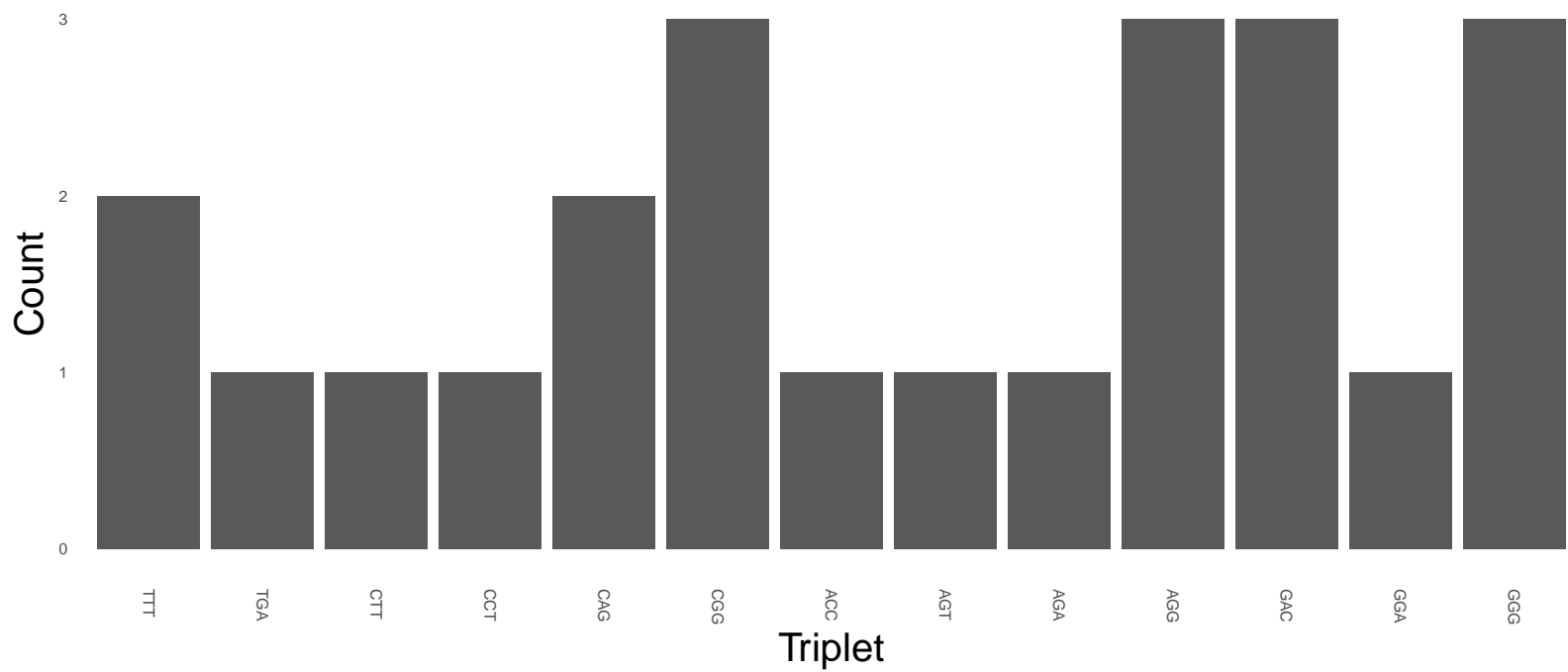
TRBV12–4*01_C87T– Final 3 nucleotides as a triplet



TRBV14*02_322CAAGA326– Final 3 nucleotides as a triplet

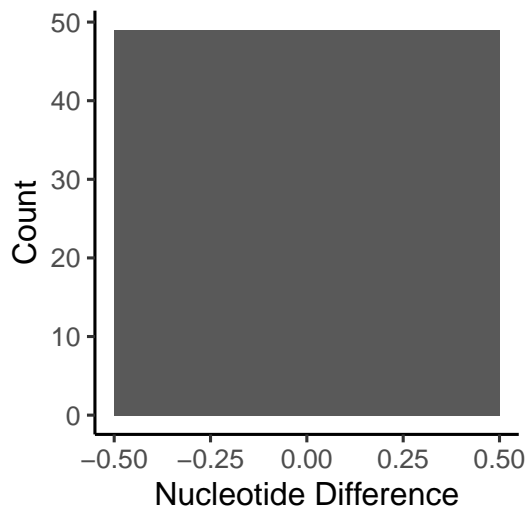


TRBV19*01_A24G– Final 3 nucleotides as a triplet



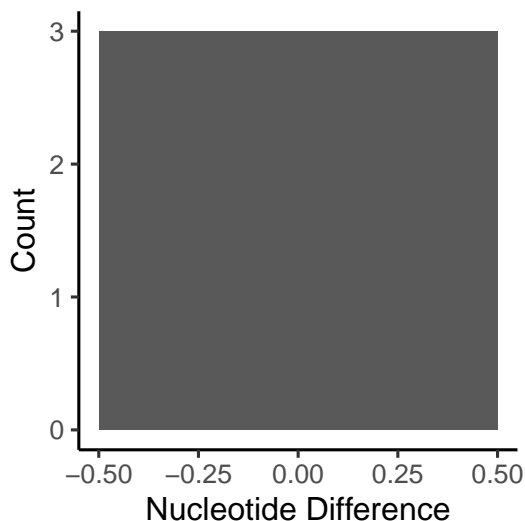
TRBV20–1*01

49 sequences assigned
49 (100%) exact matches, in which:
49 unique CDR3
9 unique J



