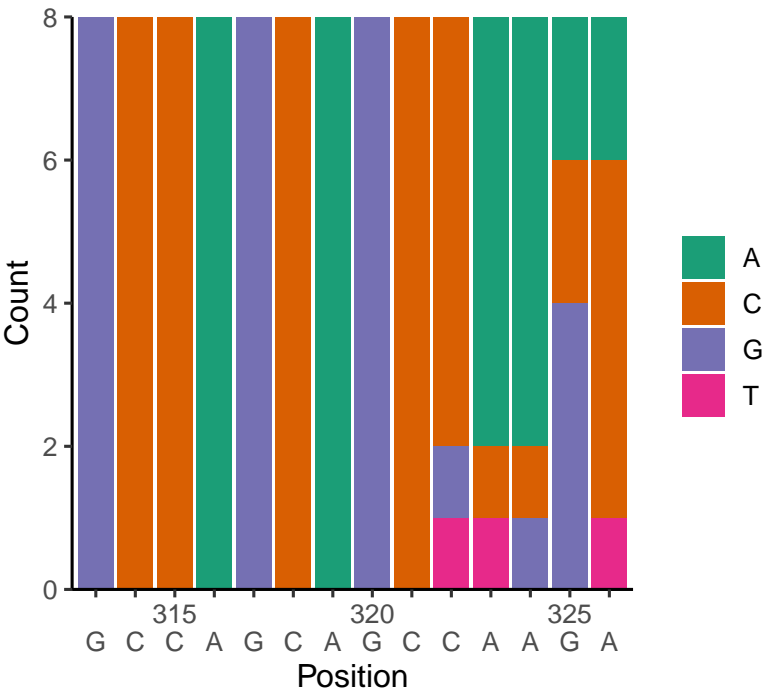
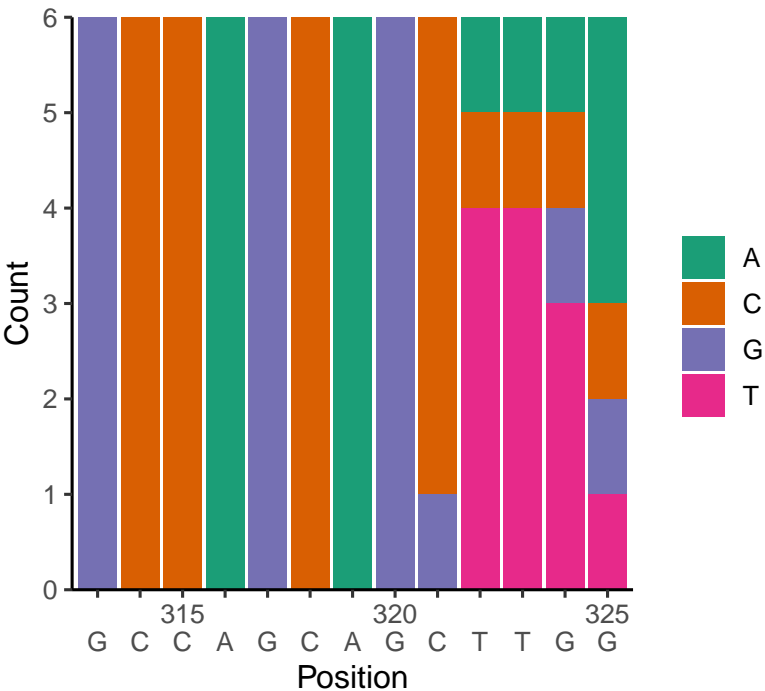


