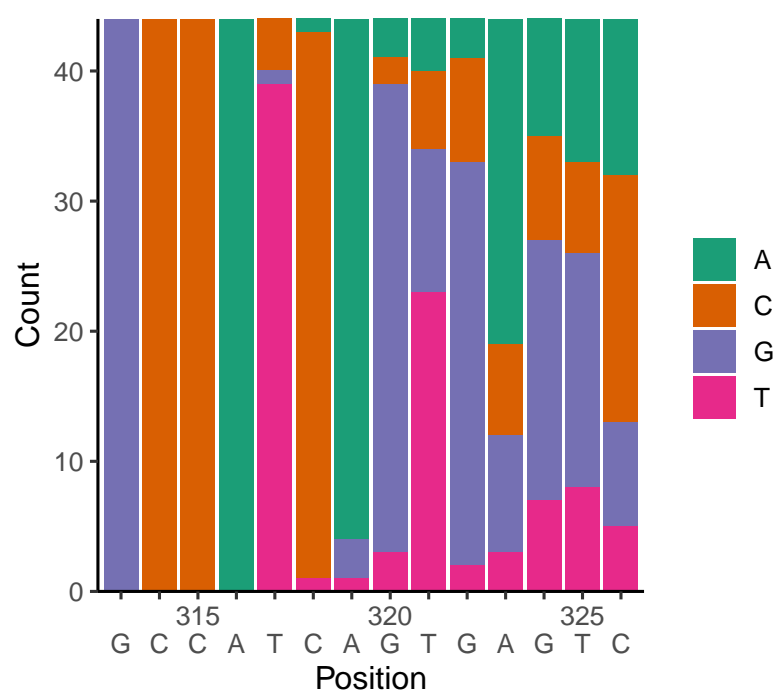
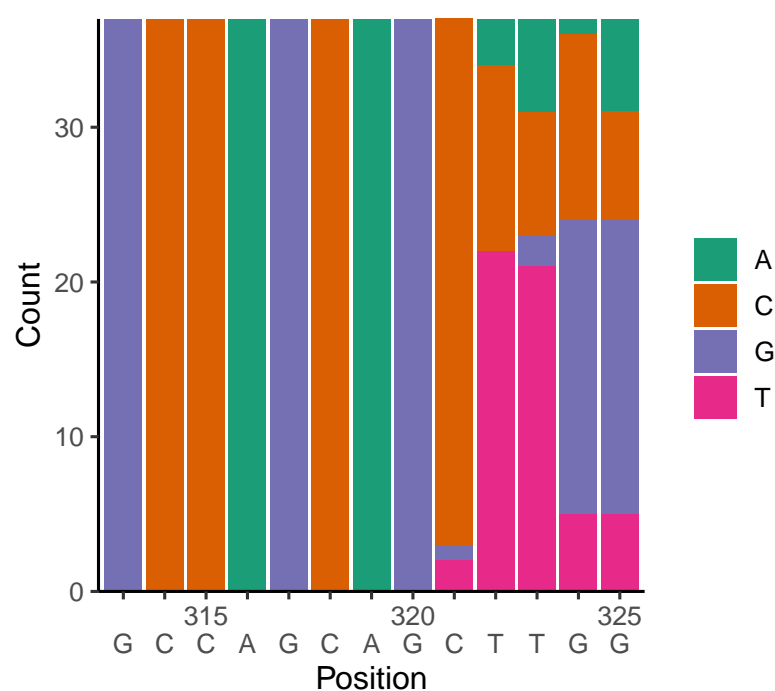


