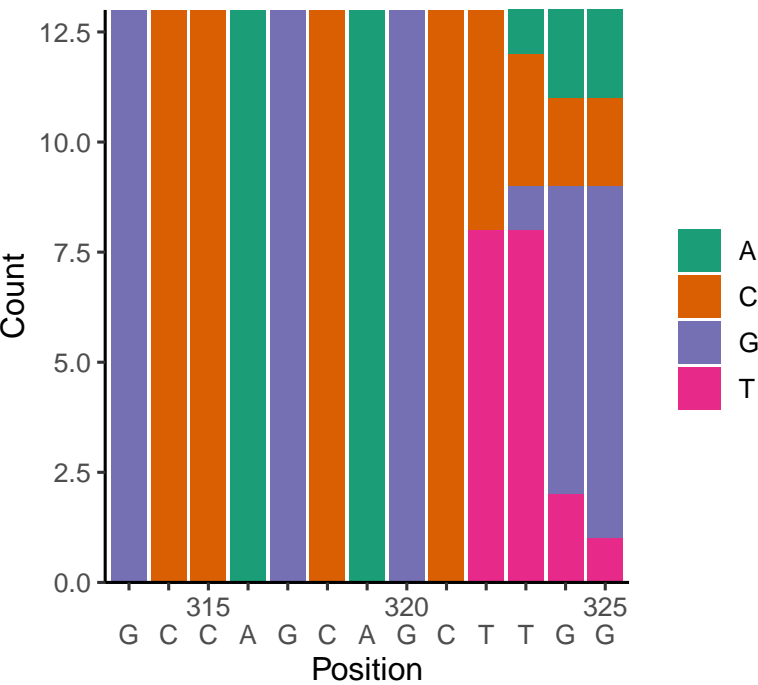
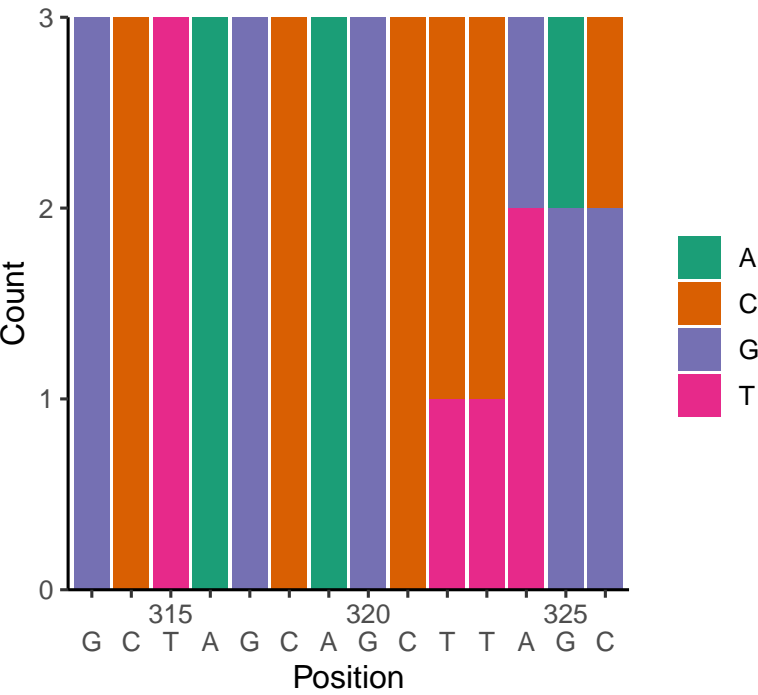