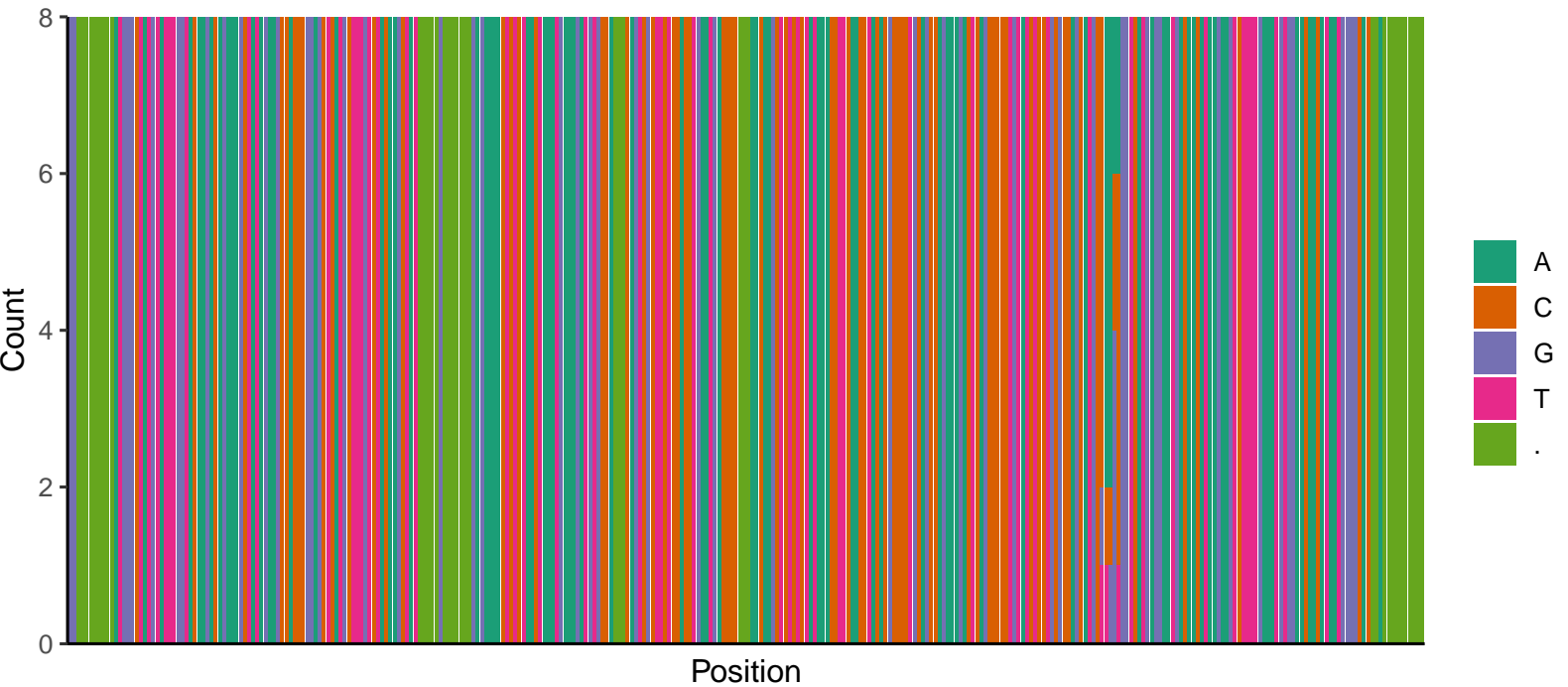
Gene TRBV4-1*01_C14T



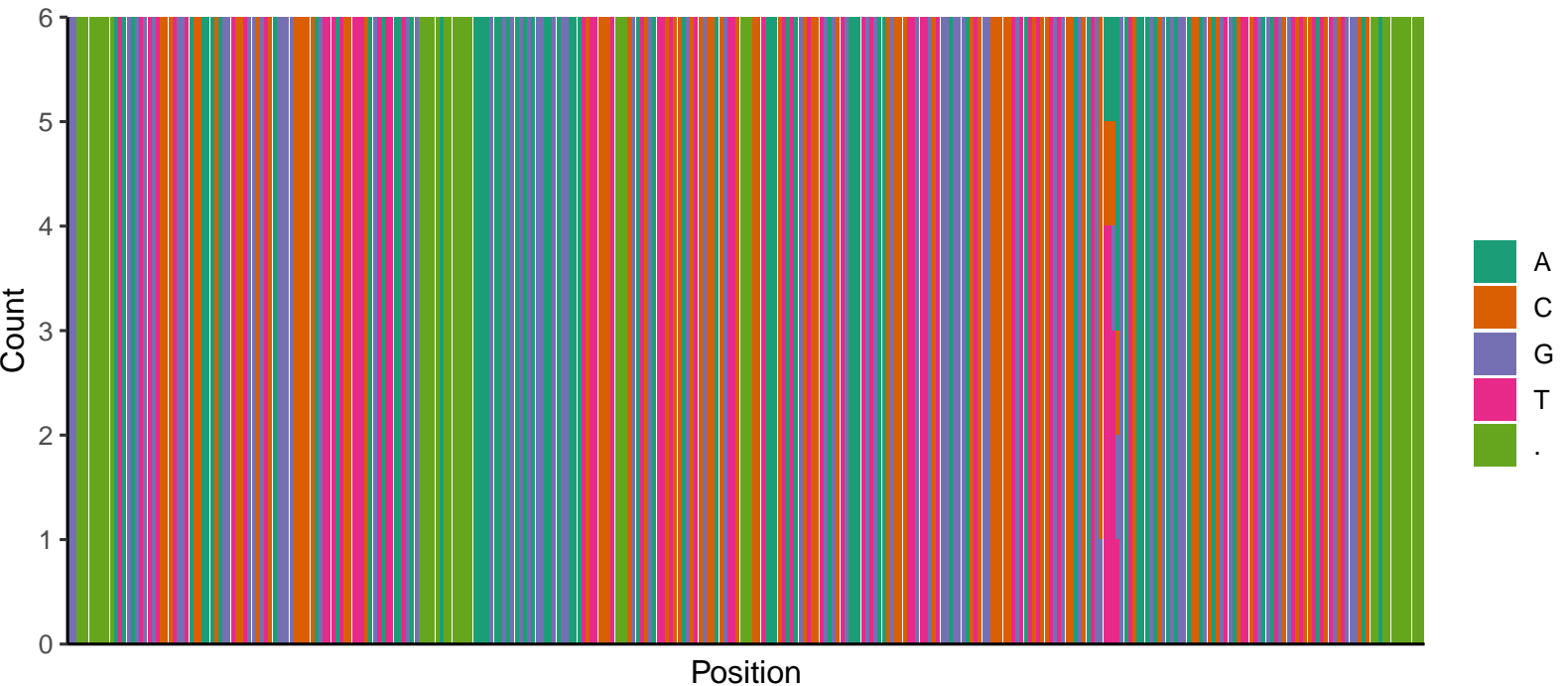
Gene TRBV5-5*02_322TTGG325



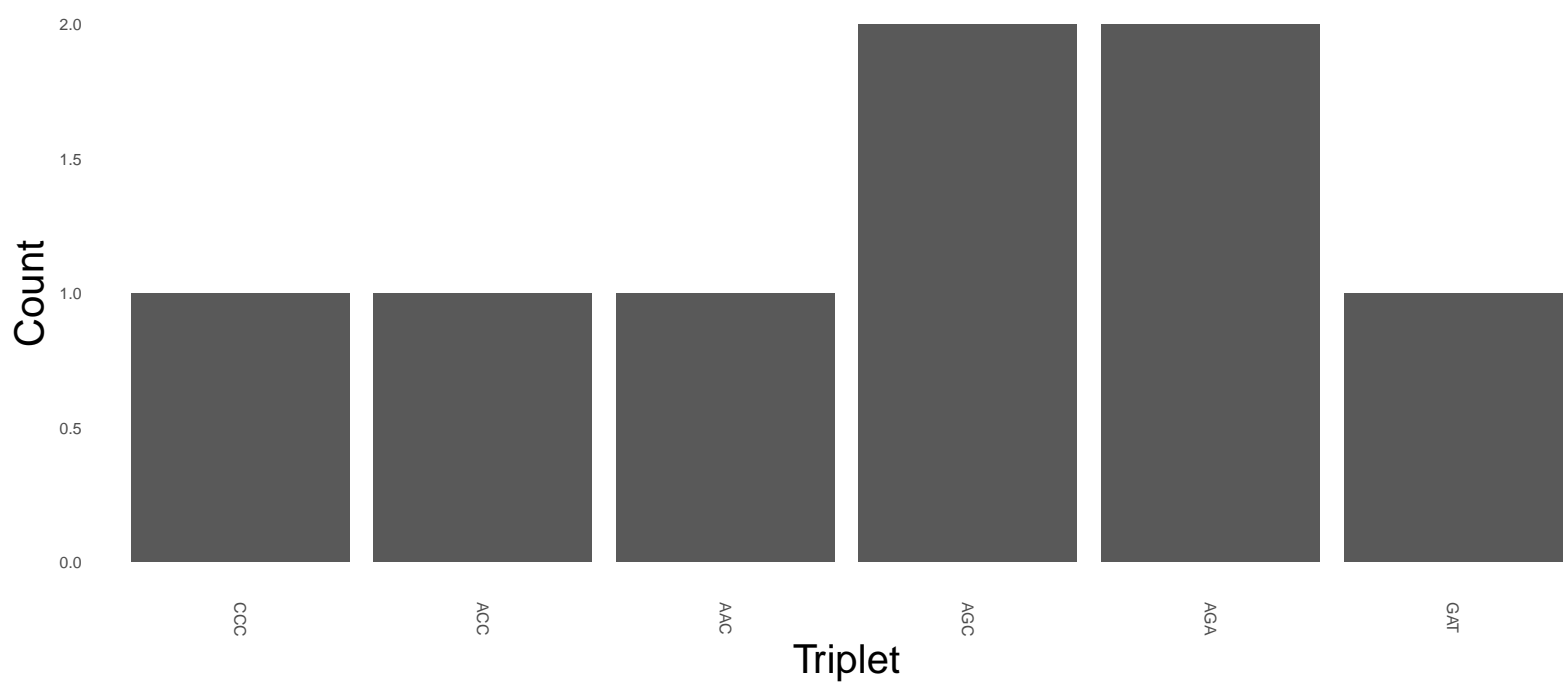
Gene TRBV4-1*01_C14T



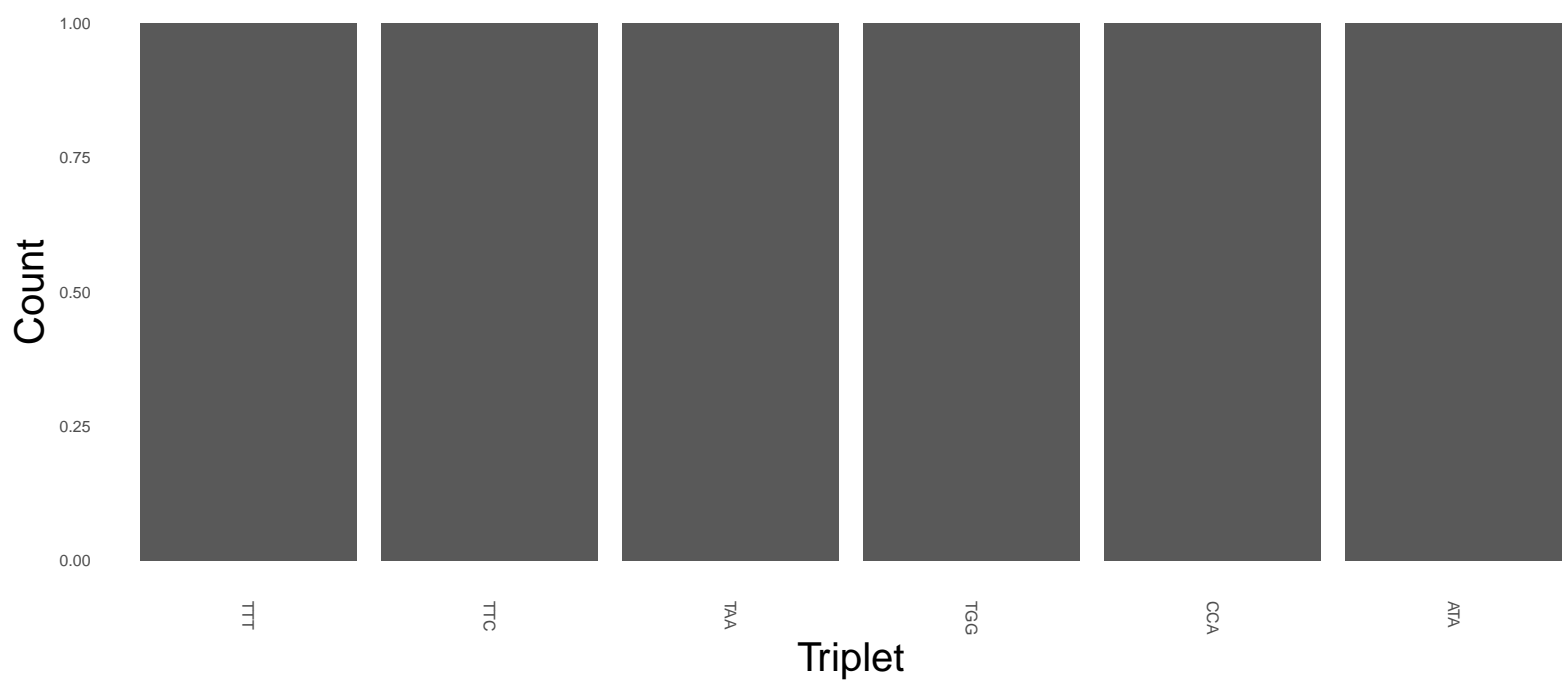
Gene TRBV5-5*02_322TTGG325