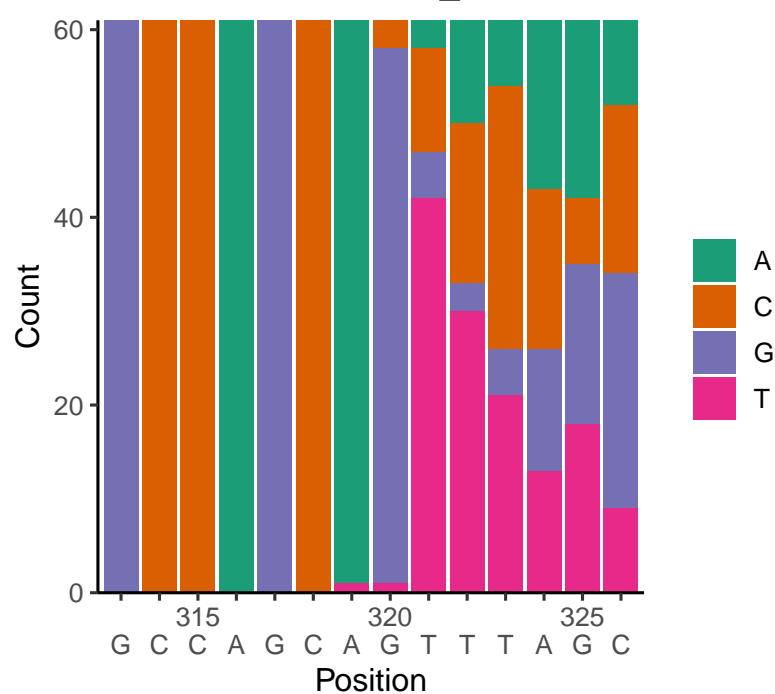
Gene TRBV10-3*03_313GCCATCAGTGA



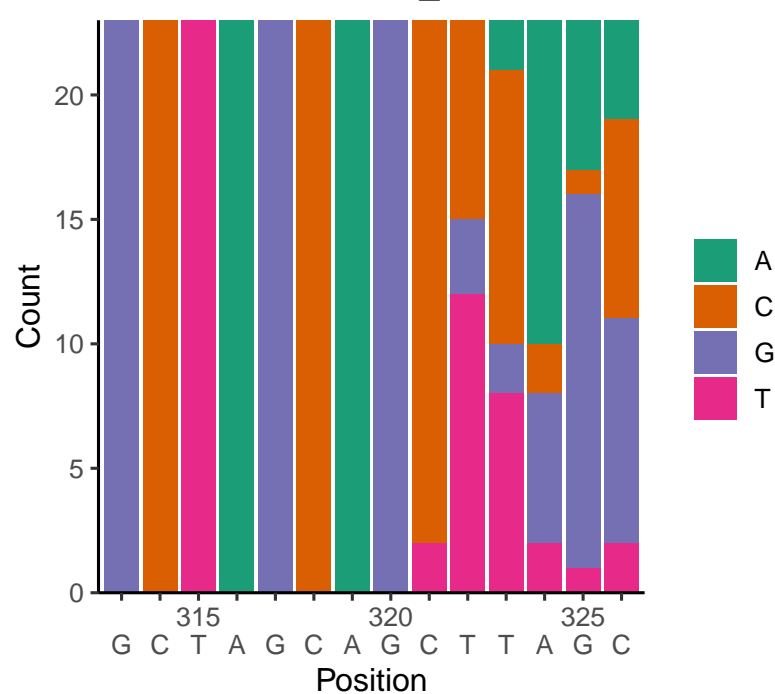
