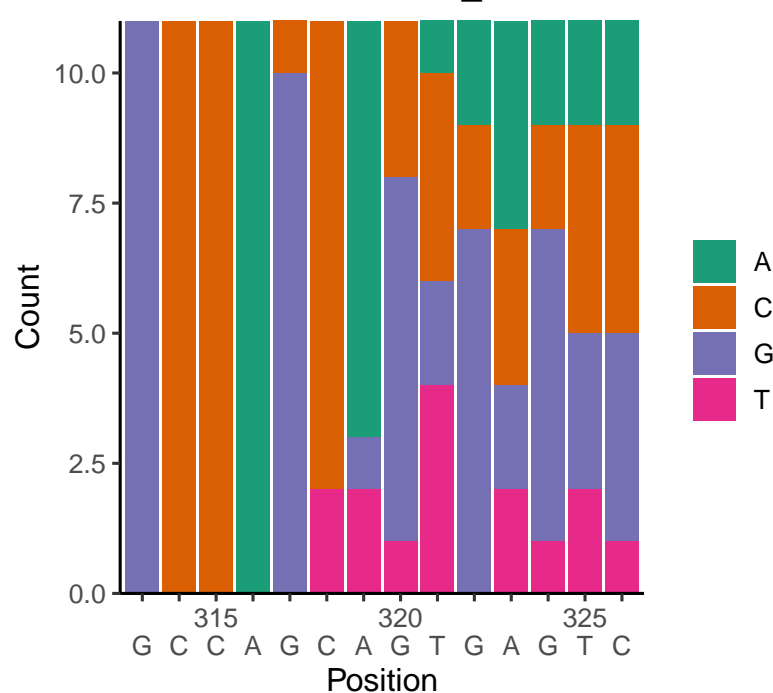
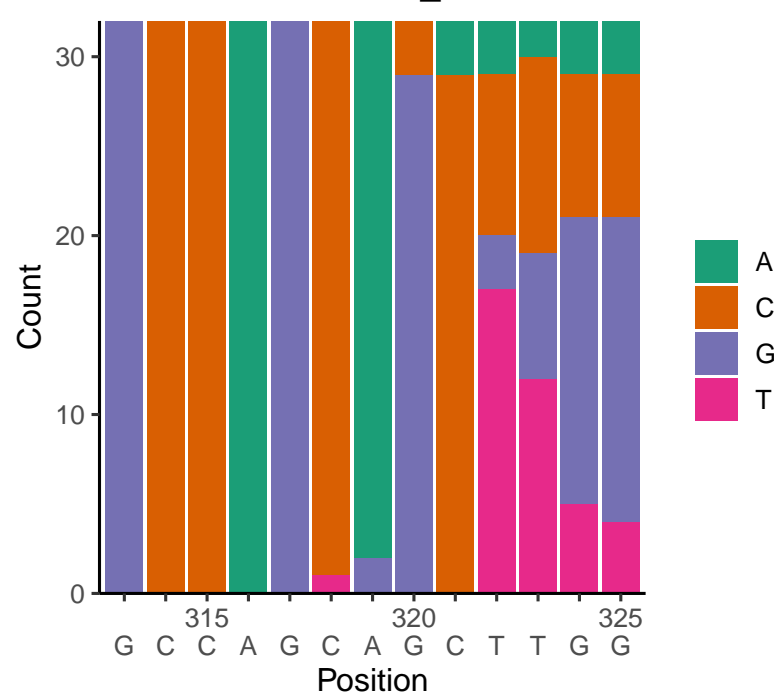