TRBV4-1*01_C14T- Final 3 nucleotides as a triplet

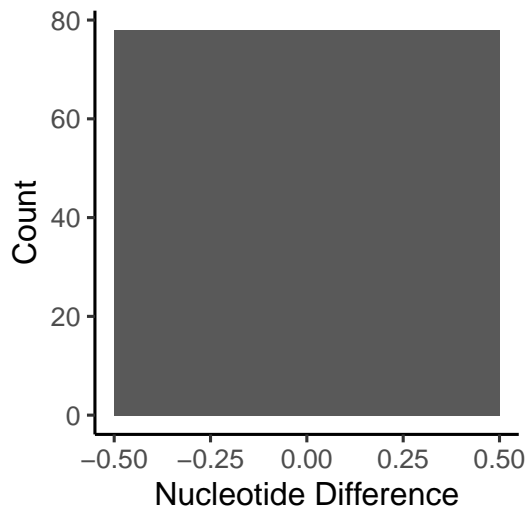


TRBV5-5*02_322TTGG325- Final 3 nucleotides as a triplet



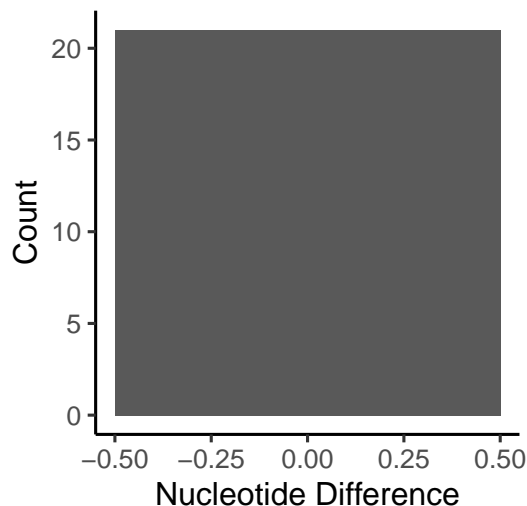
TRBV20–1*01

78 sequences assigned
78 (100%) exact matches, in which:
78 unique CDR3
12 unique J



