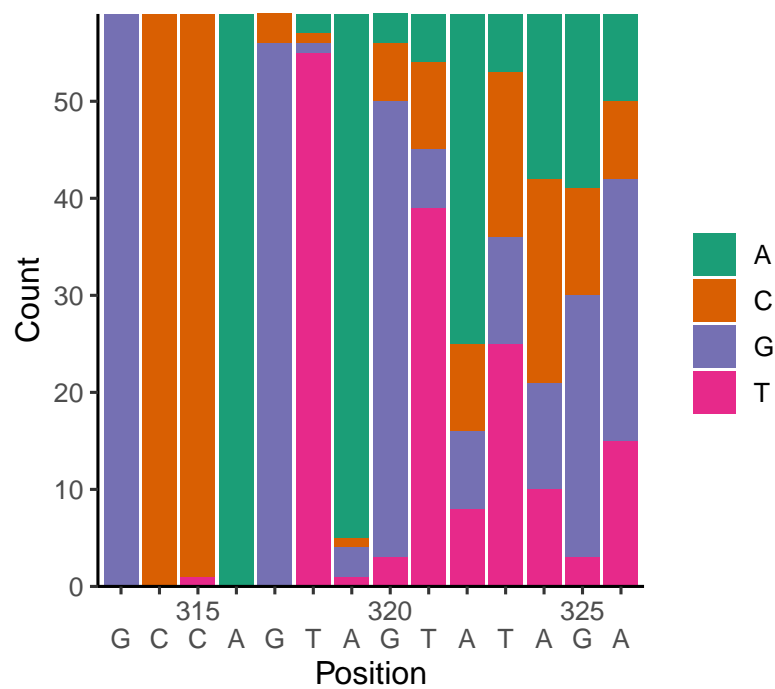
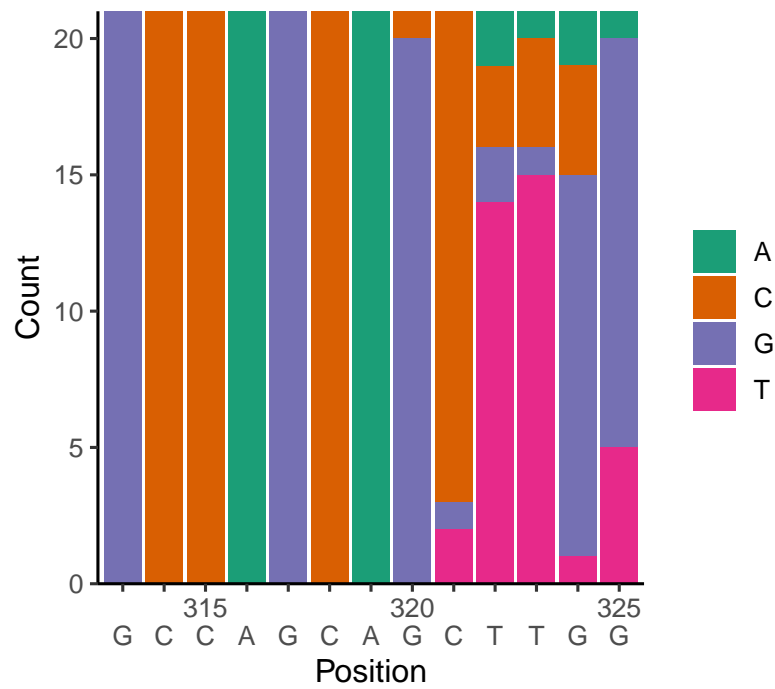