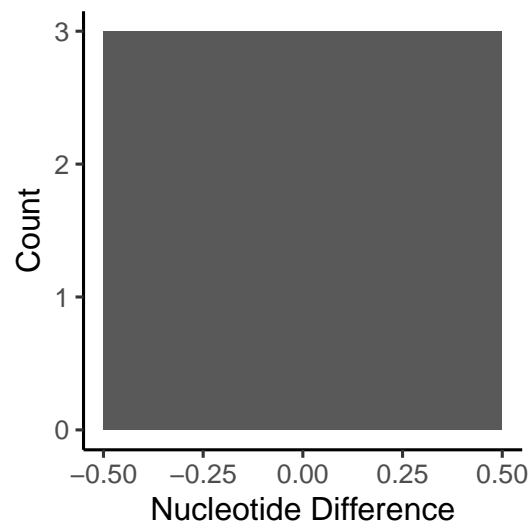
TRBV10–2*01

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



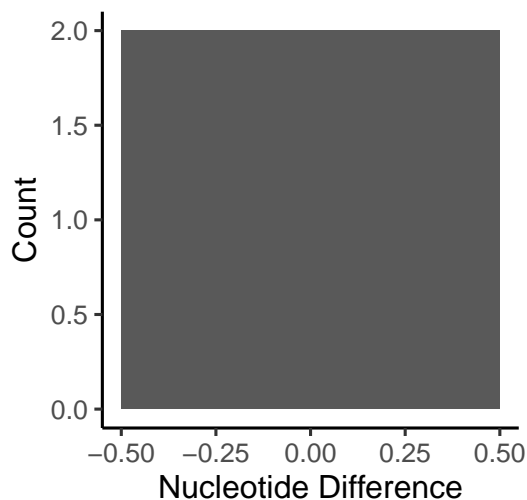
TRBV13*01

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



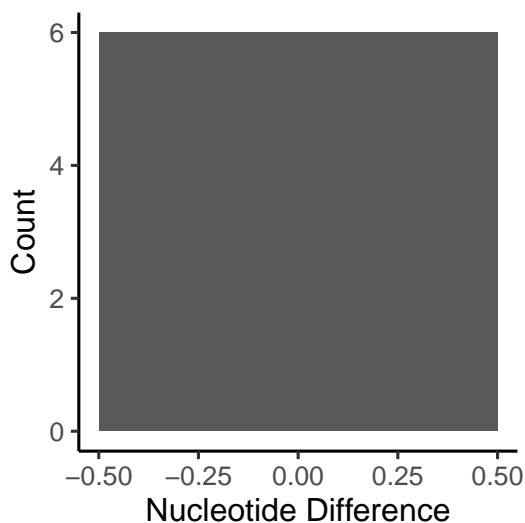
TRBV10–1*01

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



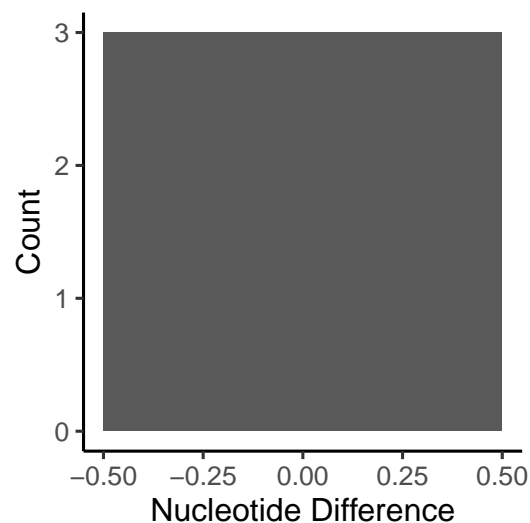
TRBV10–3*02

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



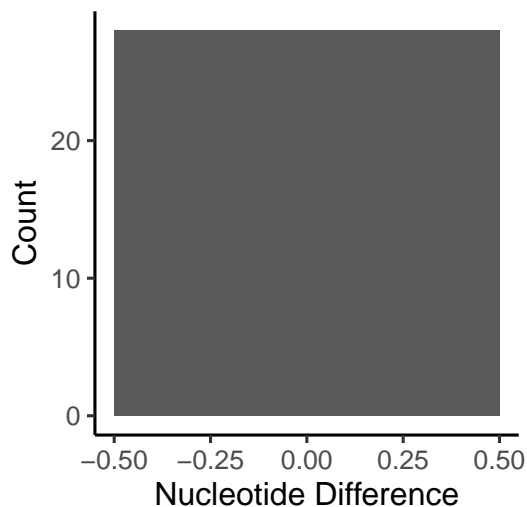
TRBV14*01

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



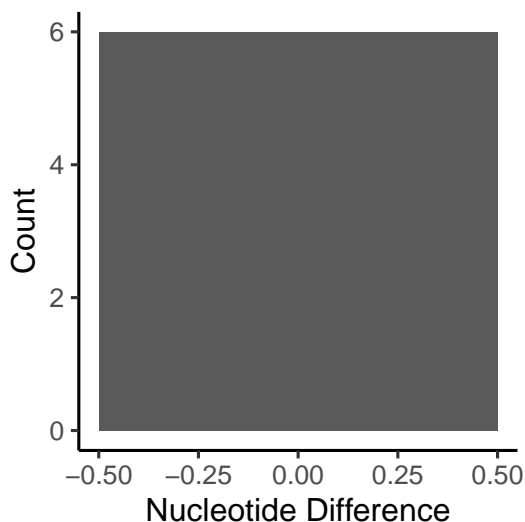
TRBV20–1*05

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



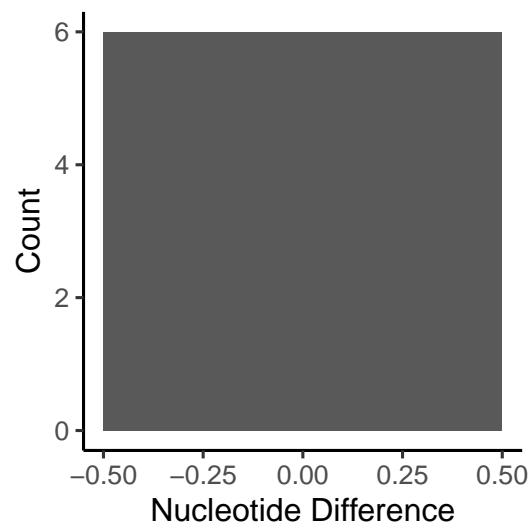
TRBV10–3*03_313GCCATG

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