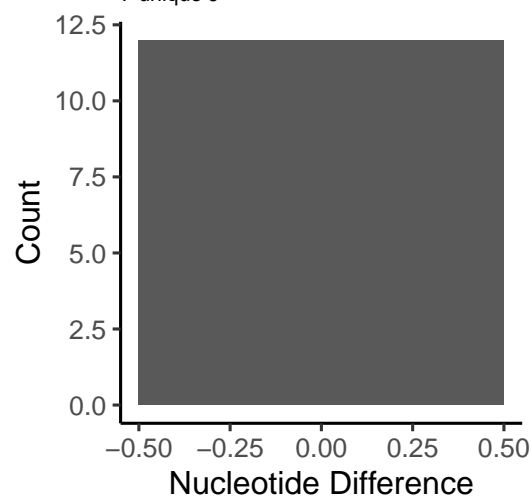
TRBV10–3*02

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



TRBV15*02

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



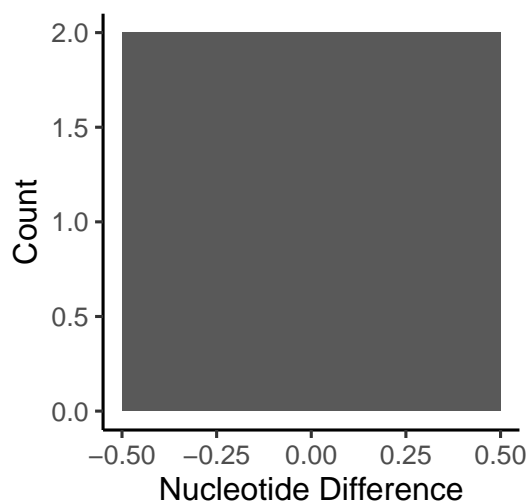
TRBV10–1*01

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



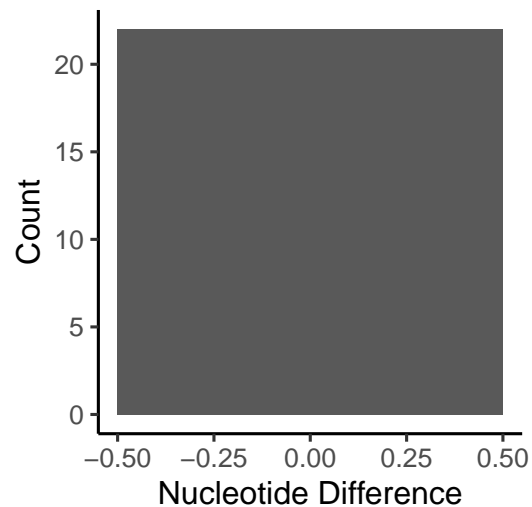
TRBV13*01

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



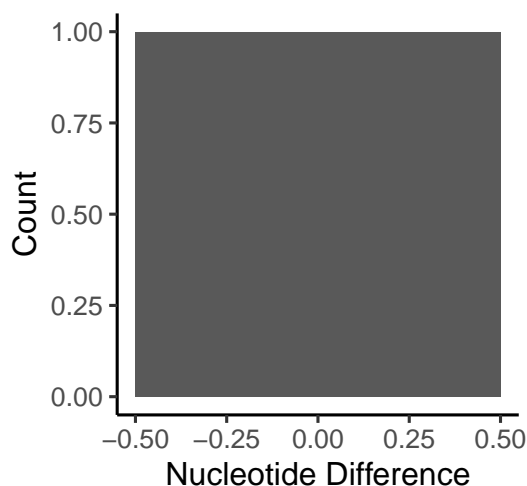
TRBV18*01

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



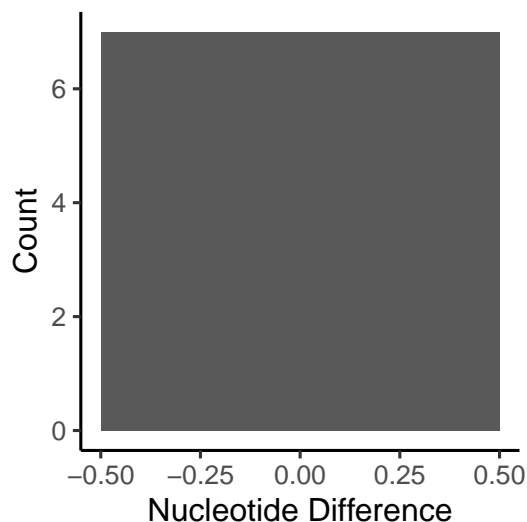
TRBV10–2*01

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



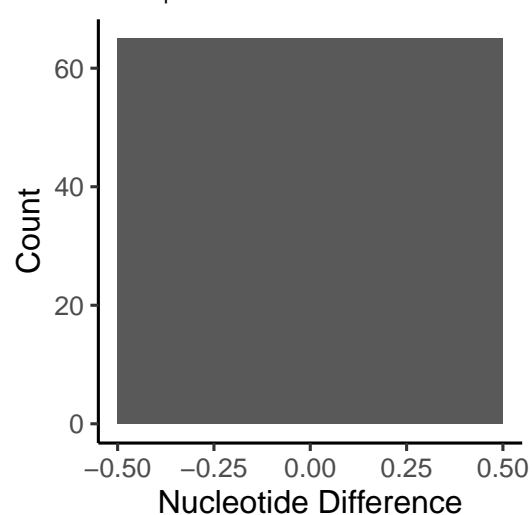
TRBV14*01

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



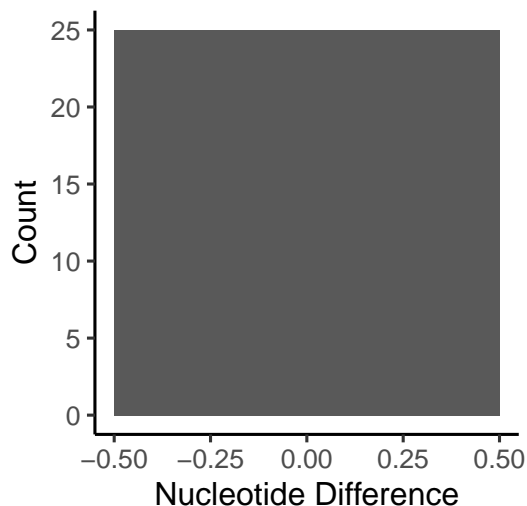
TRBV19*01

65 sequences assigned
65 (100%) exact matches, in which:
65 unique CDR3
11 unique J



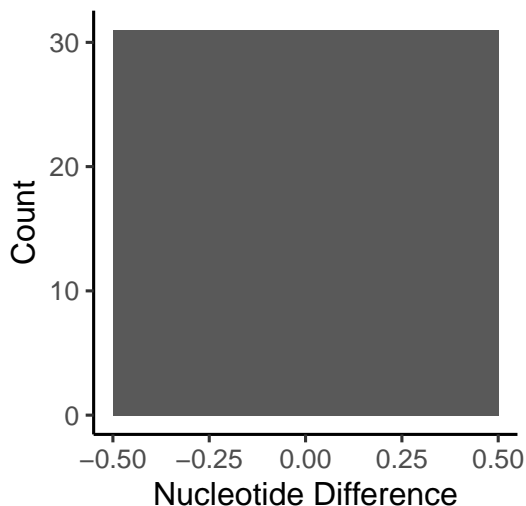
TRBV2*01

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



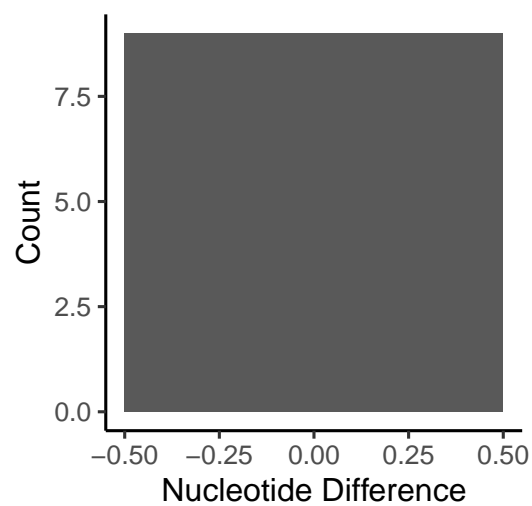
TRBV30*01

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



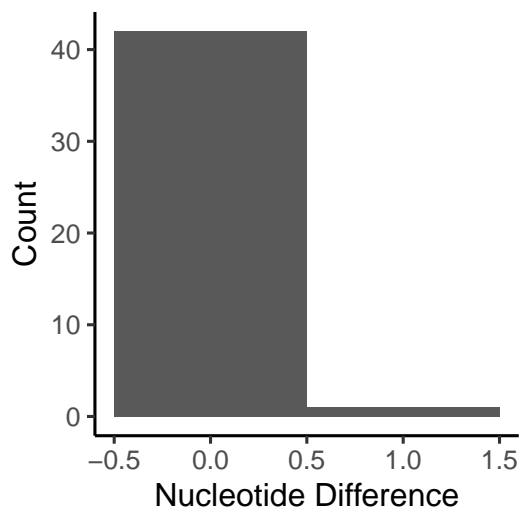
TRBV11-2*01

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



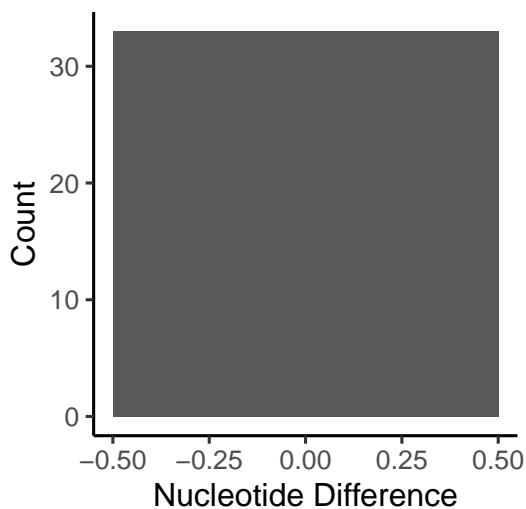
TRBV27*01

43 sequences assigned
42 (97.7%) exact matches, in which:
42 unique CDR3
12 unique J



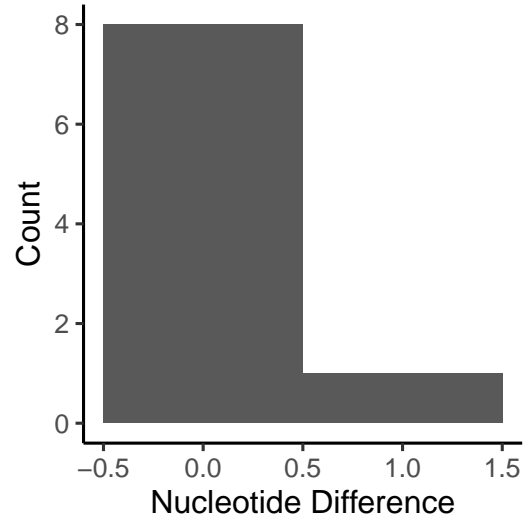
TRBV9*01

33 sequences assigned
33 (100%) exact matches, in which:
33 unique CDR3
11 unique J