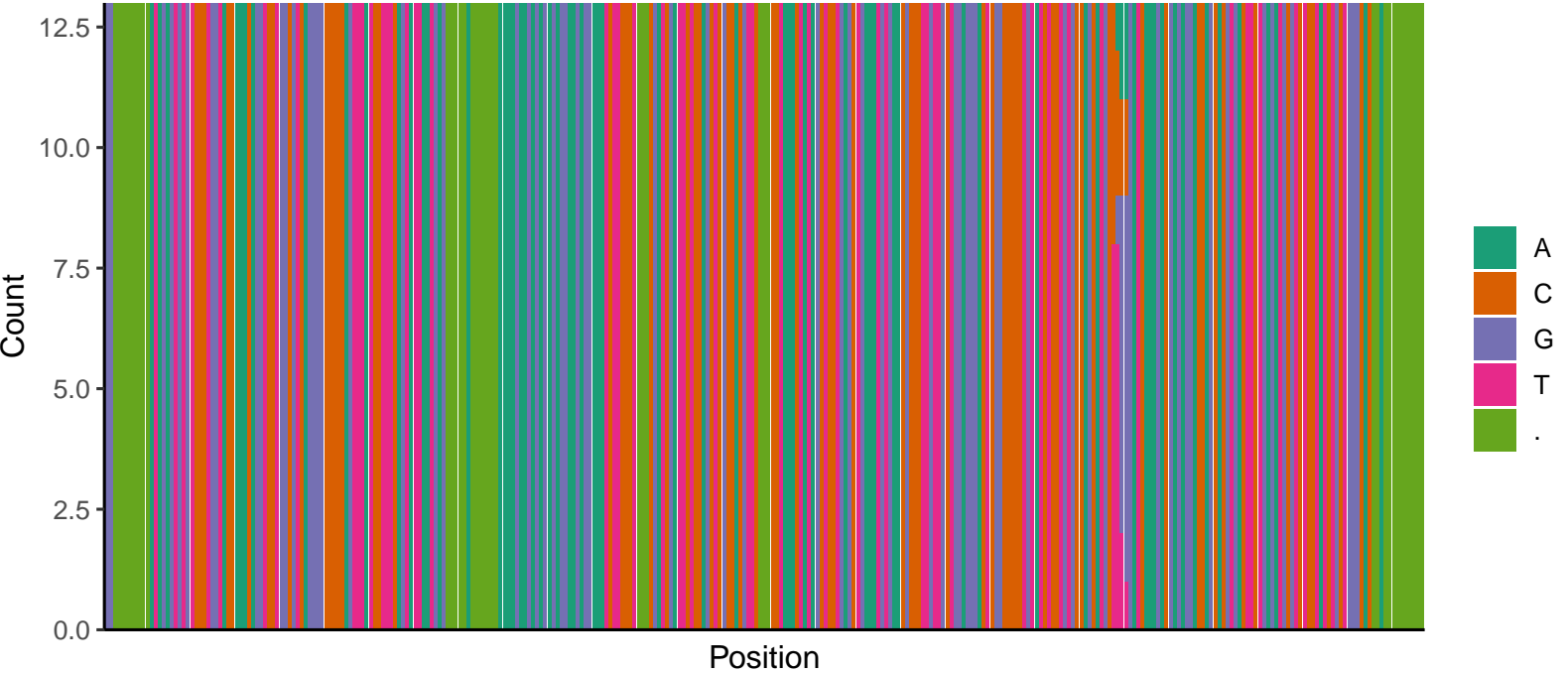
Gene TRBV5-5*02_322TTGG325



Gene TRBV7-7*01_C315T



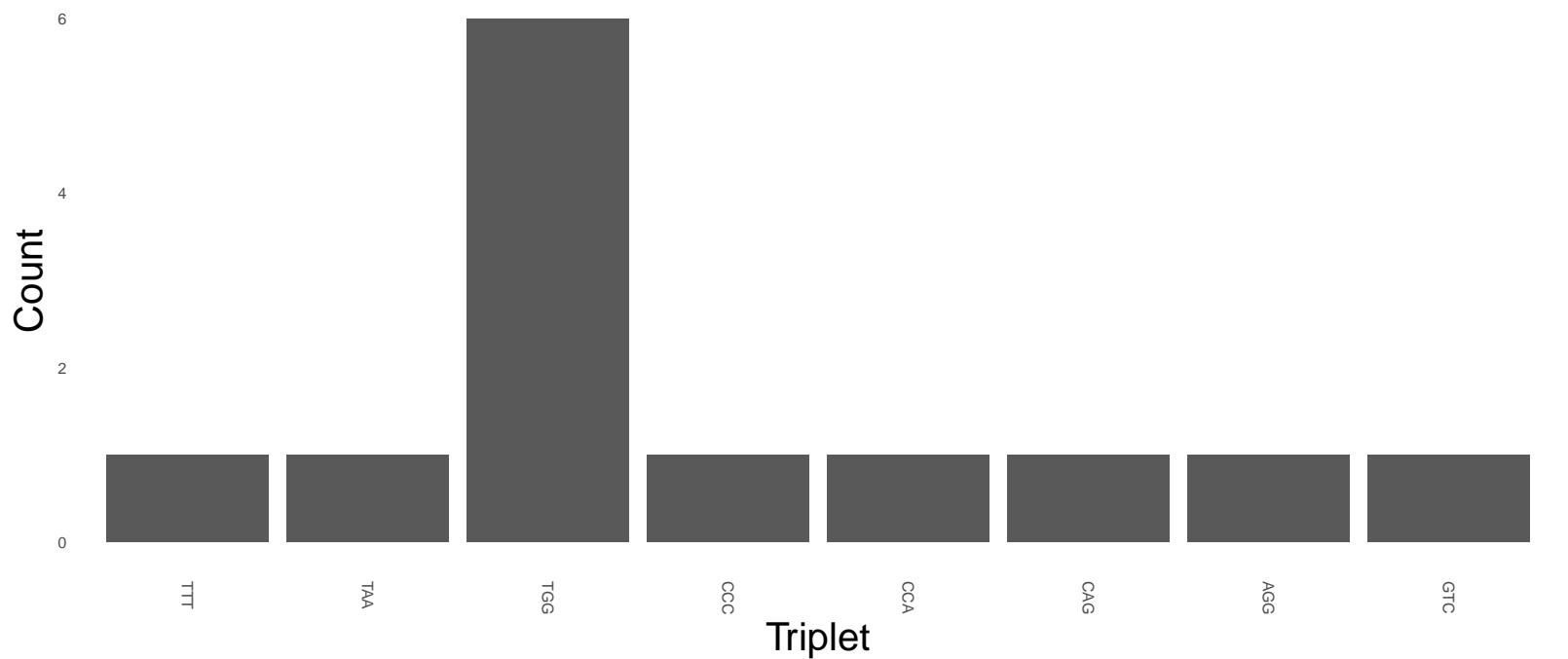
Gene TRBV5-5*02_322TTGG325



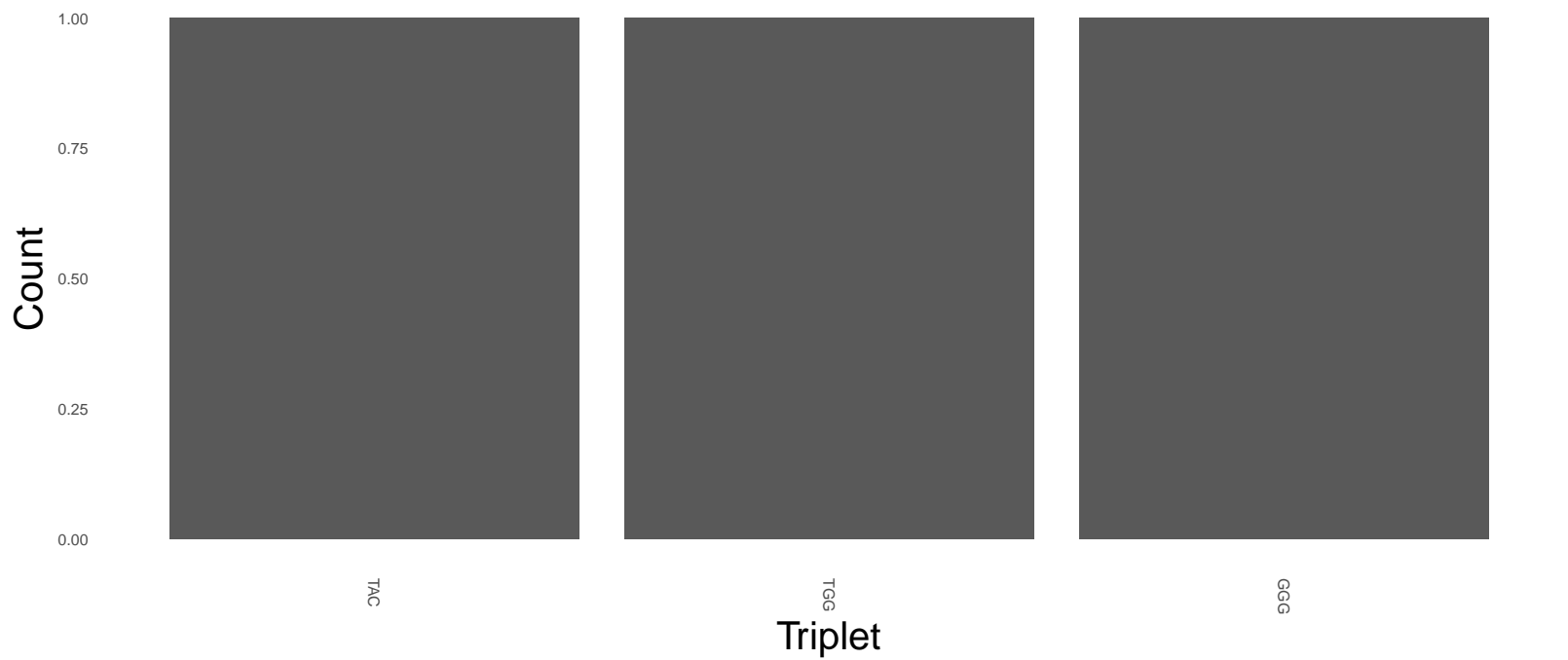
Gene TRBV7-7*01_C315T



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