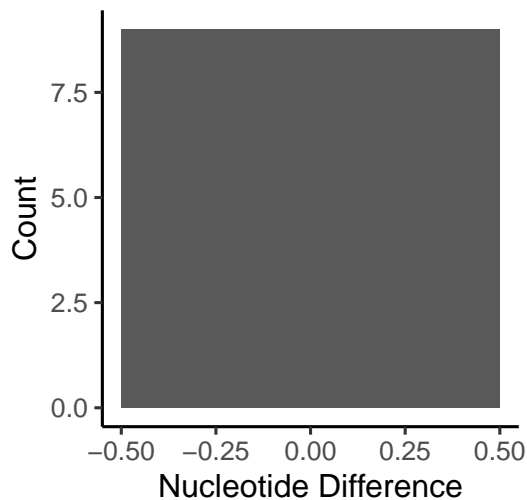
TRBV14*02_322CAAGA32

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



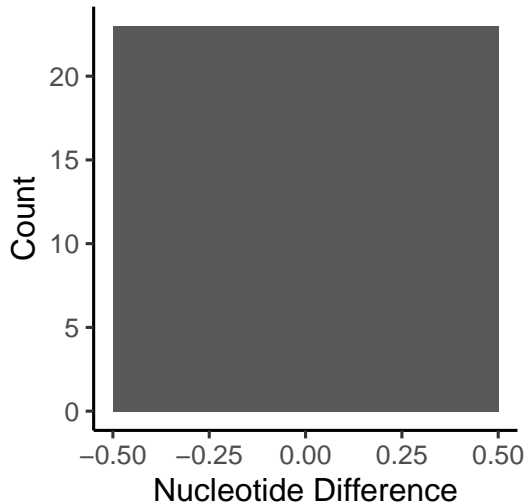
TRBV15*02

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



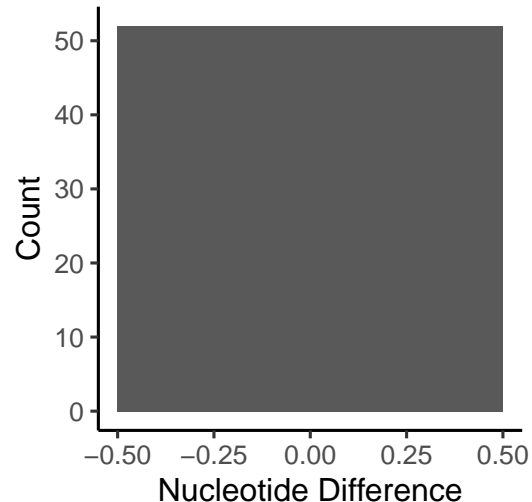
TRBV19*01_A24G

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



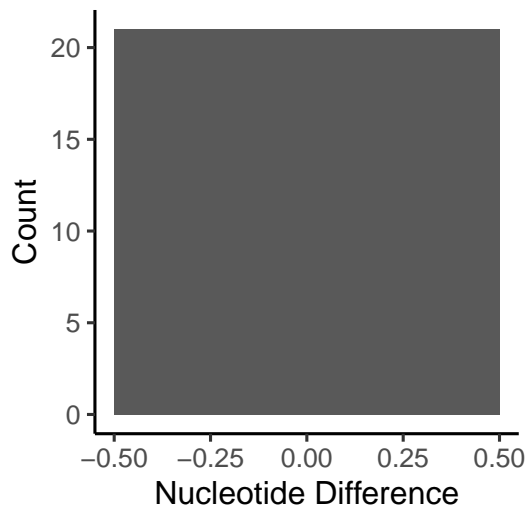
TRBV28*01

52 sequences assigned
52 (100%) exact matches, in which:
52 unique CDR3
9 unique J



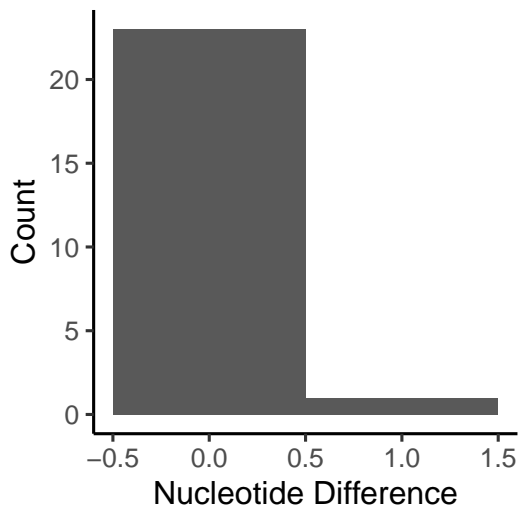
TRBV18*01

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



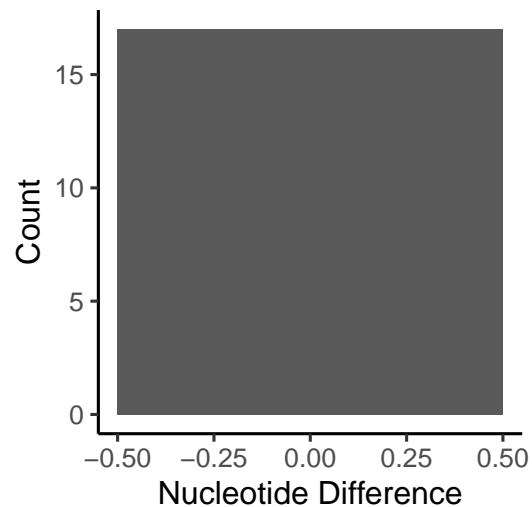
TRBV2*01

24 sequences assigned
23 (95.8%) exact matches, in which:
23 unique CDR3
10 unique J