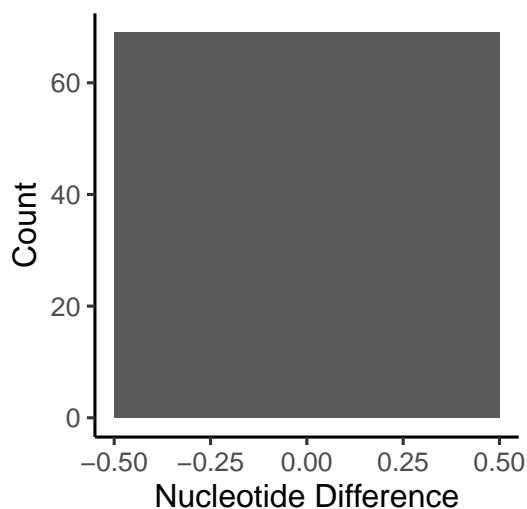
TRBV11-2*03

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



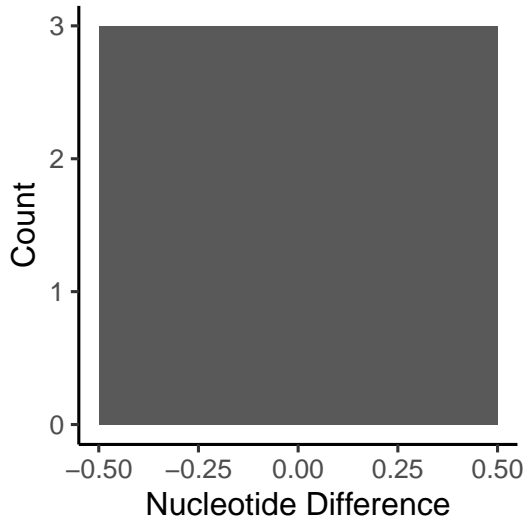
TRBV28*01

69 sequences assigned
69 (100%) exact matches, in which:
69 unique CDR3
12 unique J