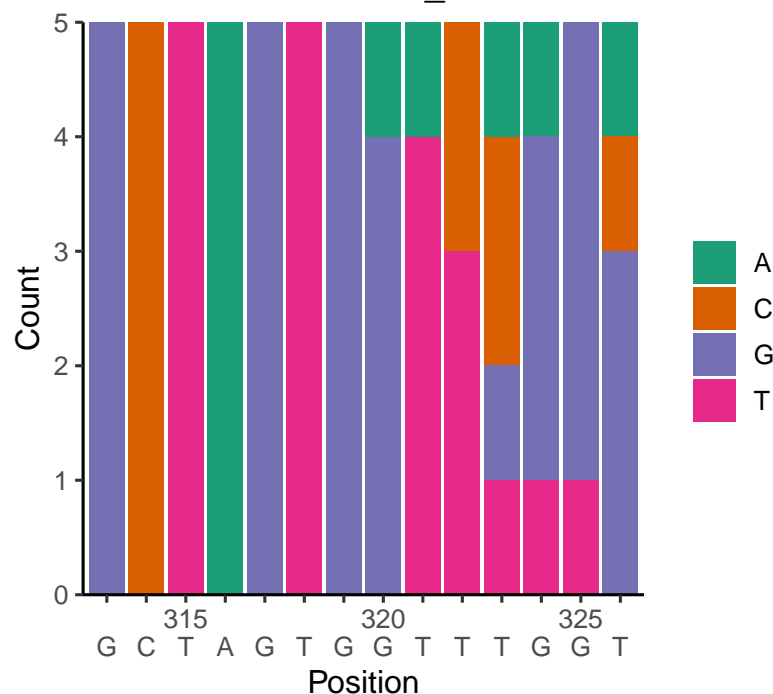
Gene TRBV10-2*01_C214T



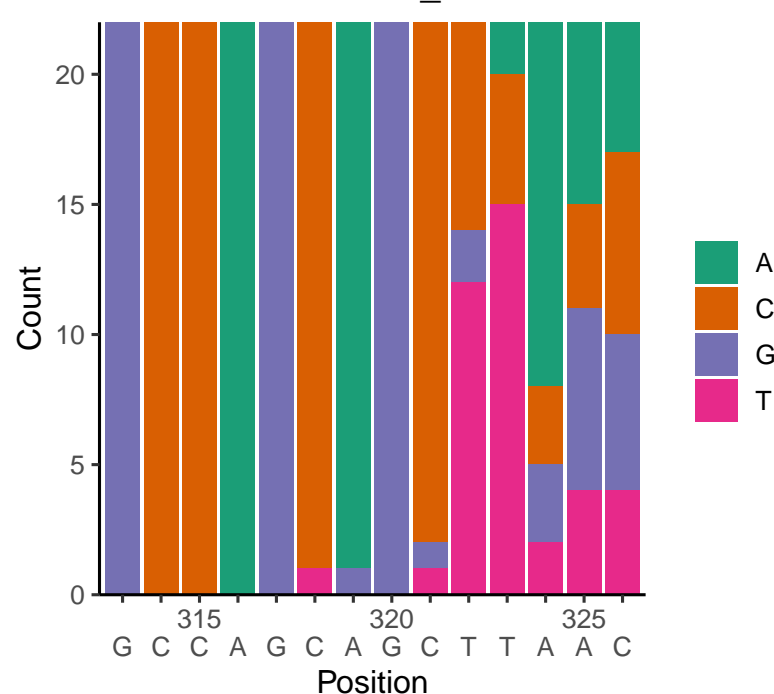
Gene TRBV5-5*02_322TTGG325



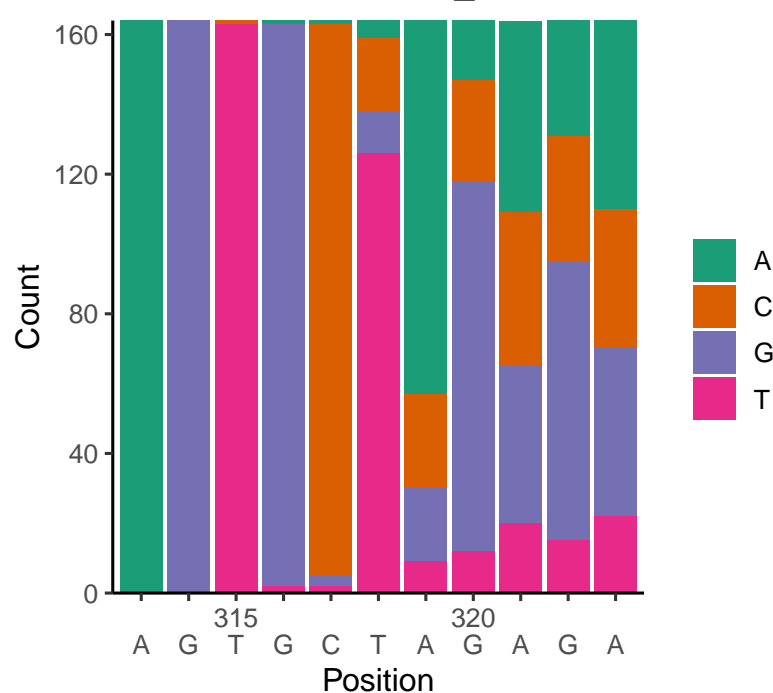
Gene TRBV12-5*01_C28G



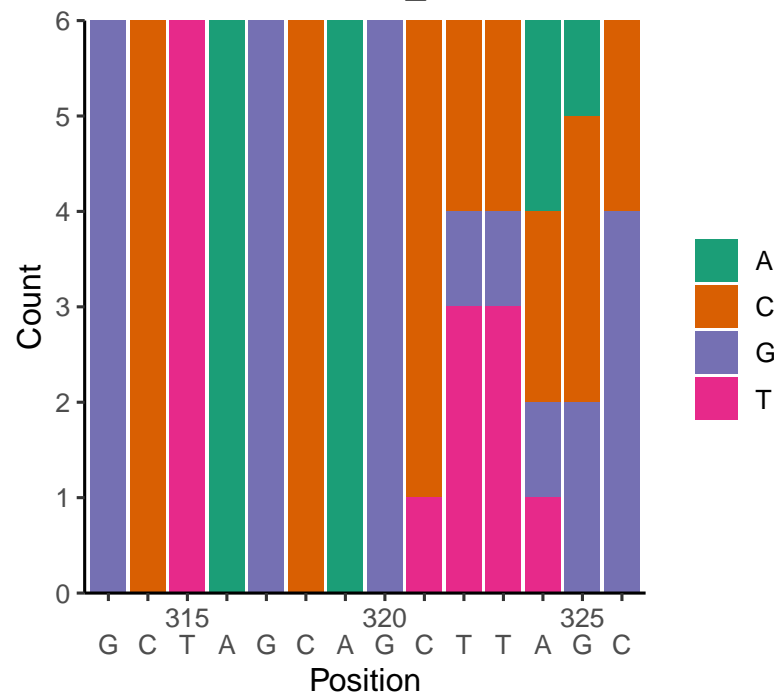
Gene TRBV7-3*01_G301A



Gene TRBV20-1*02_319AGAGA323



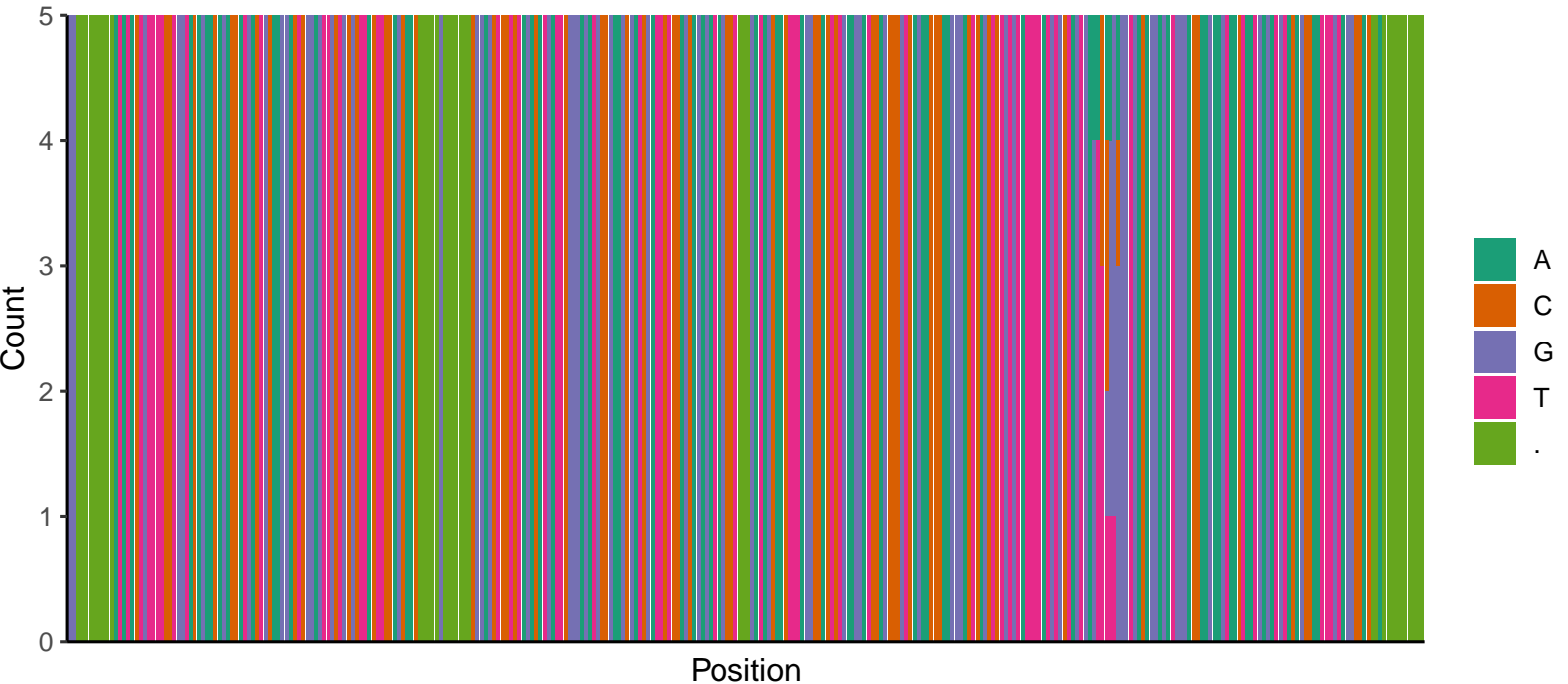
Gene TRBV7-7*01_C315T



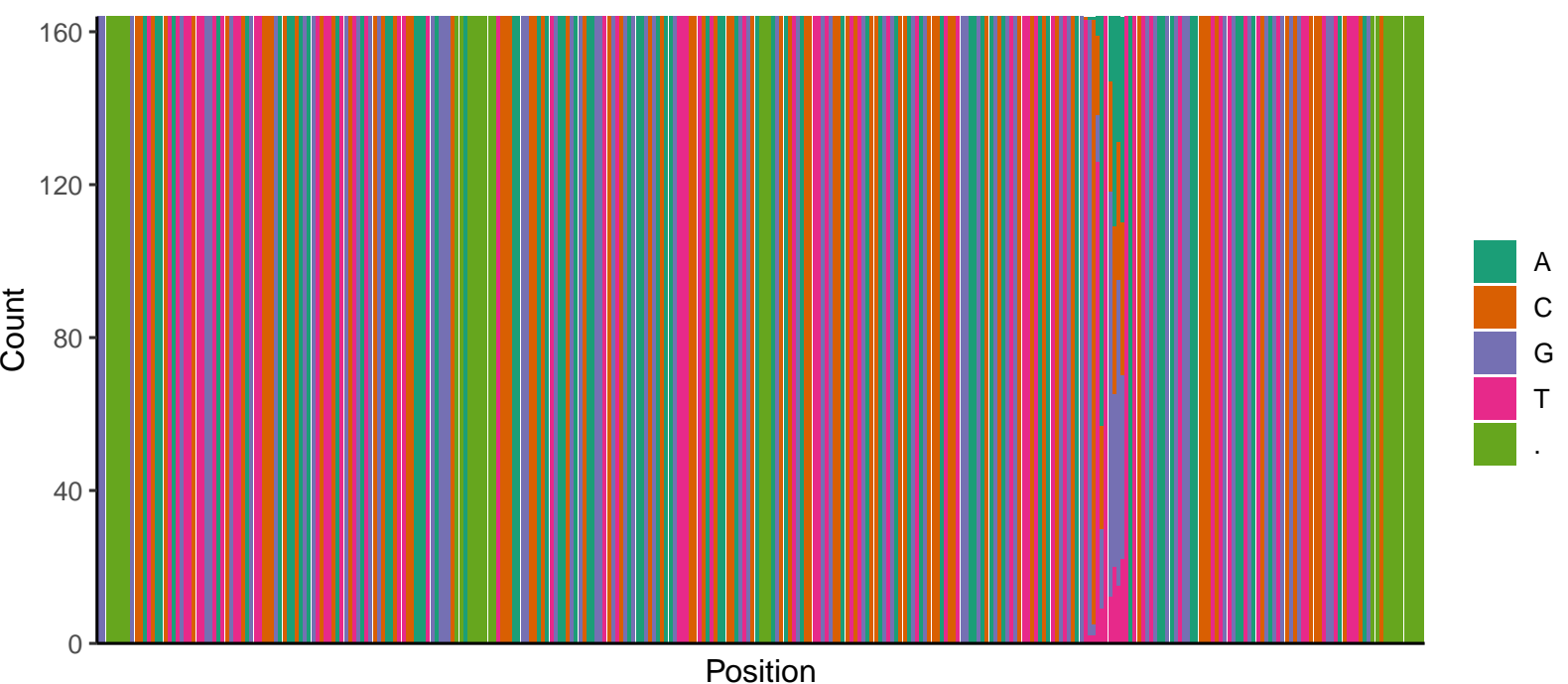
Gene TRBV10-2*01_C214T



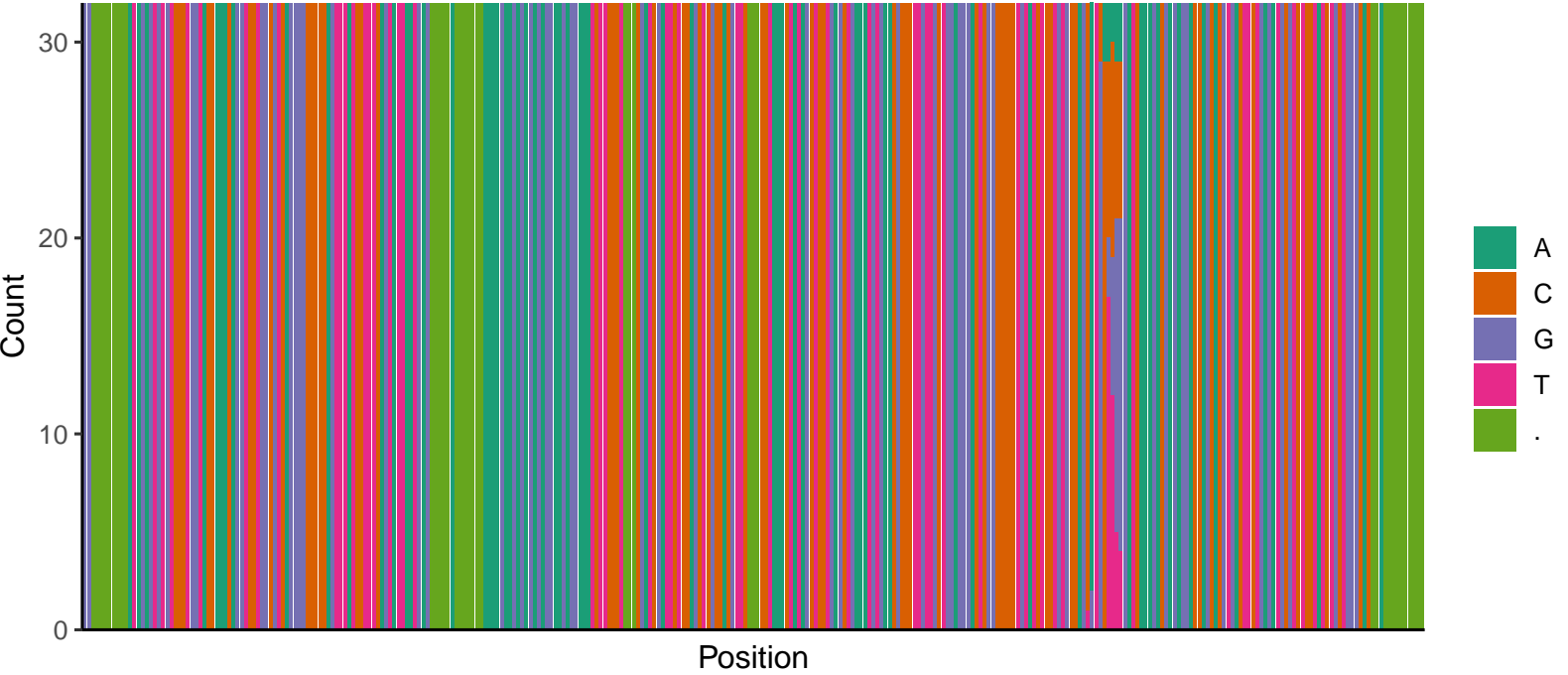
Gene TRBV12-5*01_C28G