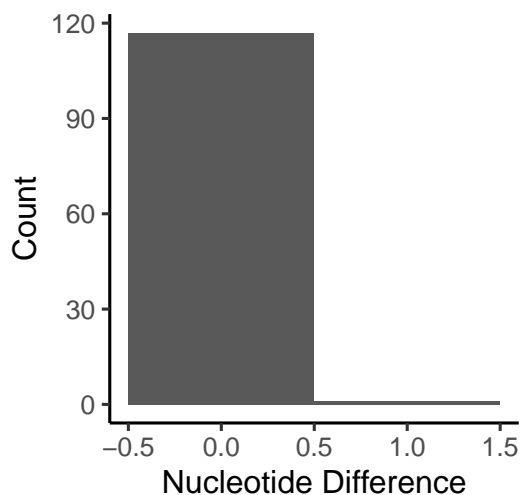


TRBV7-7*01_C315T- Final 3 nucleotides as a triplet



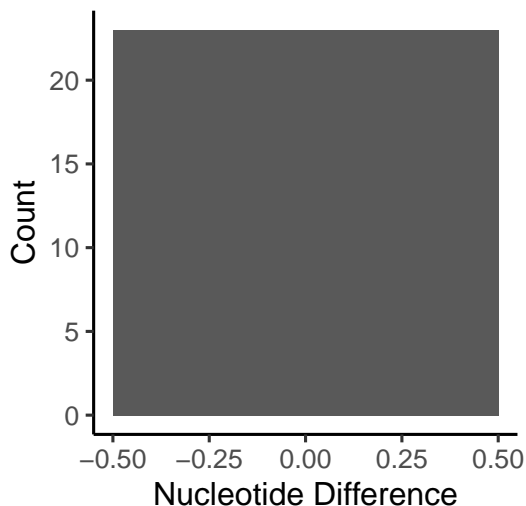
TRBV20–1*01

118 sequences assigned
117 (99.2%) exact matches, in which:
117 unique CDR3
13 unique J