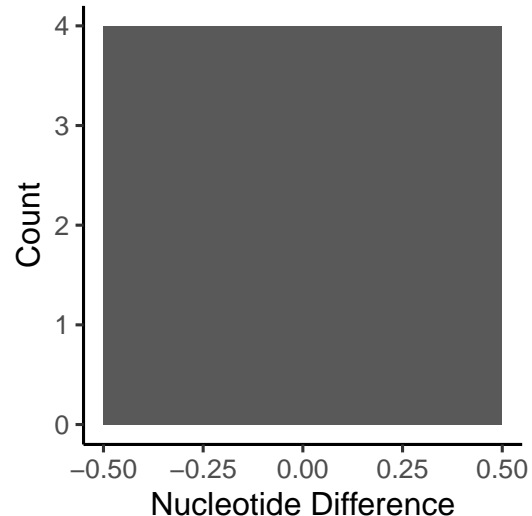
TRBV11-1*01

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



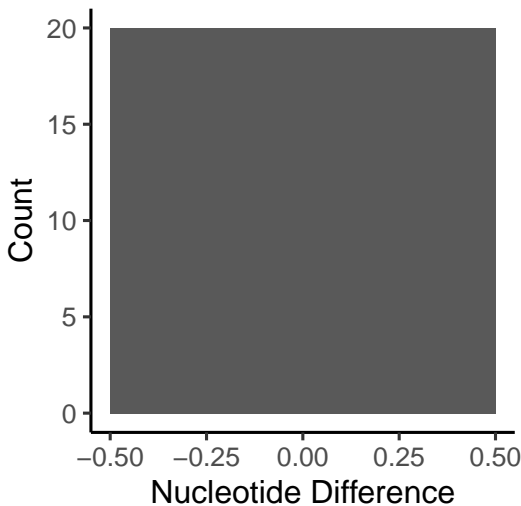
TRBV11-3*01

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



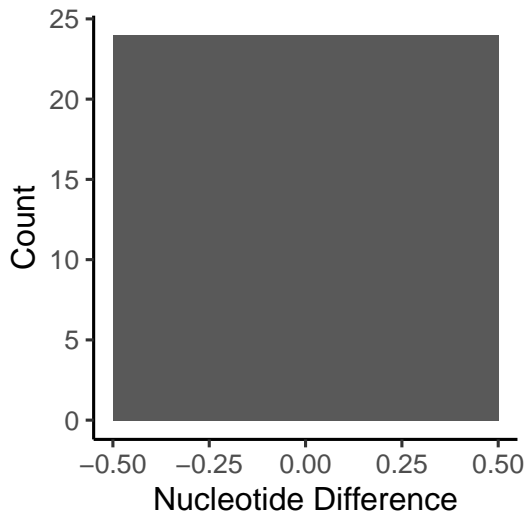
TRBV12–3*01

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



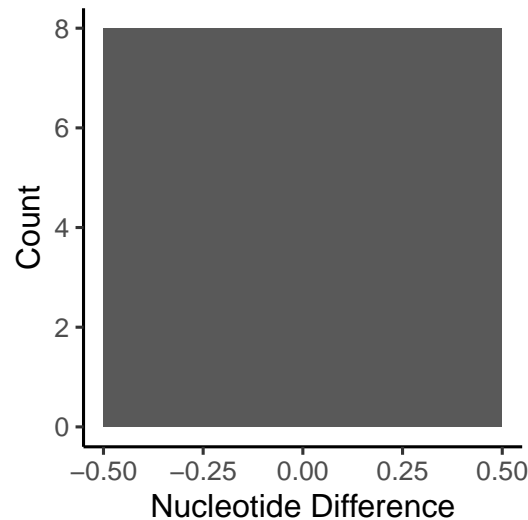
TRBV3–1*01

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



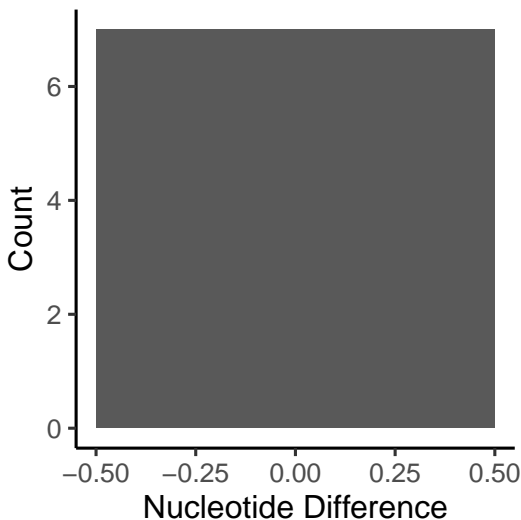
TRBV4–1*01_C14T

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



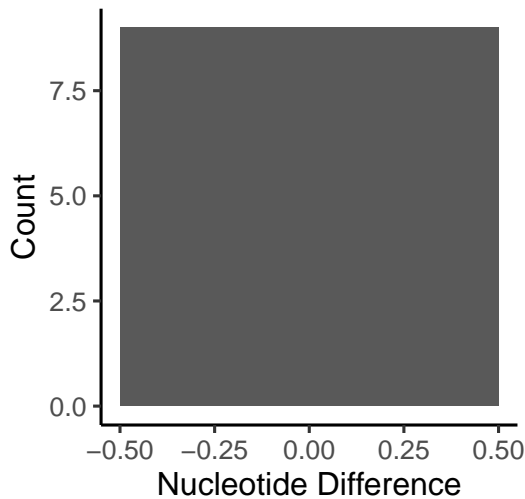
TRBV12–4*01

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



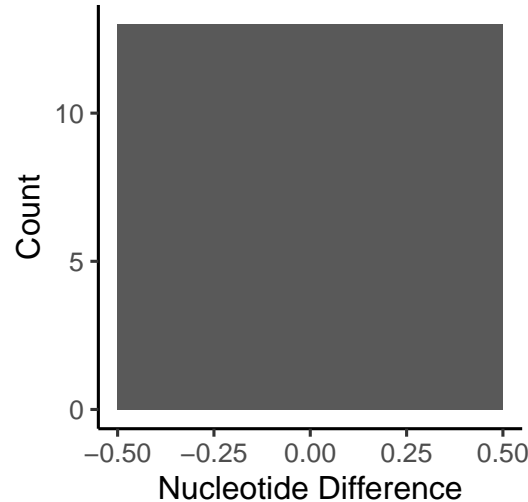
TRBV4–1*01

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



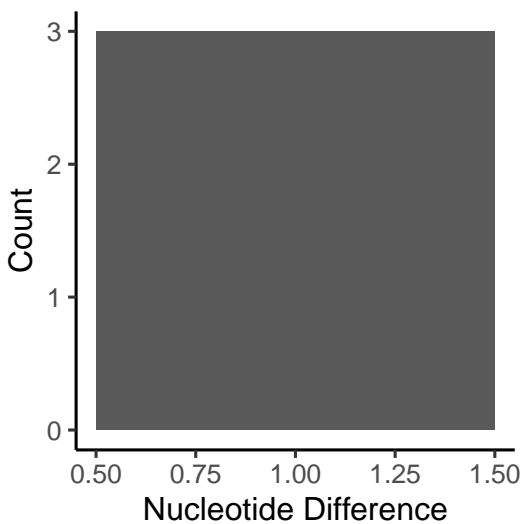
TRBV4–2*01

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



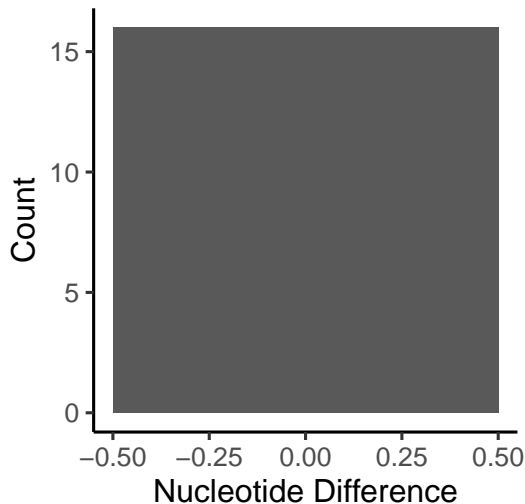
TRBV12–5*01

3 sequences assigned
No exact matches.



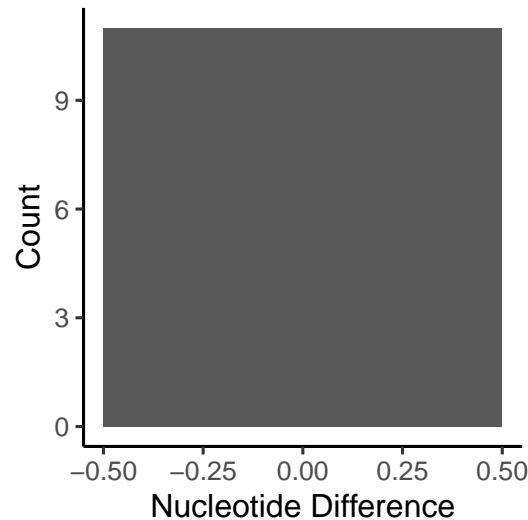
TRBV24–1*01

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



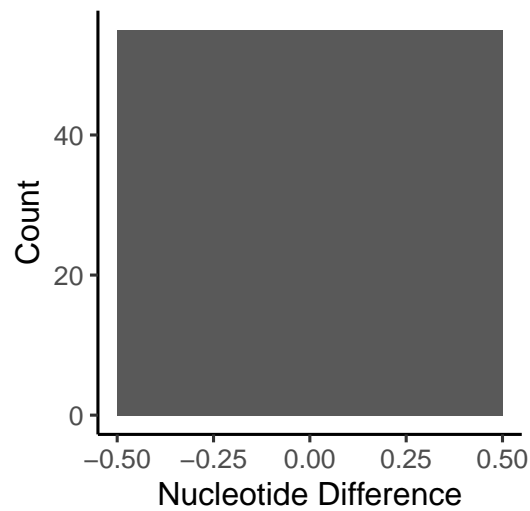
TRBV4–3*01

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



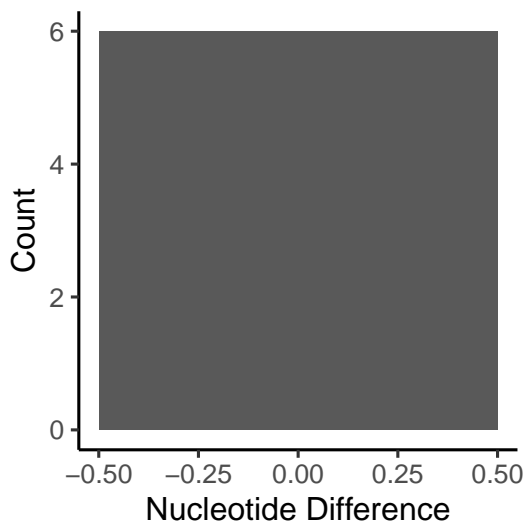
TRBV5–1*01

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



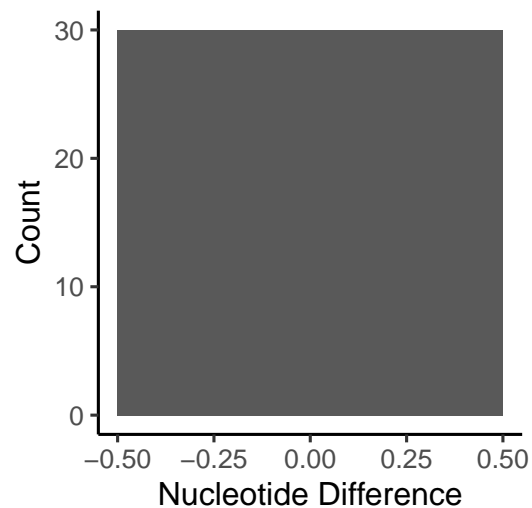
TRBV5–5*02_322TTGG32!