Gene TRBV20-1*02_319AGAGA323



Gene TRBV5-5*02_322TTGG325



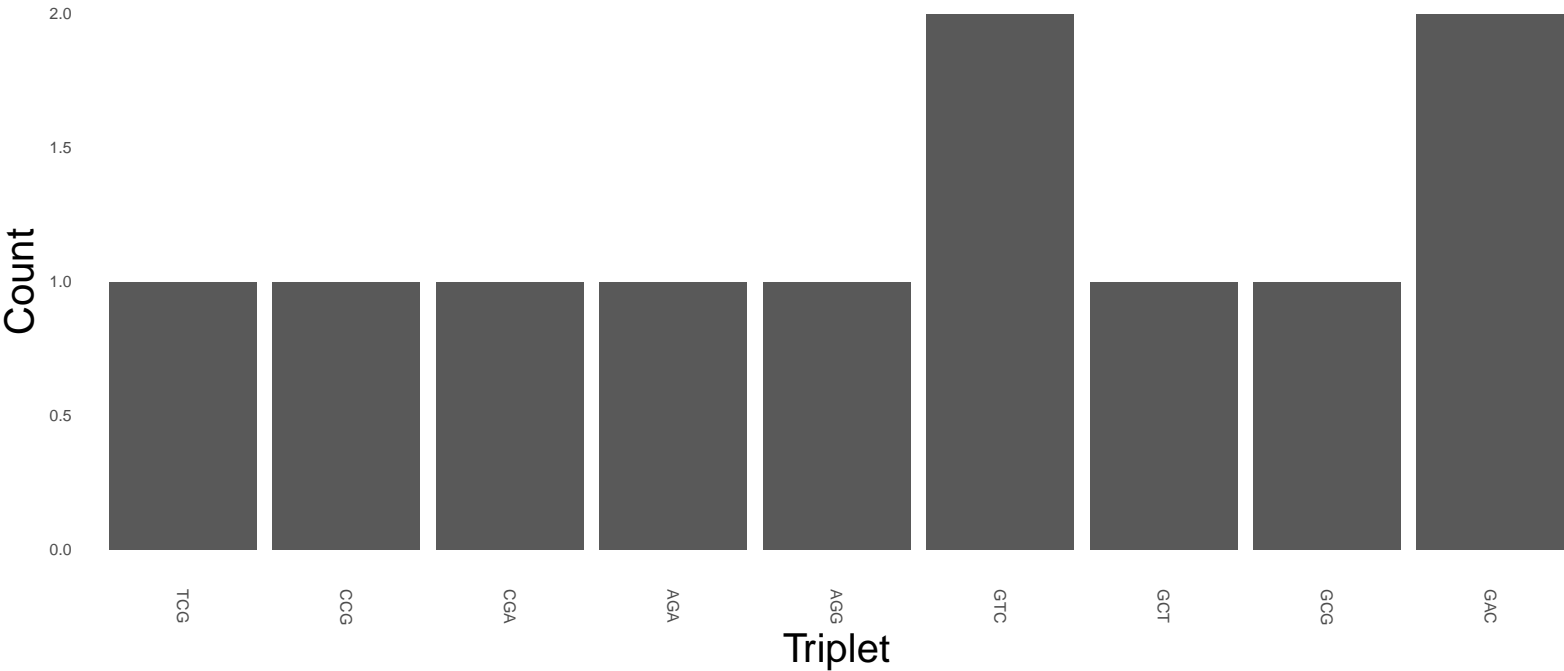
Gene TRBV7-3*01_G301A



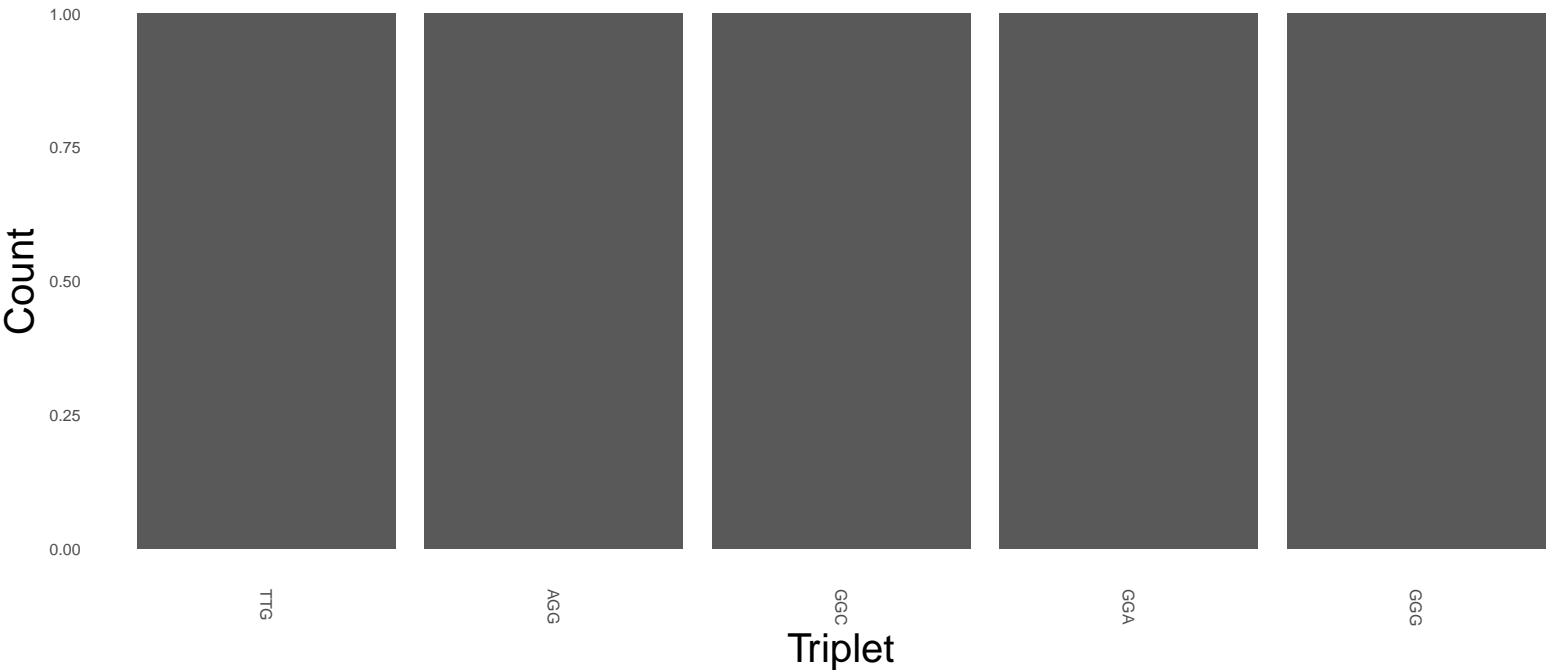
Gene TRBV7-7*01_C315T



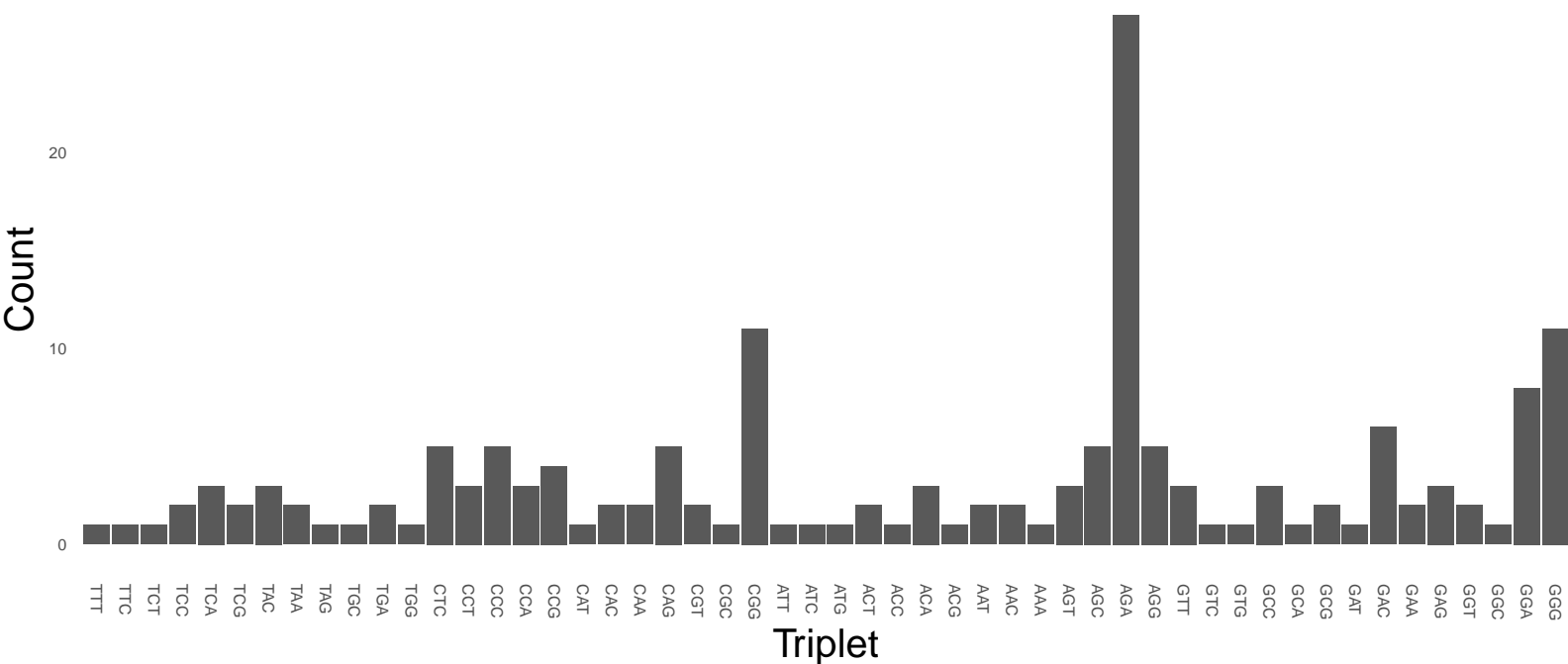
TRBV10–2*01_C214T– Final 3 nucleotides as a triplet



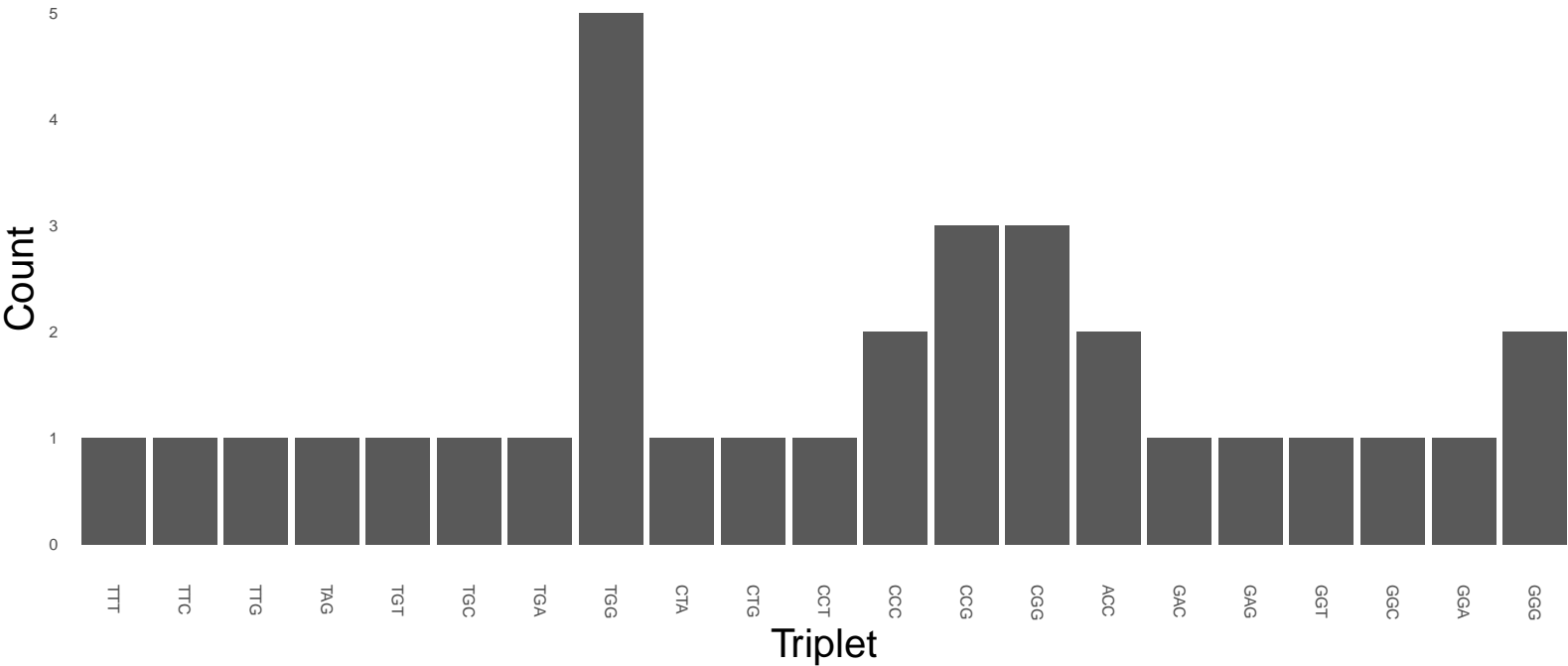
TRBV12–5*01_C28G– Final 3 nucleotides as a triplet



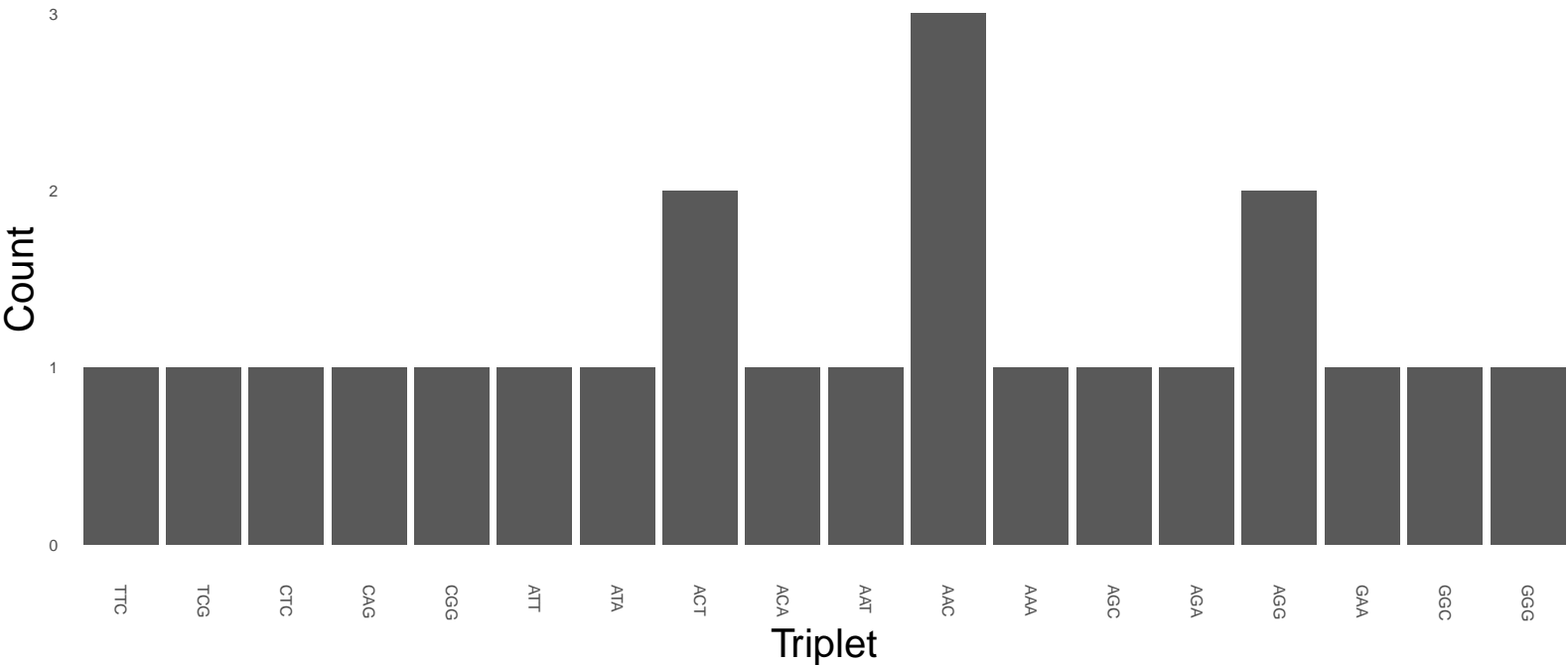
TRBV20–1*02_319AGAGA323– Final 3 nucleotides as a triplet



