

Gene TRBV10-3*03_313GCCATCAGTGA



Gene TRBV19*01_A24G



Gene TRBV12-4*01_C87T



Gene TRBV29-1*01_A280C



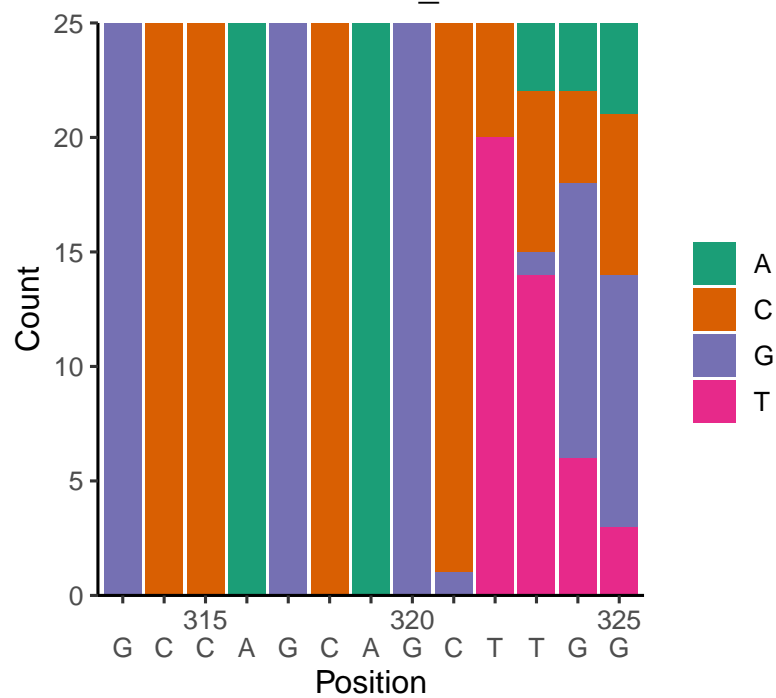
Gene TRBV14*02_322CAAGA326



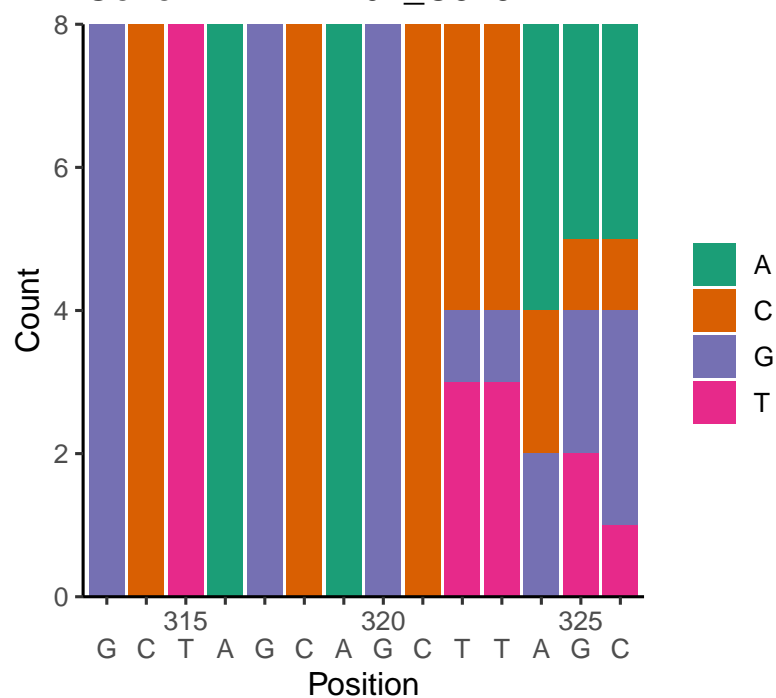
Gene TRBV4-3*01_C274T



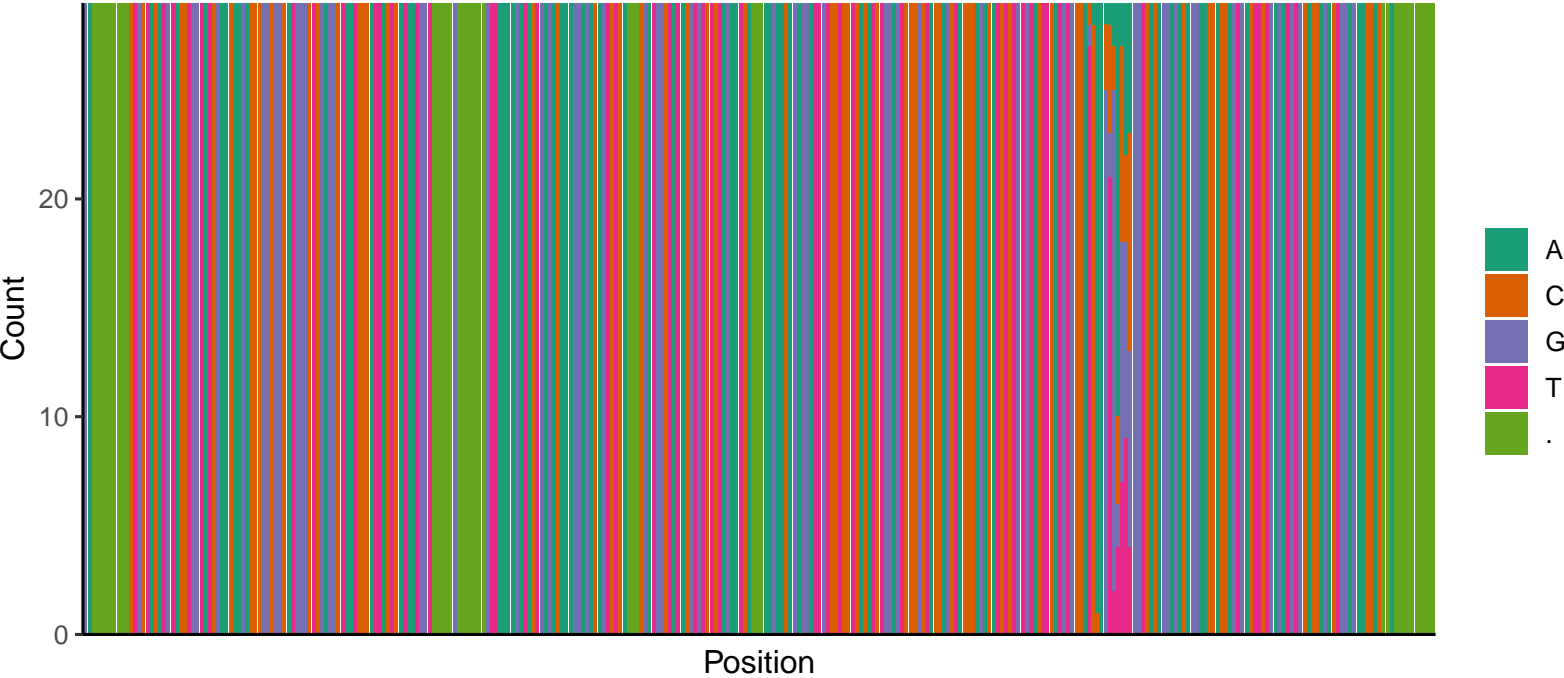
Gene TRBV5-5*02_322TTGG325



Gene TRBV7-7*01_C315T



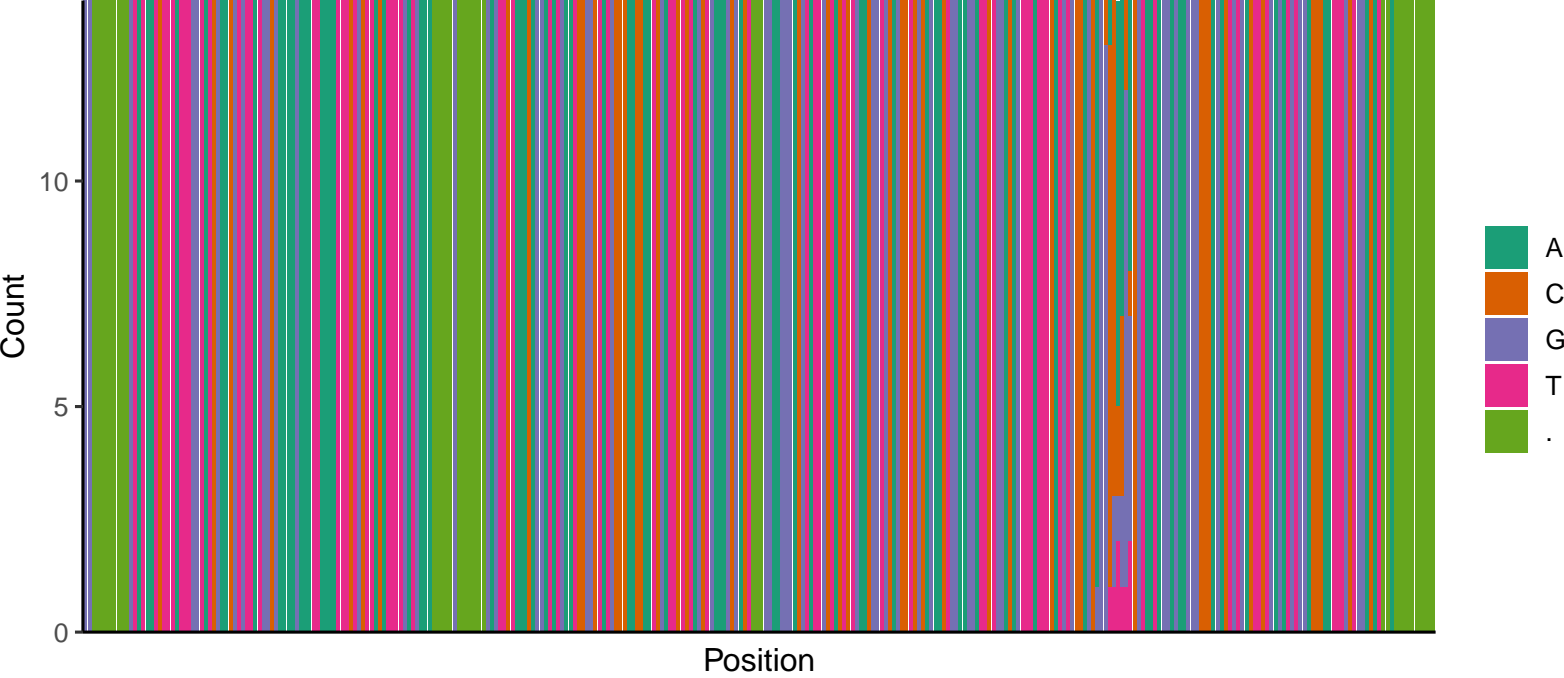
Gene TRBV10-3*03_313GCCATCAGTGAGTC326



Gene TRBV12-4*01_C87T