Gene TRBV19*01_A24G



Gene TRBV5-5*02_322TTGG325



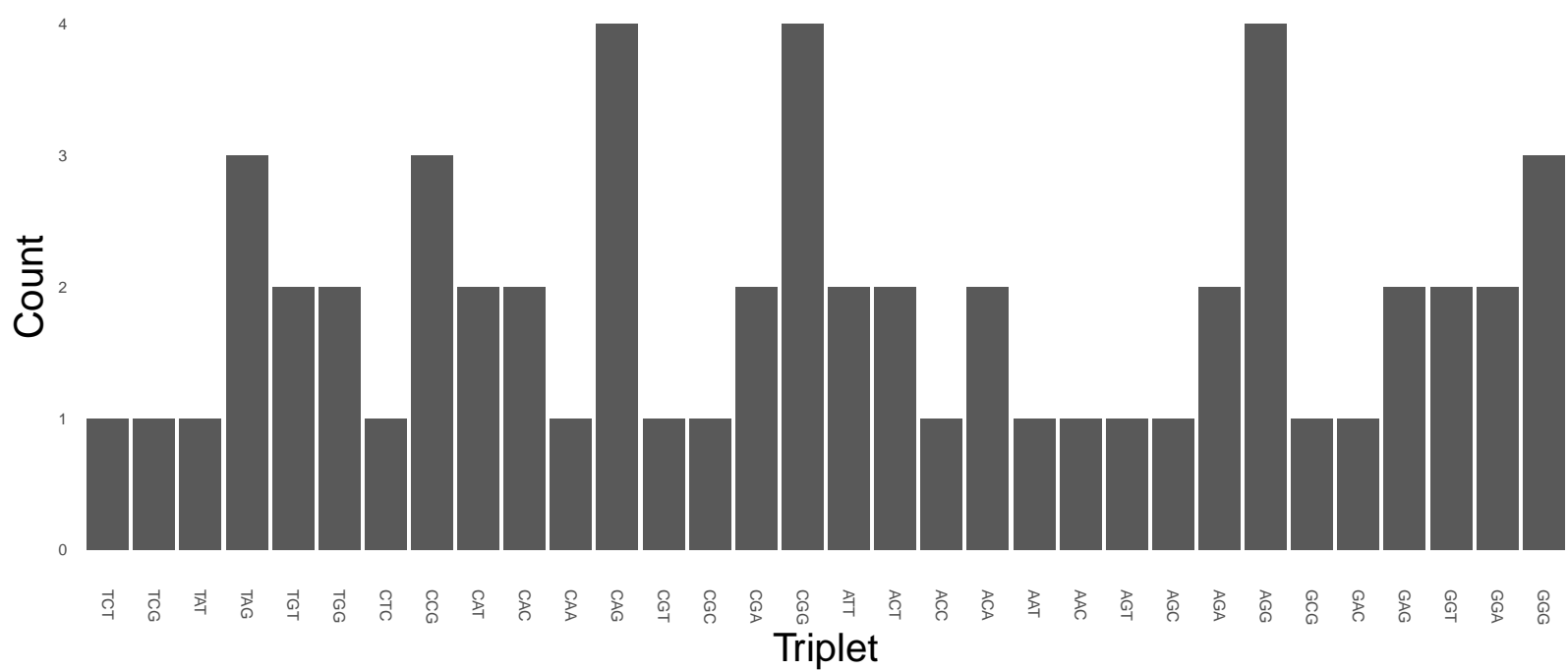
Gene TRBV19*01_A24G



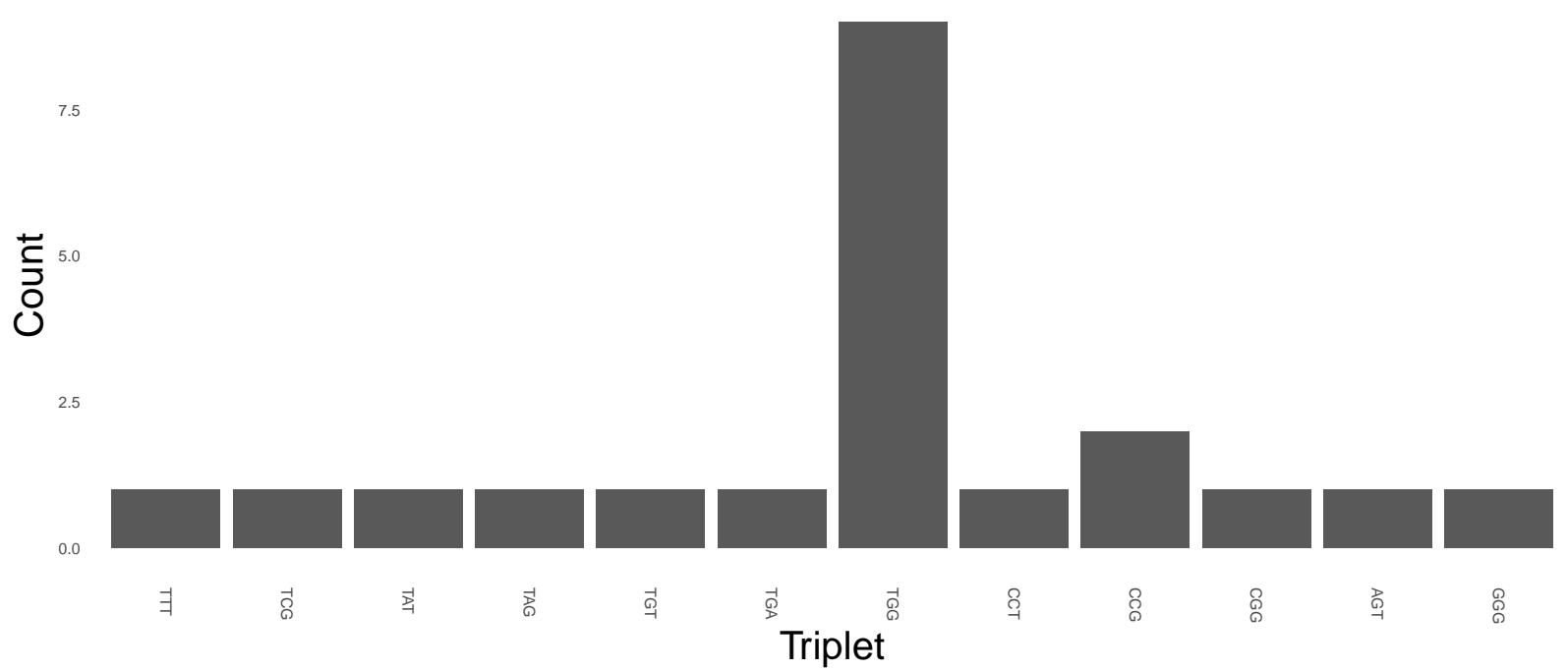
Gene TRBV5-5*02_322TTGG325



TRBV19*01_A24G– Final 3 nucleotides as a triplet

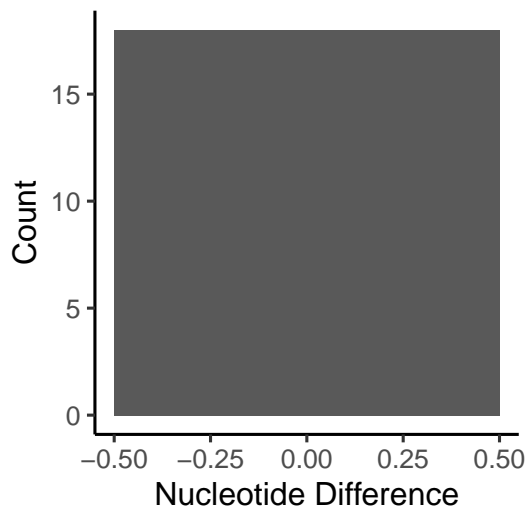


TRBV5–5*02_322TTGG325– Final 3 nucleotides as a triplet



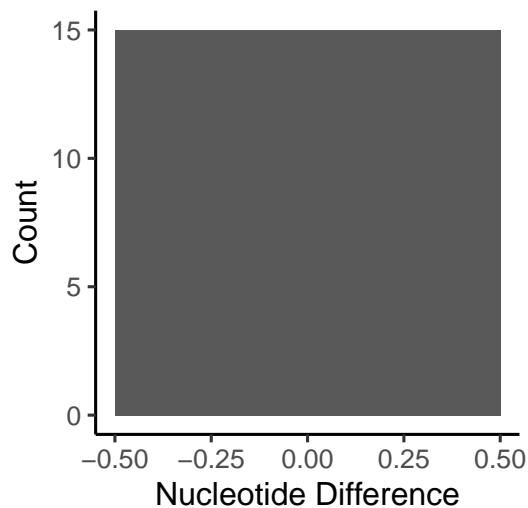
TRBV10–1*01

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



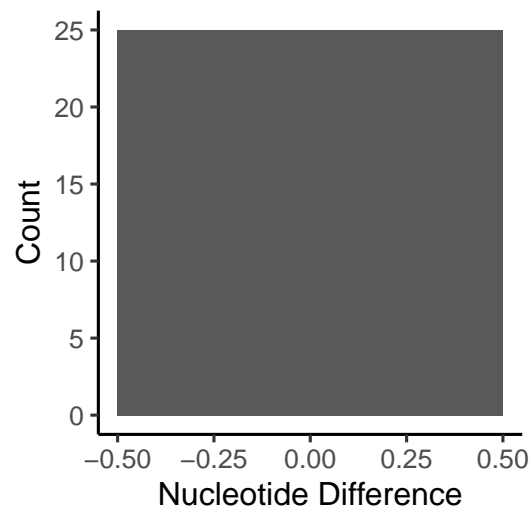
TRBV10–2*01

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



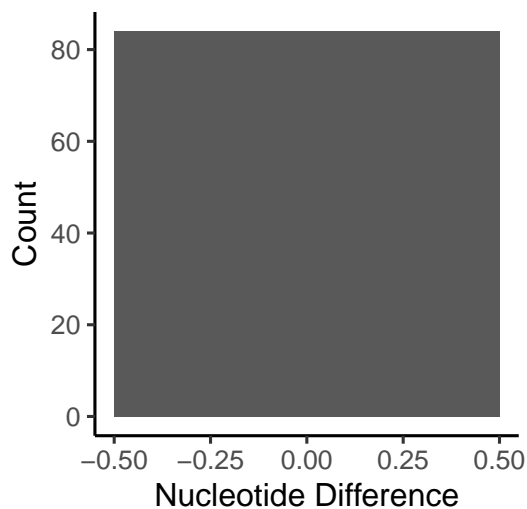
TRBV14*01

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



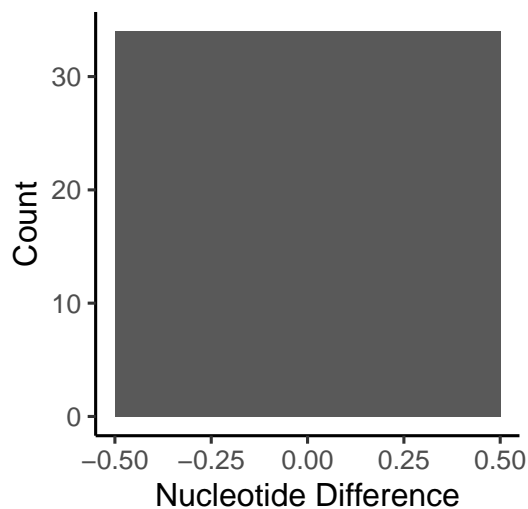
TRBV20–1*01

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



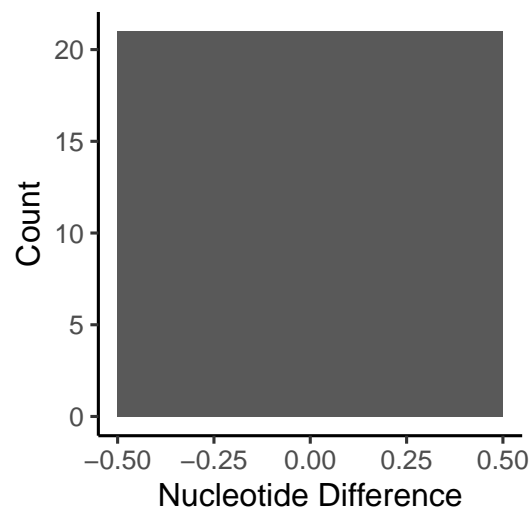
TRBV10–3*02

34 sequences assigned
34 (100%) exact matches, in which:
34 unique CDR3
11 unique J