Gene TRBV5-5*02_322TTGG325



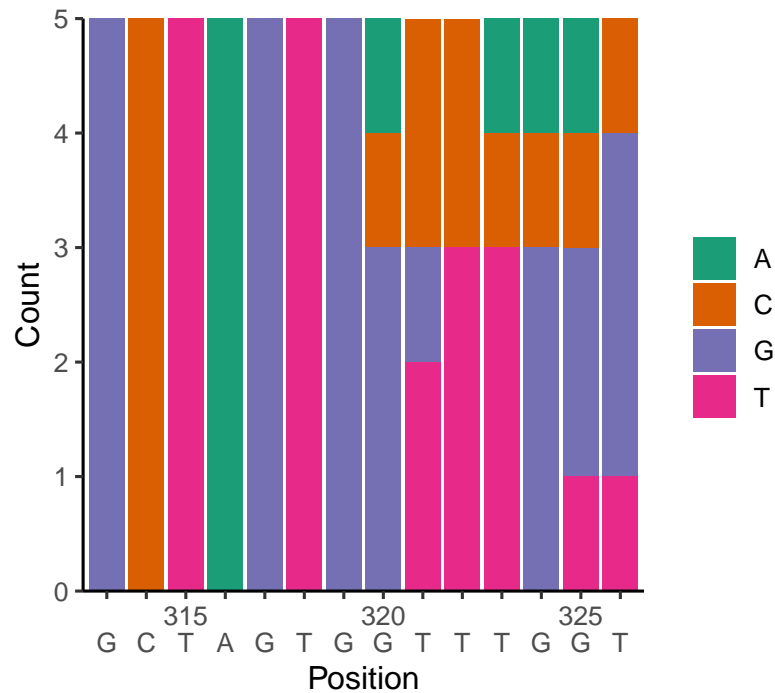
Gene TRBV12-4*01_C87T