Gene TRBV14*02_322CAAGA326



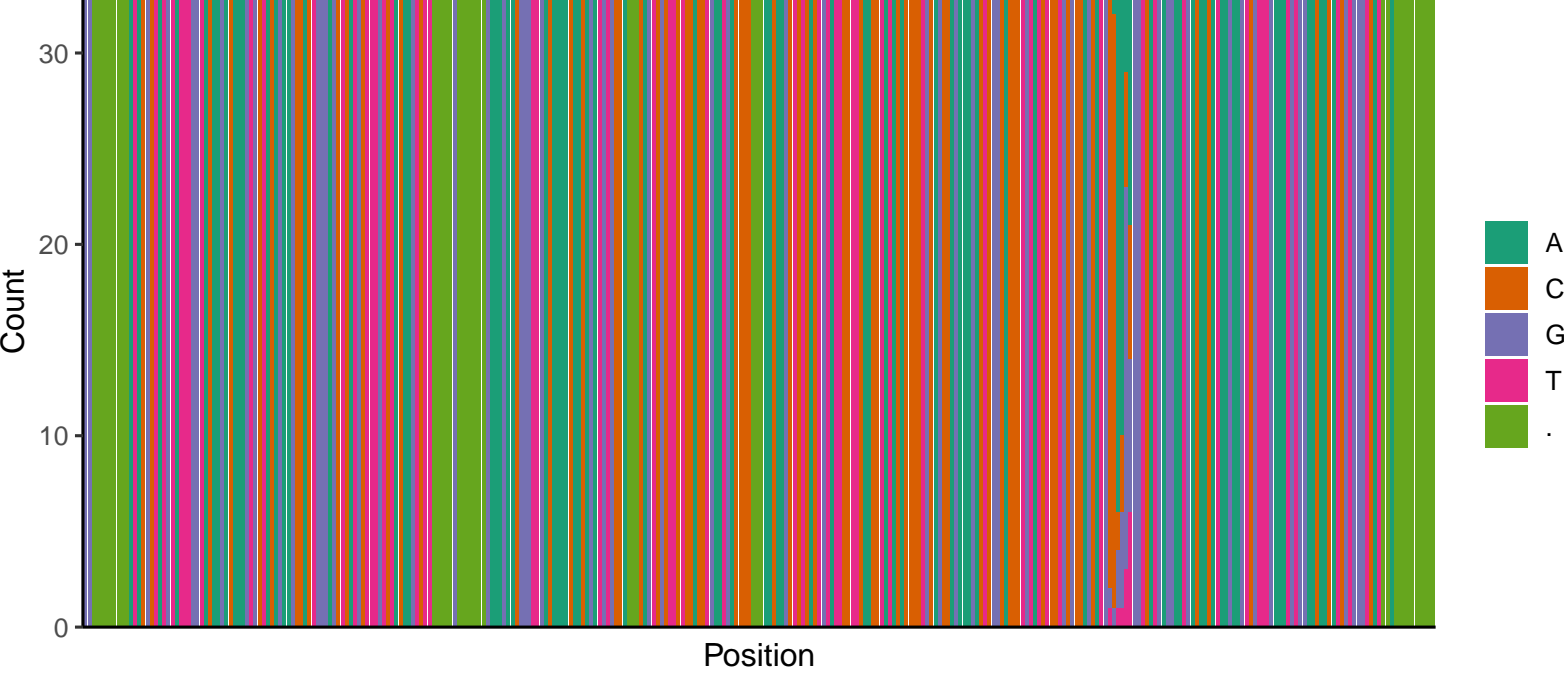
Gene TRBV19*01_A24G



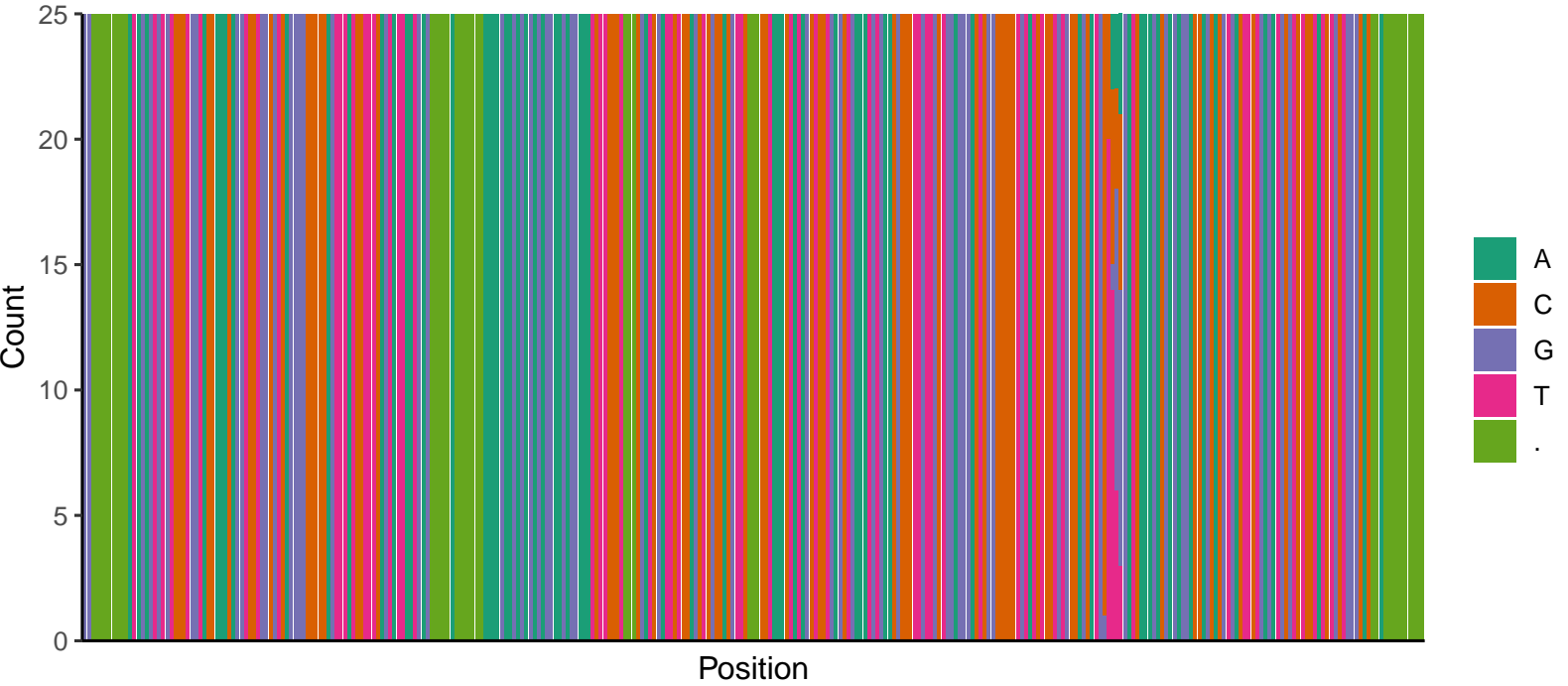
Gene TRBV29-1*01_A280C



Gene TRBV4-3*01_C274T



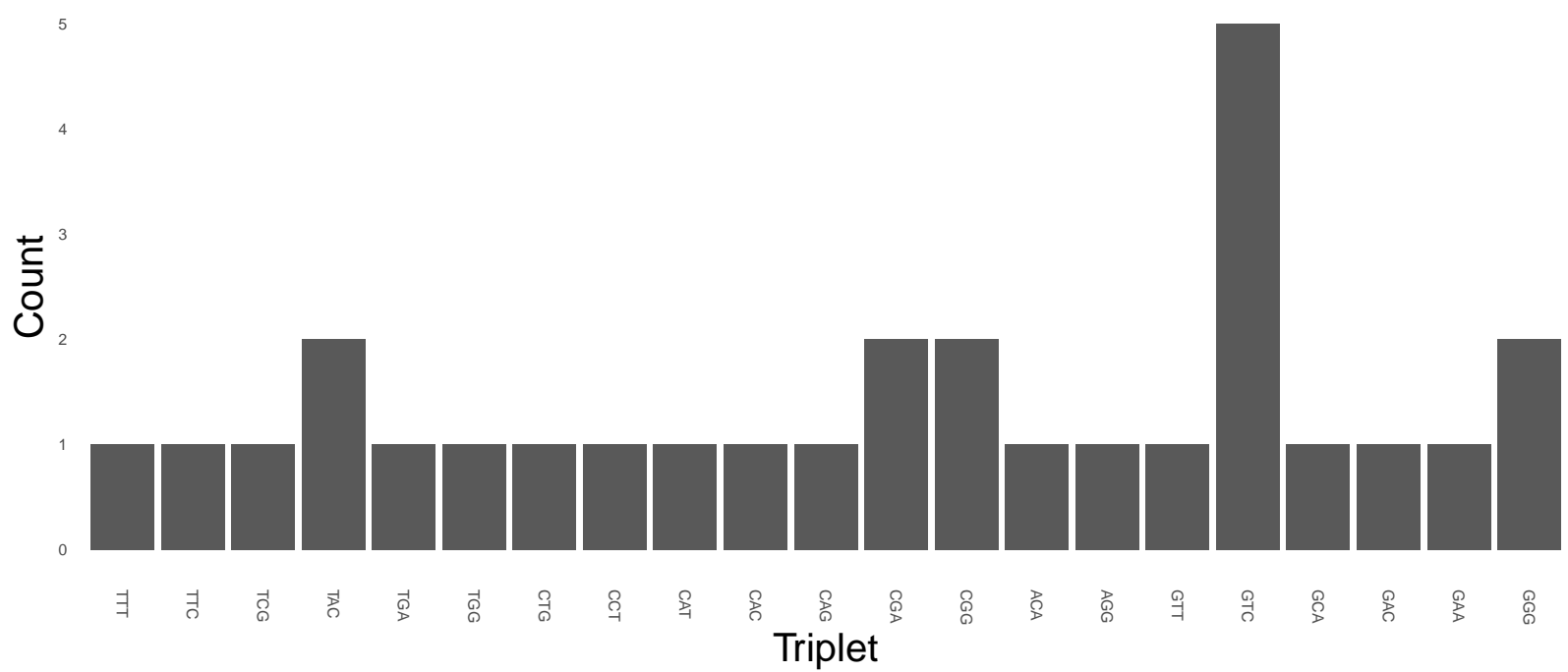
Gene TRBV5-5*02_322TTGG325



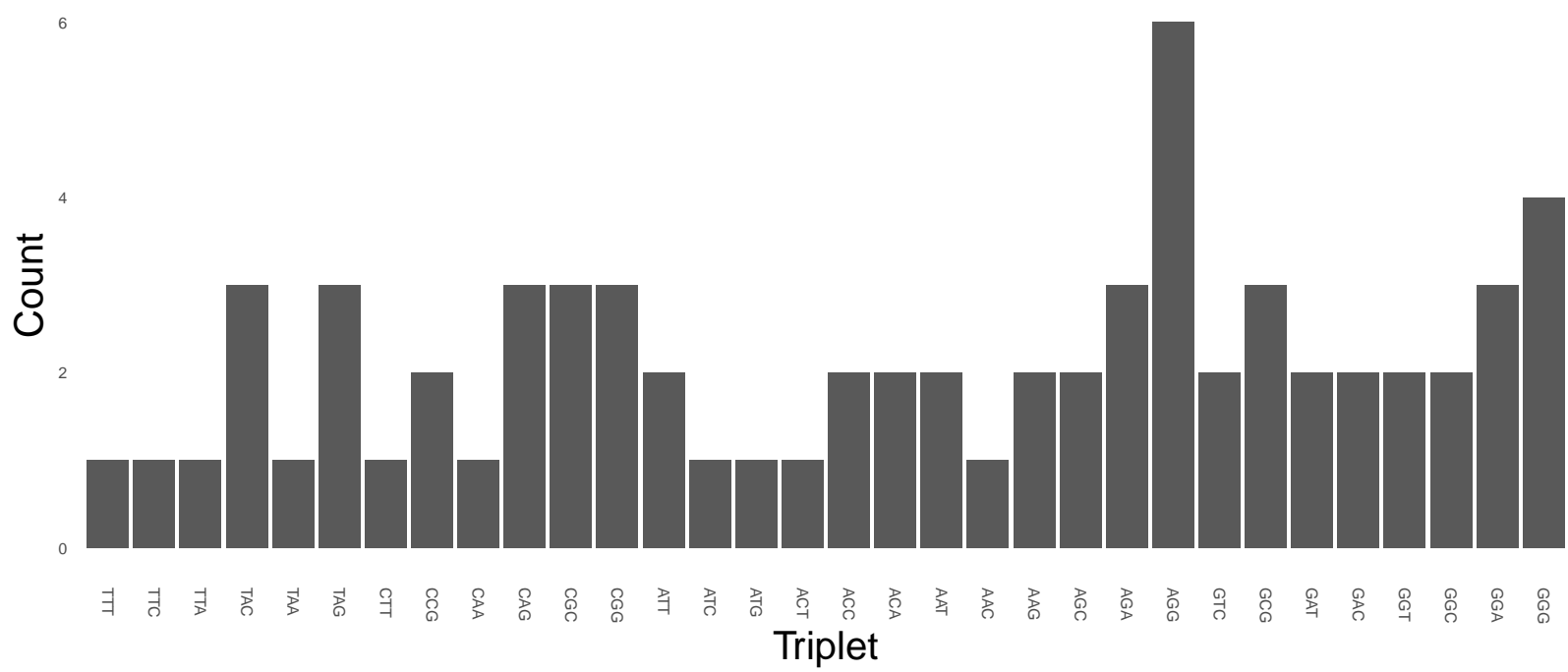
Gene TRBV7-7*01_C315T



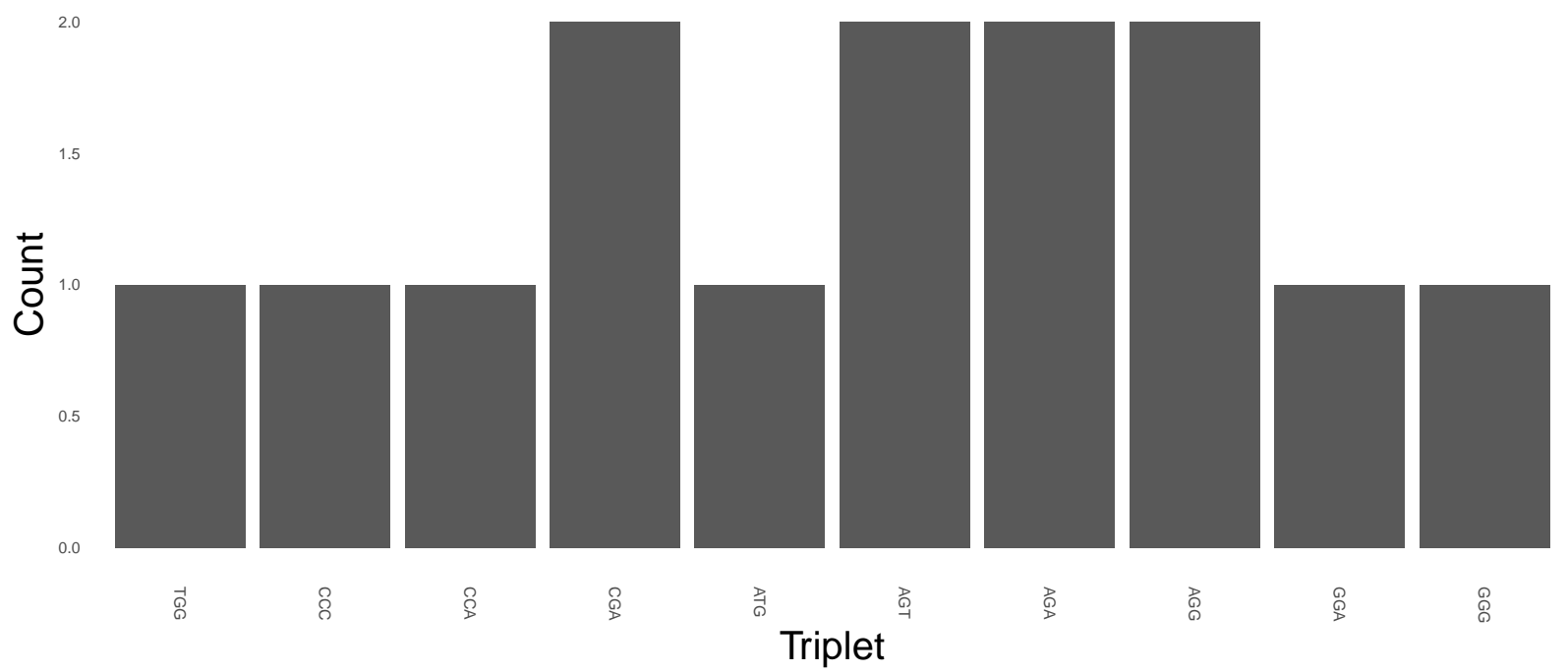
TRBV10–3*03_313GCCATCAGTGAGTC326– Final 3 nucleotides as a triplet