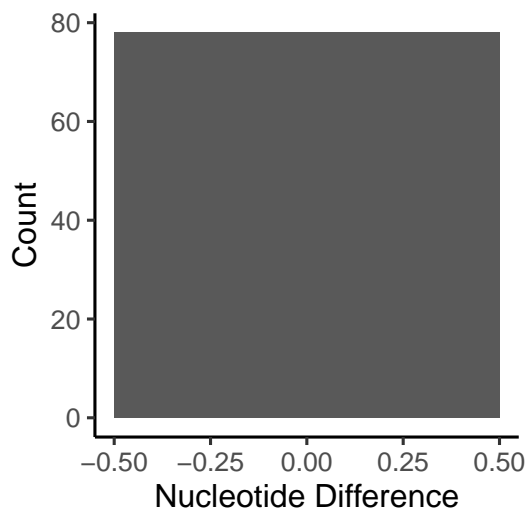
TRBV15*02

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



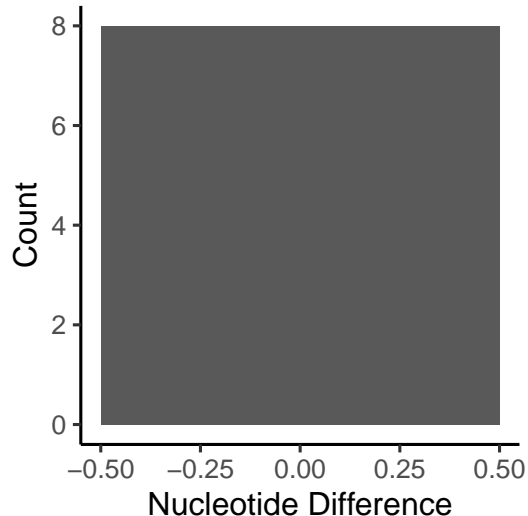
TRBV20–1*05

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



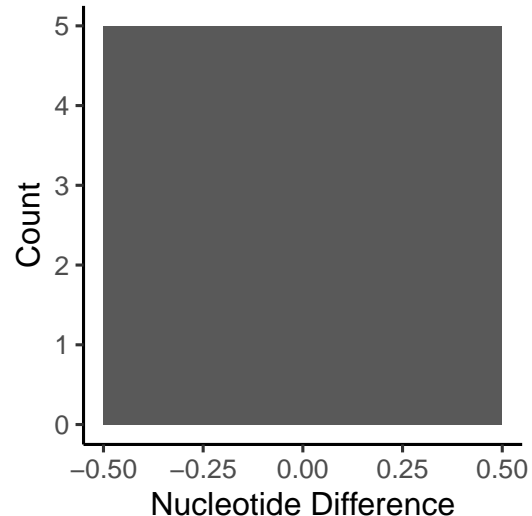
TRBV13*01

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



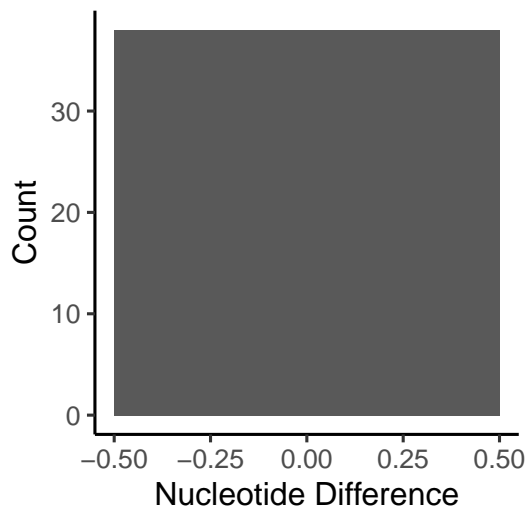
TRBV16*01

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



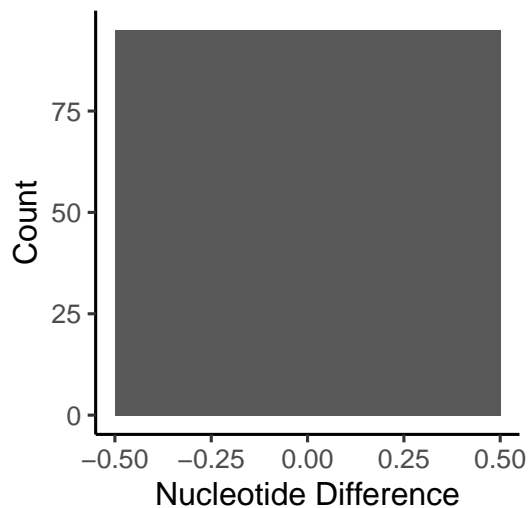
TRBV18*01

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



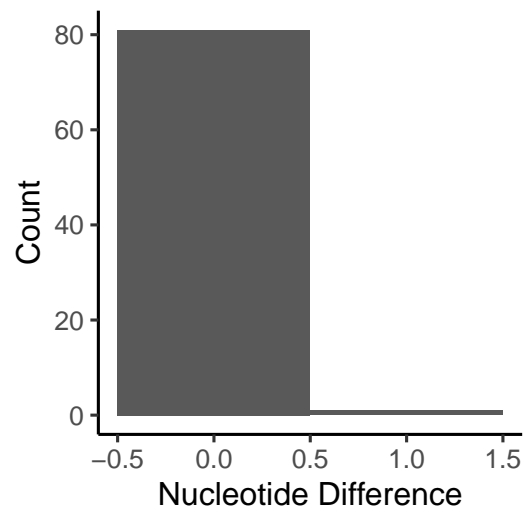
TRBV2*01

95 sequences assigned
95 (100%) exact matches, in which:
95 unique CDR3
13 unique J



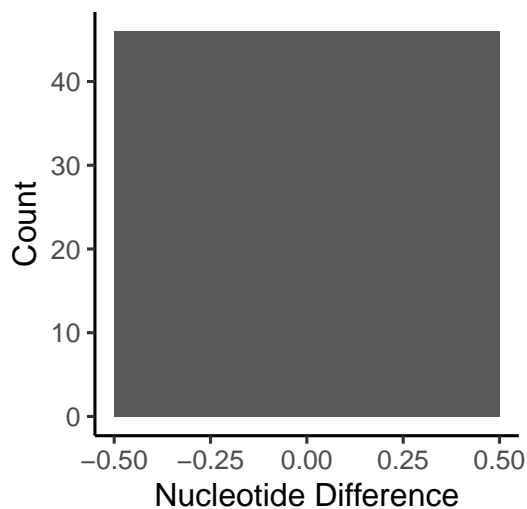
TRBV30*01

82 sequences assigned
81 (98.8%) exact matches, in which:
81 unique CDR3
14 unique J