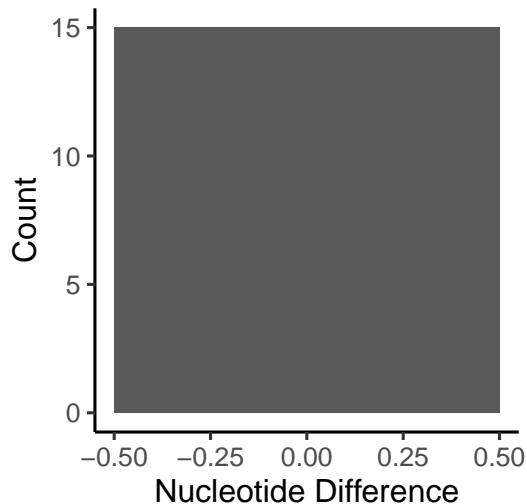
TRBV30*01

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



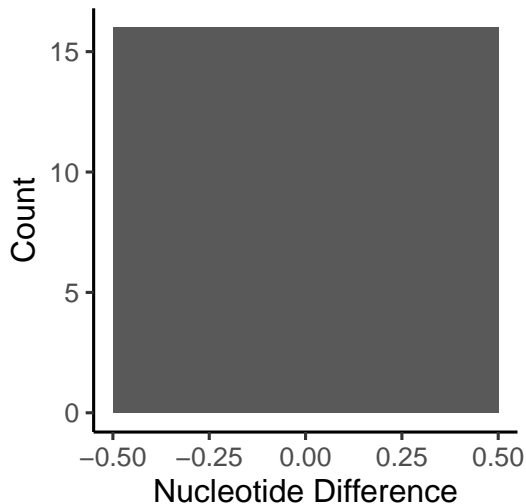
TRBV19*01

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



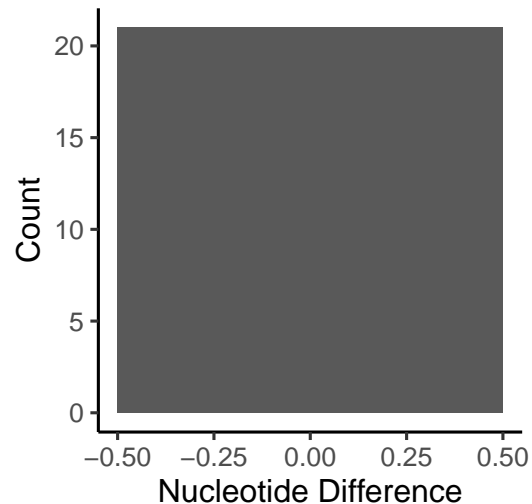
TRBV27*01

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



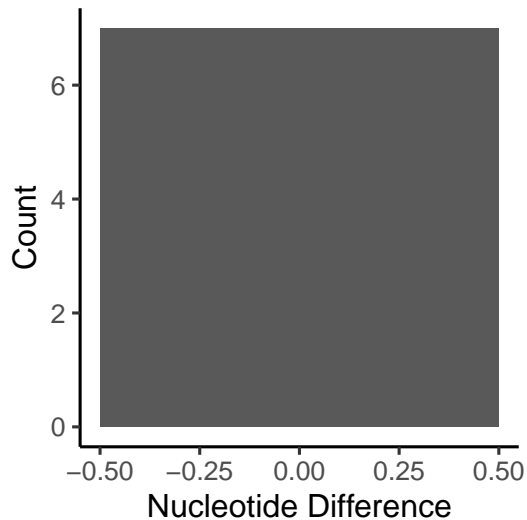
TRBV9*01

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



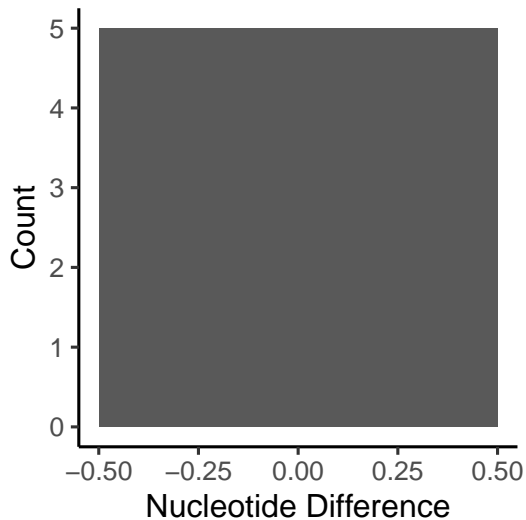
TRBV11–1*01

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



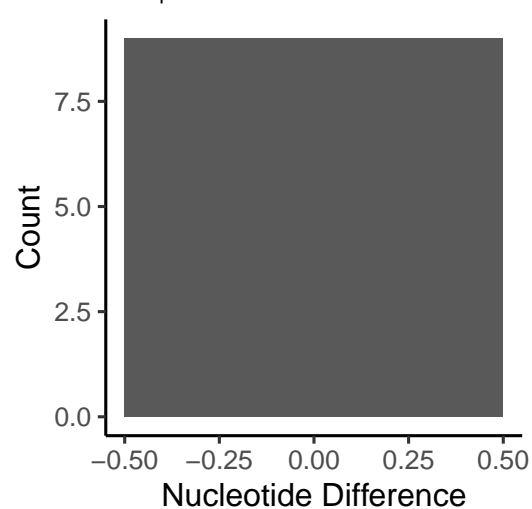
TRBV11–3*04

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



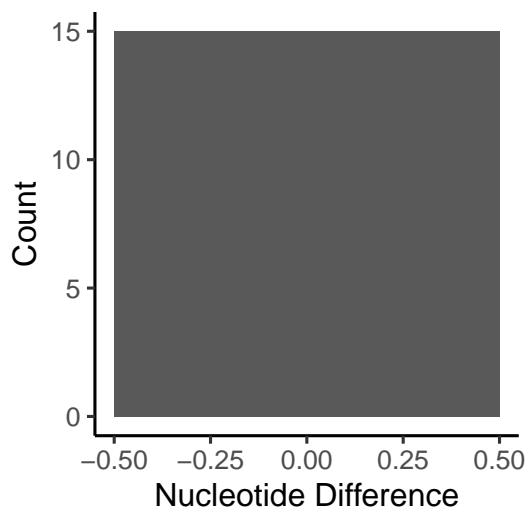
TRBV12–4*01_C87T

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



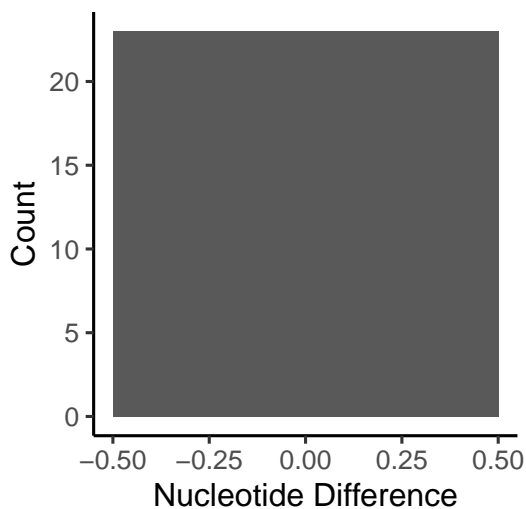
TRBV11–2*01

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



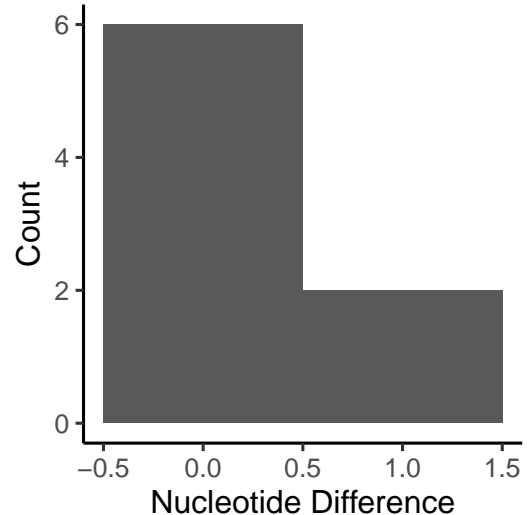
TRBV12–3*01

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



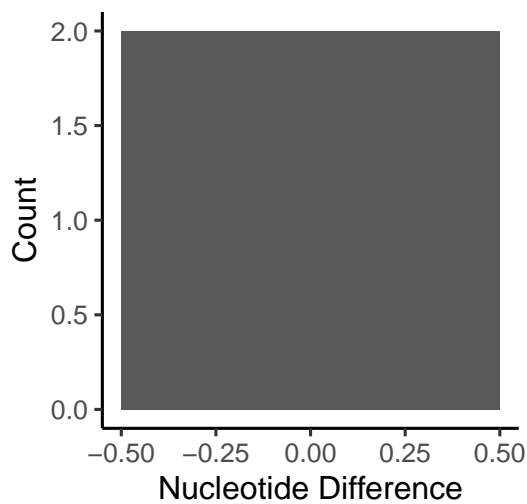