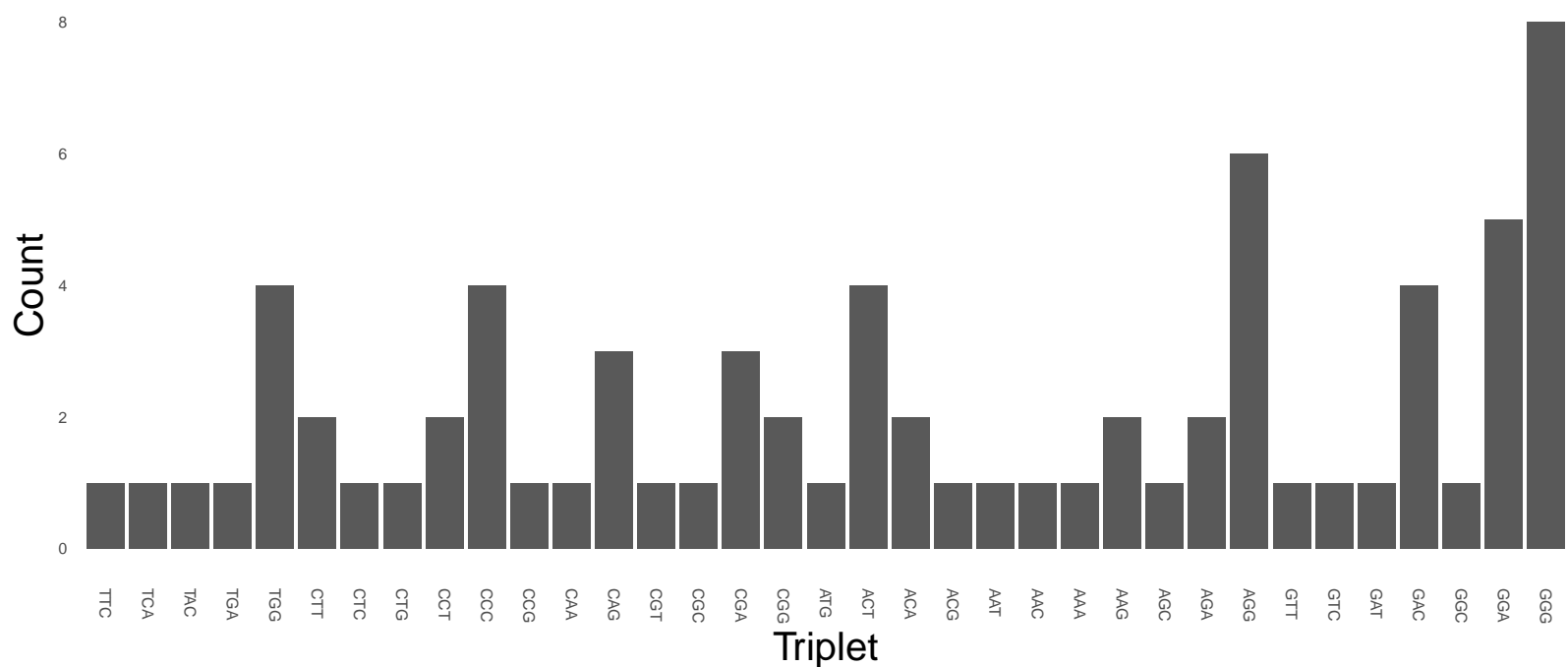
TRBV12–4*01_C87T– Final 3 nucleotides as a triplet



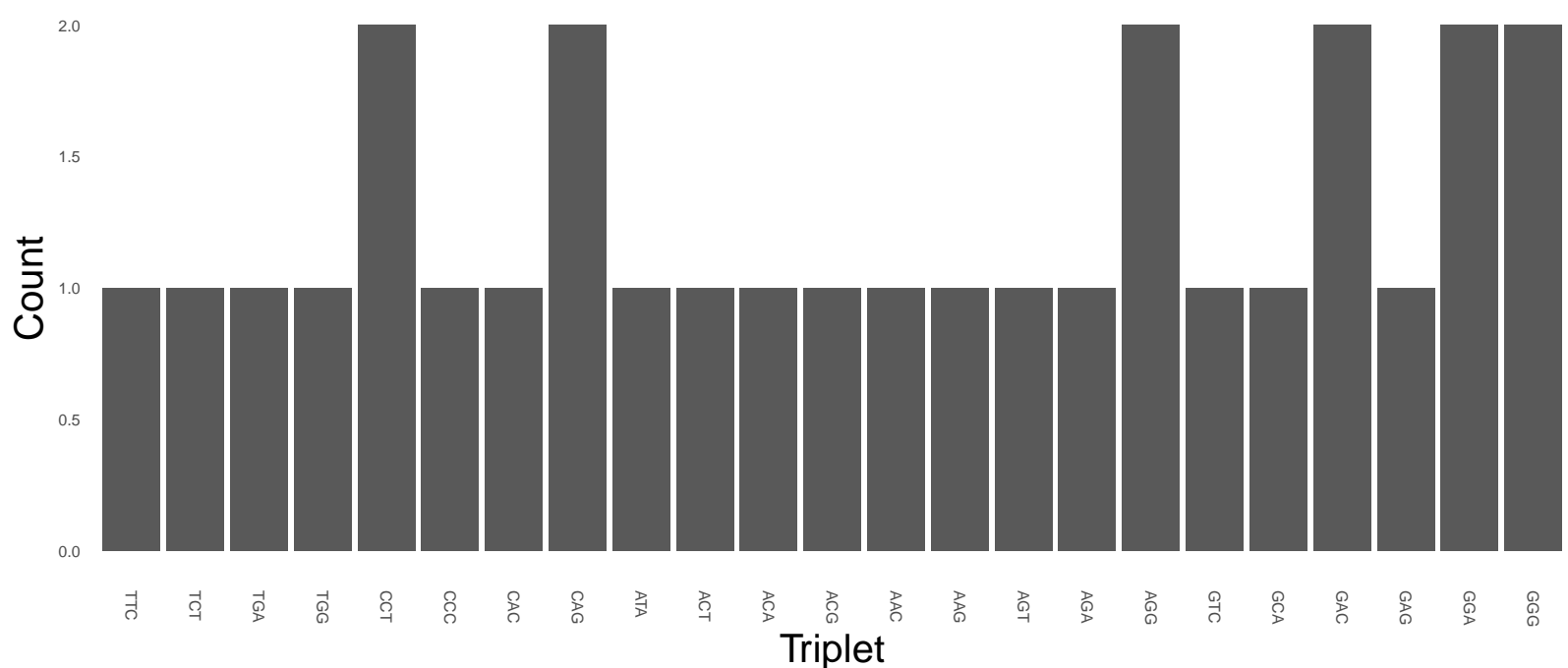
TRBV14*02_322CAAGA326– Final 3 nucleotides as a triplet



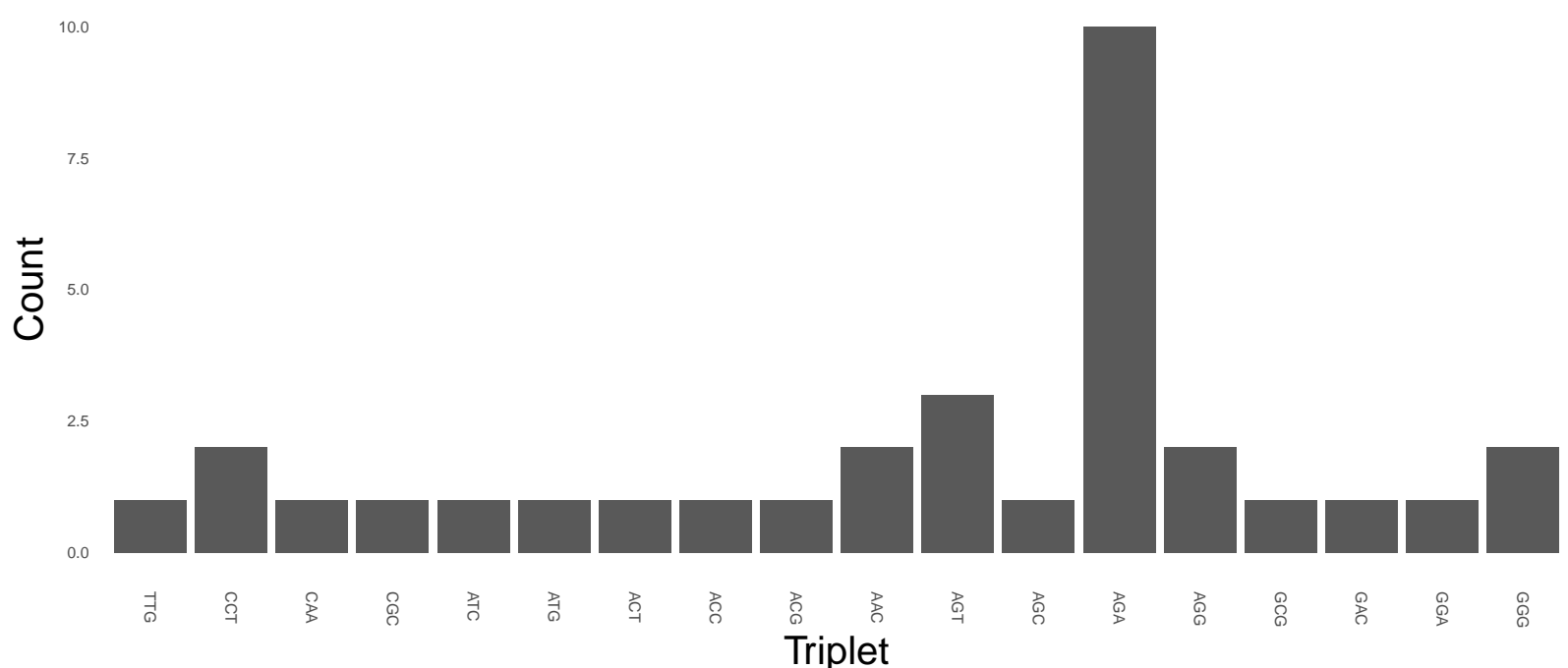
TRBV19*01_A24G– Final 3 nucleotides as a triplet



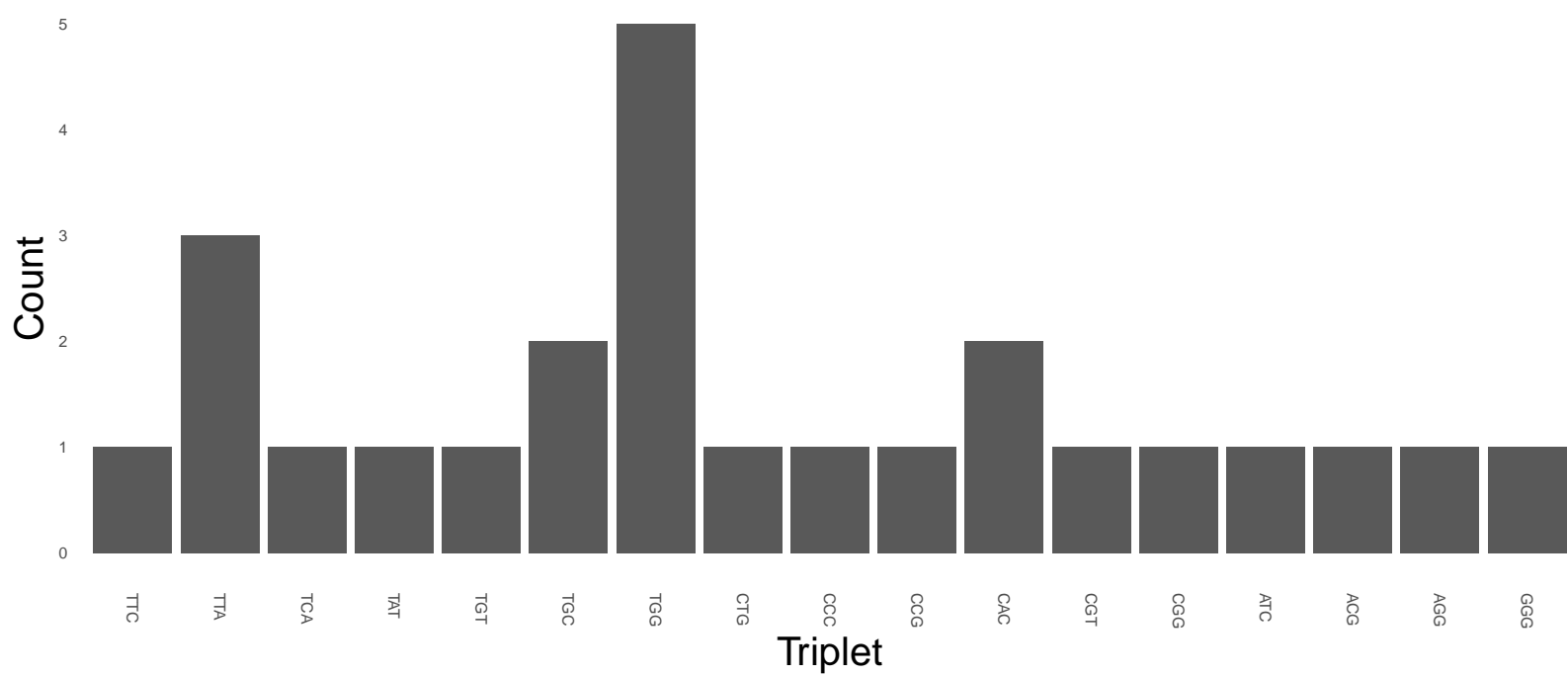
TRBV29–1*01_A280C– Final 3 nucleotides as a triplet