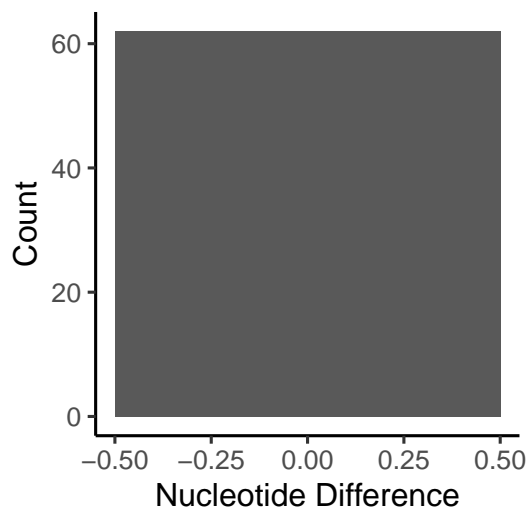
TRBV19*01

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



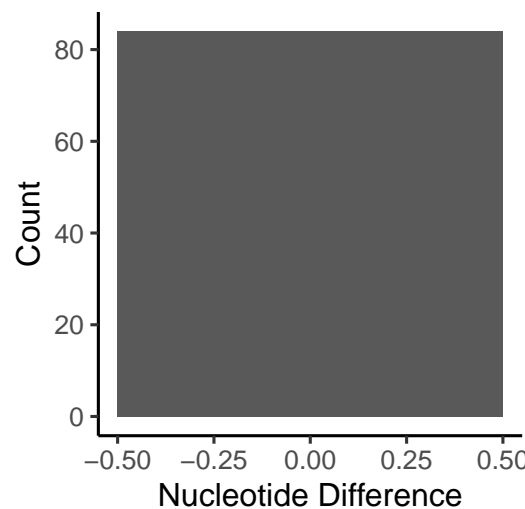
TRBV27*01

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



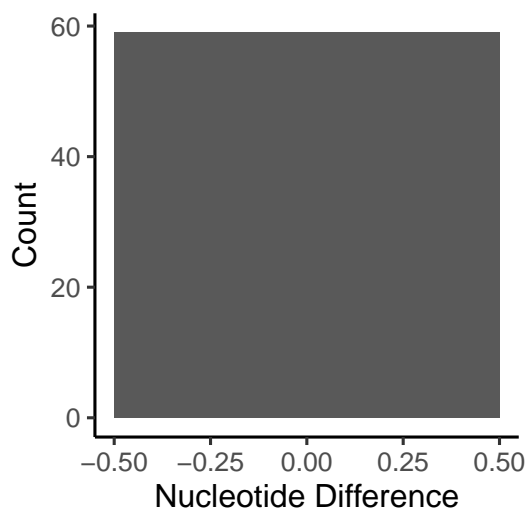
TRBV9*01

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



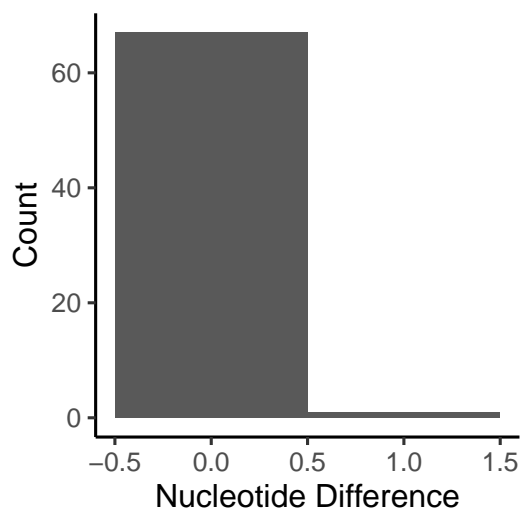
TRBV19*01_A24G

59 sequences assigned
59 (100%) exact matches, in which:
59 unique CDR3
13 unique J



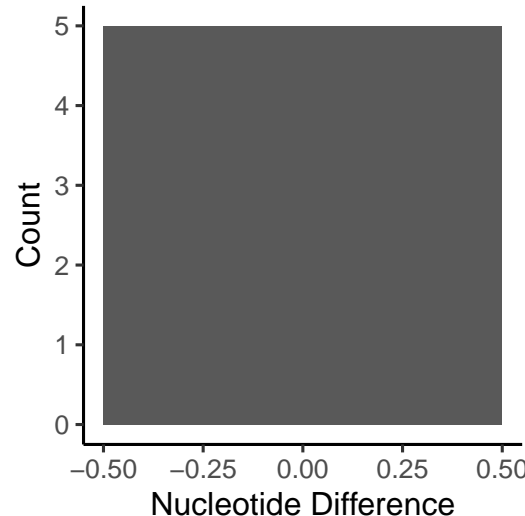
TRBV28*01

68 sequences assigned
67 (98.5%) exact matches, in which:
67 unique CDR3
13 unique J



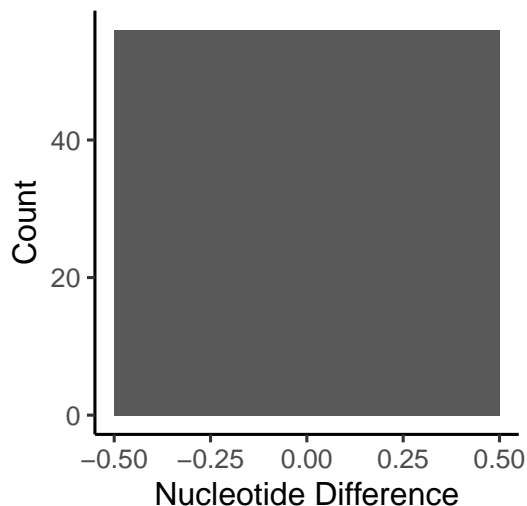
TRBV11-1*01

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



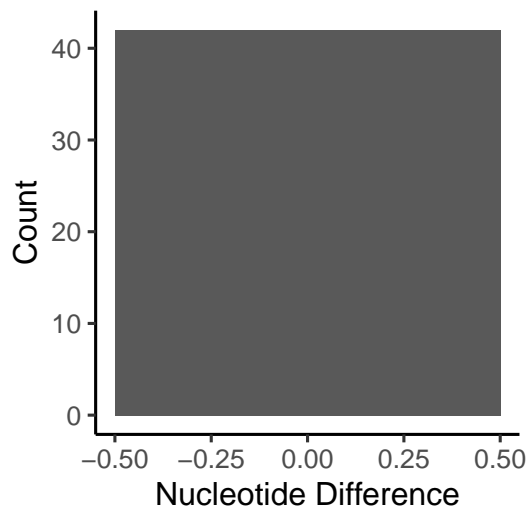
TRBV11-2*01

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



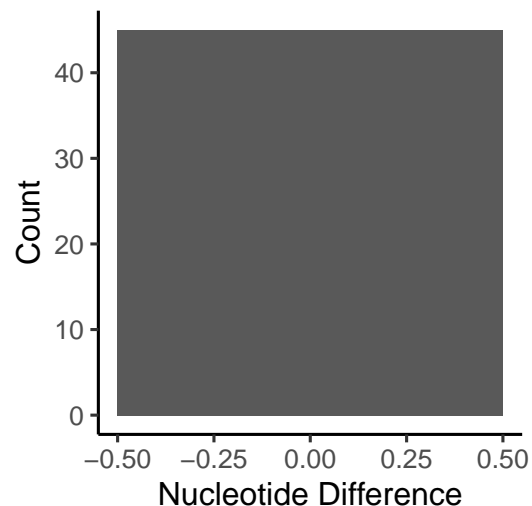
TRBV12-4*01

42 sequences assigned
42 (100%) exact matches, in which:
42 unique CDR3
8 unique J



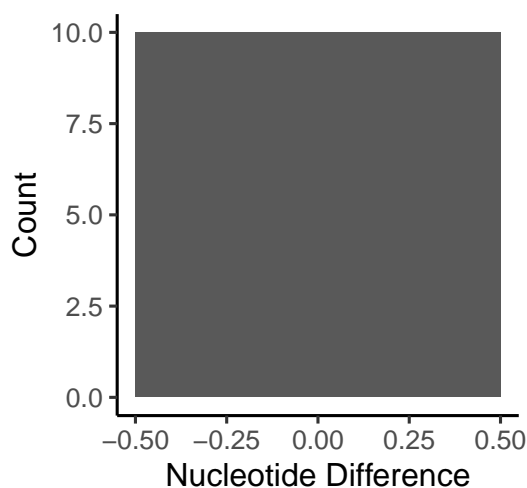
TRBV4-1*01

45 sequences assigned
45 (100%) exact matches, in which:
45 unique CDR3
12 unique J