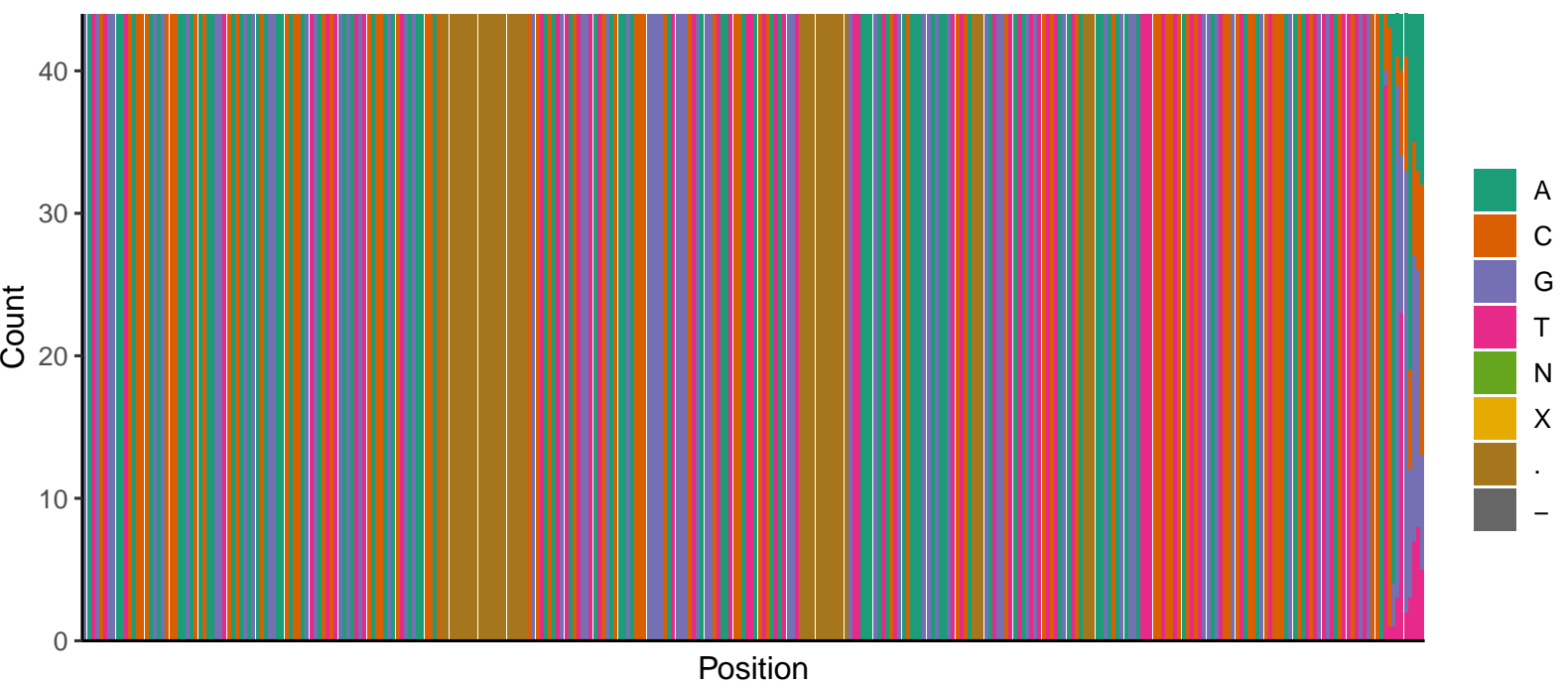
Gene TRBV7-7*01_C315T



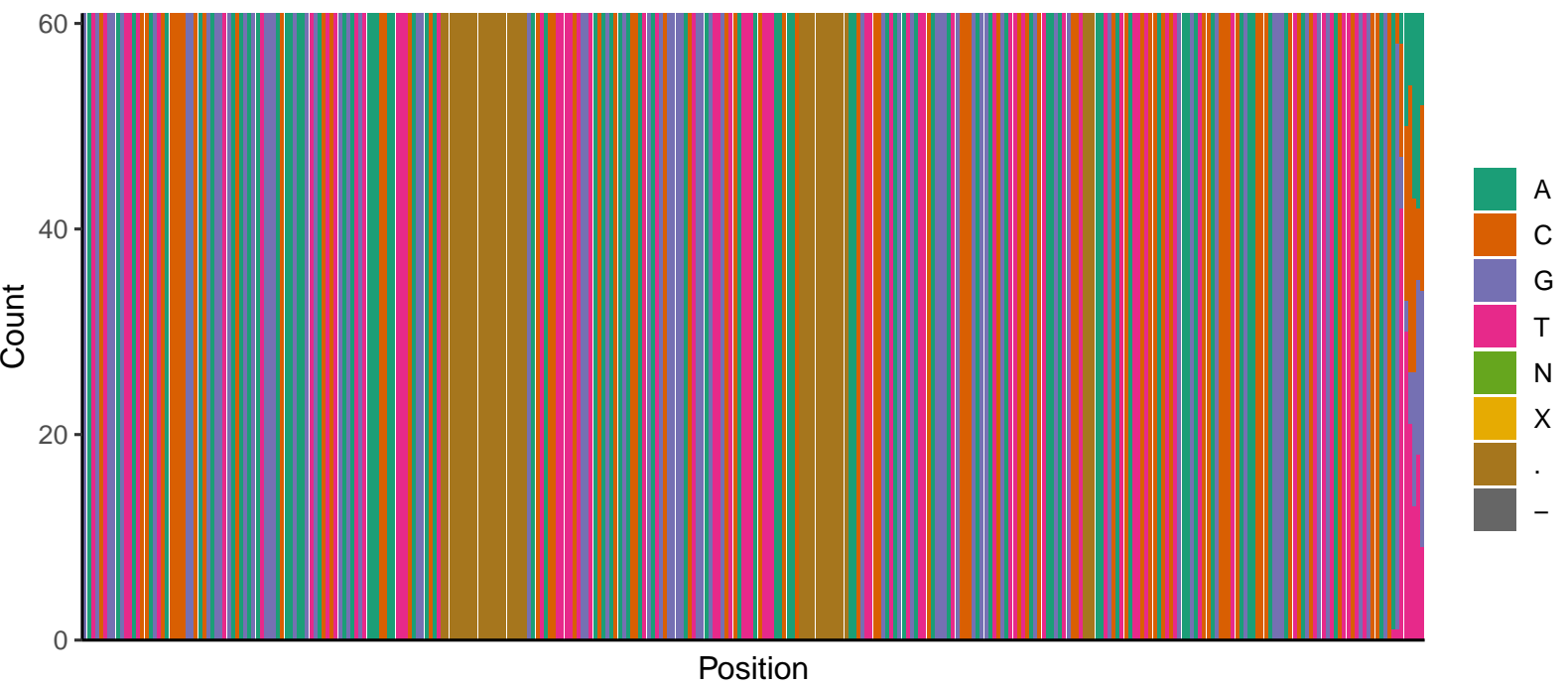