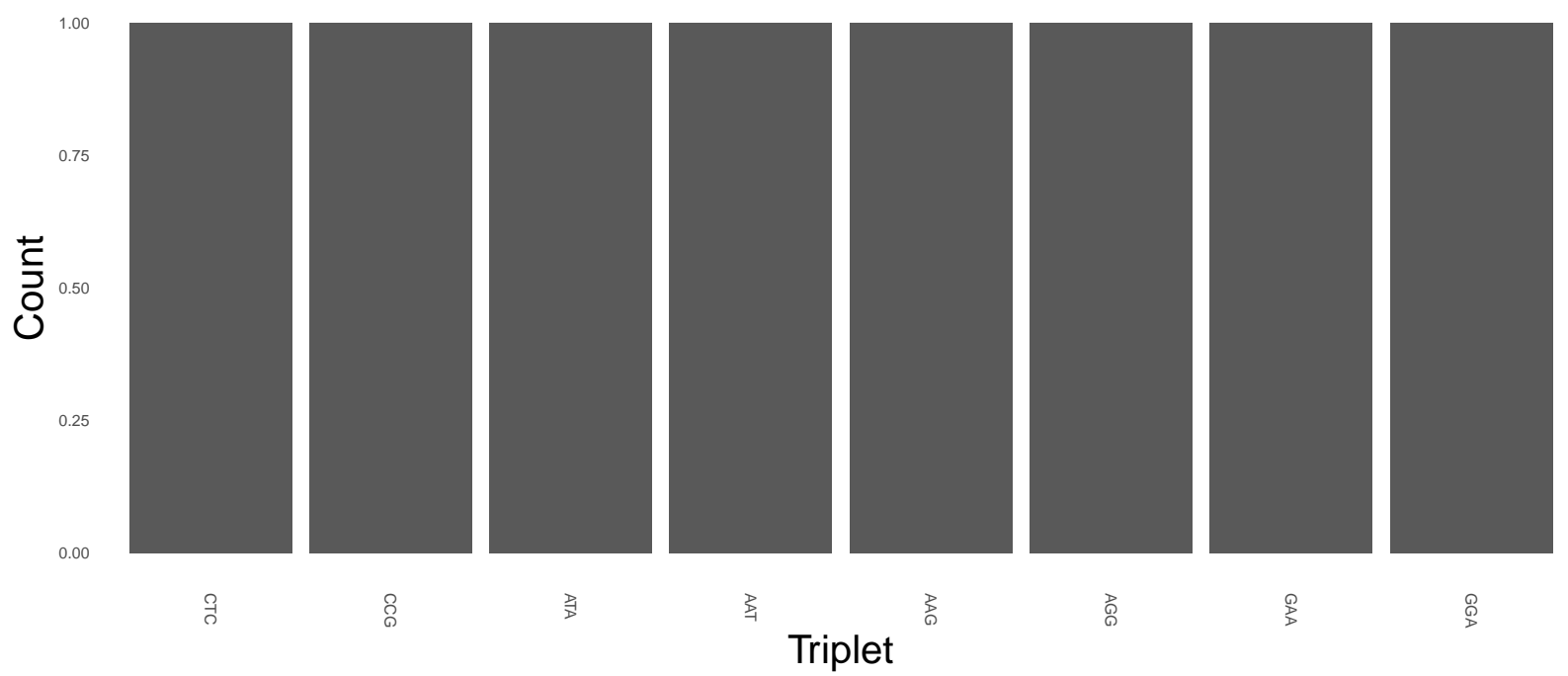
TRBV4–3*01_C274T– Final 3 nucleotides as a triplet



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

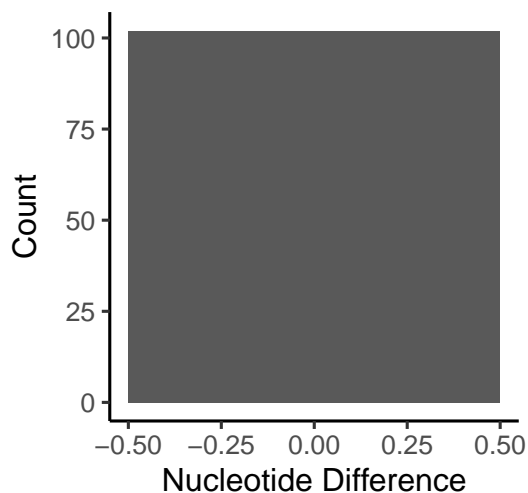


TRBV7-7*01_C315T- Final 3 nucleotides as a triplet



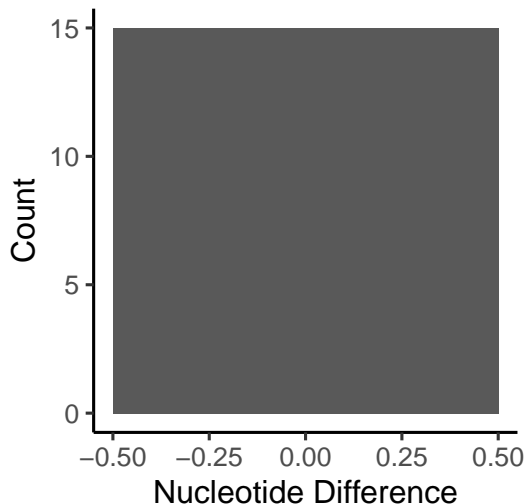
TRBV20–1*01

102 sequences assigned
102 (100%) exact matches, in which:
102 unique CDR3
10 unique J