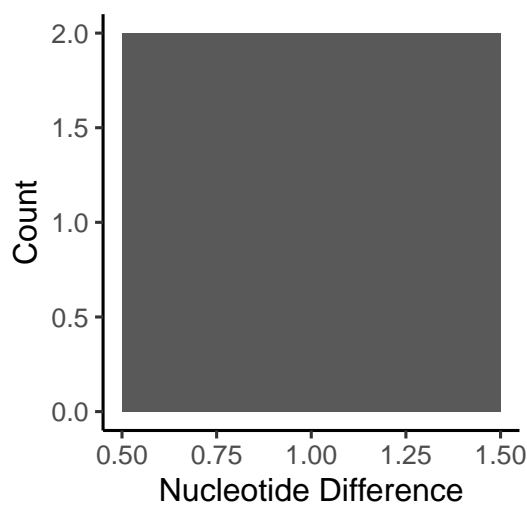
TRBV11-3*01

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



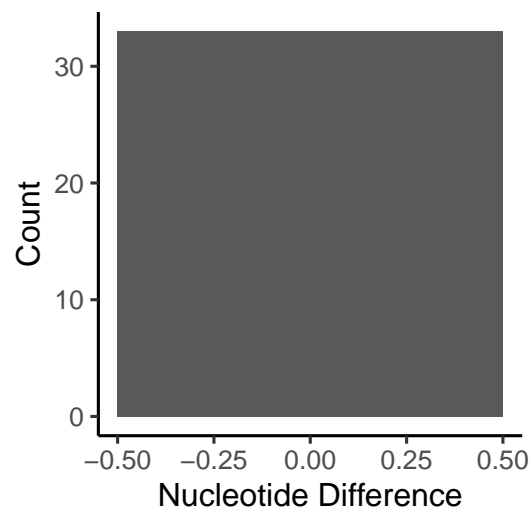
TRBV12-5*01

2 sequences assigned
No exact matches.



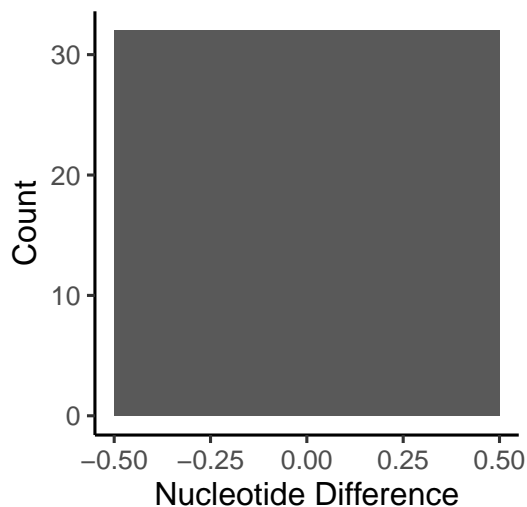
TRBV24-1*01

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



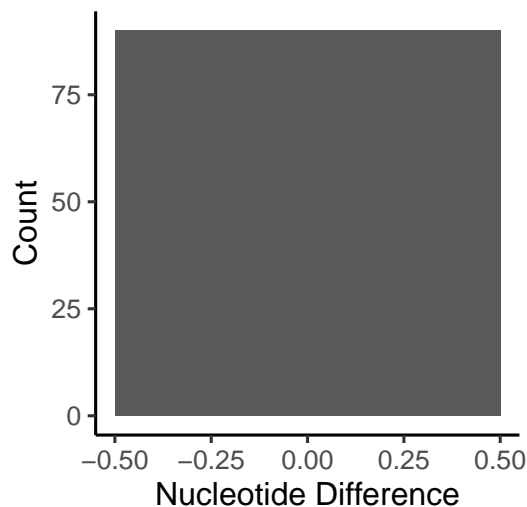
TRBV12-3*01

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



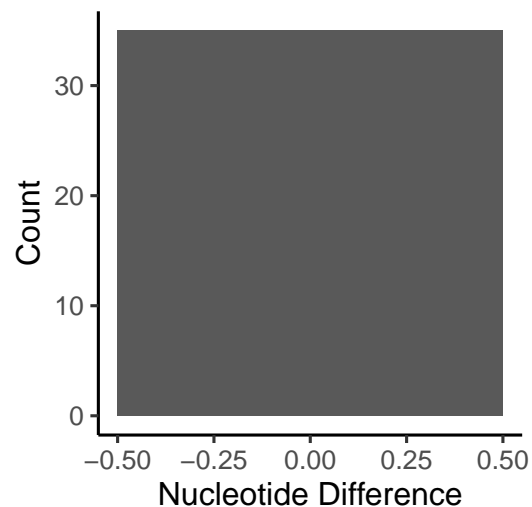
TRBV3-1*01

90 sequences assigned
90 (100%) exact matches, in which:
90 unique CDR3
14 unique J



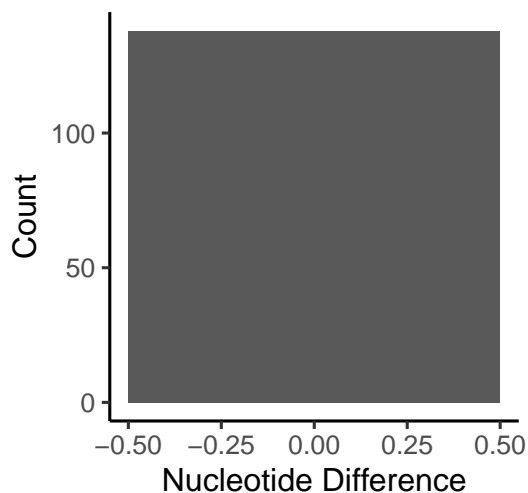
TRBV4-2*01

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



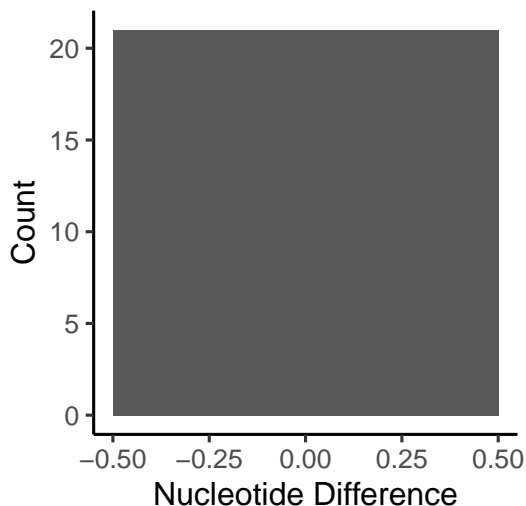
TRBV5-1*01

138 sequences assigned
138 (100%) exact matches, in which:
138 unique CDR3
14 unique J



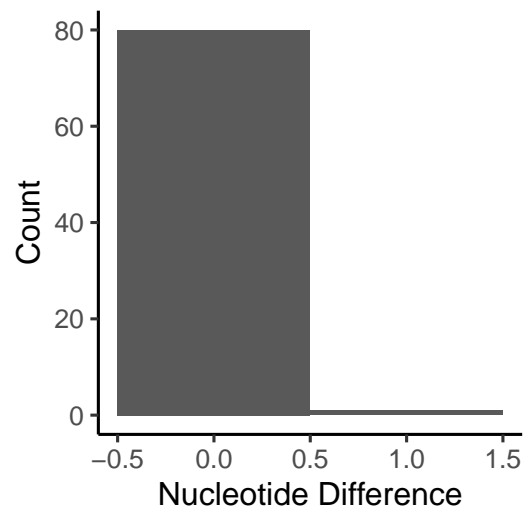
TRBV5-5*02_322TTGG32

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



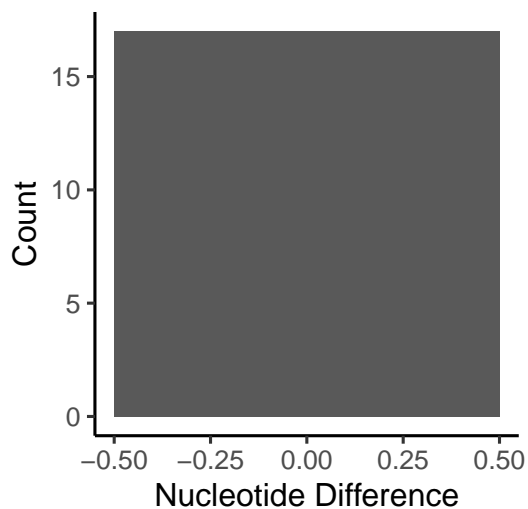
TRBV6-1*01

81 sequences assigned
80 (98.8%) exact matches, in which:
80 unique CDR3
13 unique J