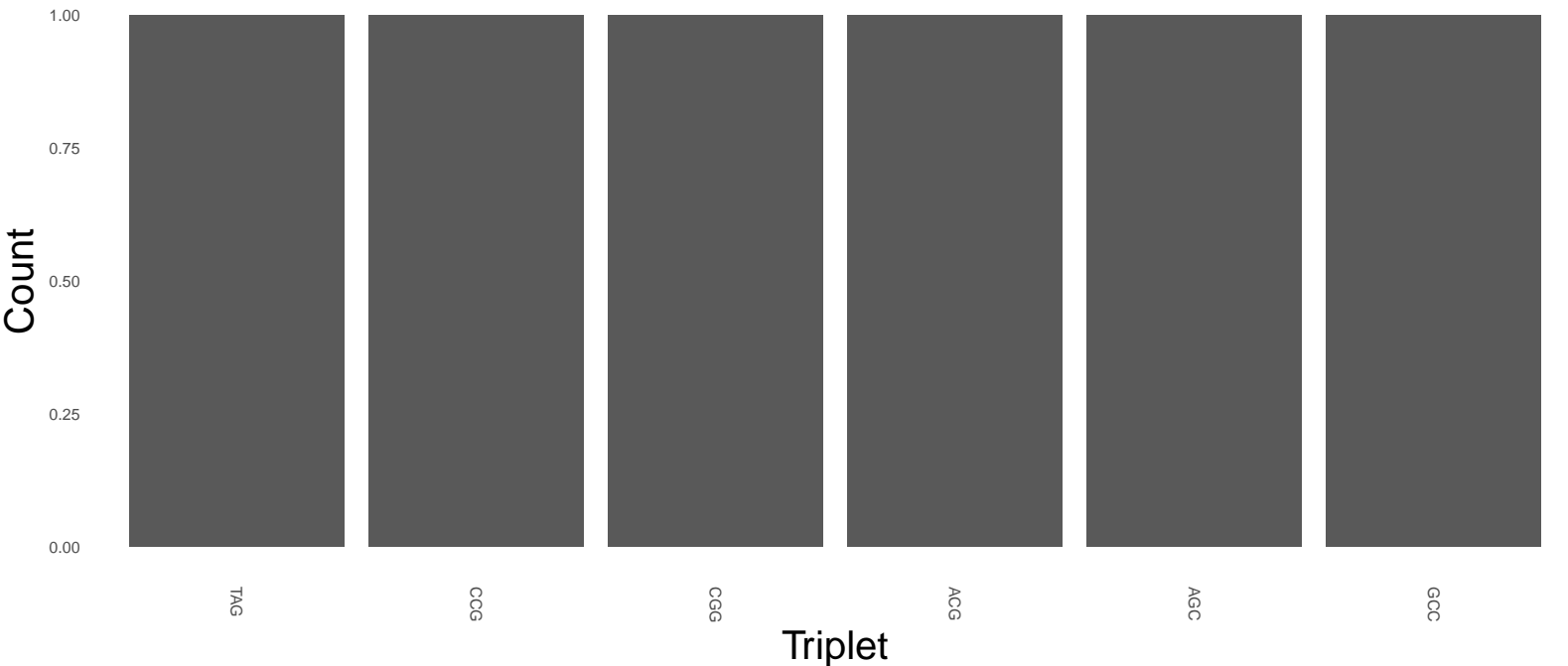
TRBV5–5*02_322TTGG325– Final 3 nucleotides as a triplet



TRBV7–3*01_G301A– Final 3 nucleotides as a triplet

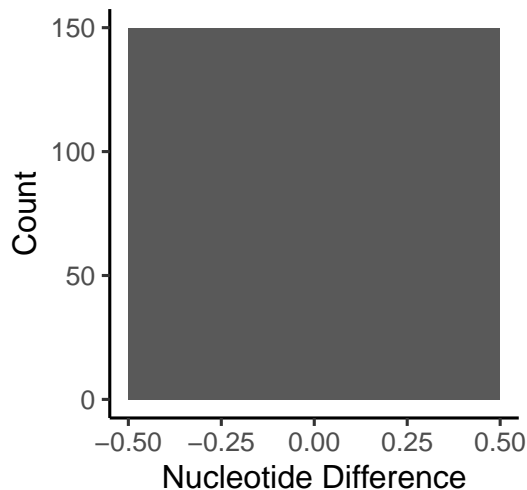


TRBV7–7*01_C315T– Final 3 nucleotides as a triplet



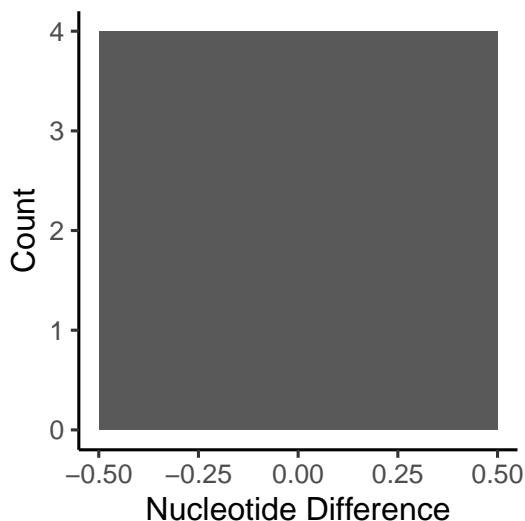
TRBV20–1*01

150 sequences assigned
150 (100%) exact matches, in which:
150 unique CDR3
13 unique J



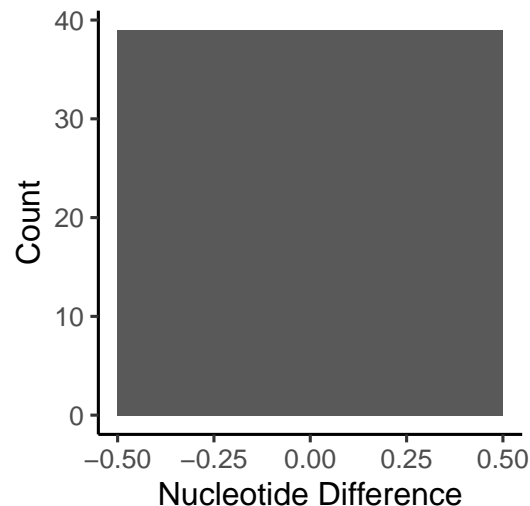
TRBV10–2*01

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



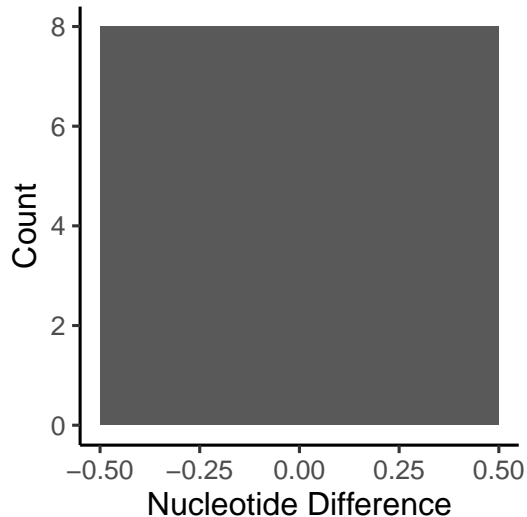
TRBV10–3*02

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



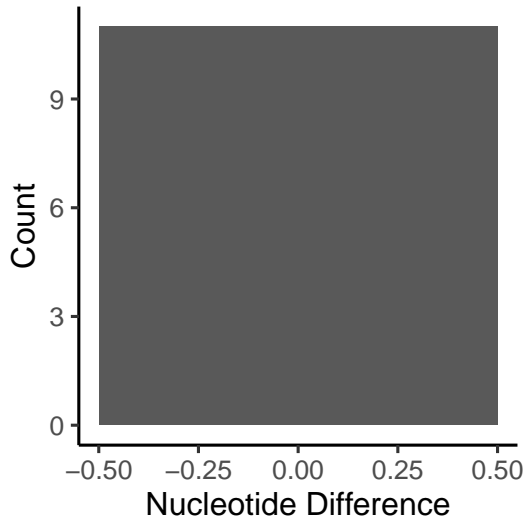
TRBV10–1*02

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



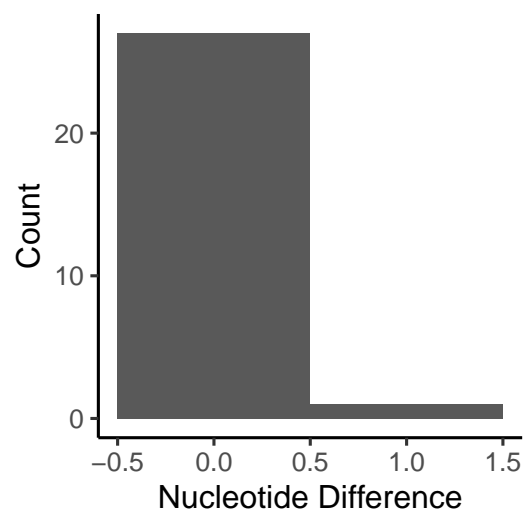
TRBV10–2*01_C214T

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



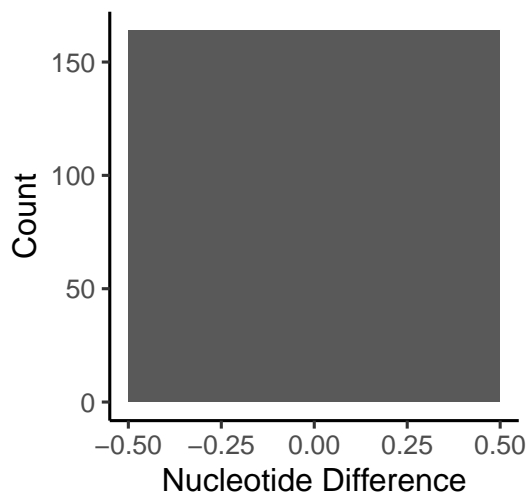
TRBV13*01

28 sequences assigned
27 (96.4%) exact matches, in which:
27 unique CDR3
11 unique J



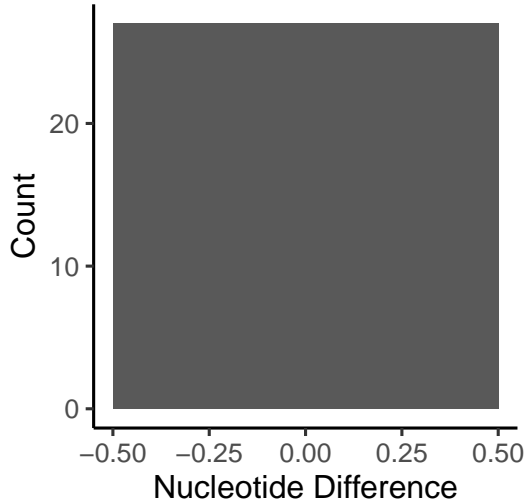
TRBV20–1*02_319AGAG

164 sequences assigned
164 (100%) exact matches, in which:
164 unique CDR3
13 unique J