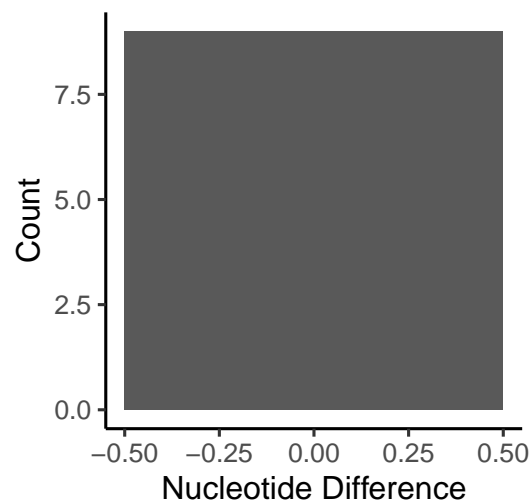
TRBV10–2*01

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



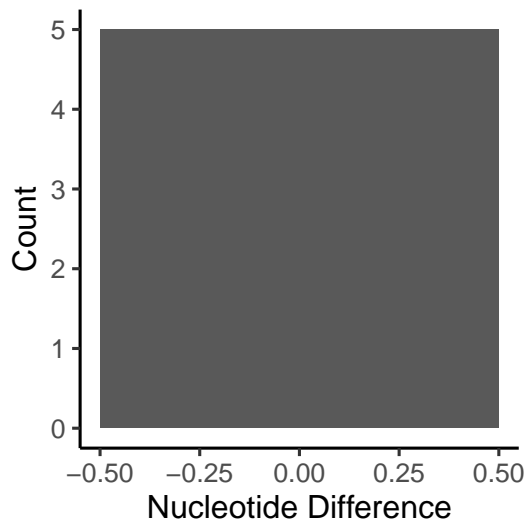
TRBV14*01

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



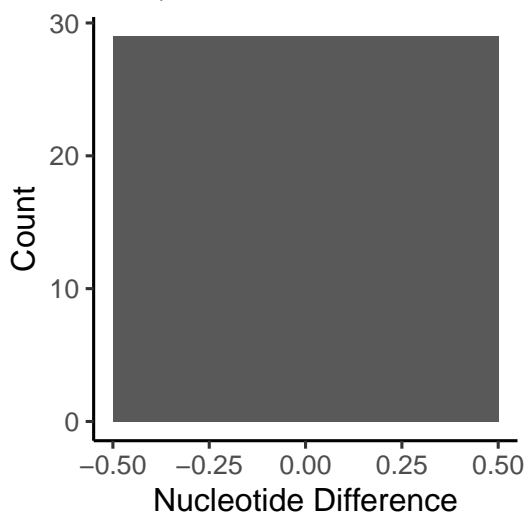
TRBV10–1*01

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



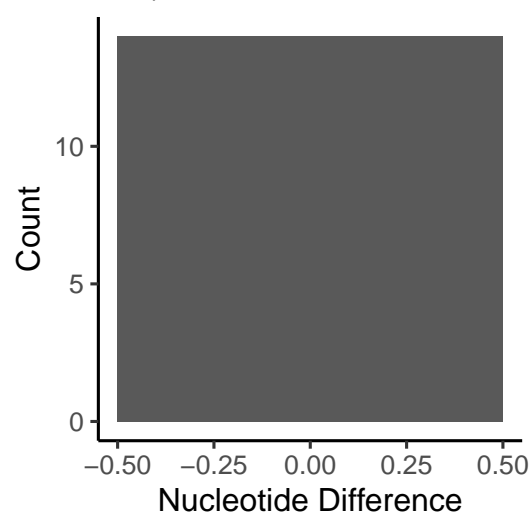
TRBV10–3*03_313GCCA1

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



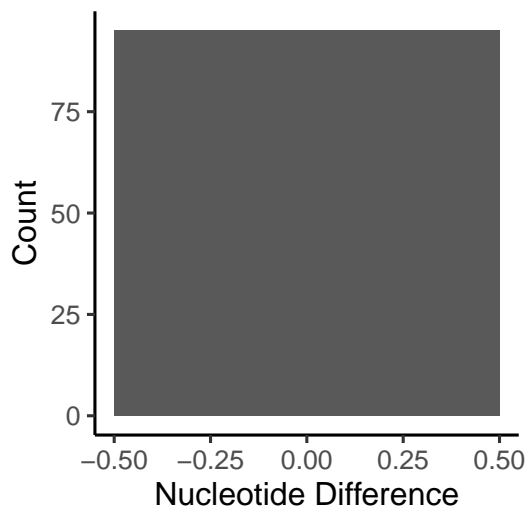
TRBV14*02_322CAAGA32

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



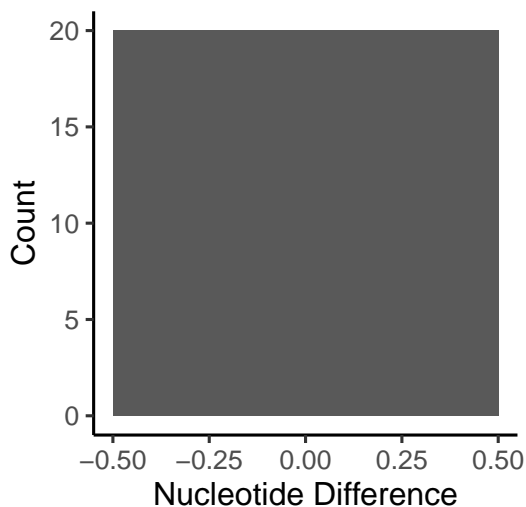
TRBV20–1*05

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



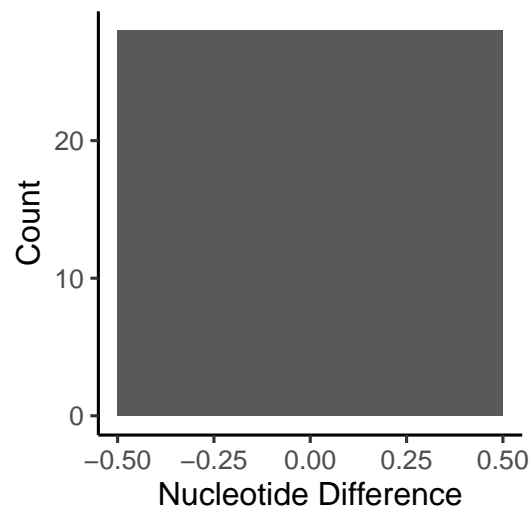
TRBV13*01

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



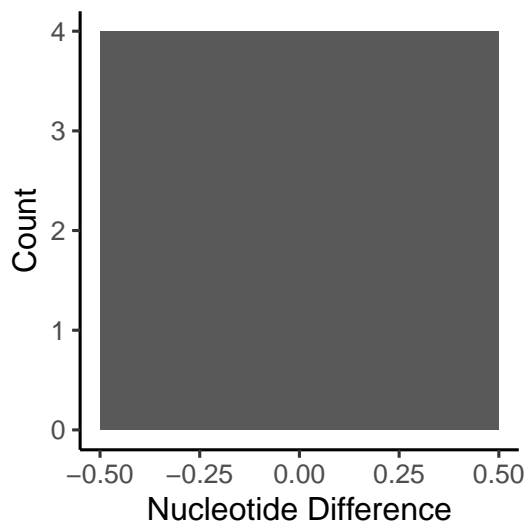
TRBV15*02

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



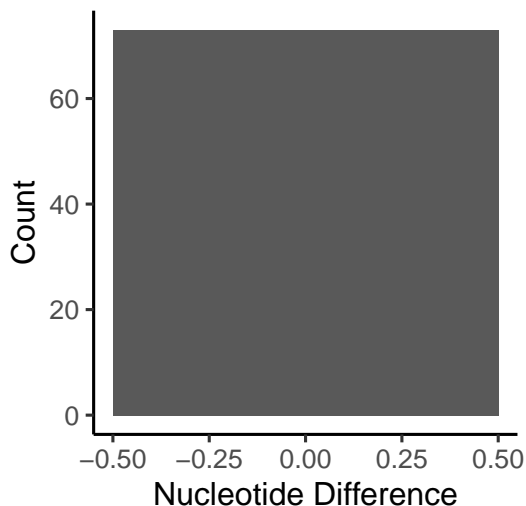
TRBV16*01

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



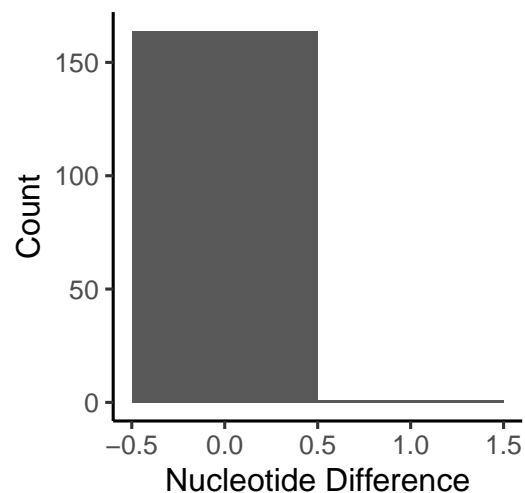
TRBV19*01_A24G

73 sequences assigned
73 (100%) exact matches, in which:
73 unique CDR3
13 unique J



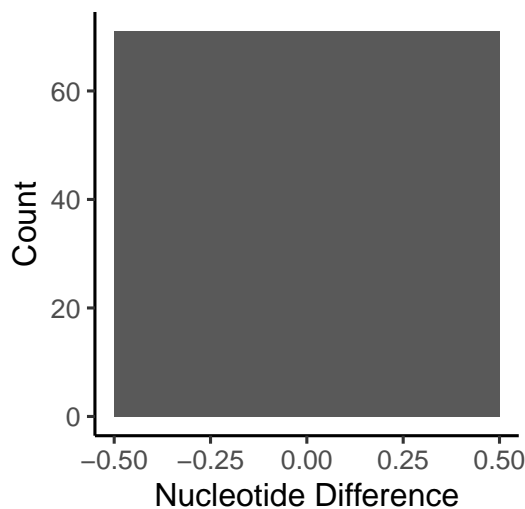
TRBV28*01

165 sequences assigned
164 (99.4%) exact matches, in which:
163 unique CDR3
14 unique J



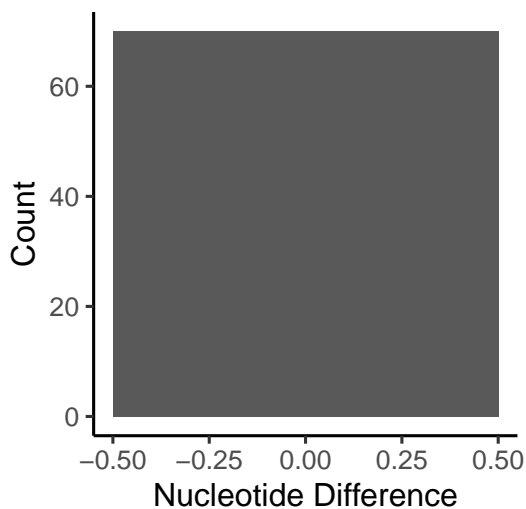
TRBV18*01

71 sequences assigned
71 (100%) exact matches, in which:
71 unique CDR3
14 unique J



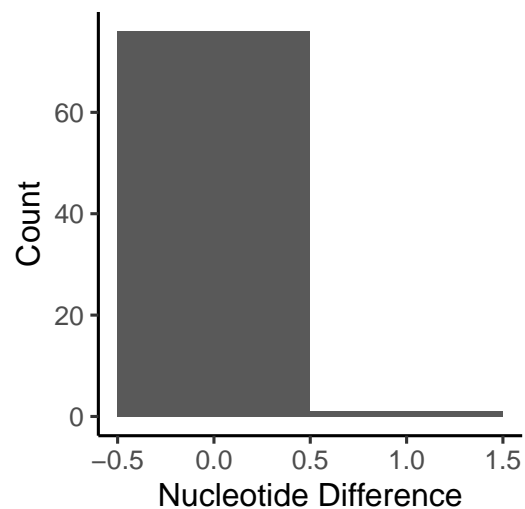
TRBV2*01

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



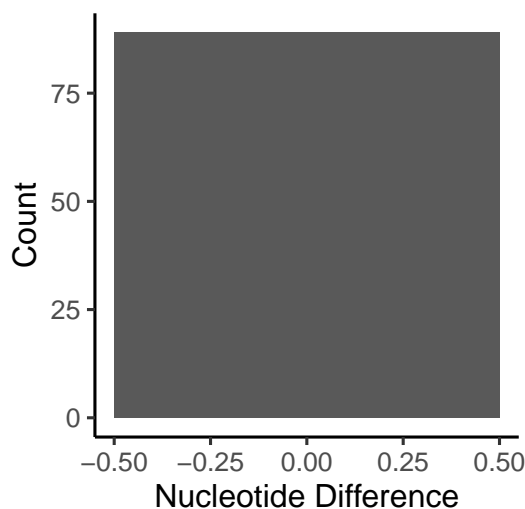
TRBV30*01

77 sequences assigned
76 (98.7%) exact matches, in which:
76 unique CDR3
13 unique J



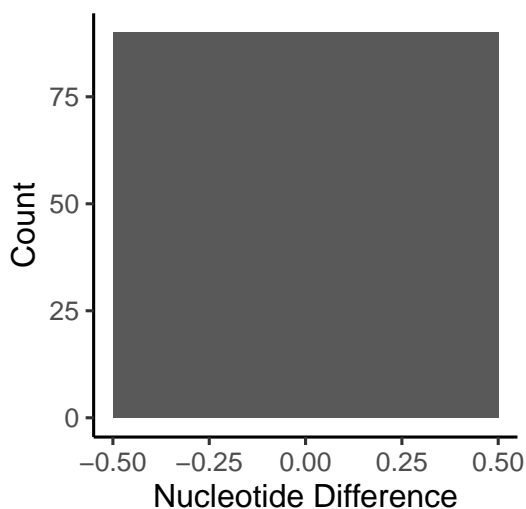
TRBV19*01

89 sequences assigned
89 (100%) exact matches, in which:
89 unique CDR3
13 unique J



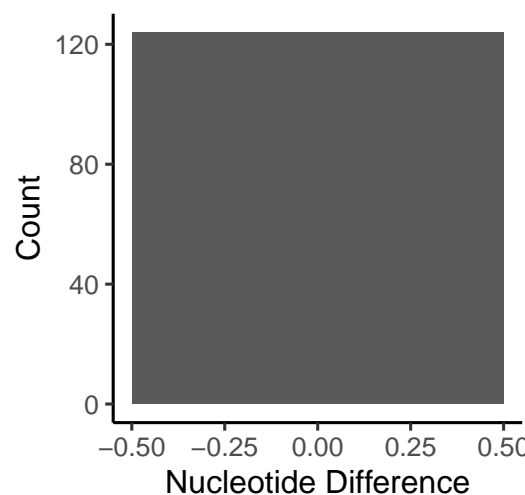
TRBV27*01

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



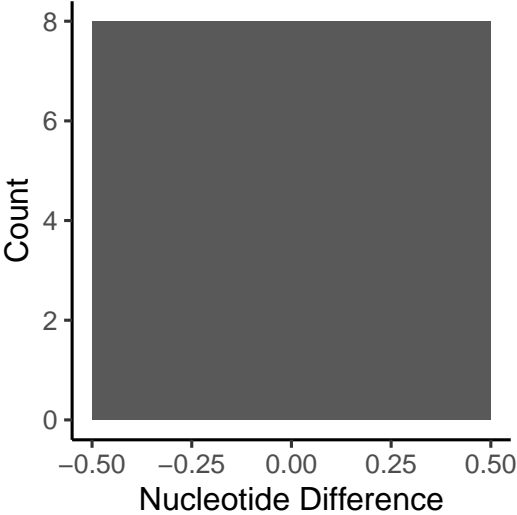
TRBV9*01

124 sequences assigned
124 (100%) exact matches, in which:
124 unique CDR3
14 unique J



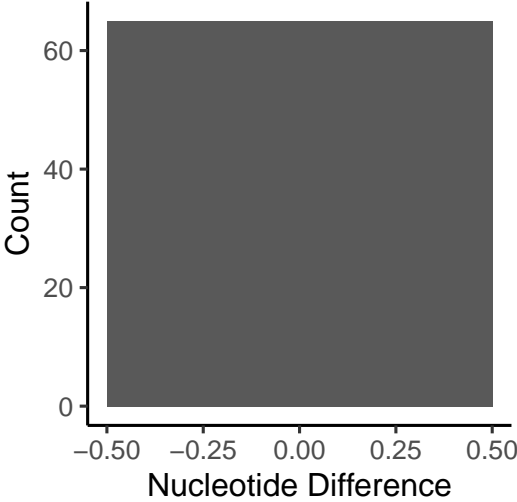
TRBV11–1*01

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



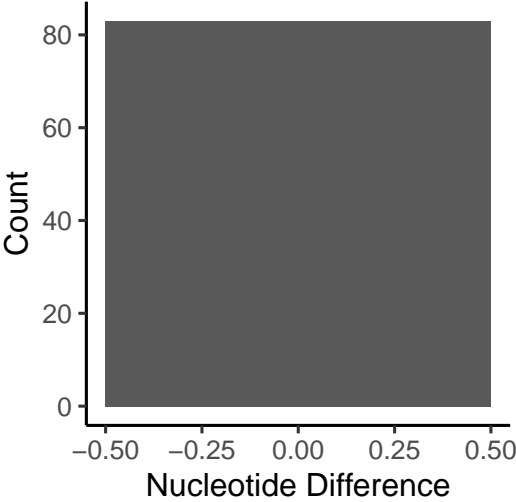
TRBV12–3*01

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



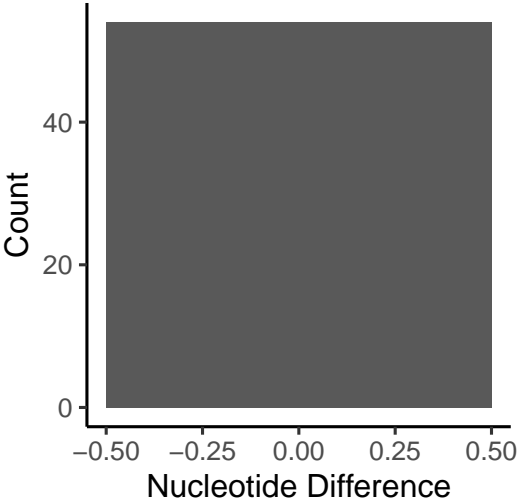
TRBV3–1*01

83 sequences assigned
83 (100%) exact matches, in which:
83 unique CDR3
13 unique J



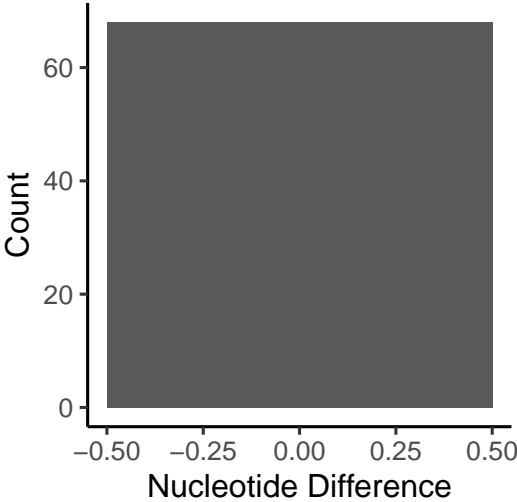
TRBV11–2*01

54 sequences assigned
54 (100%) exact matches, in which:
54 unique CDR3
13 unique J



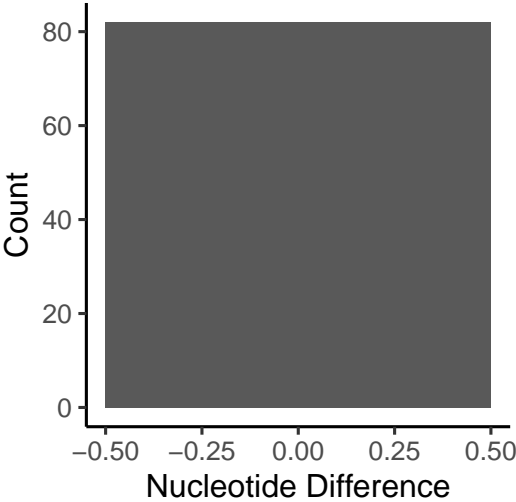
TRBV12–4*01_C87T

68 sequences assigned
68 (100%) exact matches, in which:
68 unique CDR3
11 unique J