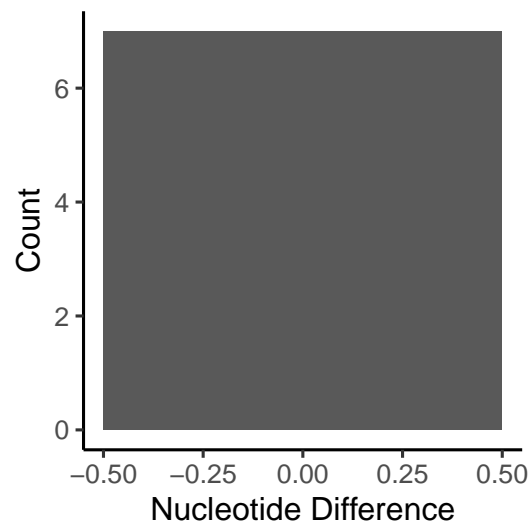
TRBV10–3*02

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



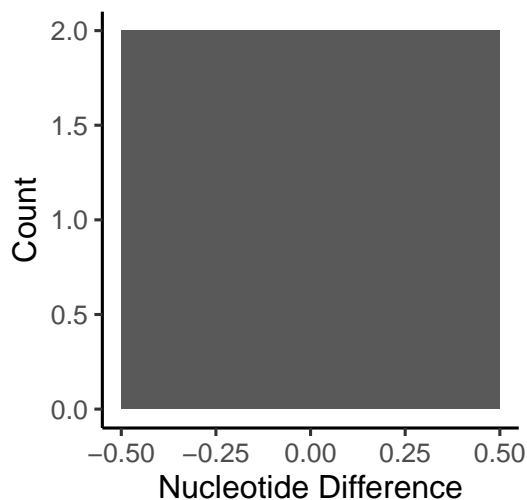
TRBV15*02

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



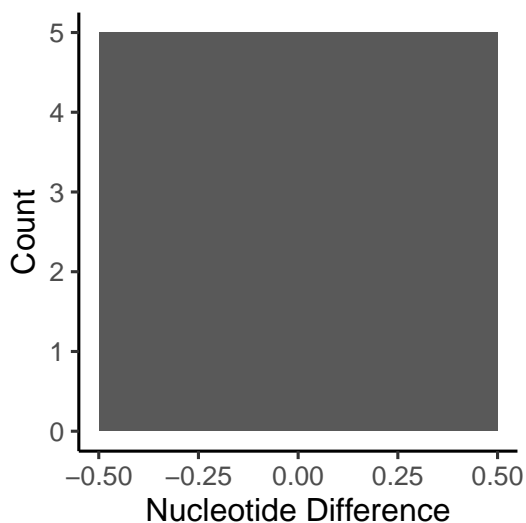
TRBV10–1*01

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



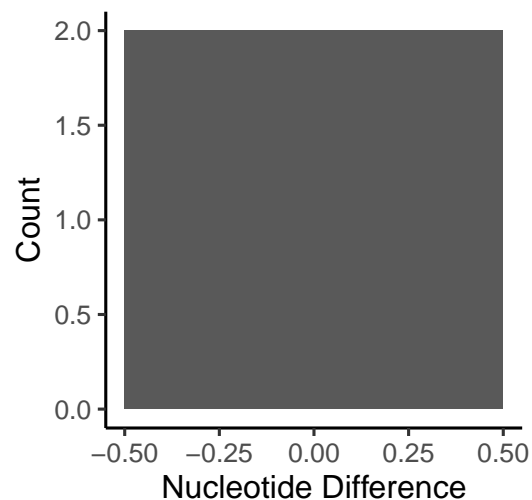
TRBV13*01

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



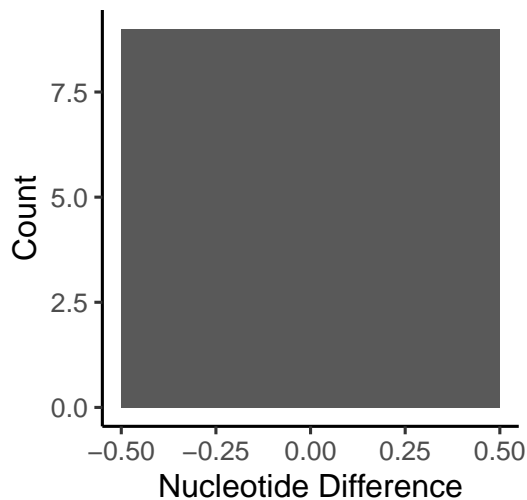
TRBV16*01

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



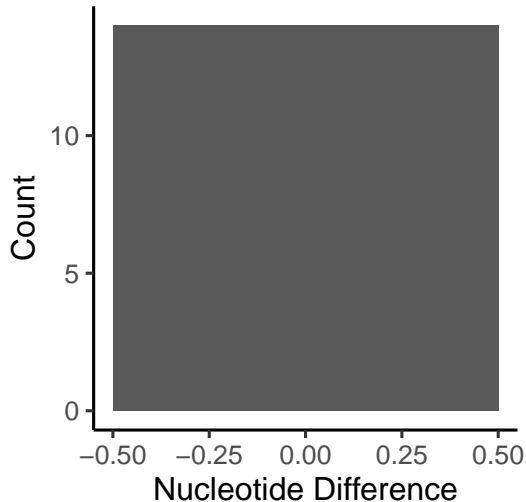
TRBV10–2*01

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



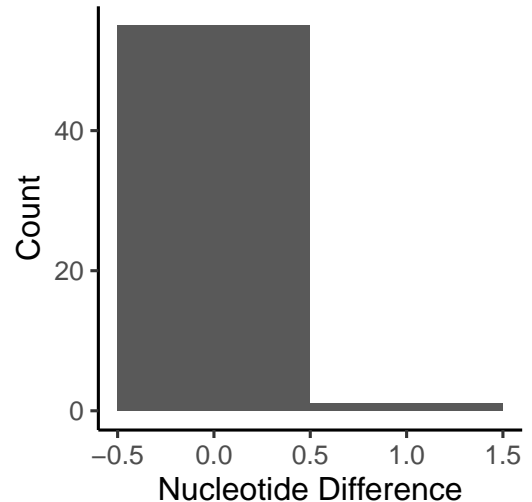
TRBV14*01

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



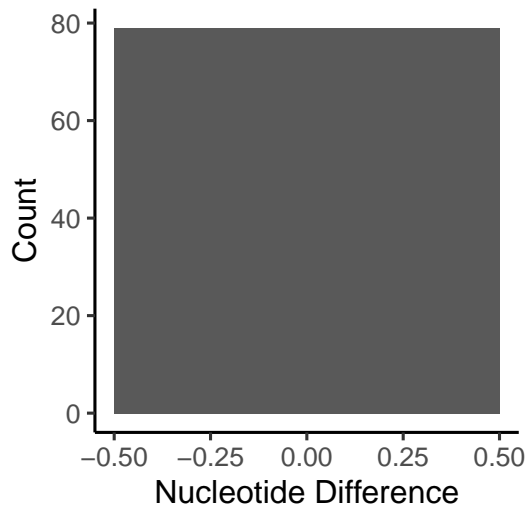
TRBV18*01

56 sequences assigned
55 (98.2%) exact matches, in which:
55 unique CDR3
12 unique J



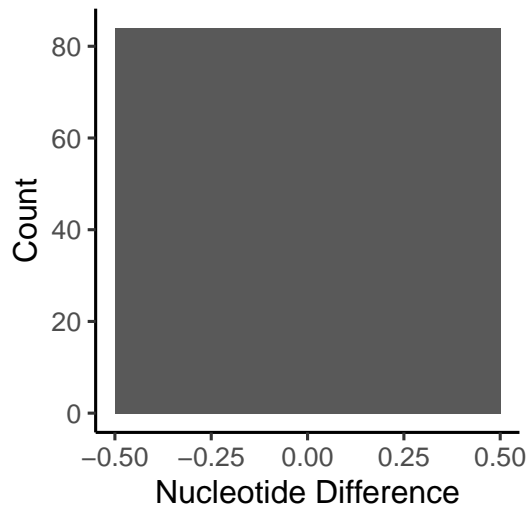
TRBV19*01

79 sequences assigned
79 (100%) exact matches, in which:
79 unique CDR3
11 unique J



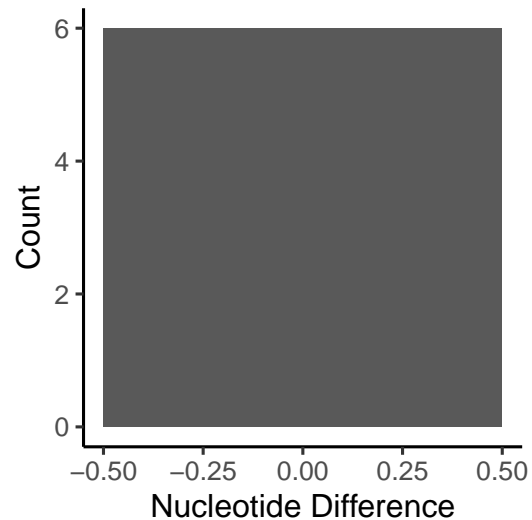
TRBV28*01

84 sequences assigned
84 (100%) exact matches, in which:
84 unique CDR3
13 unique J



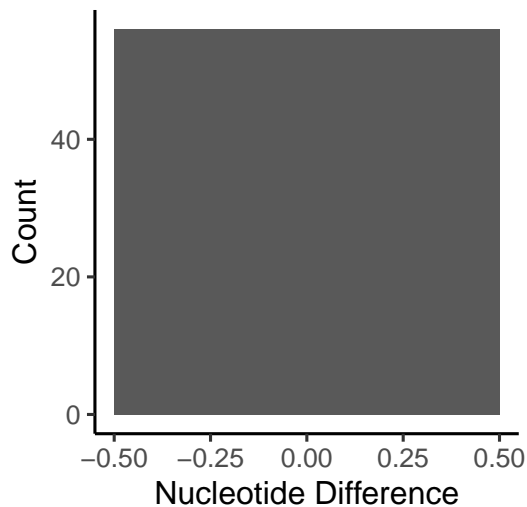
TRBV11-1*01

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



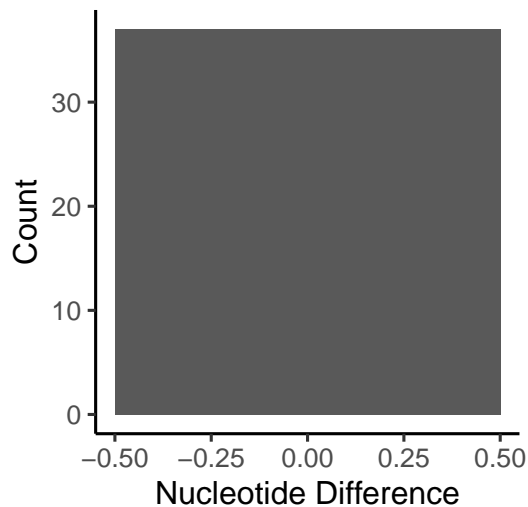
TRBV2*01

56 sequences assigned
56 (100%) exact matches, in which:
56 unique CDR3
10 unique J



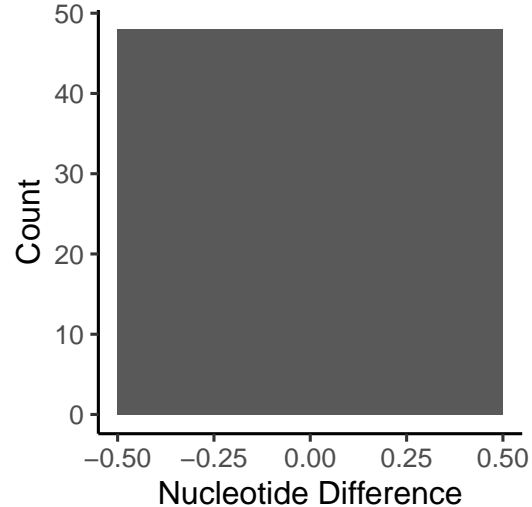
TRBV30*01

37 sequences assigned
37 (100%) exact matches, in which:
37 unique CDR3
11 unique J