TRBV12–5*01

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



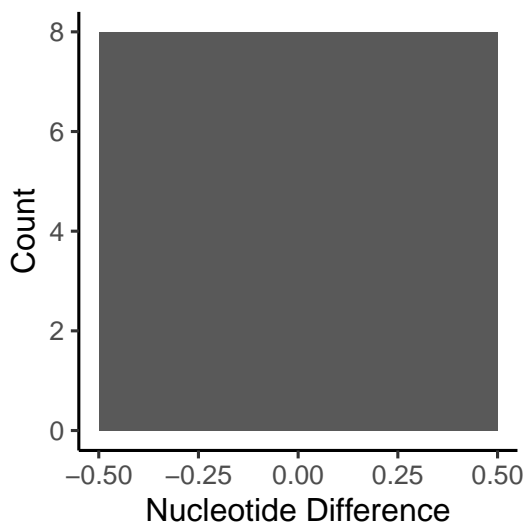
TRBV11–3*01

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



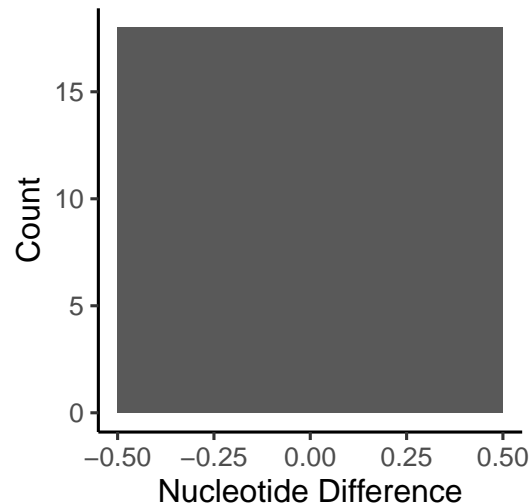
TRBV12–4*01

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



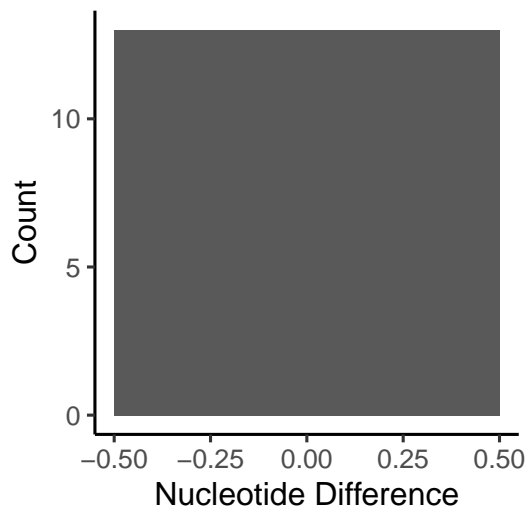
TRBV3–1*01

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



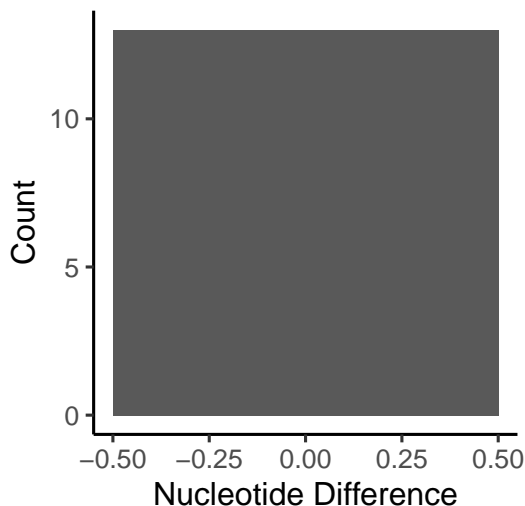
TRBV24–1*01

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



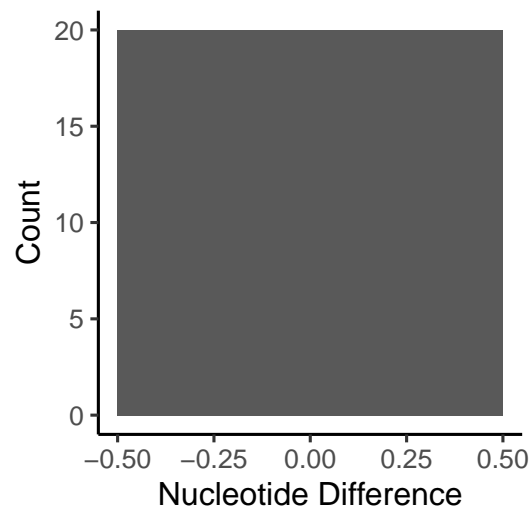
TRBV4–3*01

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



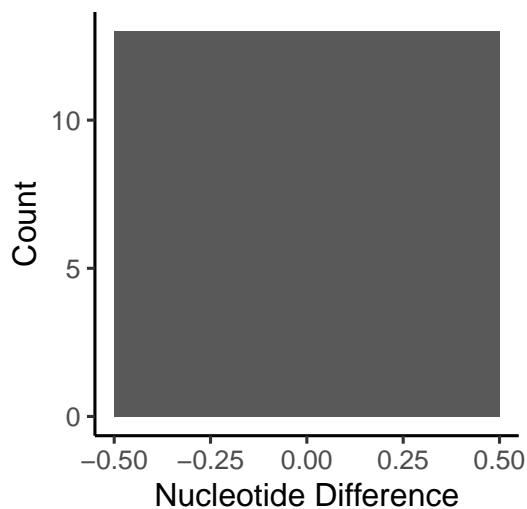
TRBV5–4*01

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



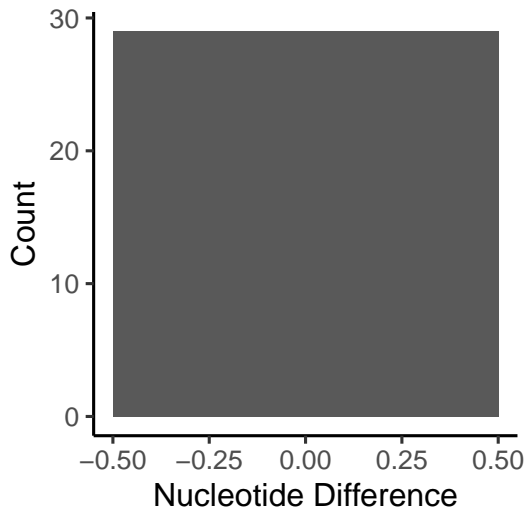
TRBV4–1*01

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



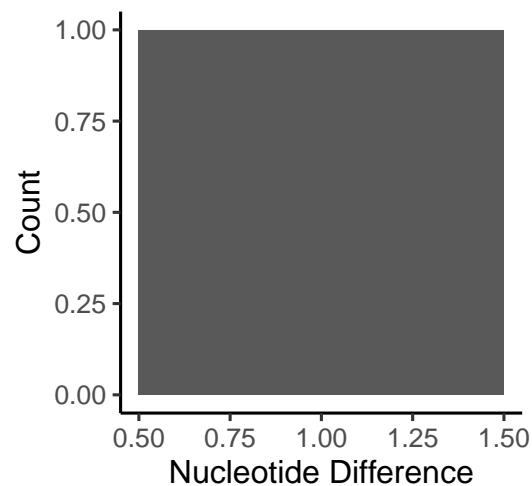
TRBV5–1*01

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



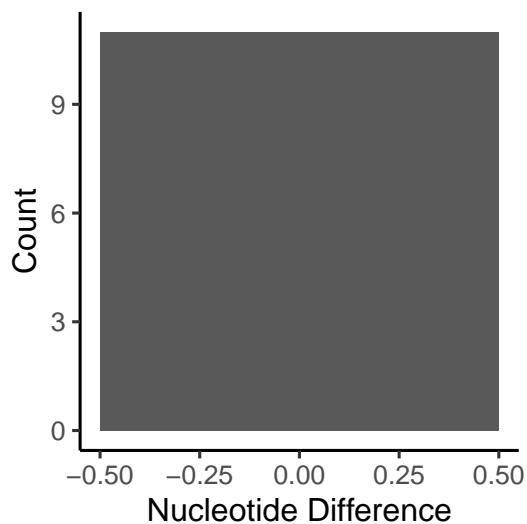
TRBV5–5*01

1 sequences assigned
No exact matches.