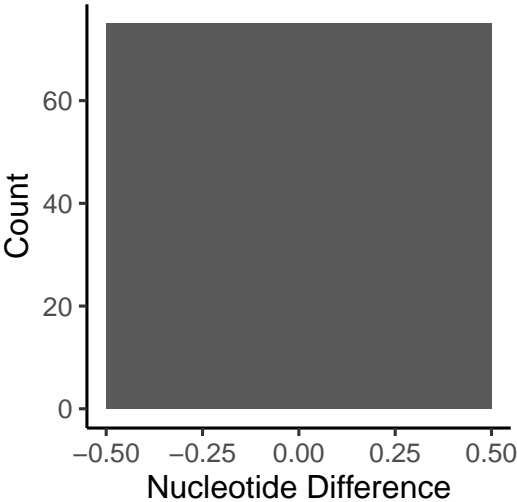
TRBV4–1*01

82 sequences assigned
82 (100%) exact matches, in which:
82 unique CDR3
14 unique J



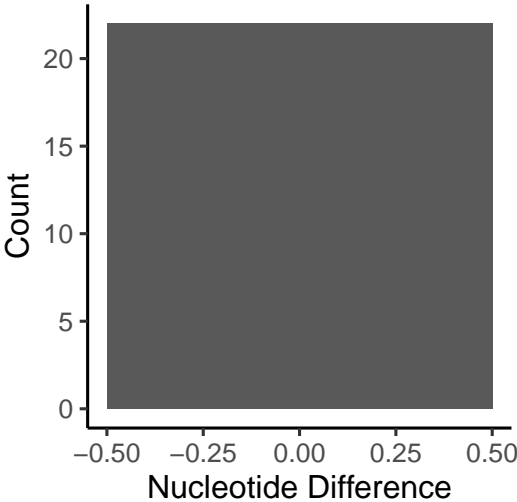
TRBV11–3*04

75 sequences assigned
75 (100%) exact matches, in which:
75 unique CDR3
12 unique J



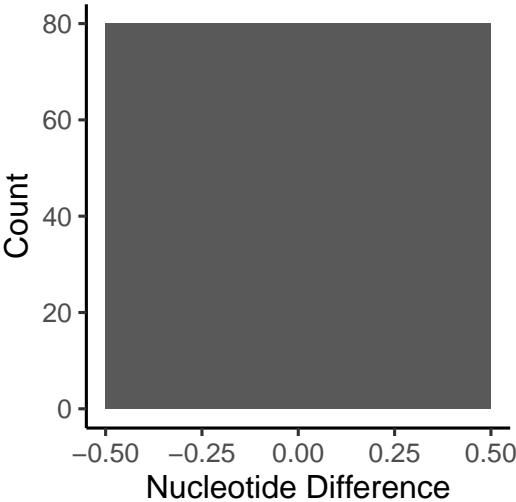
TRBV12–5*01

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



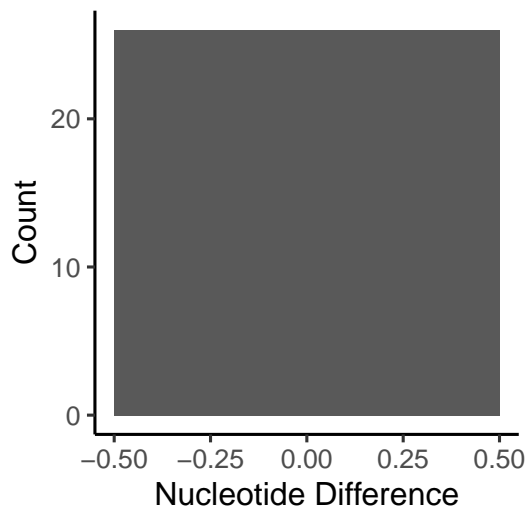
TRBV24–1*01

80 sequences assigned
80 (100%) exact matches, in which:
80 unique CDR3
10 unique J



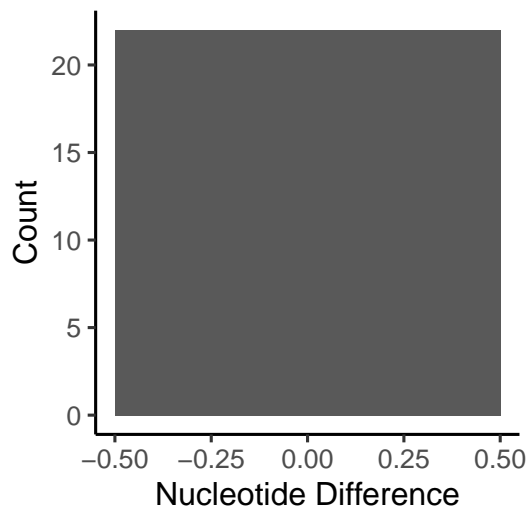
TRBV4–2*01

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



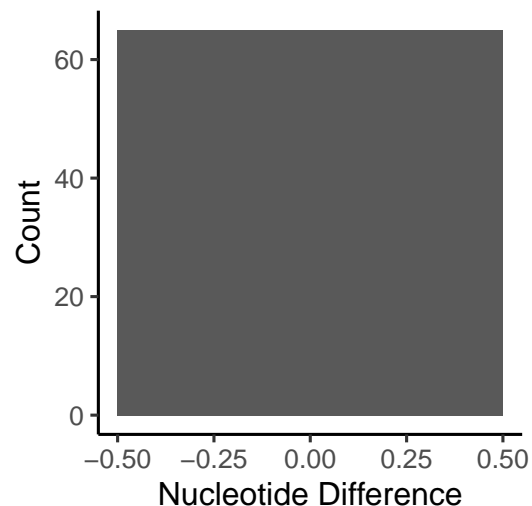
TRBV25–1*01

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



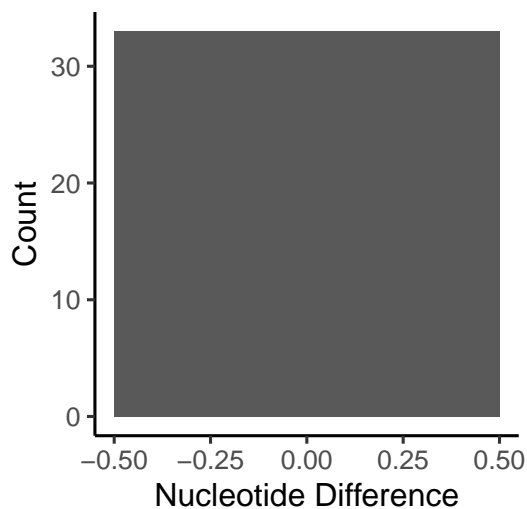
TRBV5–6*01

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



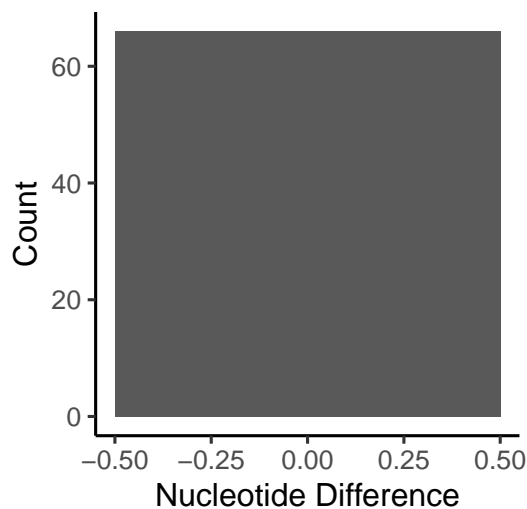
TRBV4–3*01_C274T

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



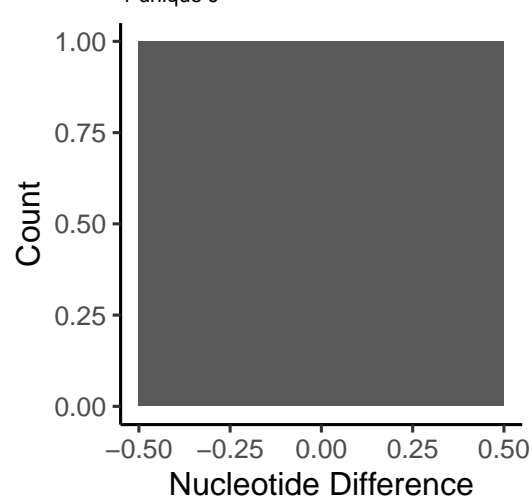
TRBV5–4*01

66 sequences assigned
66 (100%) exact matches, in which:
66 unique CDR3
12 unique J



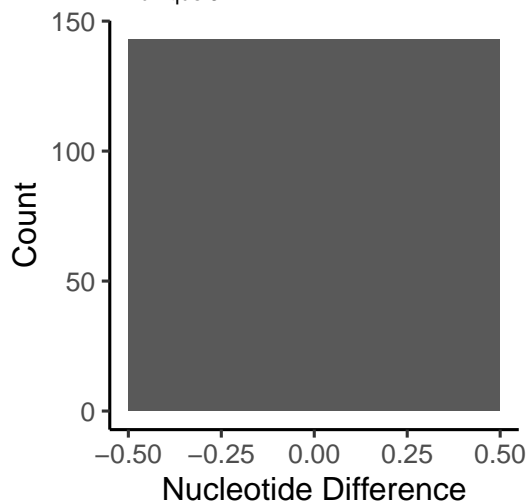
TRBV5–7*01

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



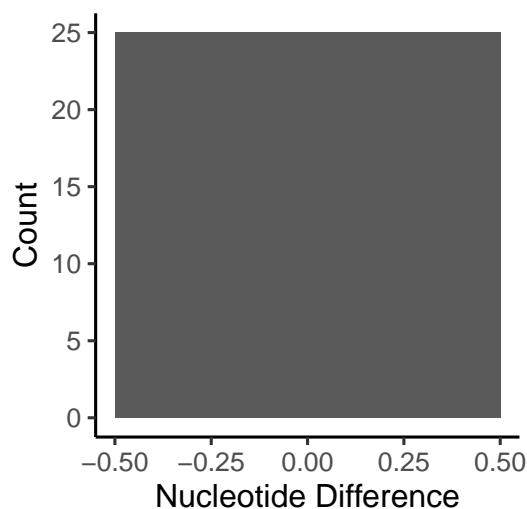
TRBV5–1*01

143 sequences assigned
143 (100%) exact matches, in which:
143 unique CDR3
14 unique J



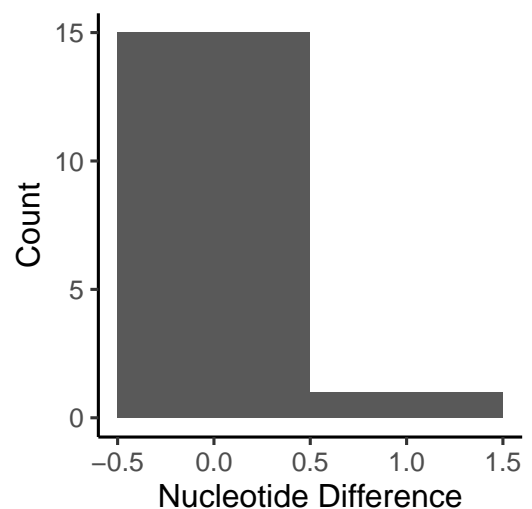
TRBV5–5*02_322TTGG32

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



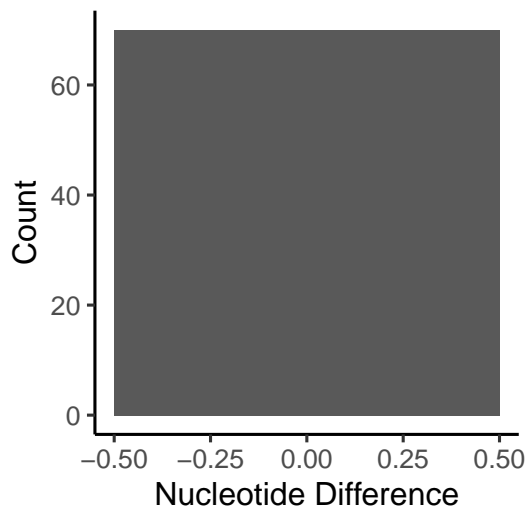
TRBV5–8*01

16 sequences assigned
15 (93.8%) exact matches, in which:
15 unique CDR3
9 unique J



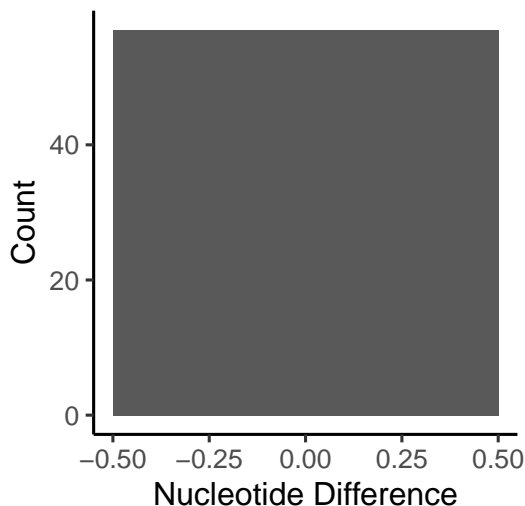
TRBV6-1*01

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



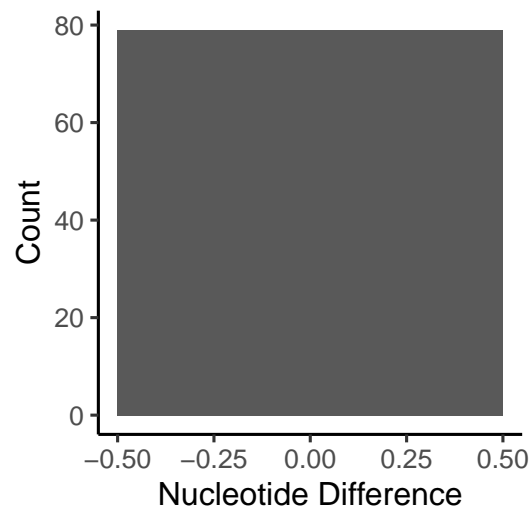
TRBV6-6*01

57 sequences assigned
57 (100%) exact matches, in which:
57 unique CDR3
14 unique J