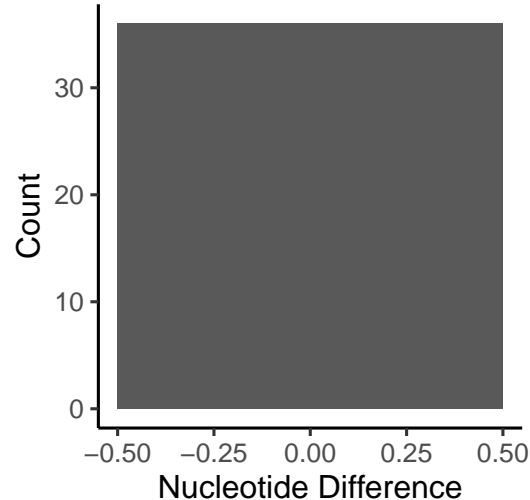
TRBV10–3*01

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



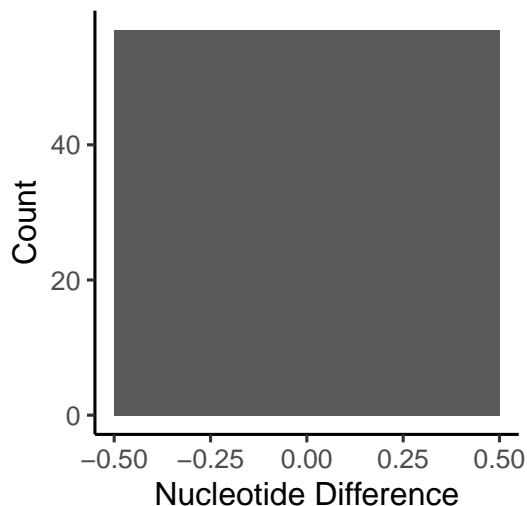
TRBV14*01

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



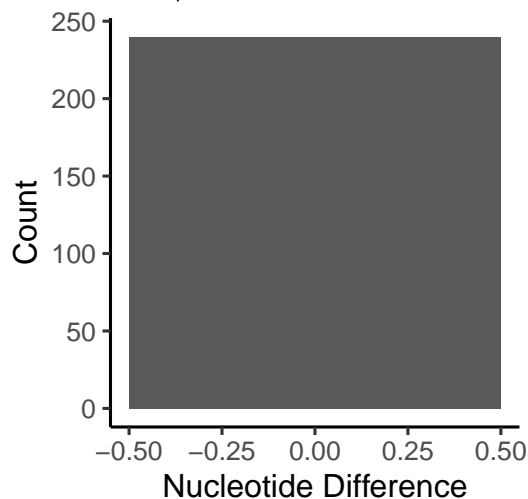
TRBV15*02

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



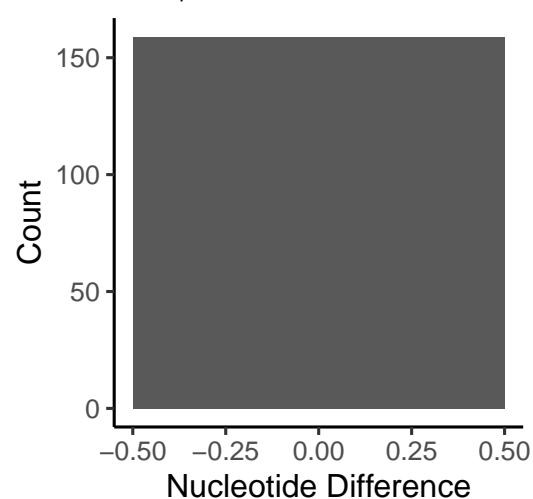
TRBV19*01

240 sequences assigned
240 (100%) exact matches, in which:
240 unique CDR3
13 unique J



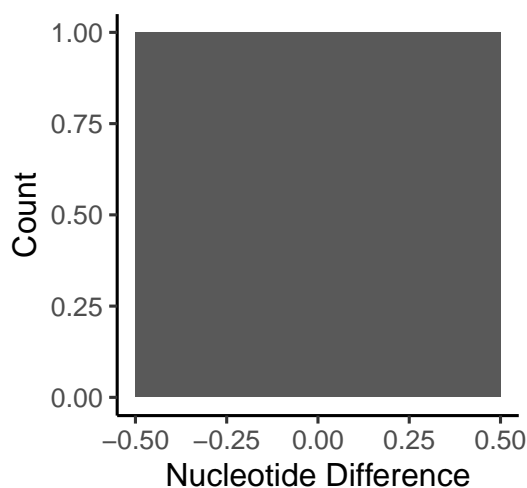
TRBV28*01

159 sequences assigned
159 (100%) exact matches, in which:
159 unique CDR3
12 unique J



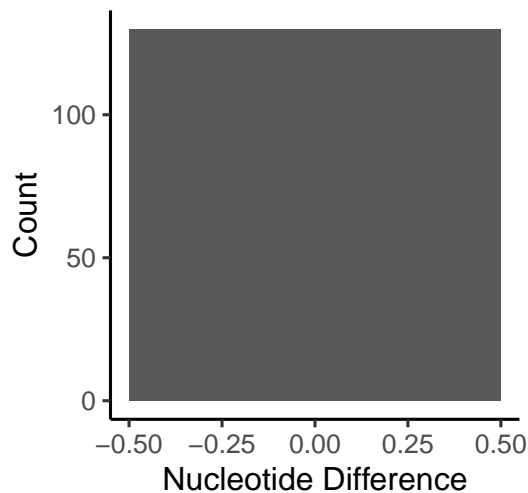
TRBV16*01

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



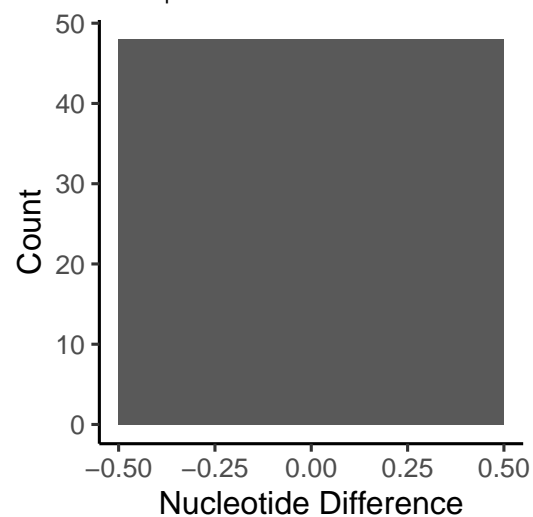
TRBV2*01

130 sequences assigned
130 (100%) exact matches, in which:
130 unique CDR3
13 unique J



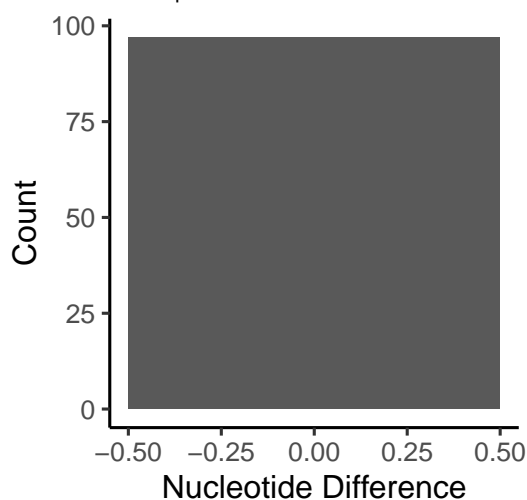
TRBV30*01

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



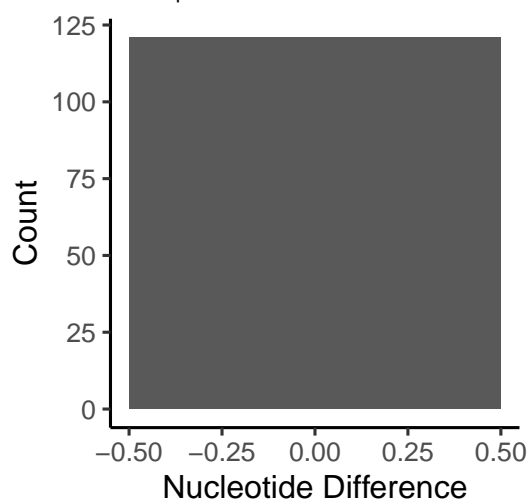
TRBV18*01

97 sequences assigned
97 (100%) exact matches, in which:
97 unique CDR3
13 unique J



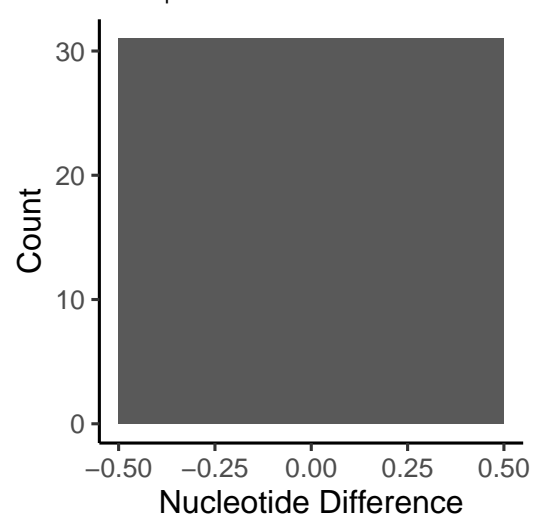
TRBV27*01

121 sequences assigned
121 (100%) exact matches, in which:
121 unique CDR3
12 unique J



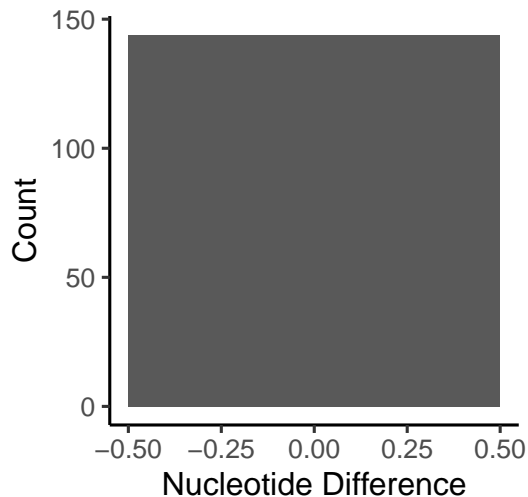
TRBV30*02

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



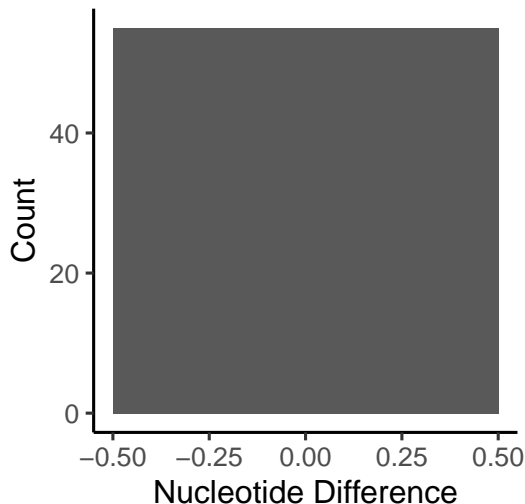
TRBV9*01

144 sequences assigned
144 (100%) exact matches, in which:
144 unique CDR3
12 unique J