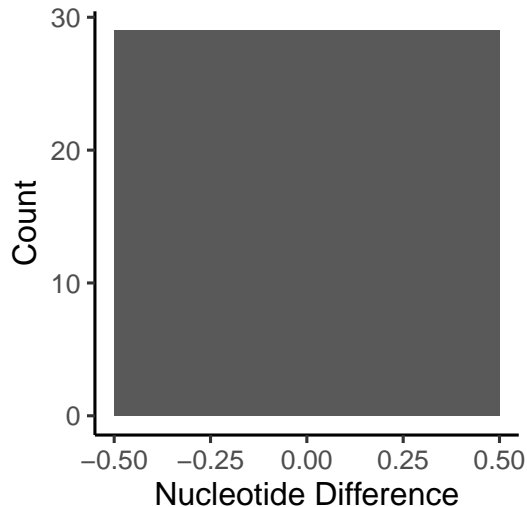
TRBV11-2*01

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



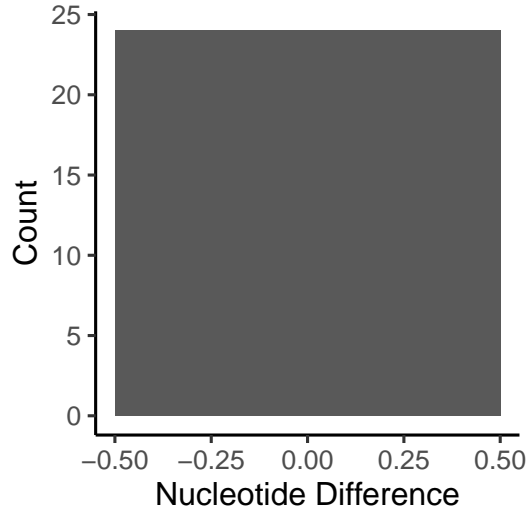
TRBV27*01

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



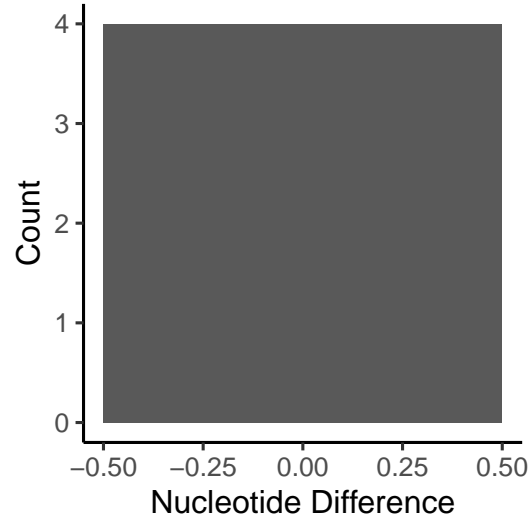
TRBV9*01

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



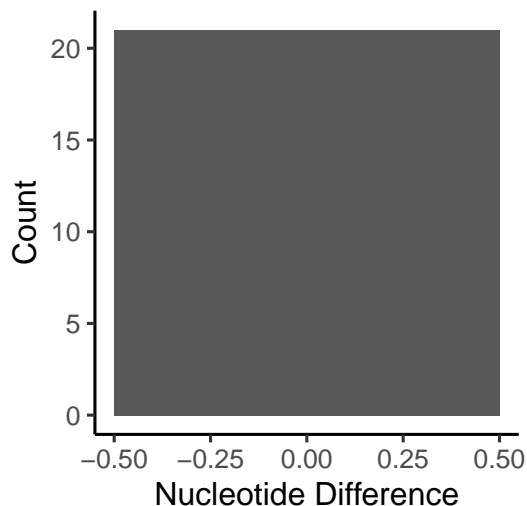
TRBV11-3*01

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



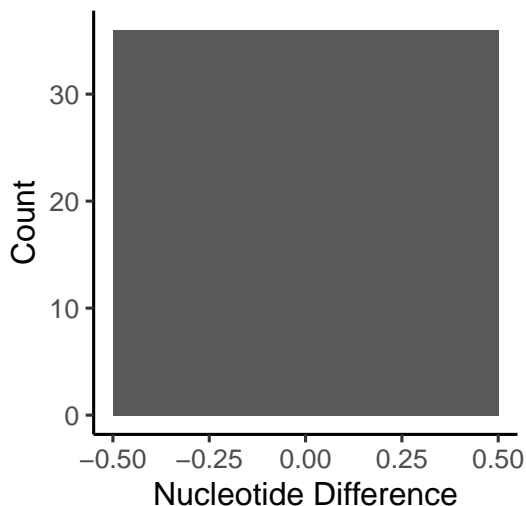
TRBV12-3*01

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



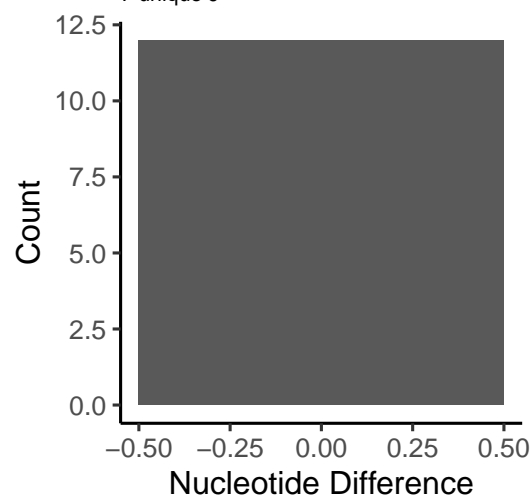
TRBV3-1*01

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



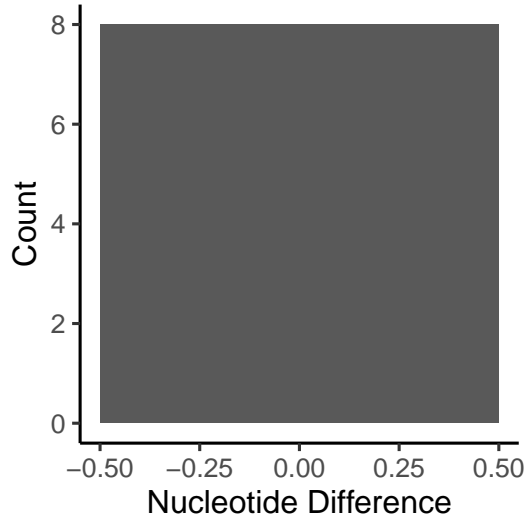
TRBV4-2*01

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



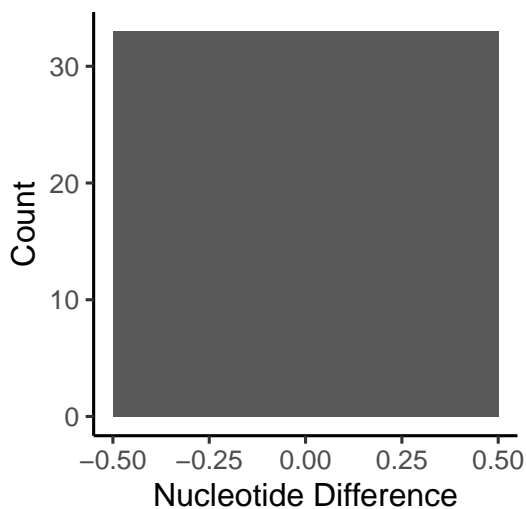