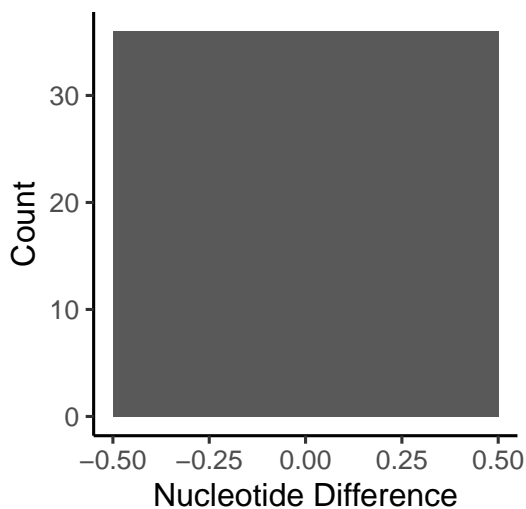
TRBV25-1*01

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



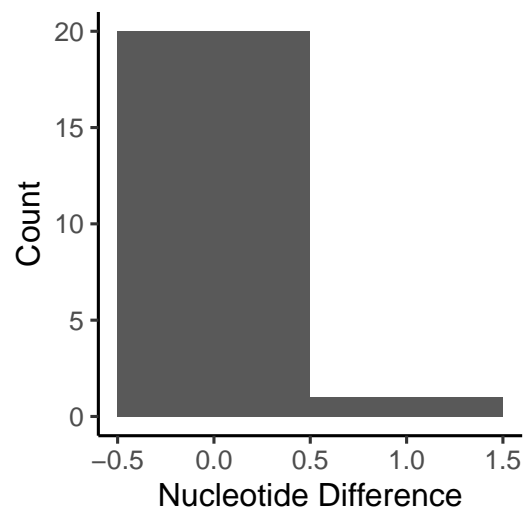
TRBV5-6*01

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



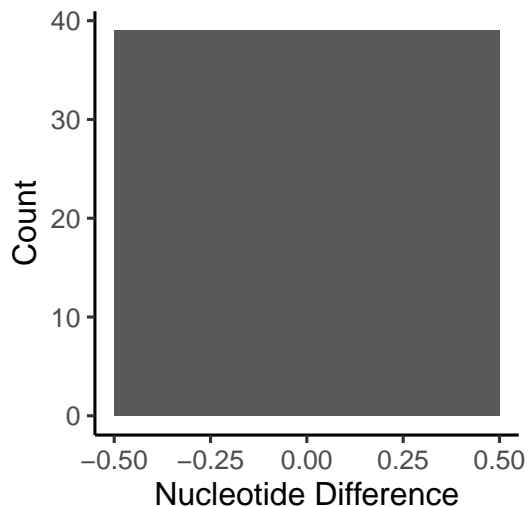
TRBV6-4*01

21 sequences assigned
20 (95.2%) exact matches, in which:
20 unique CDR3
9 unique J



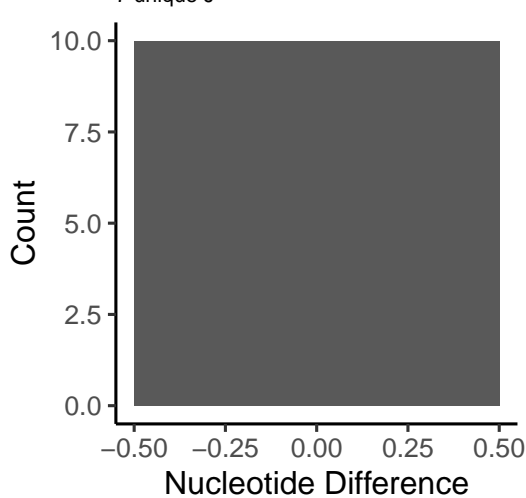
TRBV5-4*01

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



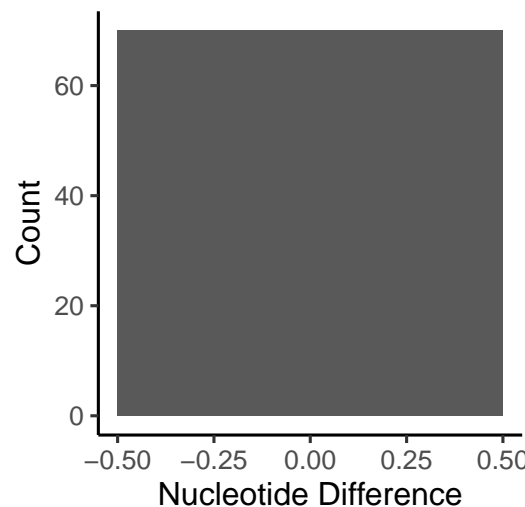
TRBV5-8*01

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



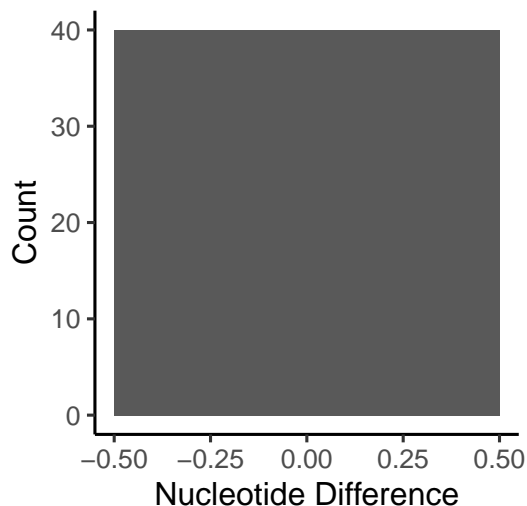
TRBV6-5*01

70 sequences assigned
70 (100%) exact matches, in which:
70 unique CDR3
13 unique J



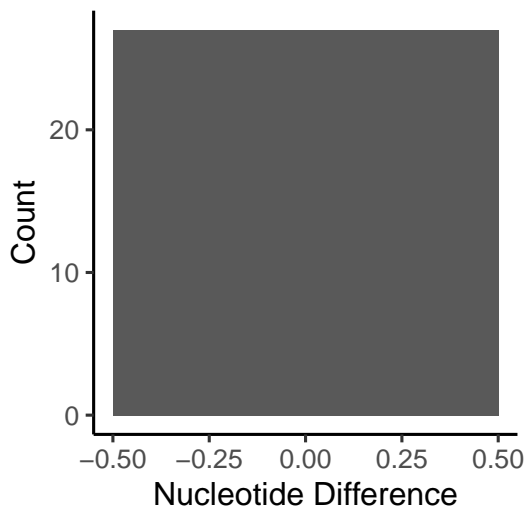
TRBV6-6*01

40 sequences assigned
40 (100%) exact matches, in which:
40 unique CDR3
12 unique J