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



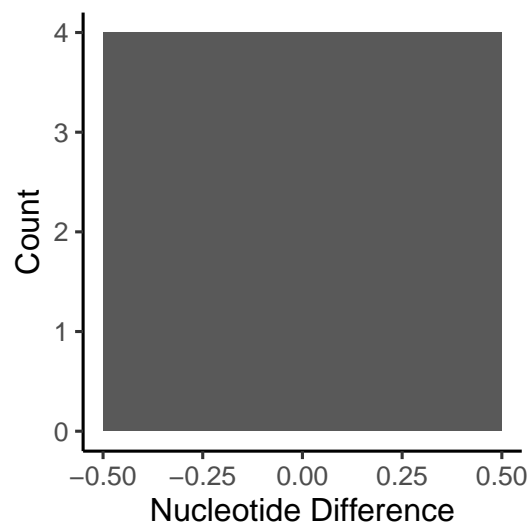
TRBV6–1*01

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



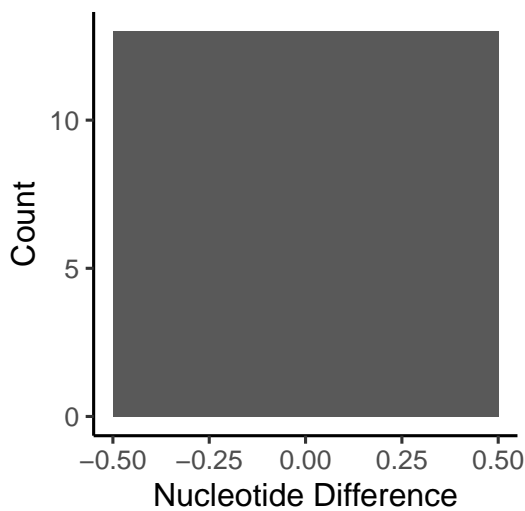
TRBV25–1*01

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



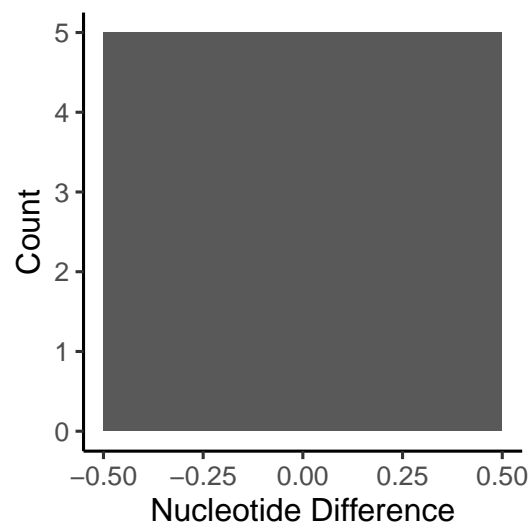
TRBV5–6*01

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



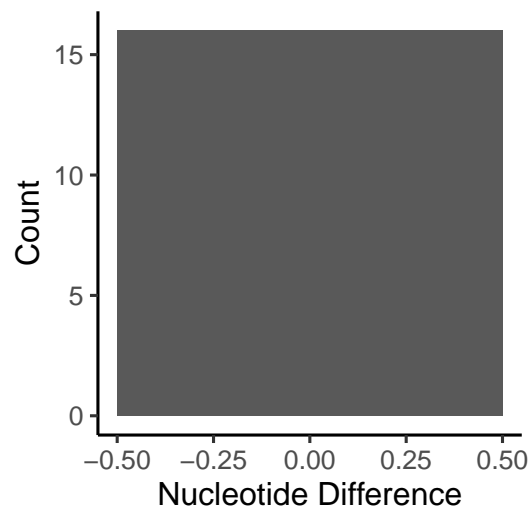
TRBV6–4*01

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



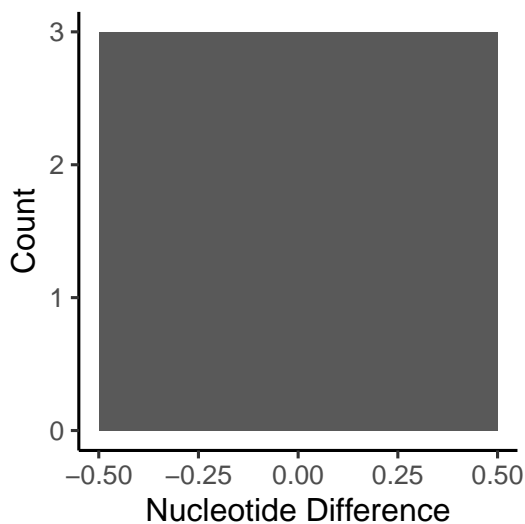
TRBV5–4*01

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



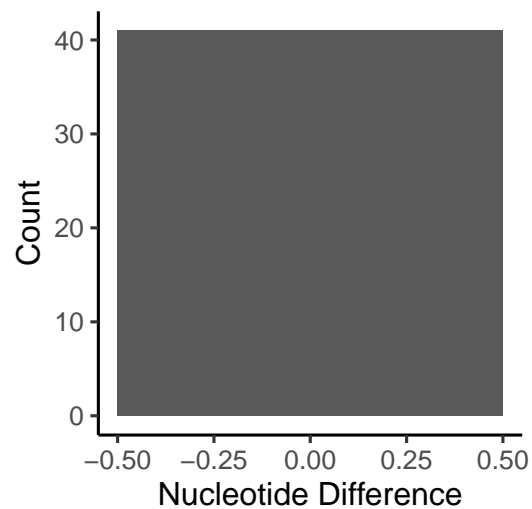
TRBV5–8*01

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



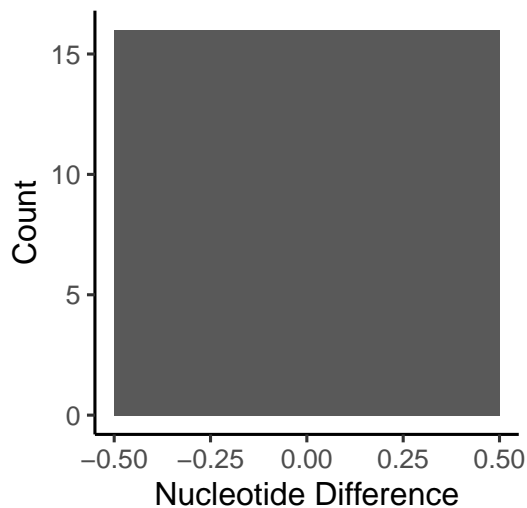
TRBV6–5*01

41 sequences assigned
41 (100%) exact matches, in which:
41 unique CDR3
11 unique J