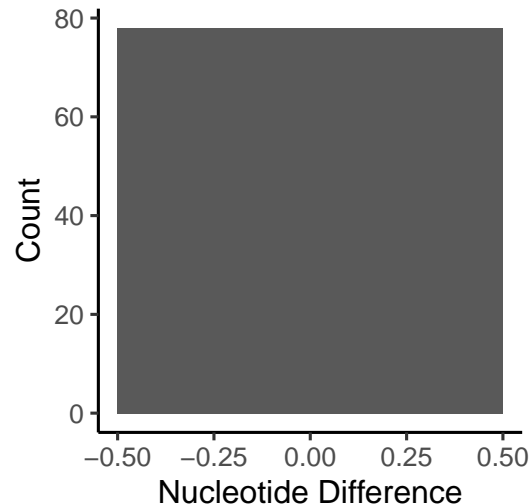
TRBV11-2*03

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



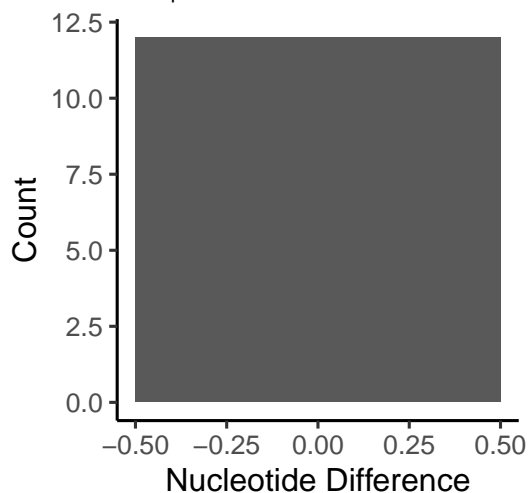
TRBV12-4*01

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



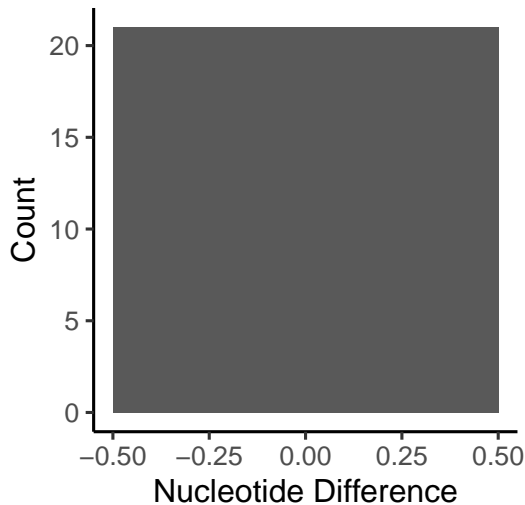
TRBV11-1*01

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



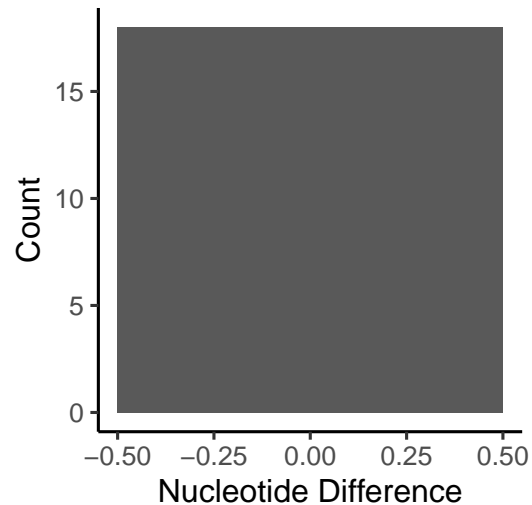
TRBV11-3*01

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



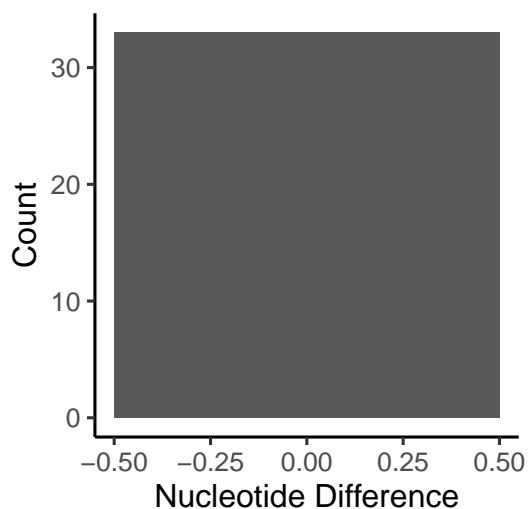
TRBV12-5*01

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



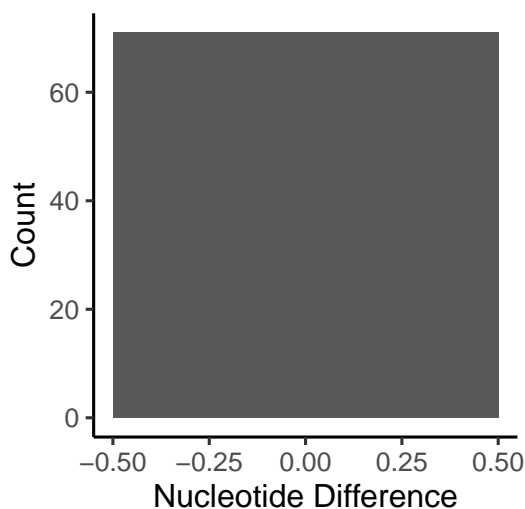
TRBV11-2*01

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



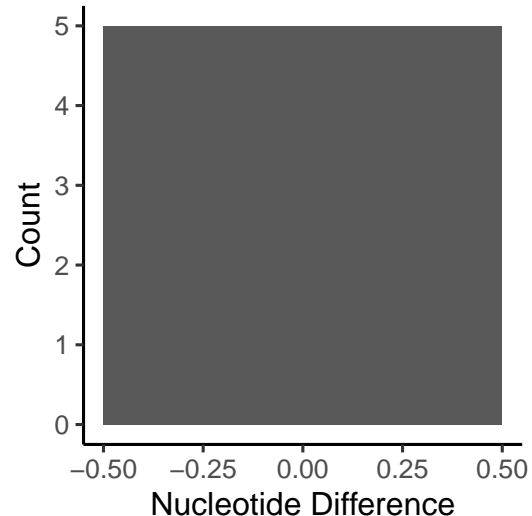
TRBV12-3*01

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



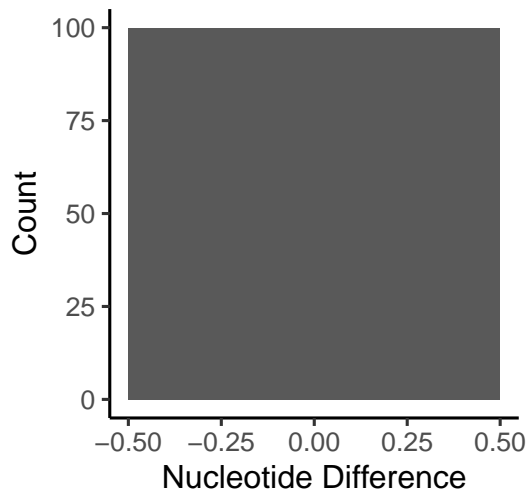
TRBV12-5*01_C28G

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



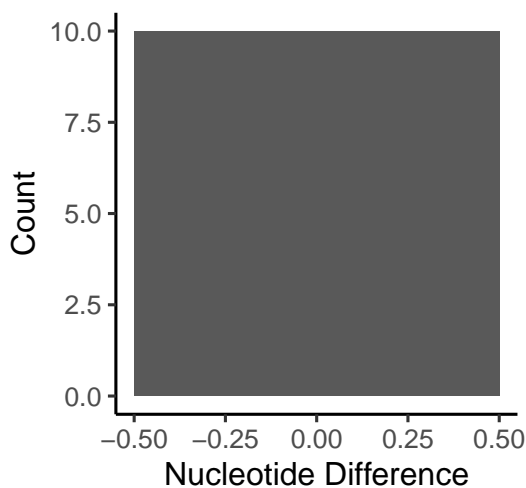
TRBV3–1*01

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



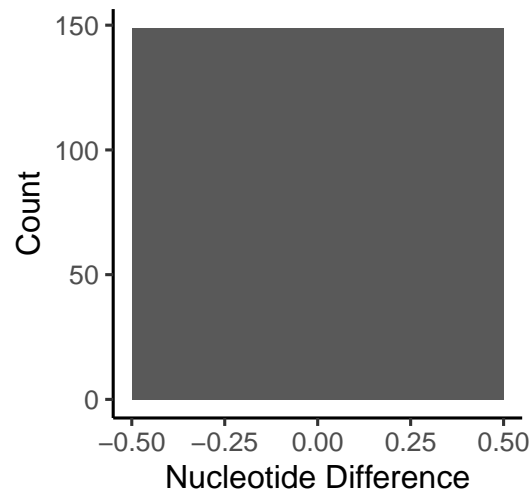
TRBV24–1*02

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



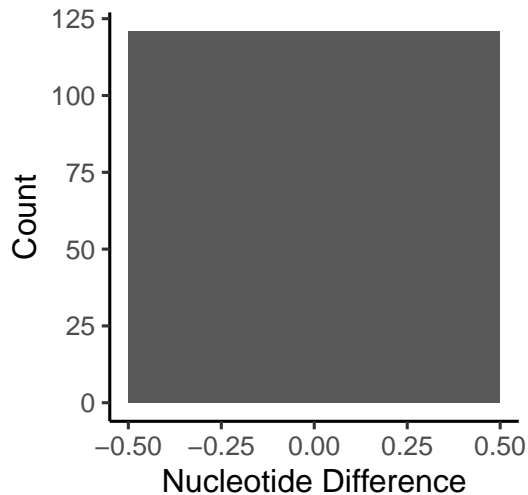
TRBV5–1*01

149 sequences assigned
149 (100%) exact matches, in which:
149 unique CDR3
13 unique J



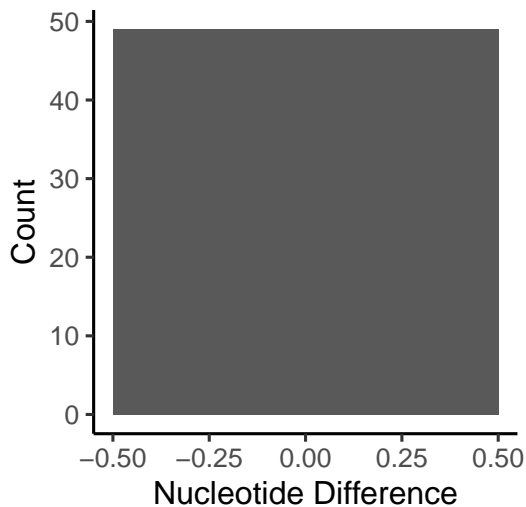
TRBV4–1*01

121 sequences assigned
121 (100%) exact matches, in which:
121 unique CDR3
13 unique J



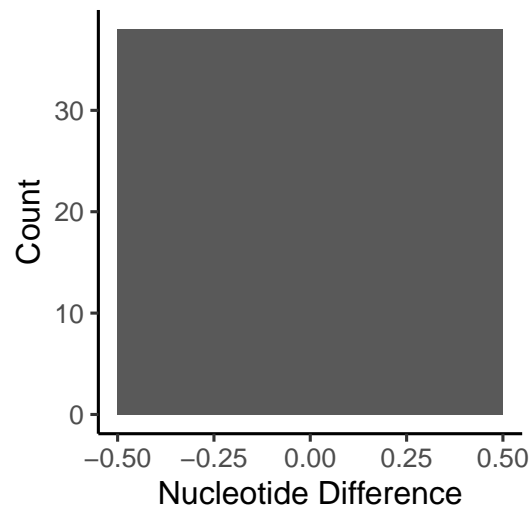
TRBV4–2*01

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



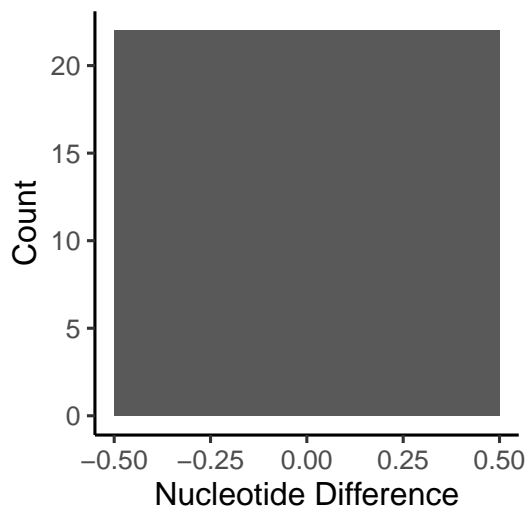
TRBV25–1*01

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



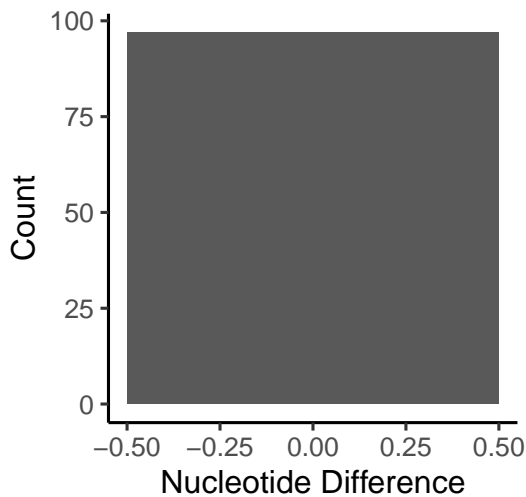
TRBV24–1*01

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



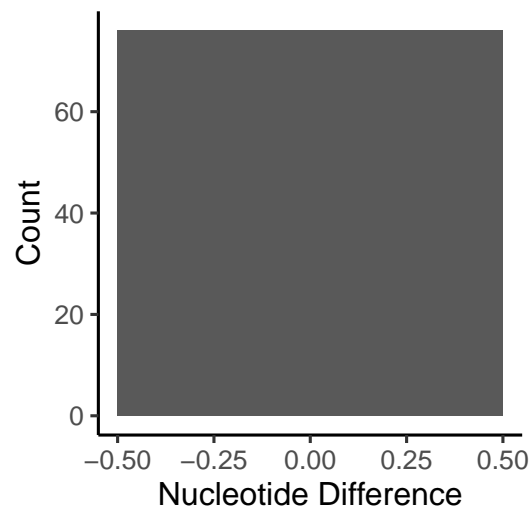
TRBV4–3*01

97 sequences assigned
97 (100%) exact matches, in which:
97 unique CDR3
13 unique J



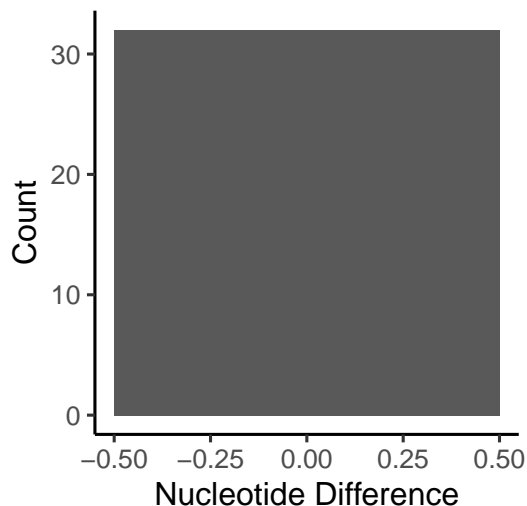
TRBV5–4*01

76 sequences assigned
76 (100%) exact matches, in which:
76 unique CDR3
13 unique J