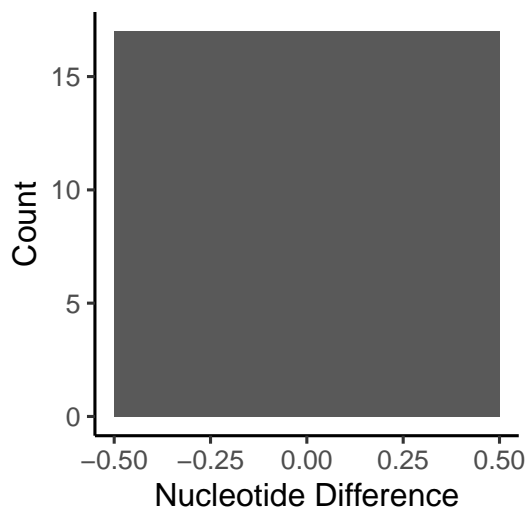
TRBV6-23*01

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



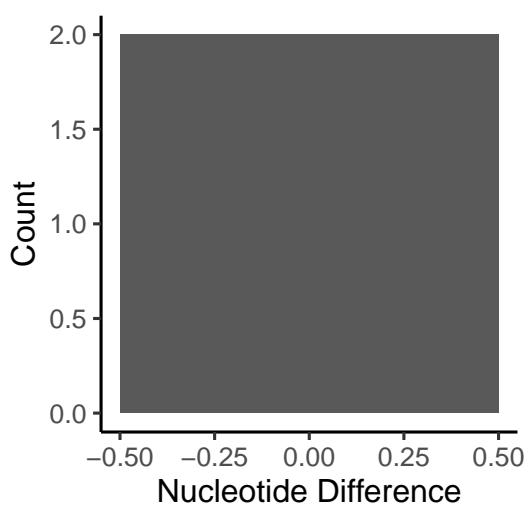
TRBV6-4*01

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



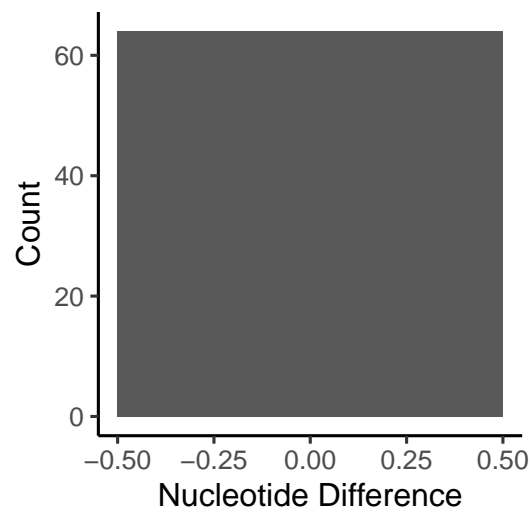
TRBV6-7*01

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



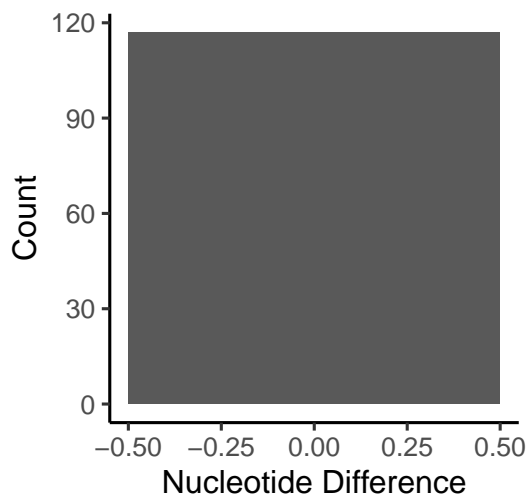
TRBV7-2*01

64 sequences assigned
64 (100%) exact matches, in which:
64 unique CDR3
12 unique J



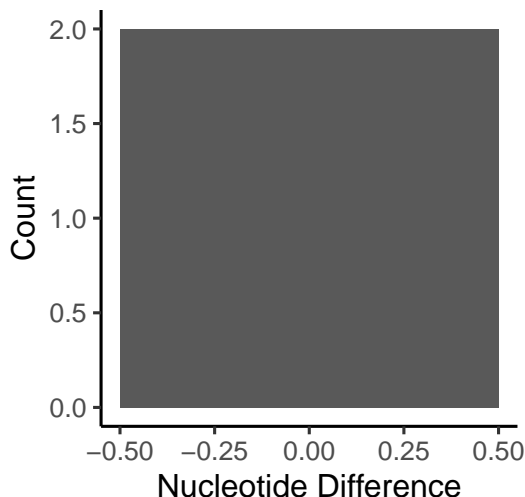
TRBV6-5*01

117 sequences assigned
117 (100%) exact matches, in which:
117 unique CDR3
14 unique J



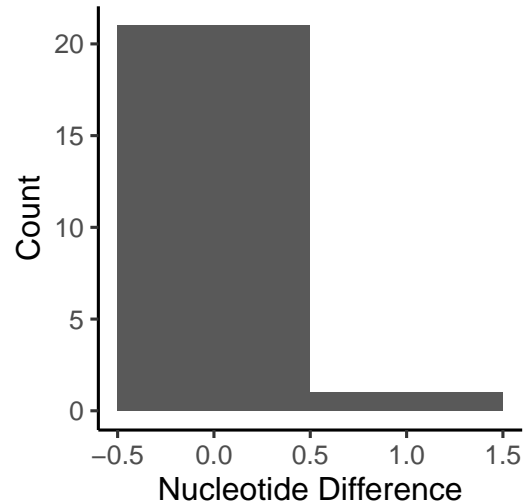
TRBV6-8*01

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



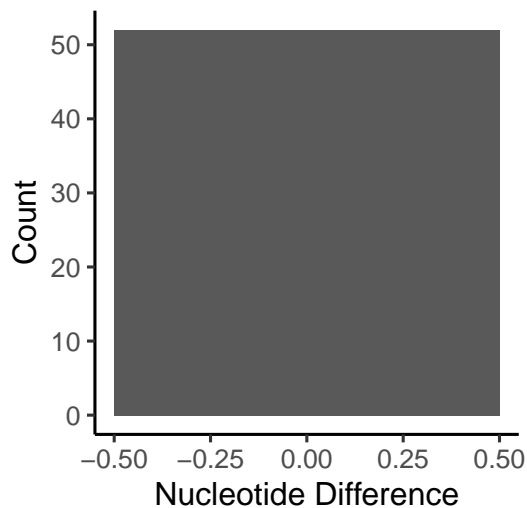
TRBV7-2*02

22 sequences assigned
21 (95.5%) exact matches, in which:
21 unique CDR3
10 unique J



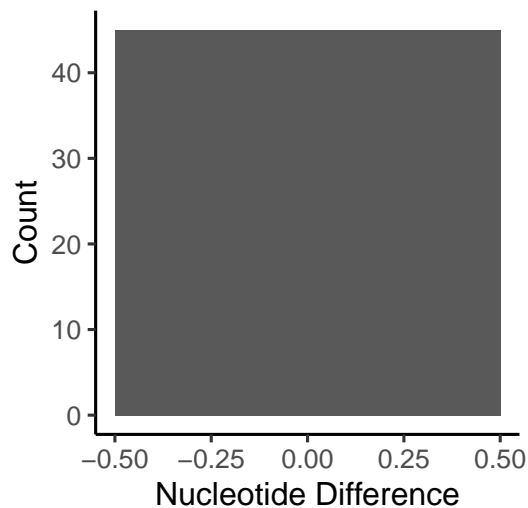
TRBV7-3*01

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



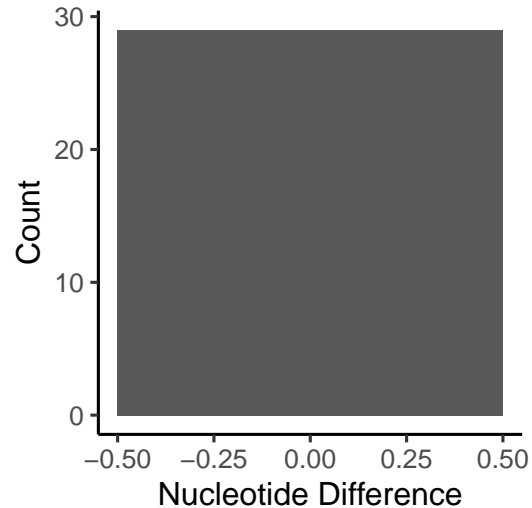
TRBV7-8*01

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



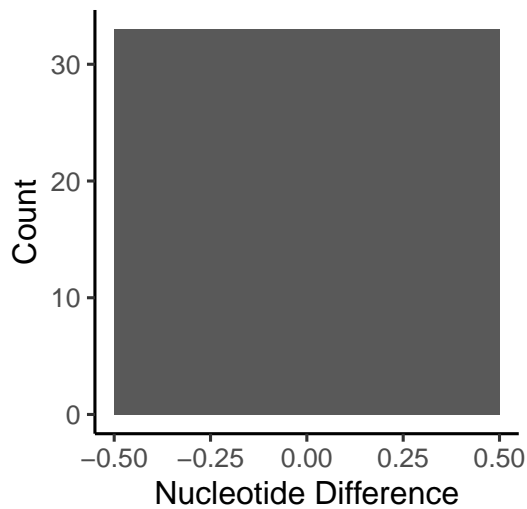
TRBV29-1*01_A280C

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



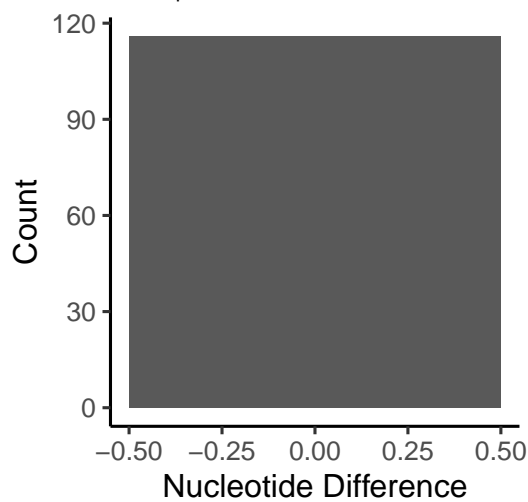
TRBV7-6*01

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



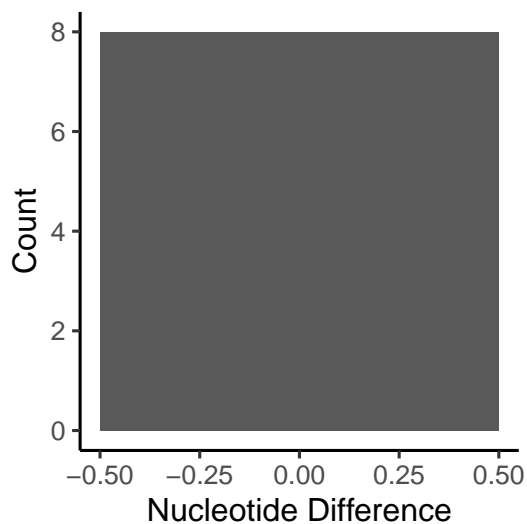
TRBV7-9*03

116 sequences assigned
116 (100%) exact matches, in which:
116 unique CDR3
13 unique J



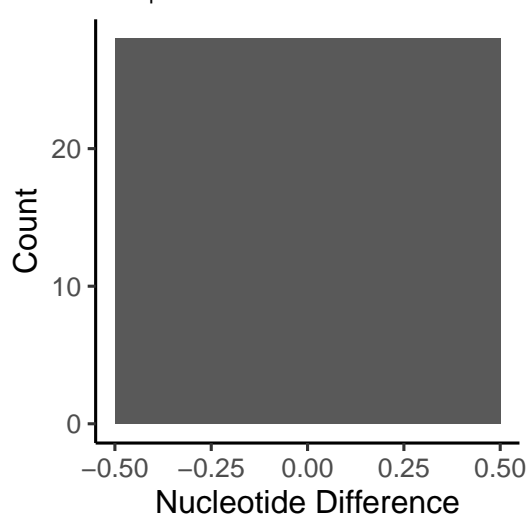
TRBV7-7*01_C315T

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

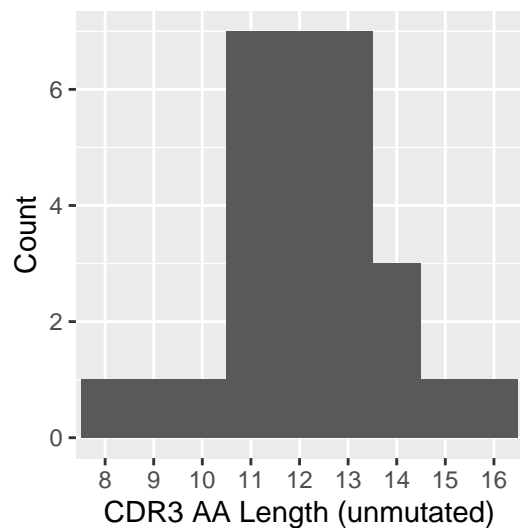


TRBV29-1*01

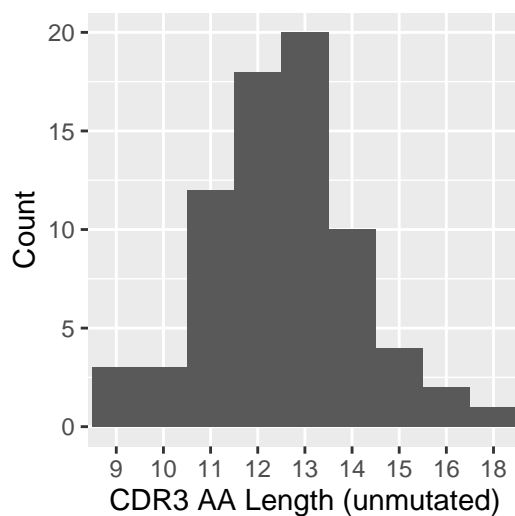
28 sequences assigned
28 (100%) exact matches, in which:
28 unique CDR3
9 unique J