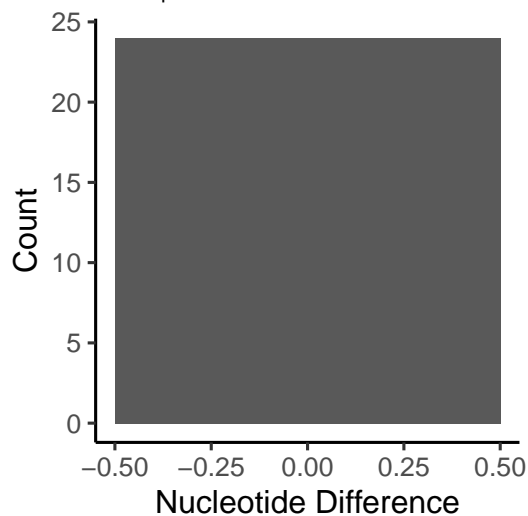
TRBV6-6*01

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



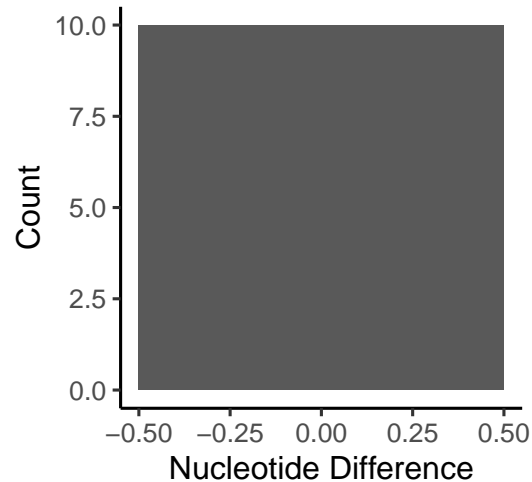
TRBV7-2*01

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



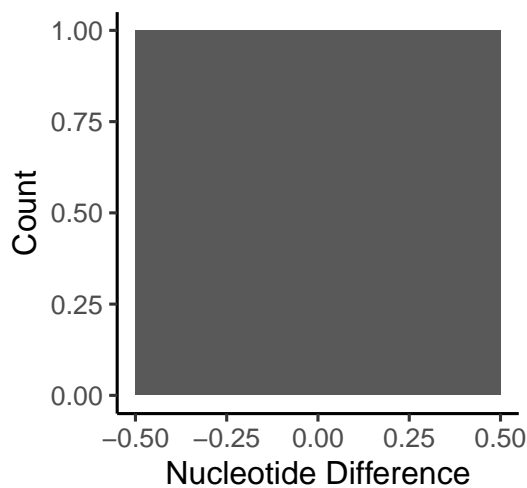
TRBV7-6*01

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



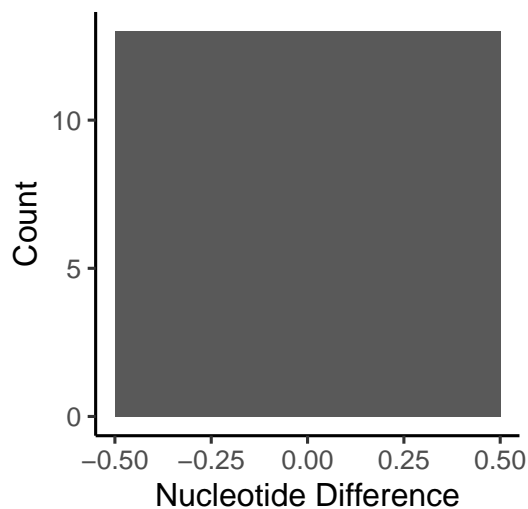
TRBV6-7*01

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



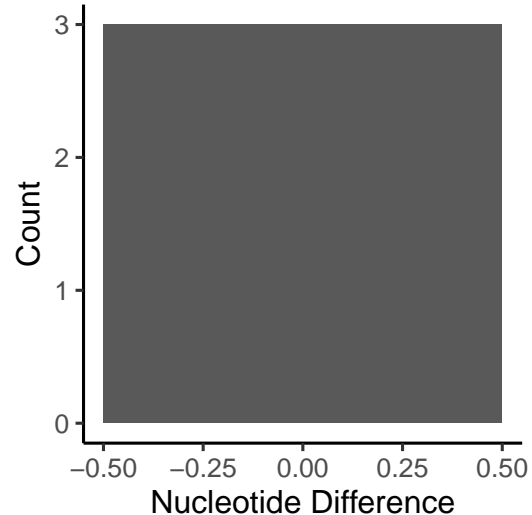
TRBV7-2*02

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



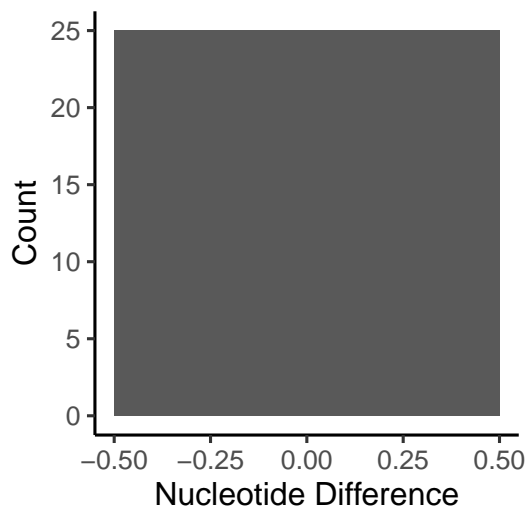
TRBV7-7*01

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



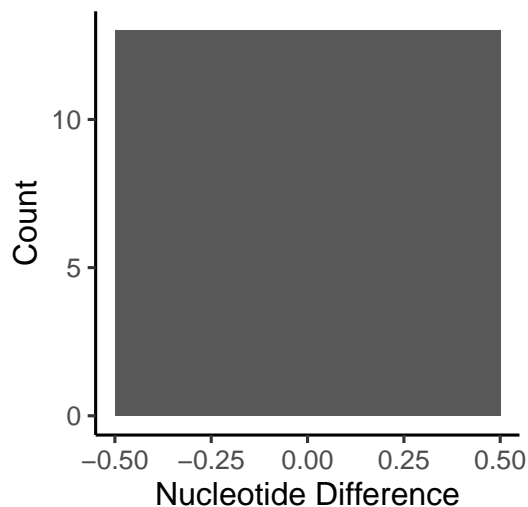
TRBV6-23*01

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



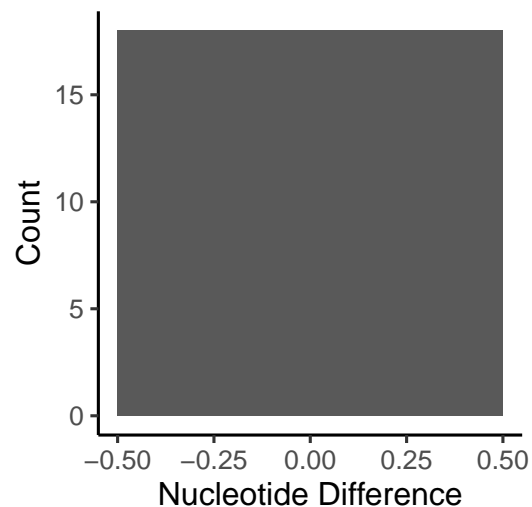
TRBV7-3*01

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



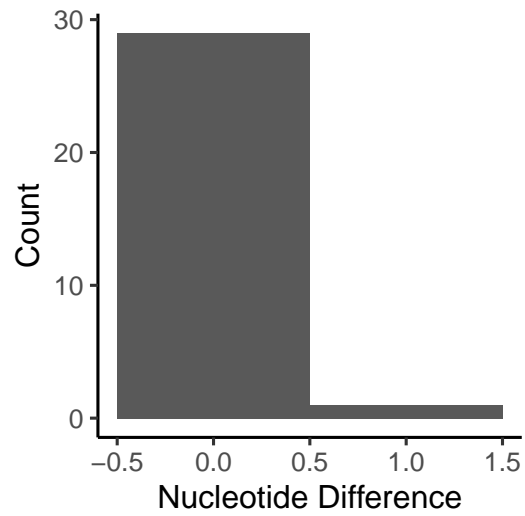
TRBV7-8*01

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



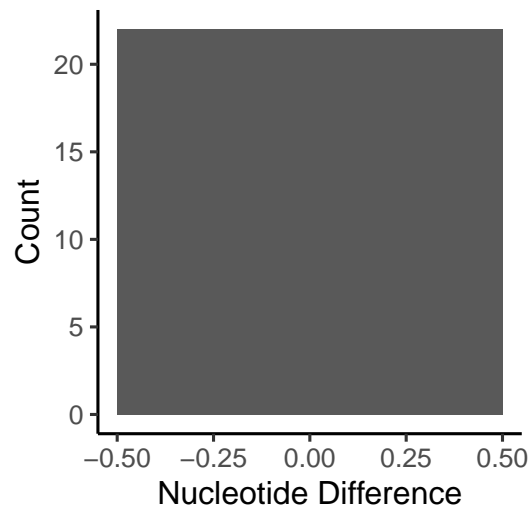
TRBV7-9*03

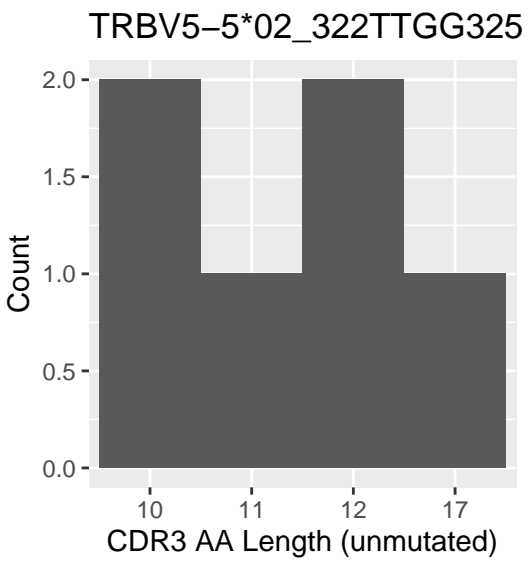
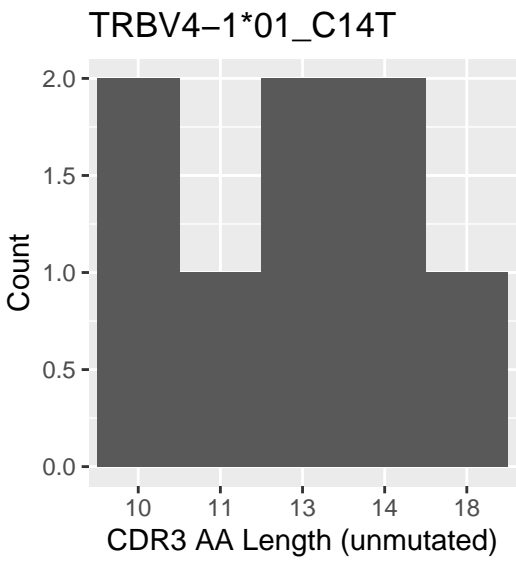
30 sequences assigned
29 (96.7%) exact matches, in which:
29 unique CDR3
10 unique J



TRBV29-1*01

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







Warning – no inferred sequences found.

Warning – no inferred sequences found.