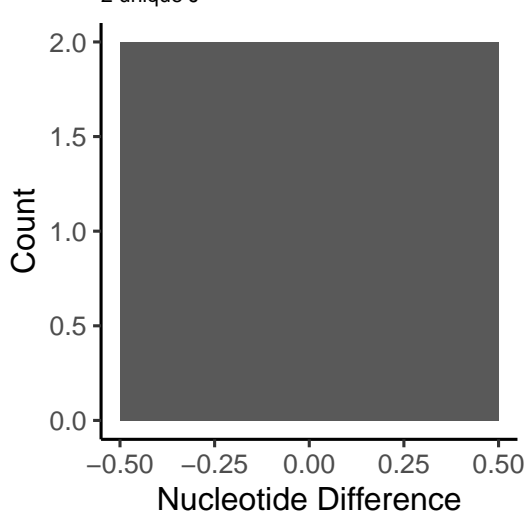
TRBV4–2*01

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



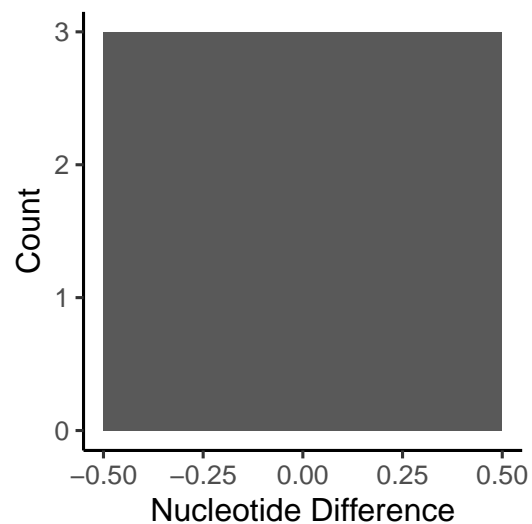
TRBV25–1*01

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



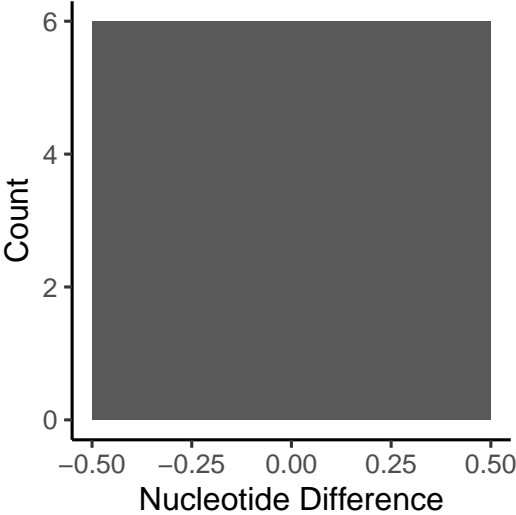
TRBV5–5*02

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



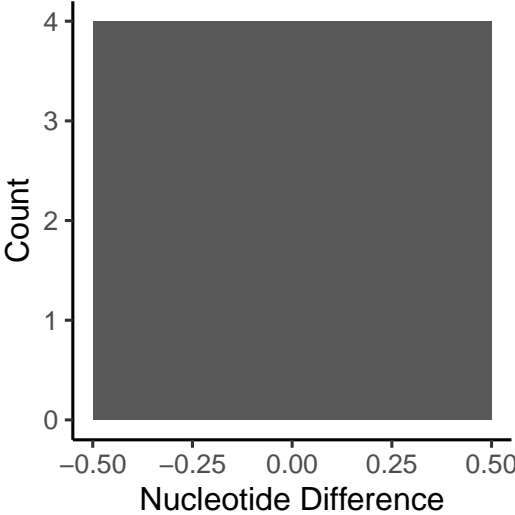
TRBV5-6*01

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



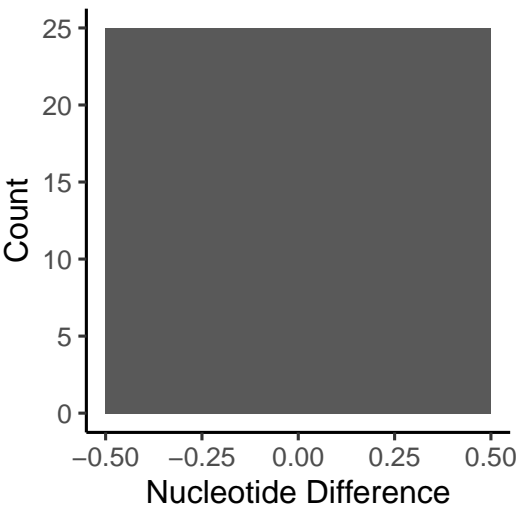
TRBV6-4*01

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



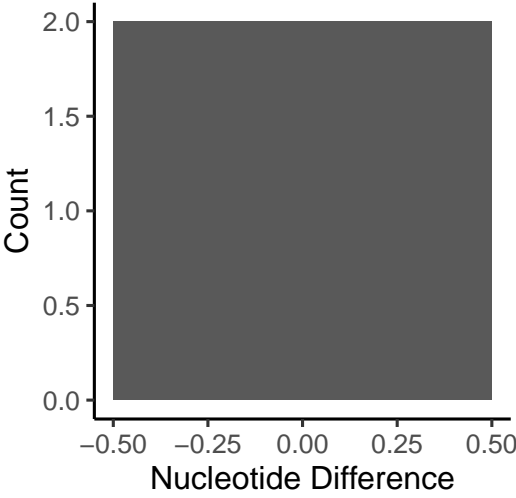
TRBV6-23*01

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



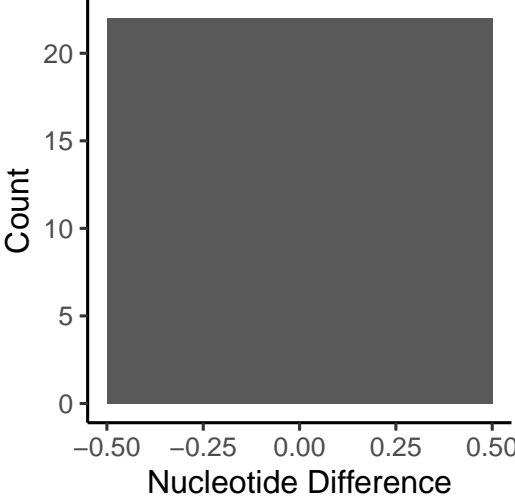
TRBV5-8*01

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



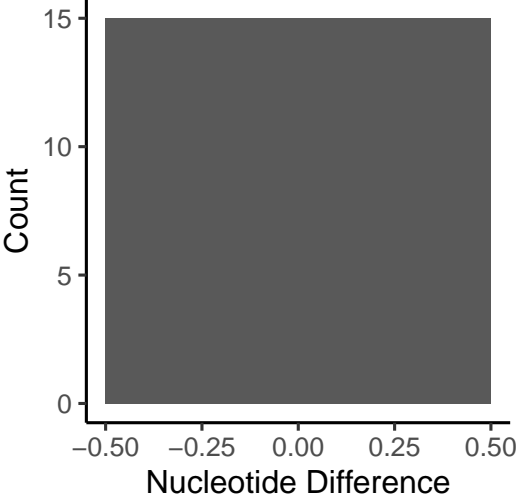
TRBV6-5*01

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



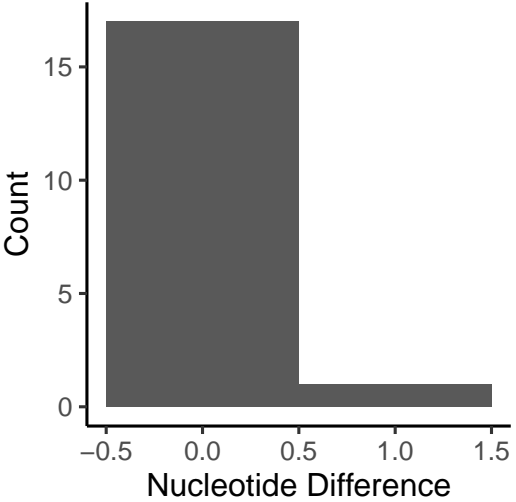
TRBV7-2*01

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



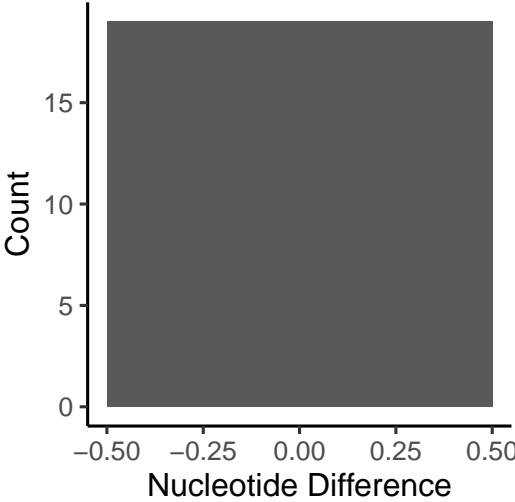
TRBV6-1*01

18 sequences assigned
17 (94.4%) exact matches, in which:
17 unique CDR3
10 unique J



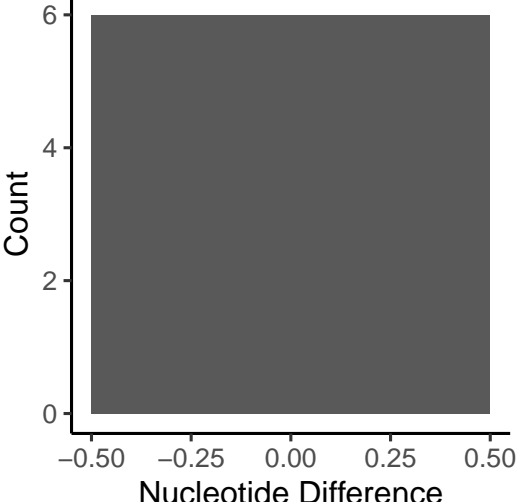
TRBV6-6*01

19 sequences assigned
19 (100%) exact matches, in which:
19 unique CDR3
9 unique J