Gene TRBV12-5*01_C28G



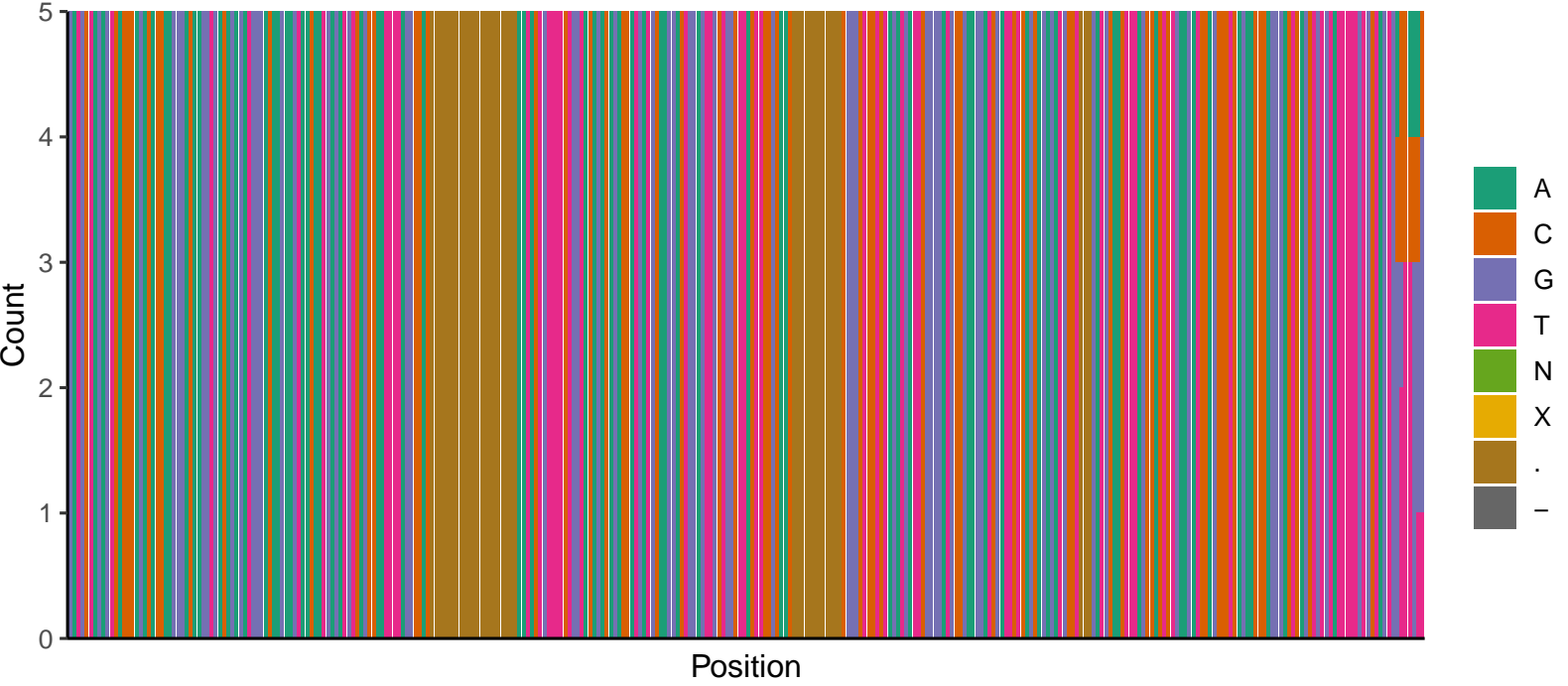
Gene TRBV10-3*03_313GCCATCAGTGAGTC326