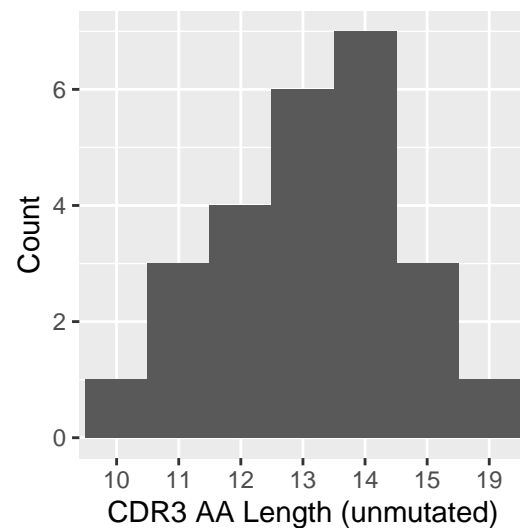
TRBV10-3*03_313GCCATCA



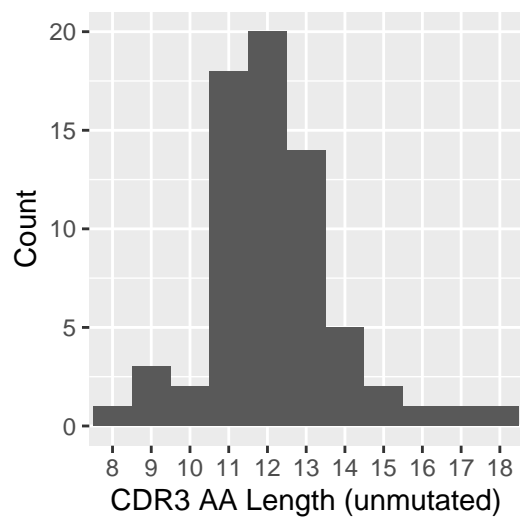
TRBV19*01_A24G



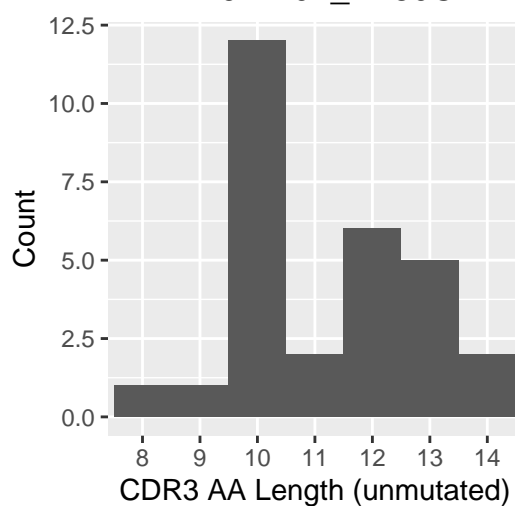
TRBV5-5*02_322TTGG325



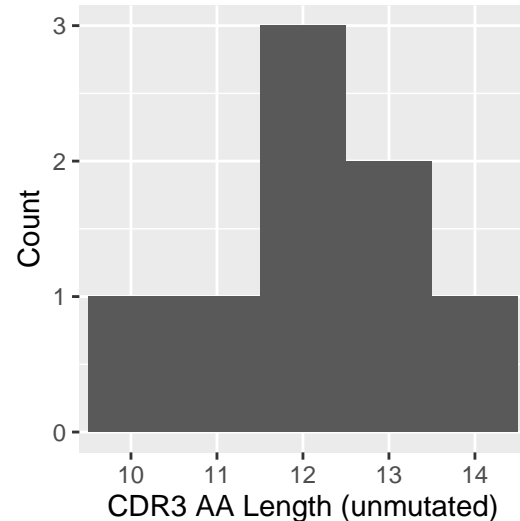
TRBV12-4*01_C87T



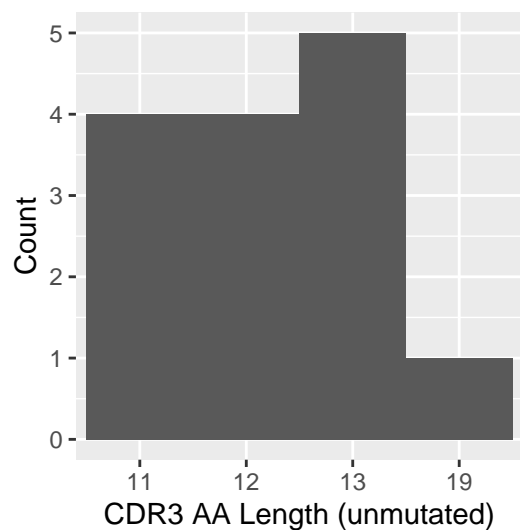
TRBV29-1*01_A280C



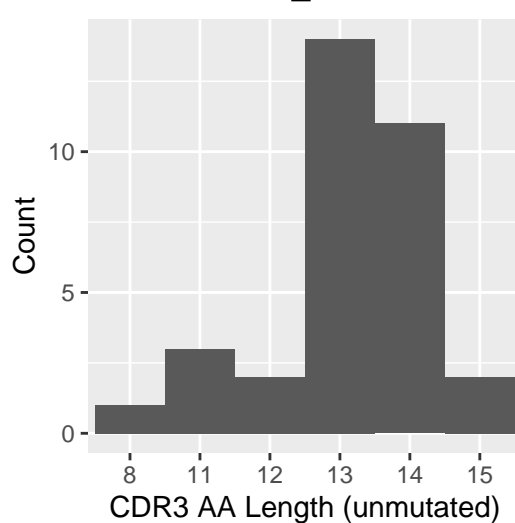
TRBV7-7*01_C315T

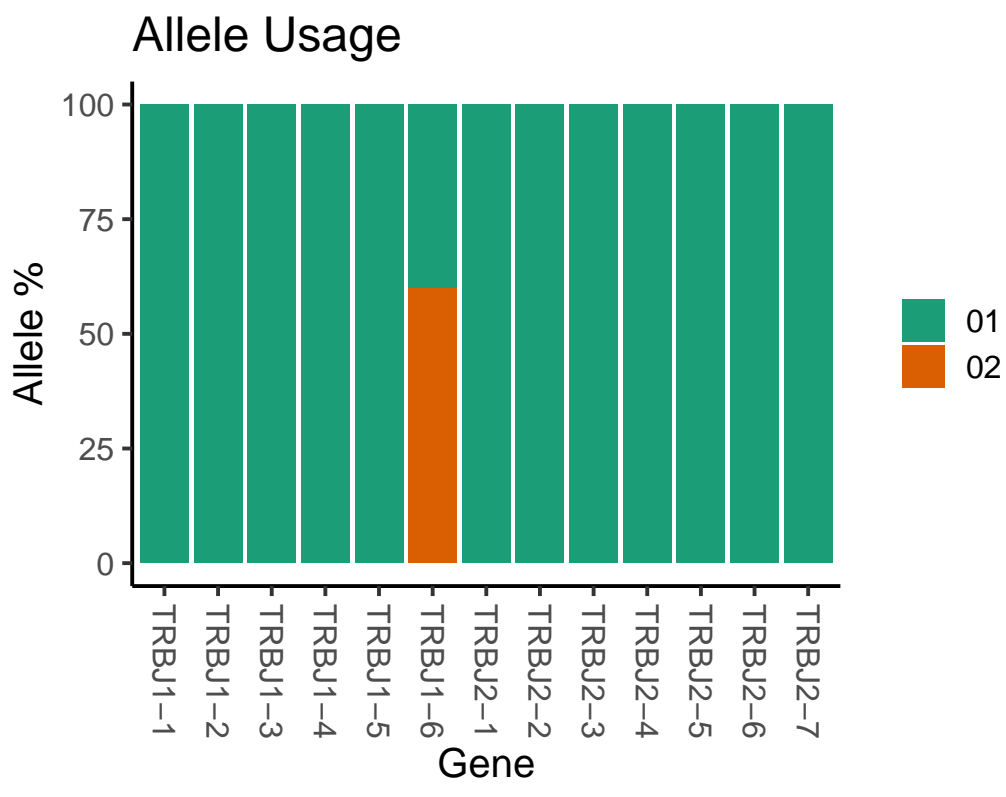


TRBV14*02_322CAAGA326

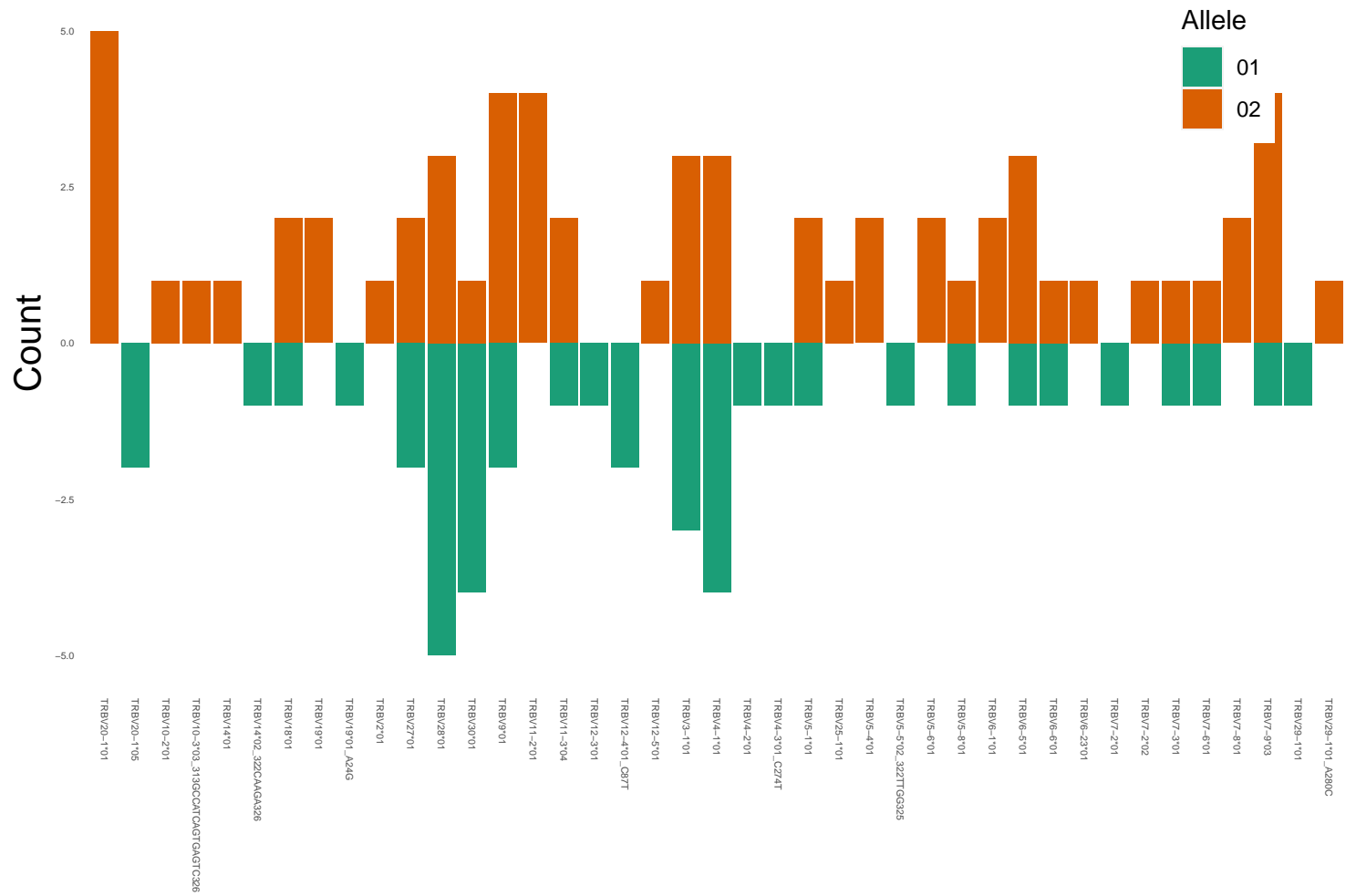


TRBV4-3*01_C274T





Sequence Count by TRBJ1–6 allele usage



Warning – no inferred sequences found.

Warning – no inferred sequences found.