TRBV12-4*01

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



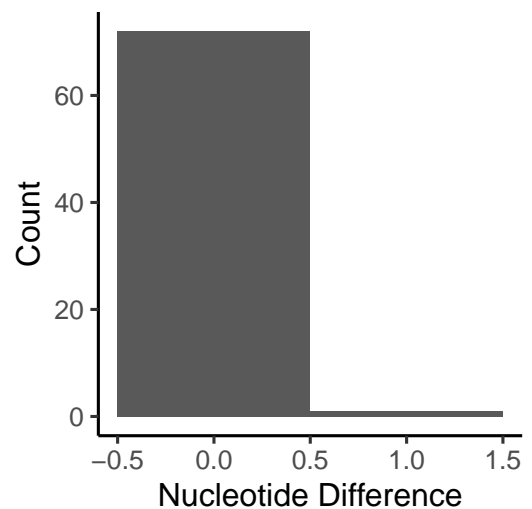
TRBV4-1*01

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



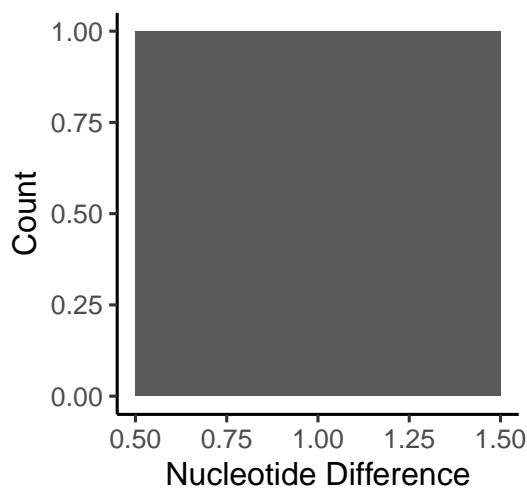
TRBV5-1*01

73 sequences assigned
72 (98.6%) exact matches, in which:
72 unique CDR3
10 unique J



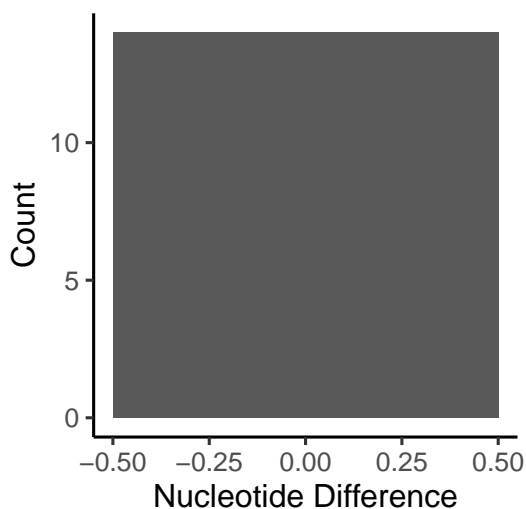
TRBV12-5*01

1 sequences assigned
No exact matches.



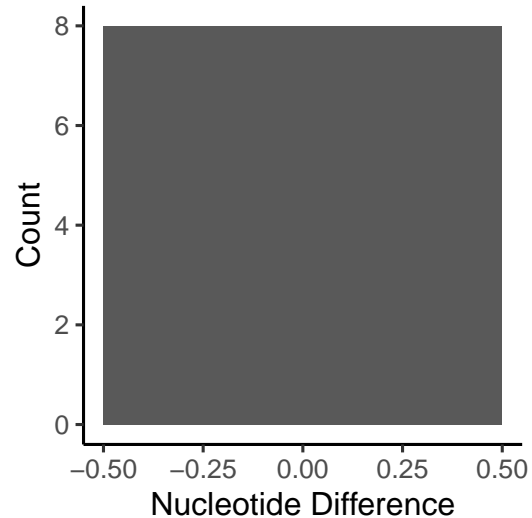
TRBV24-1*01

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



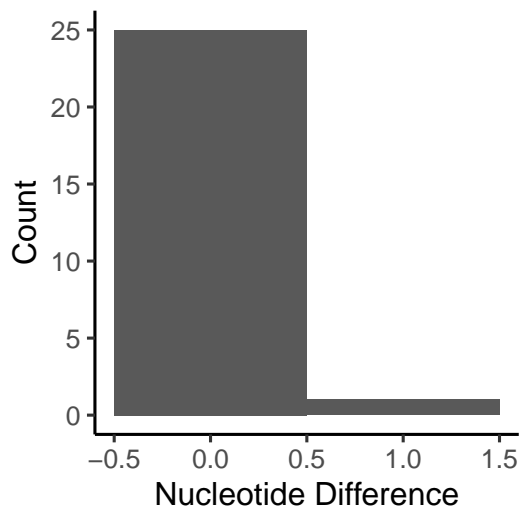
TRBV25-1*01

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



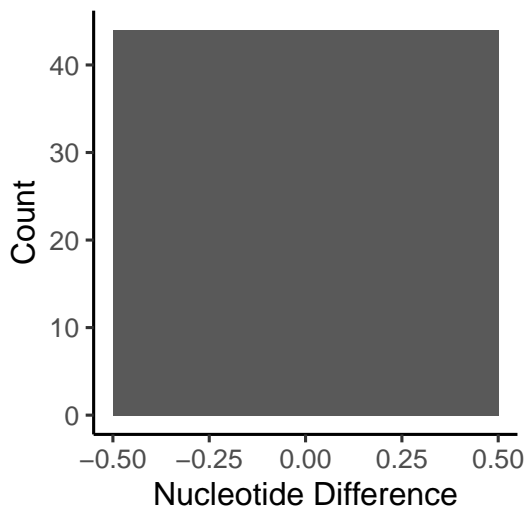
TRBV5-4*01

26 sequences assigned
25 (96.2%) exact matches, in which:
25 unique CDR3
6 unique J