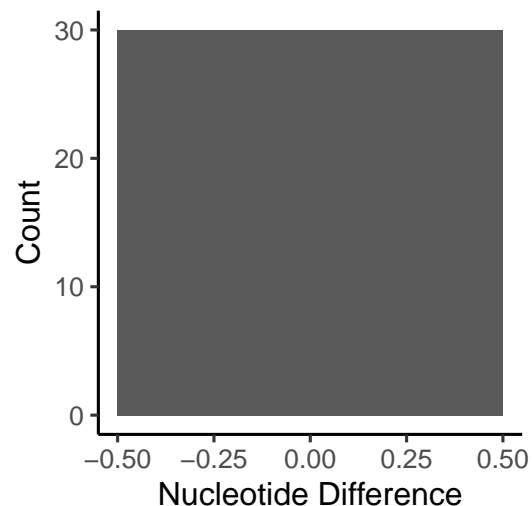
TRBV7-3*01

27 sequences assigned
27 (100%) exact matches, in which:
27 unique CDR3
10 unique J



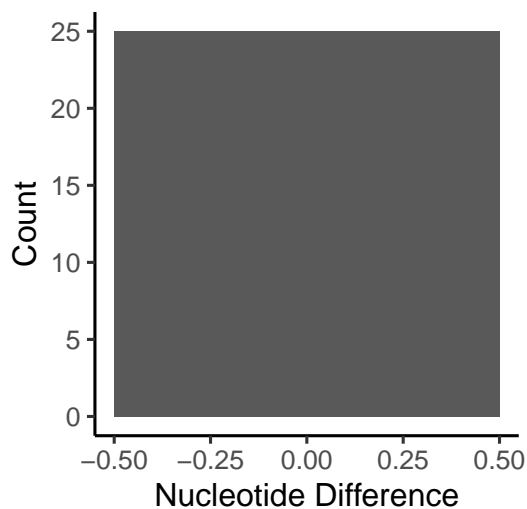
TRBV7-8*01

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



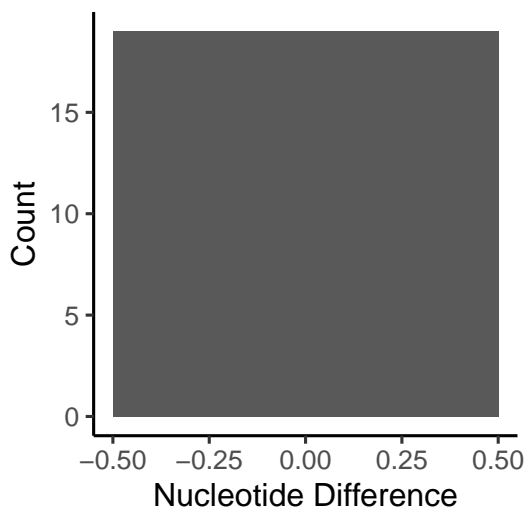
TRBV6-23*01

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



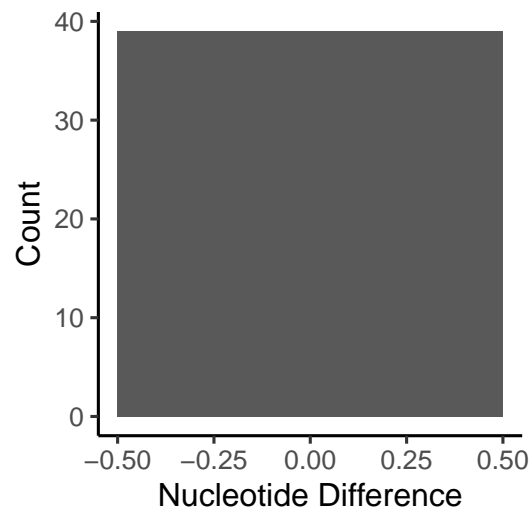
TRBV7-6*01

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



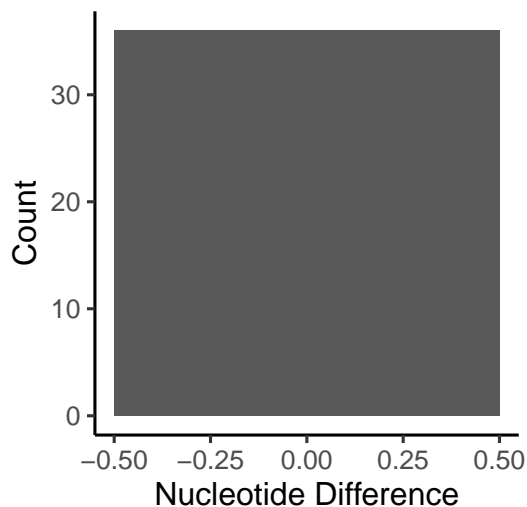
TRBV7-9*01

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



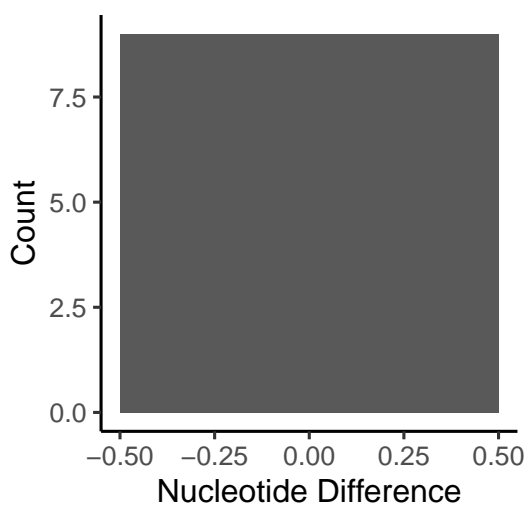
TRBV7-2*02

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