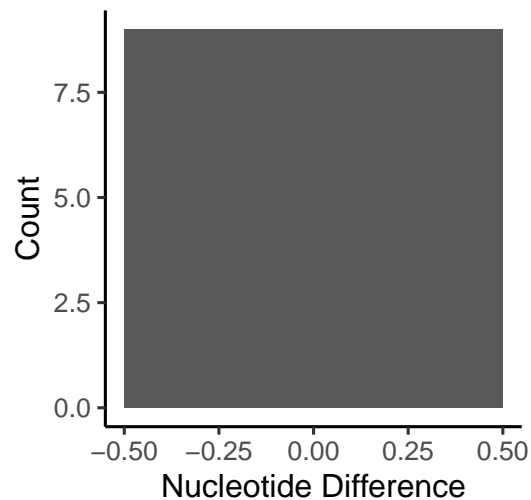
TRBV7-2*02

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



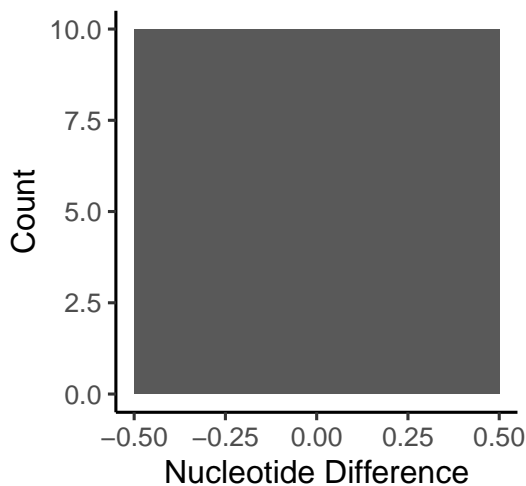
TRBV7-3*01

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



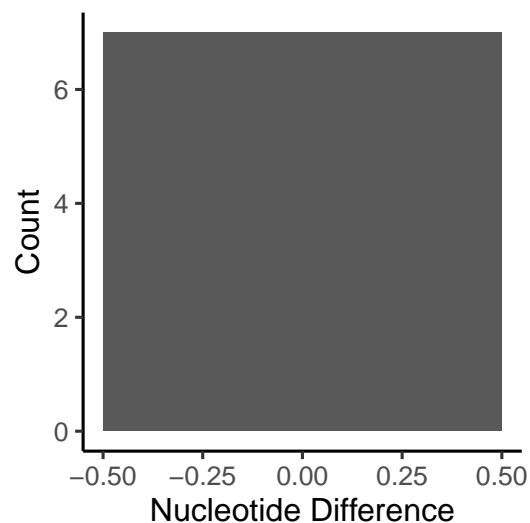
TRBV7-8*01

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



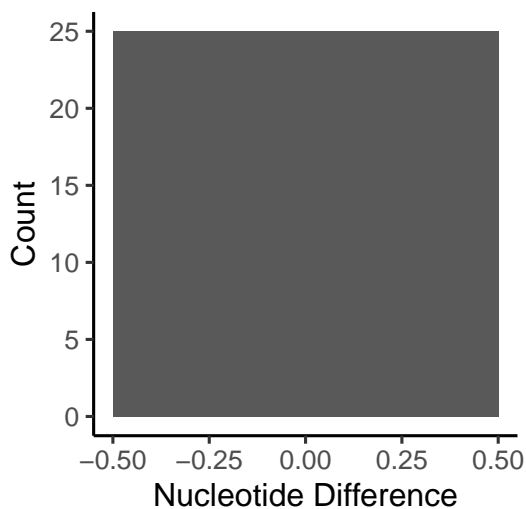
TRBV7-6*01

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



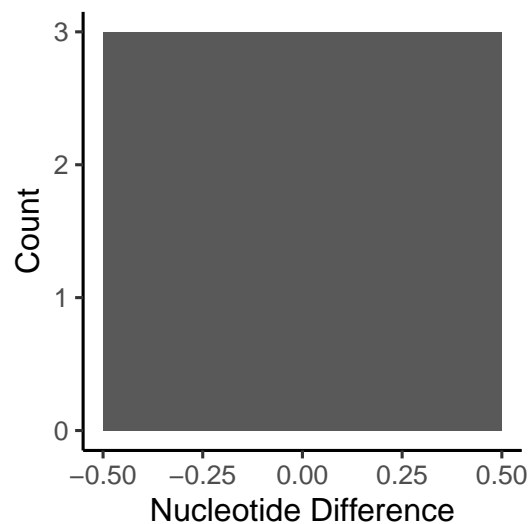
TRBV7-9*03

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