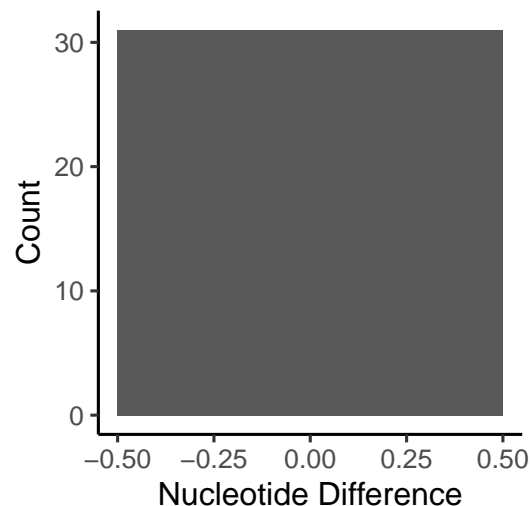
TRBV6-1*01

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



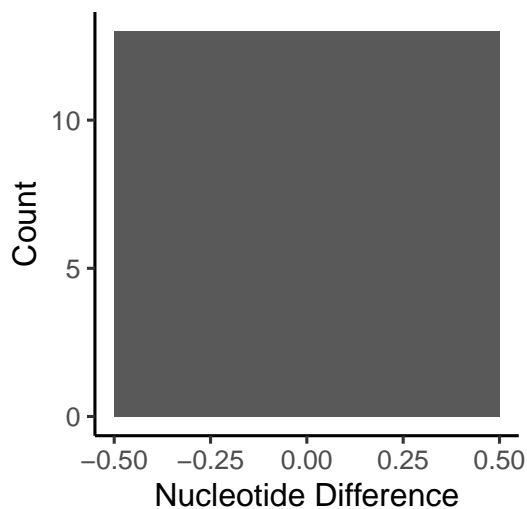
TRBV6-6*01

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



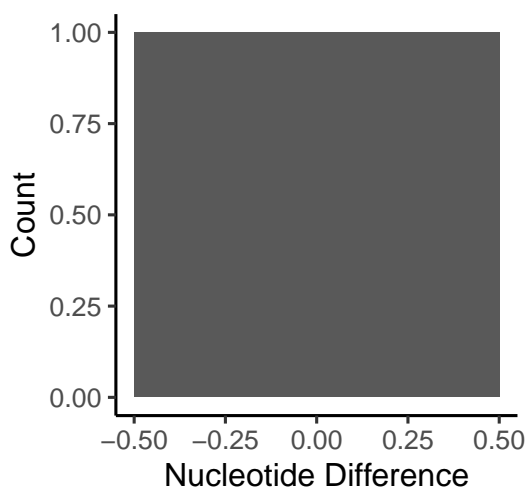
TRBV5-5*02_322TTGG32

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



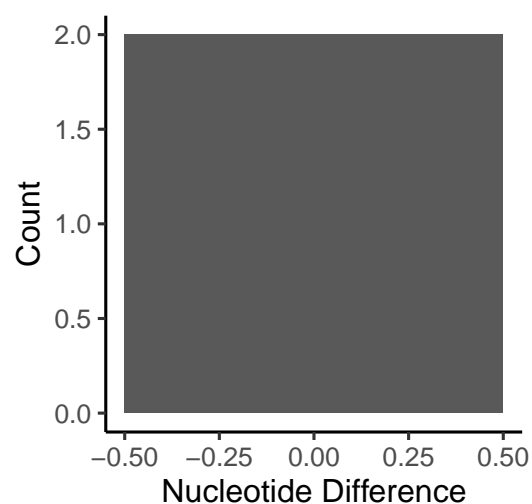
TRBV6-4*01

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



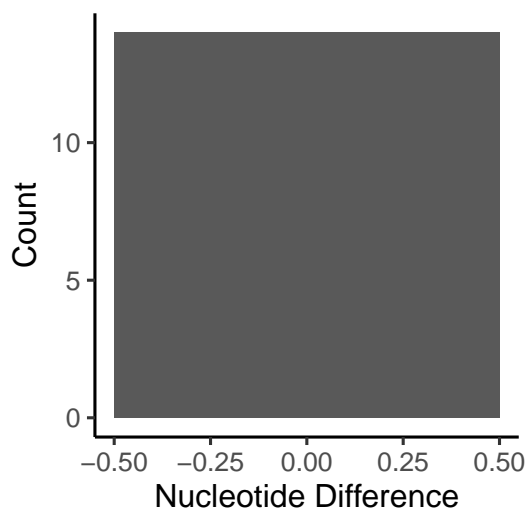
TRBV6-8*01

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



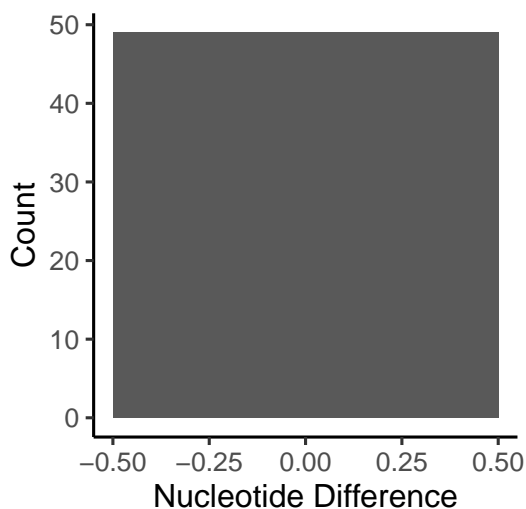
TRBV5-6*01

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



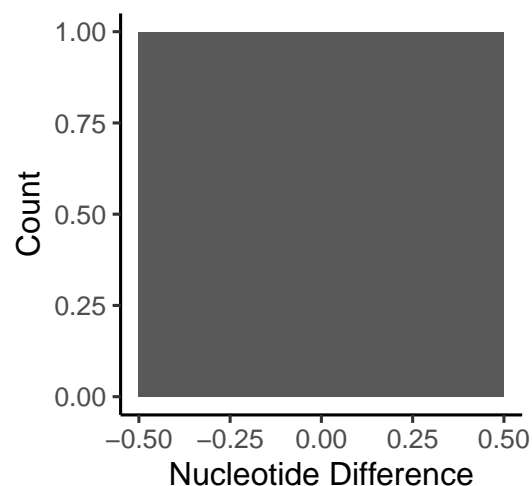
TRBV6-5*01

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



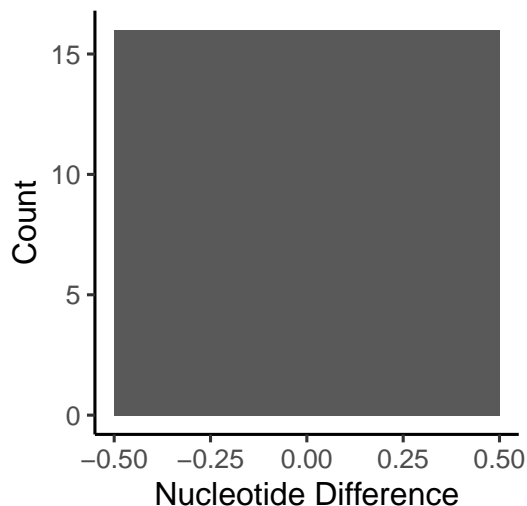
TRBV6-9*01

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



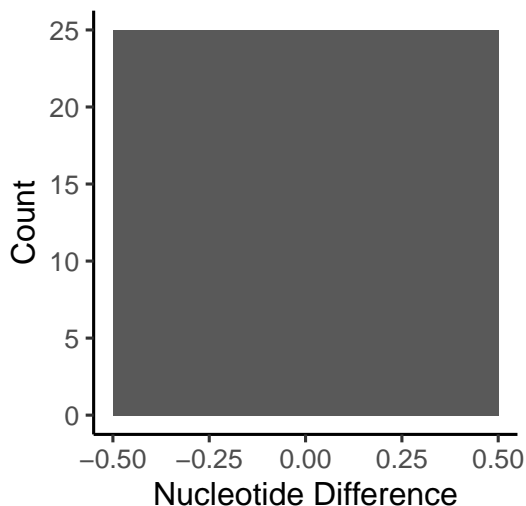
TRBV6–23*01

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