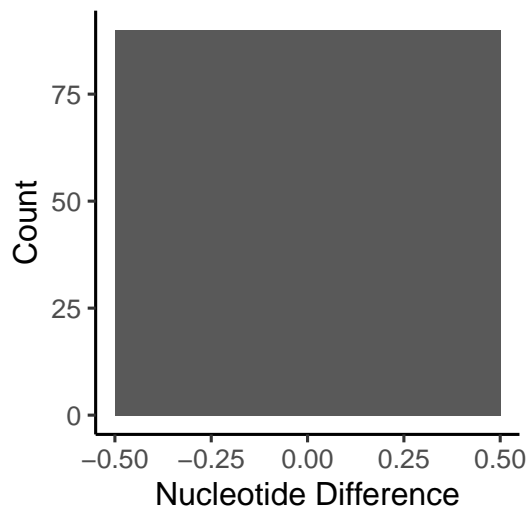
TRBV5-5*02_322TTGG32

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



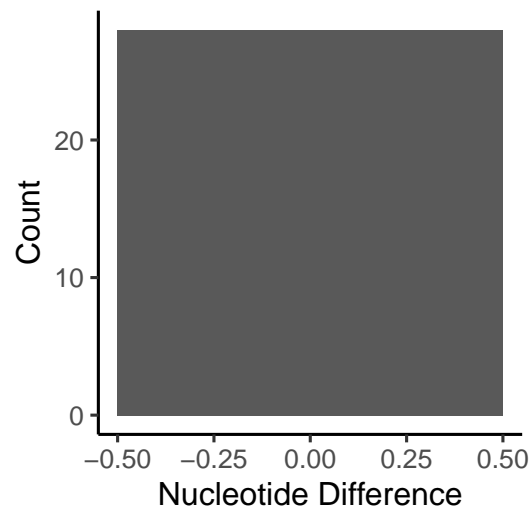
TRBV6-1*01

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



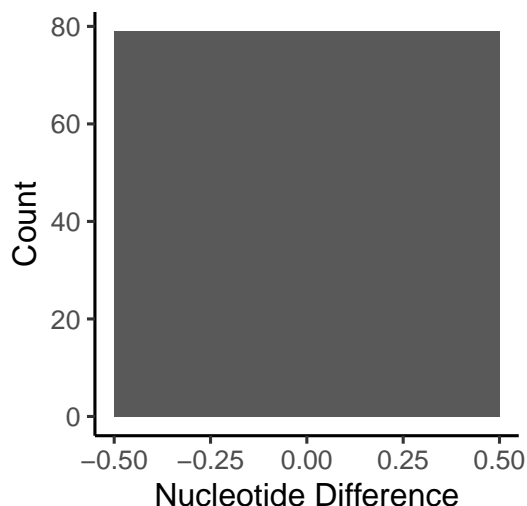
TRBV6-6*01

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



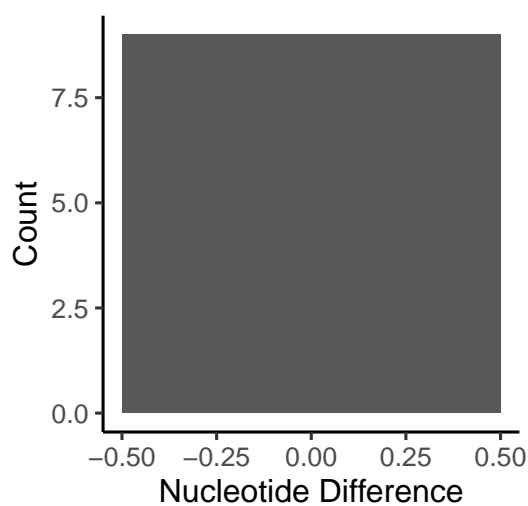
TRBV5-6*01

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



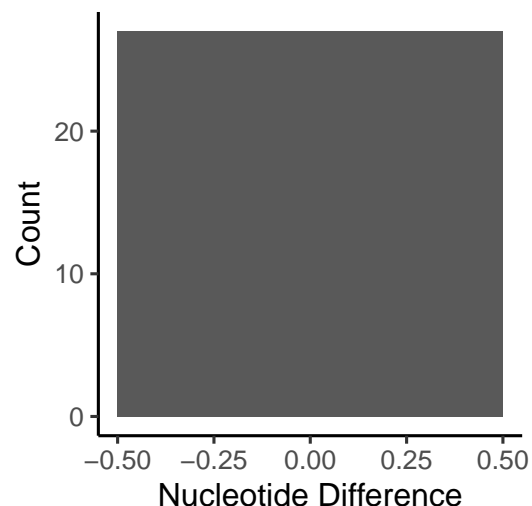
TRBV6-4*01

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



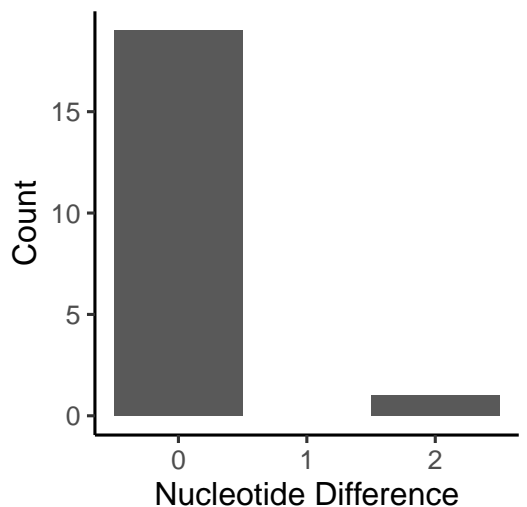
TRBV6-6*02

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



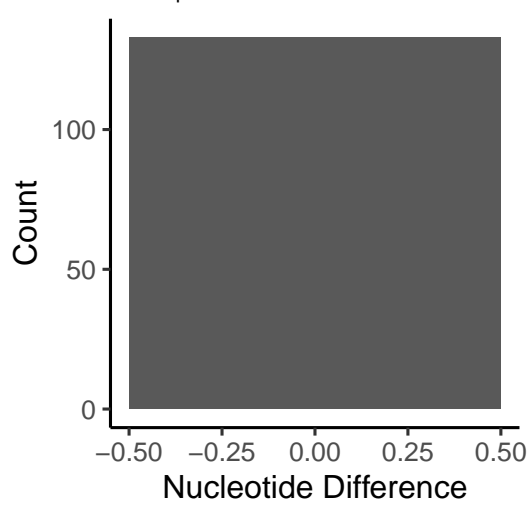
TRBV5-8*01

20 sequences assigned
19 (95%) exact matches, in which:
19 unique CDR3
9 unique J



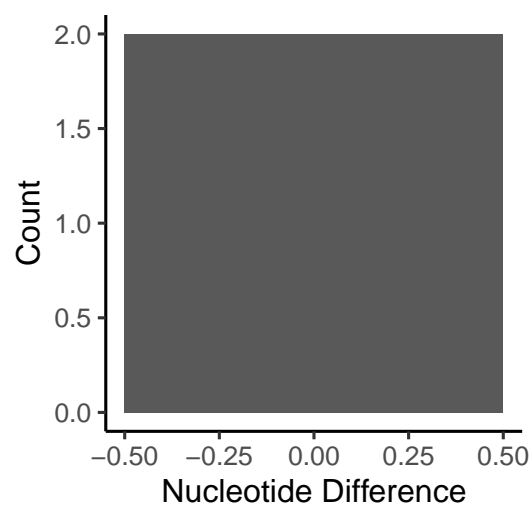
TRBV6-5*01

133 sequences assigned
133 (100%) exact matches, in which:
133 unique CDR3
13 unique J



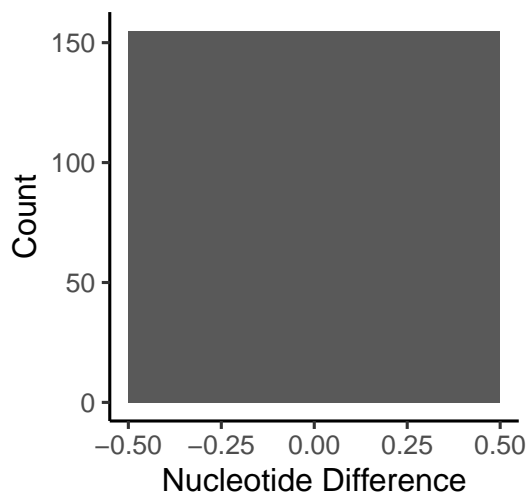
TRBV6-7*01

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



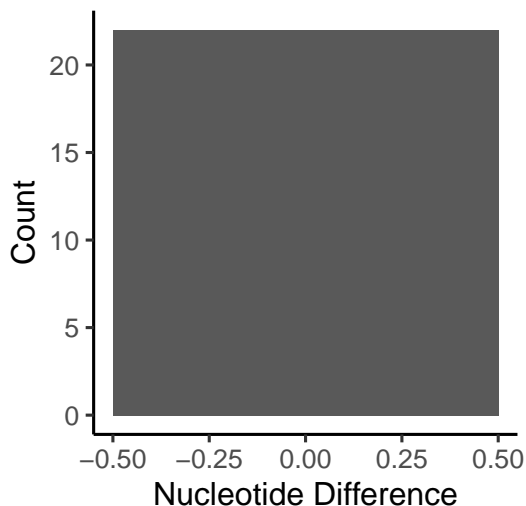
TRBV6–23*01

155 sequences assigned
155 (100%) exact matches, in which:
155 unique CDR3
12 unique J



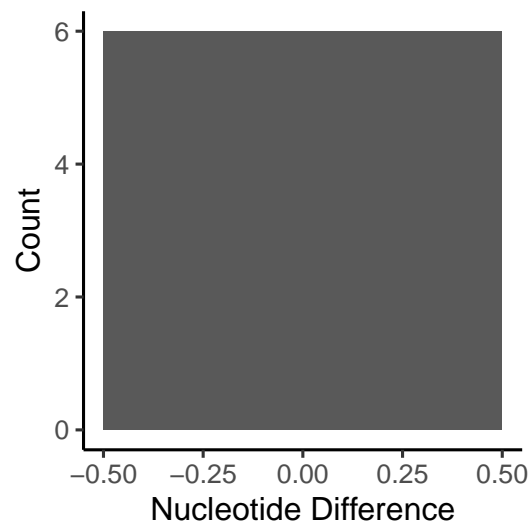
TRBV7–3*01_G301A

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



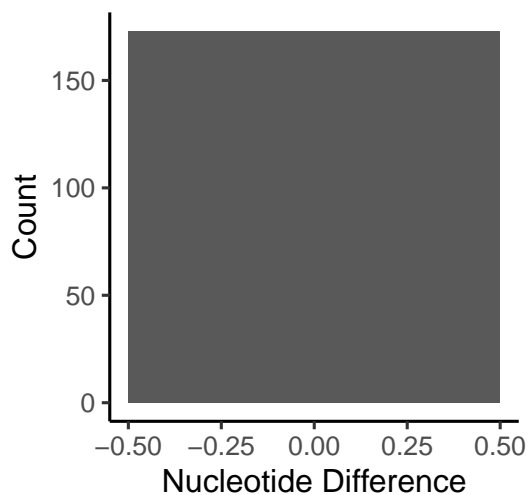
TRBV7–7*01_C315T

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



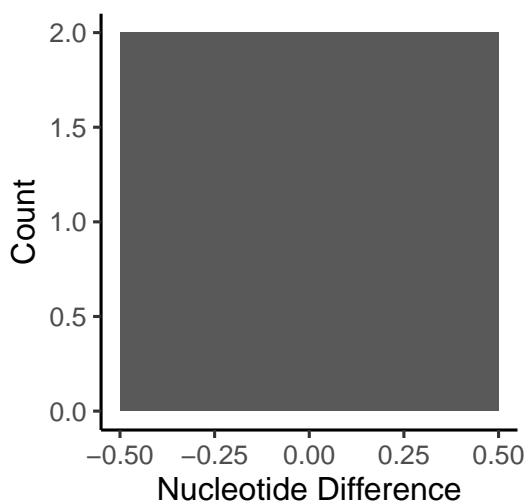
TRBV7–2*01

173 sequences assigned
173 (100%) exact matches, in which:
173 unique CDR3
13 unique J