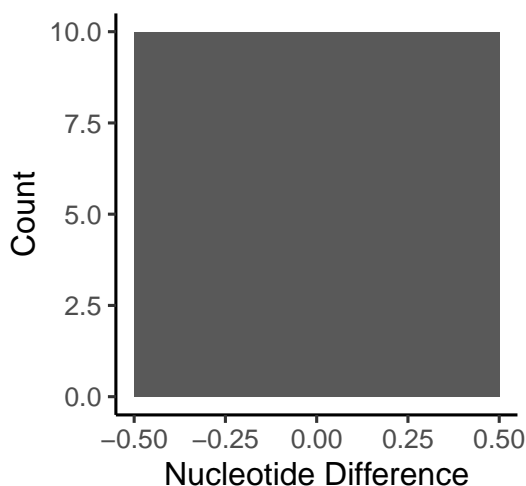
TRBV7-7*01

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

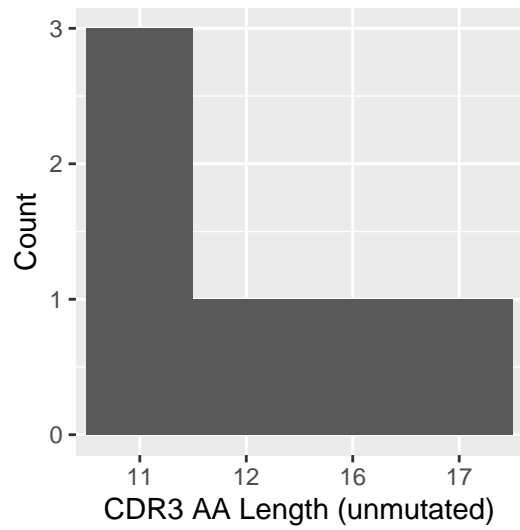


TRBV29-1*01

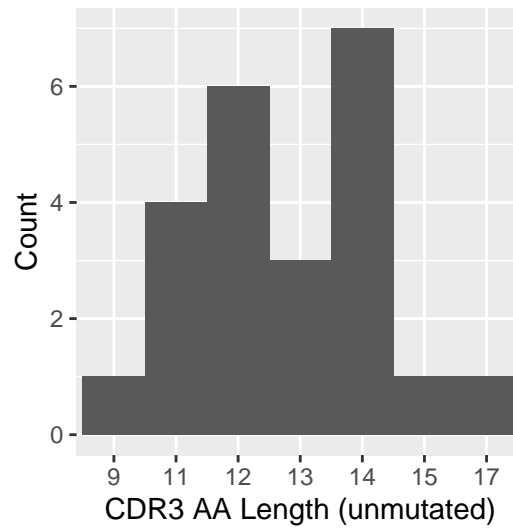
10 sequences assigned
10 (100%) exact matches, in which:
10 unique CDR3
6 unique J



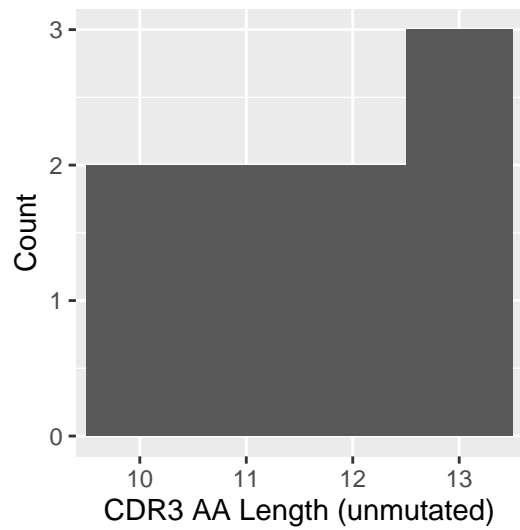
TRBV10-3*03_313GCCATCA



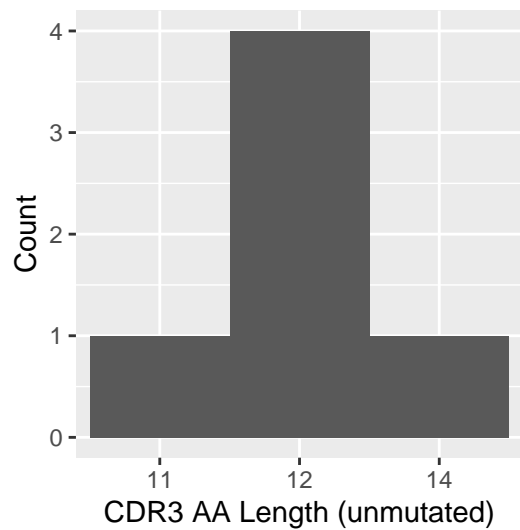
TRBV19*01_A24G

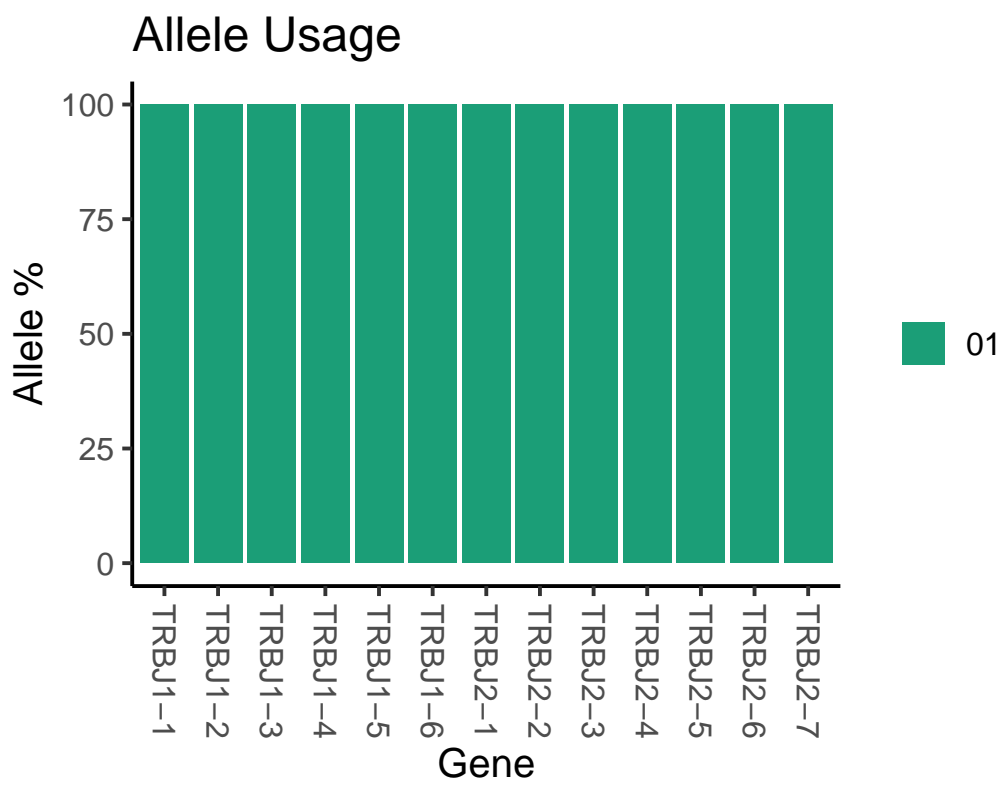


TRBV12-4*01_C87T



TRBV14*02_322CAAGA326





Warning – no inferred sequences found.

Warning – no inferred sequences found.