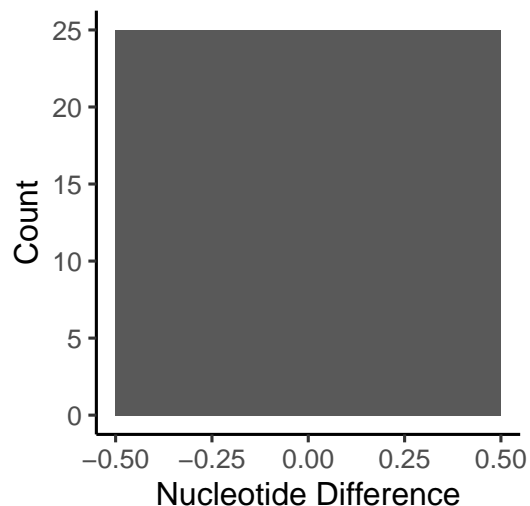
TRBV7–3*01

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



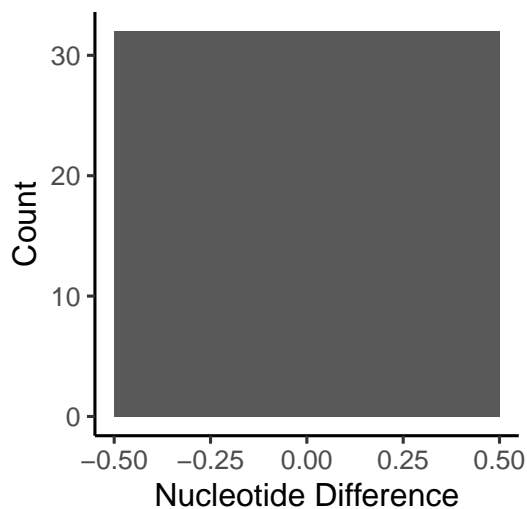
TRBV7–8*01

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



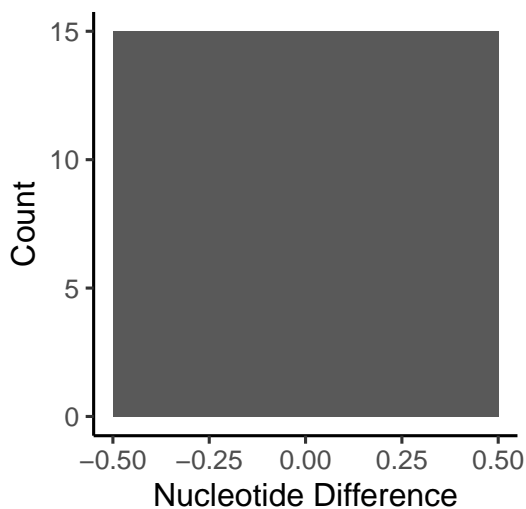
TRBV7–2*01

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



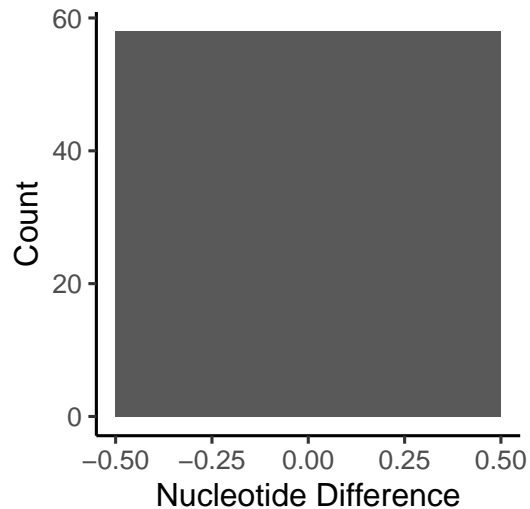
TRBV7–6*01

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



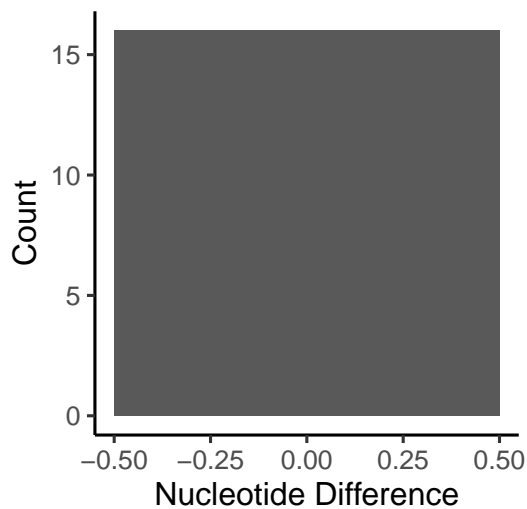
TRBV7–9*03

58 sequences assigned
58 (100%) exact matches, in which:
58 unique CDR3
11 unique J



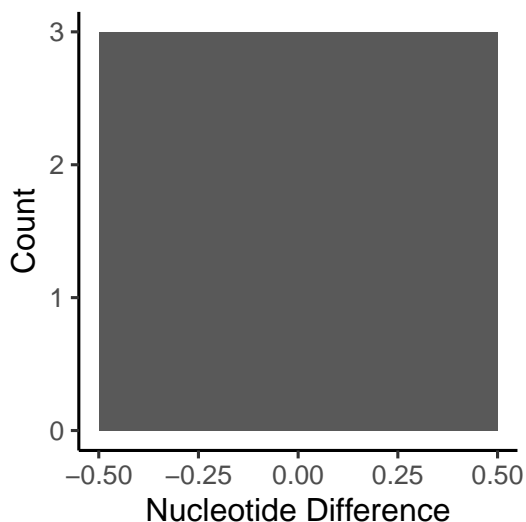
TRBV7–2*02

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



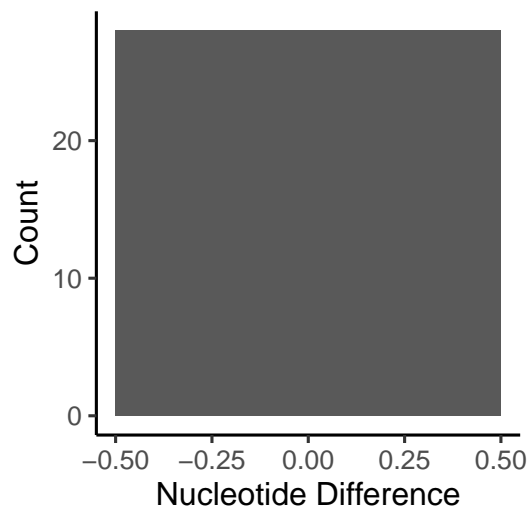
TRBV7–7*01_C315T

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

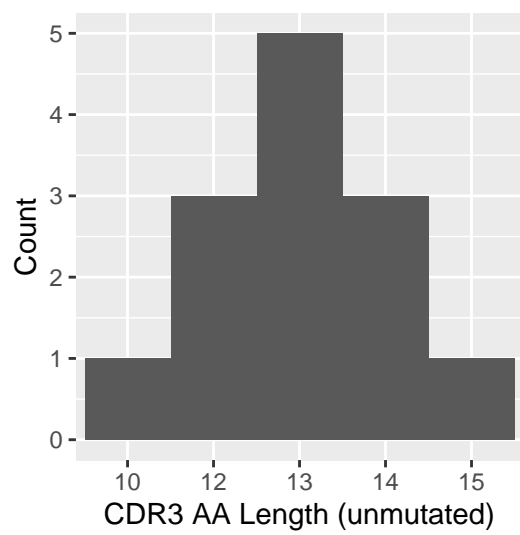


TRBV29–1*01

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



TRBV5-5*02_322TTGG325



TRBV7-7*01_C315T