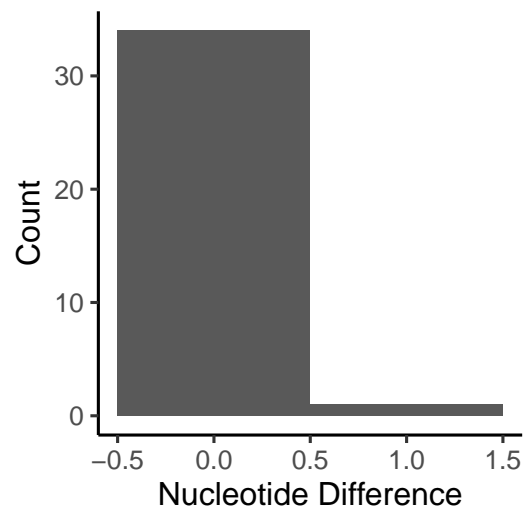
TRBV7-7*01

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



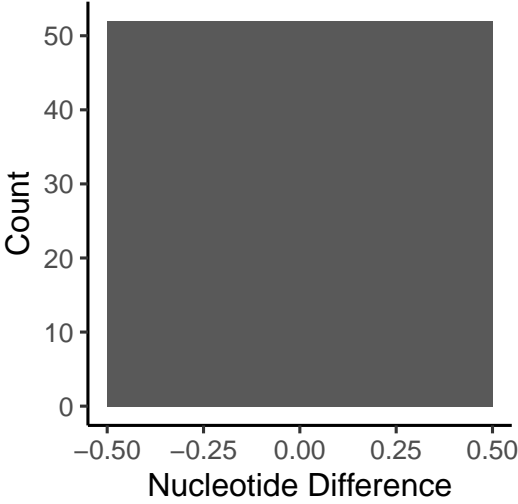
TRBV7-9*03

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

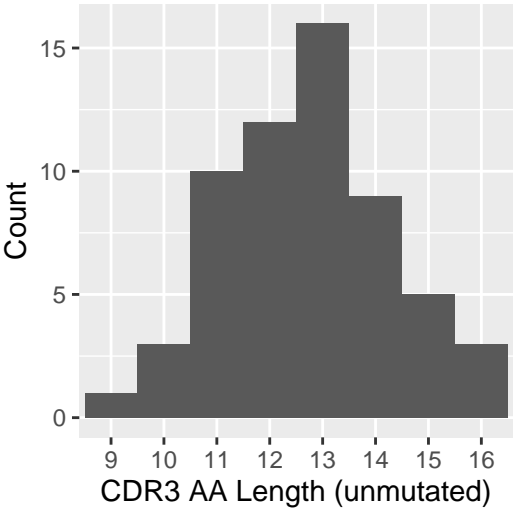


TRBV29-1*01

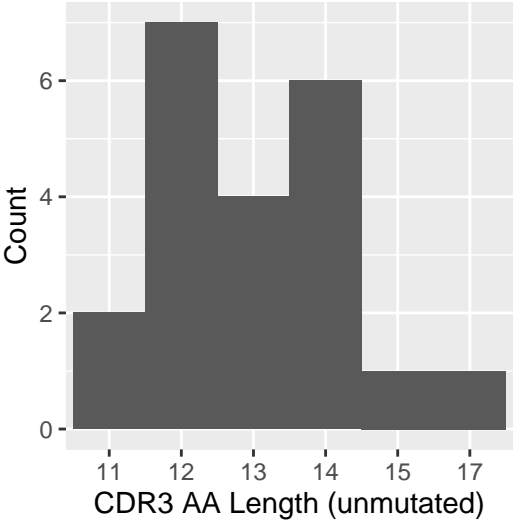
52 sequences assigned
52 (100%) exact matches, in which:
52 unique CDR3
12 unique J

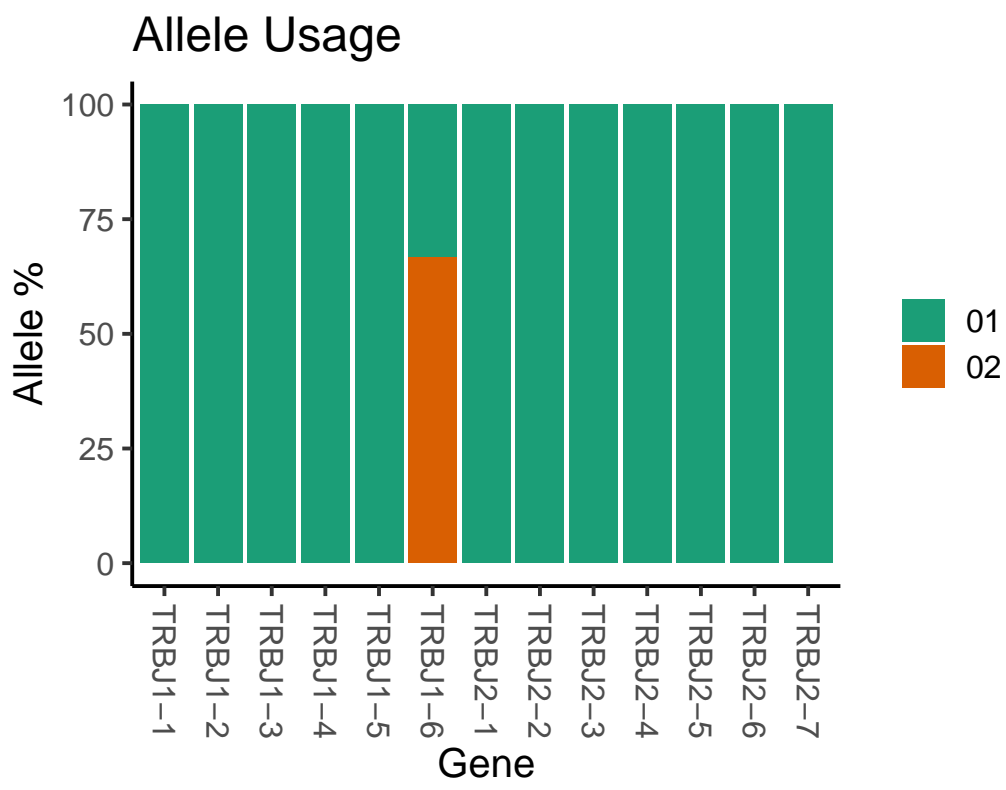


TRBV19*01_A24G

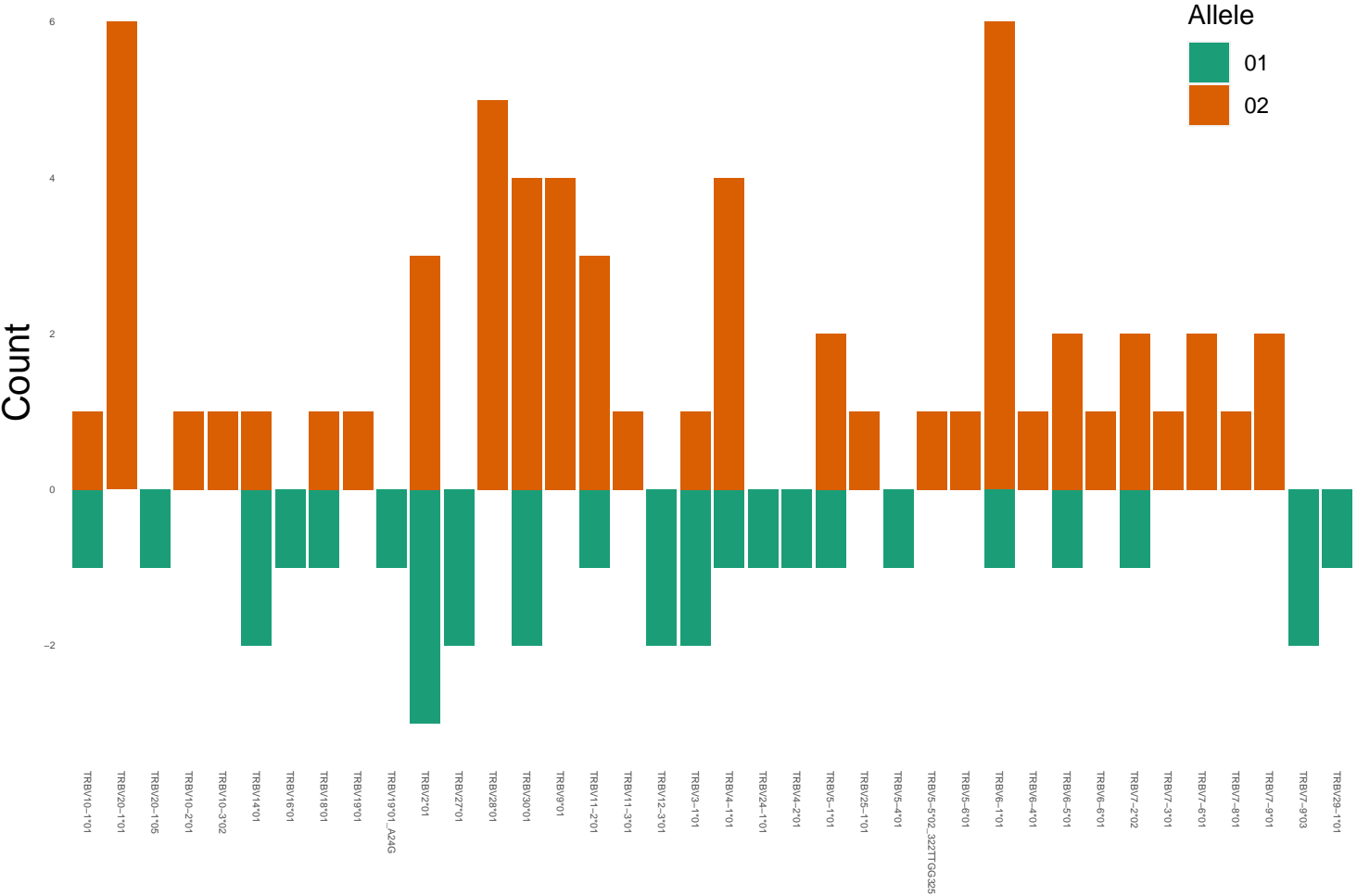


TRBV5-5*02_322TTGG325





Sequence Count by TRBJ1–6 allele usage



Warning – no inferred sequences found.

Warning – no inferred sequences found.