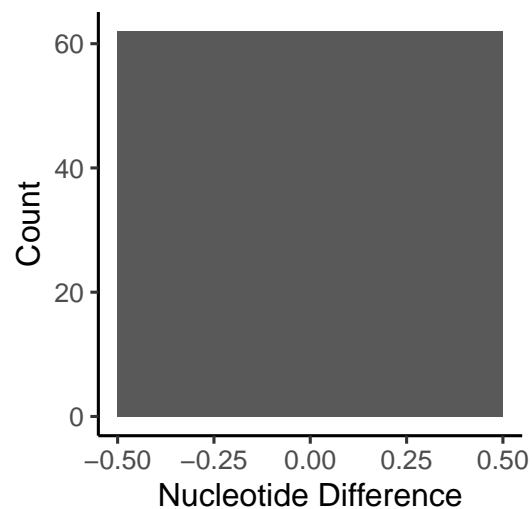
TRBV7–4*01

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



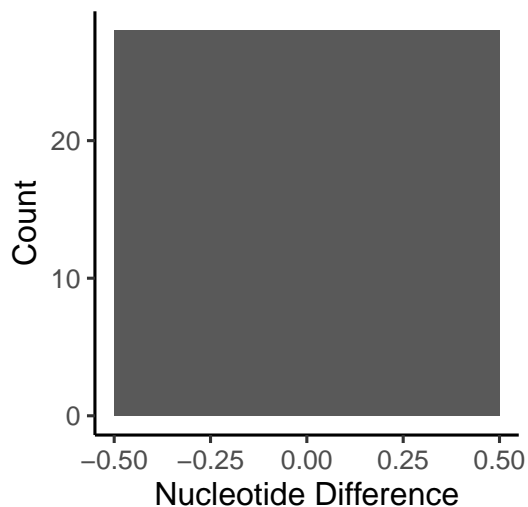
TRBV7–8*01

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



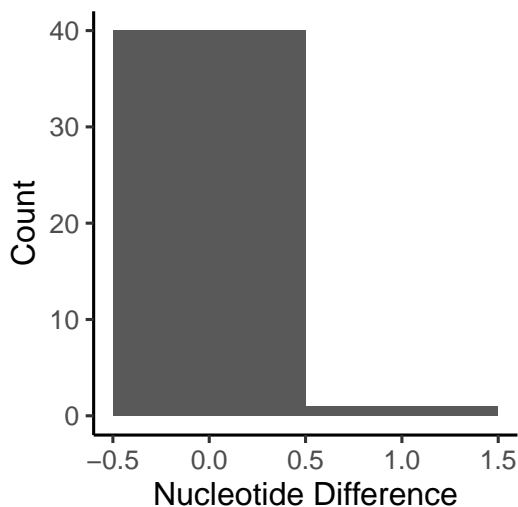
TRBV7–3*01

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



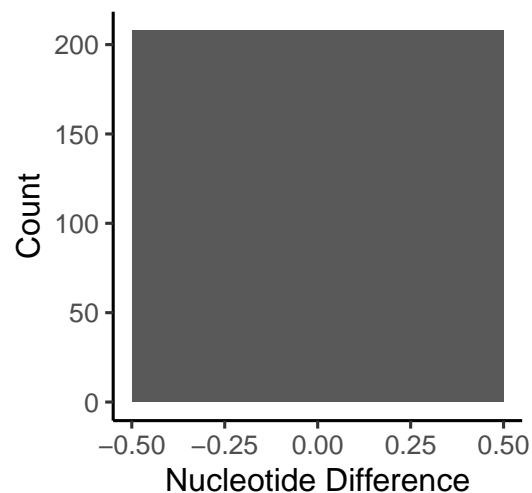
TRBV7–6*01

41 sequences assigned
40 (97.6%) exact matches, in which:
40 unique CDR3
12 unique J



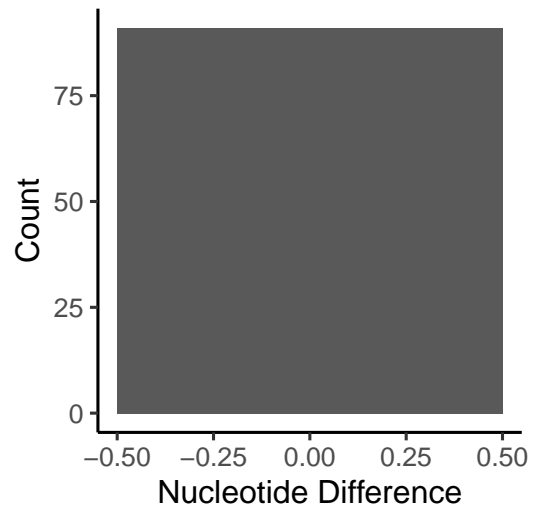
TRBV7–9*03

208 sequences assigned
208 (100%) exact matches, in which:
208 unique CDR3
13 unique J

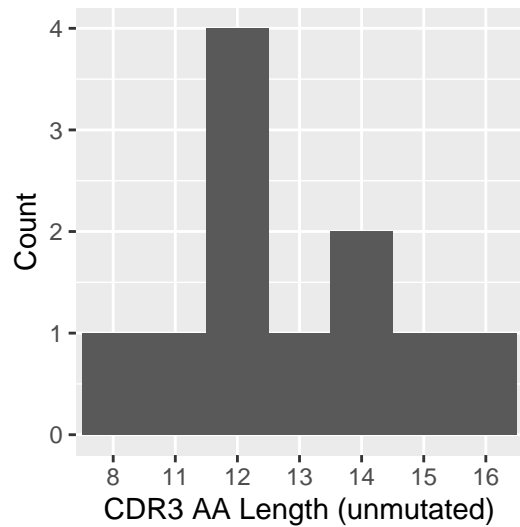


TRBV29-1*01

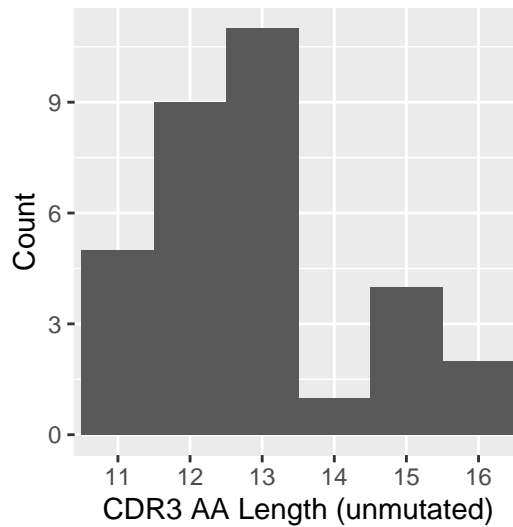
91 sequences assigned
91 (100%) exact matches, in which:
91 unique CDR3
12 unique J



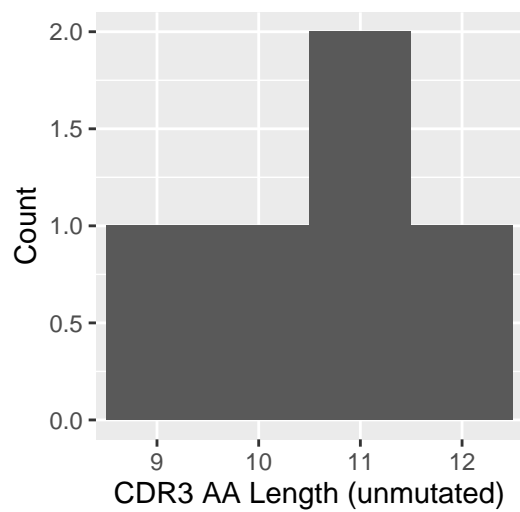
TRBV10-2*01_C214T



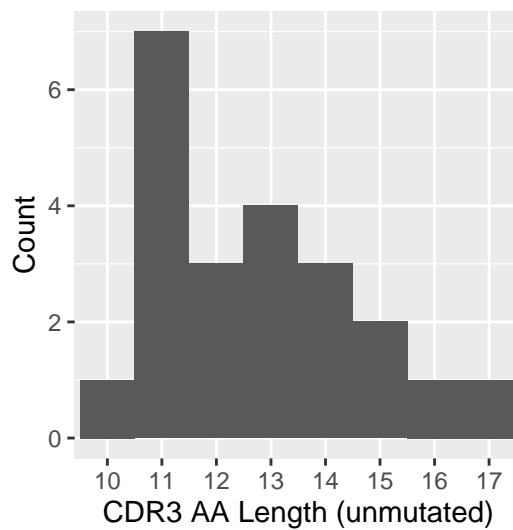
TRBV5-5*02_322TTGG325



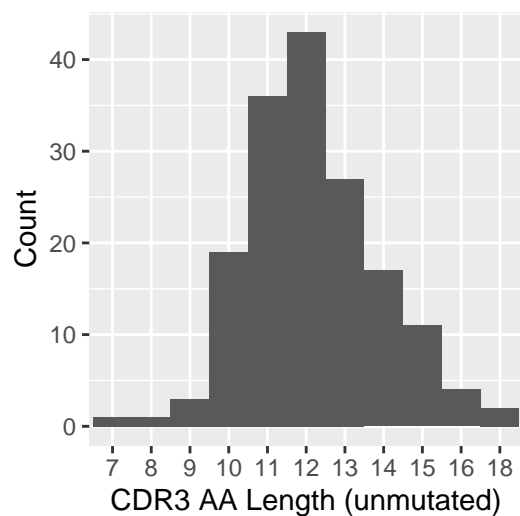
TRBV12-5*01_C28G



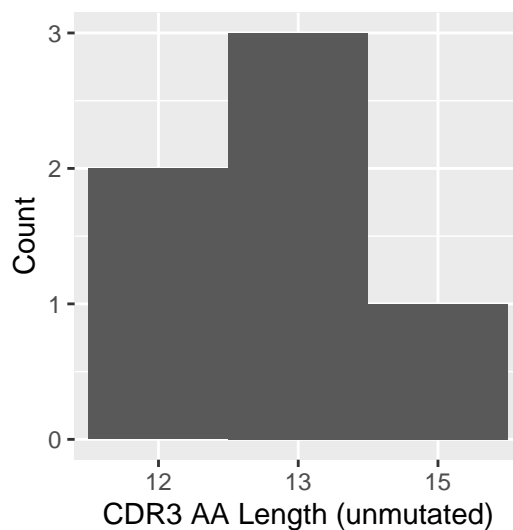
TRBV7-3*01_G301A

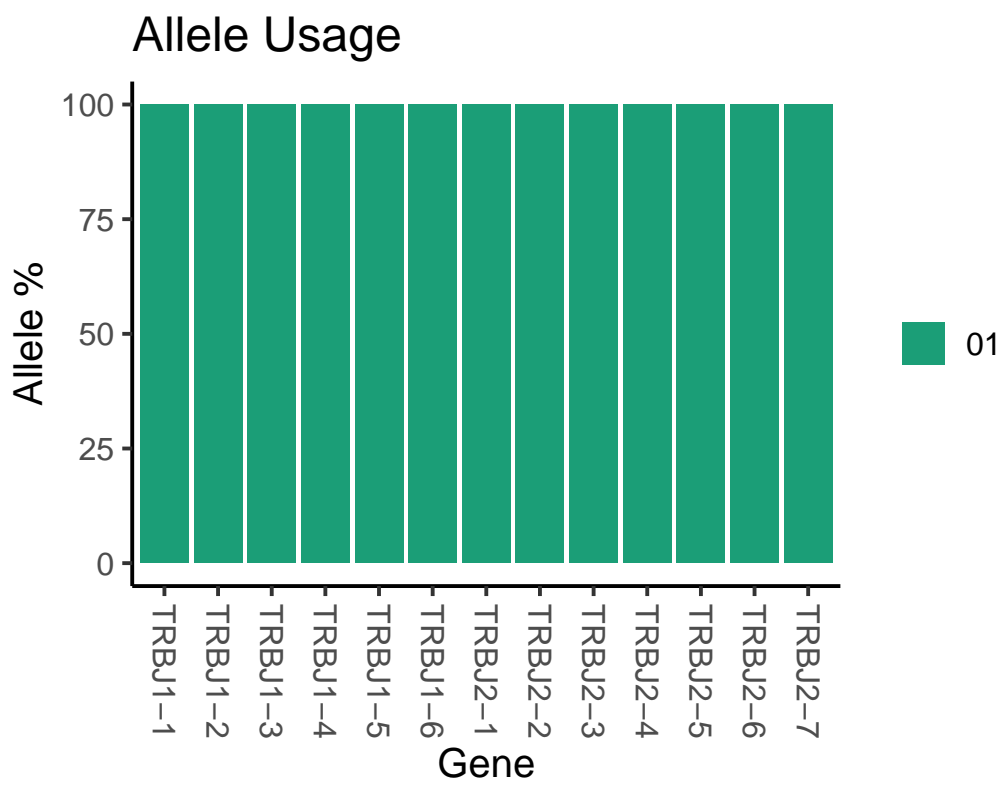


TRBV20-1*02_319AGAGA32



TRBV7-7*01_C315T





Warning – no inferred sequences found.

Warning – no inferred sequences found.