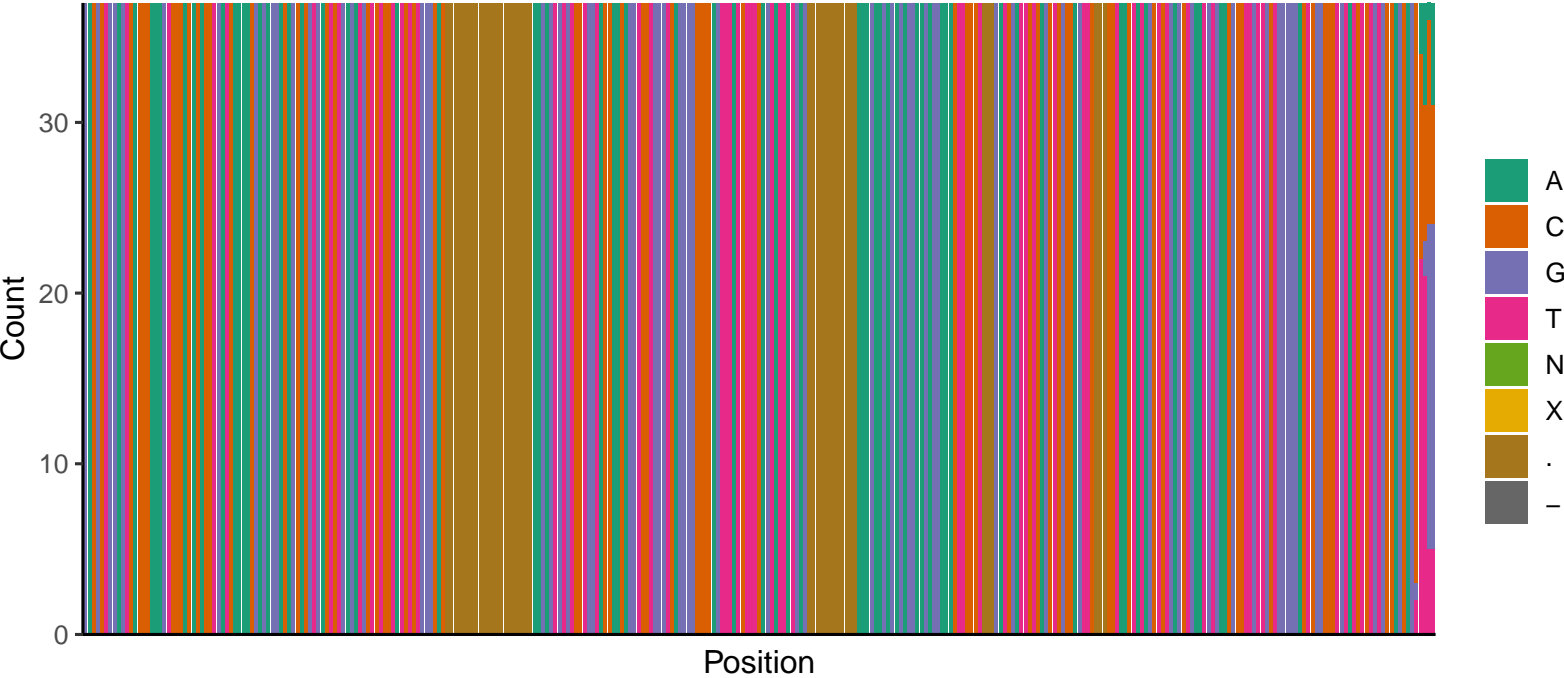
Gene TRBV12-4*01_C87T



Gene TRBV12-5*01_C28G



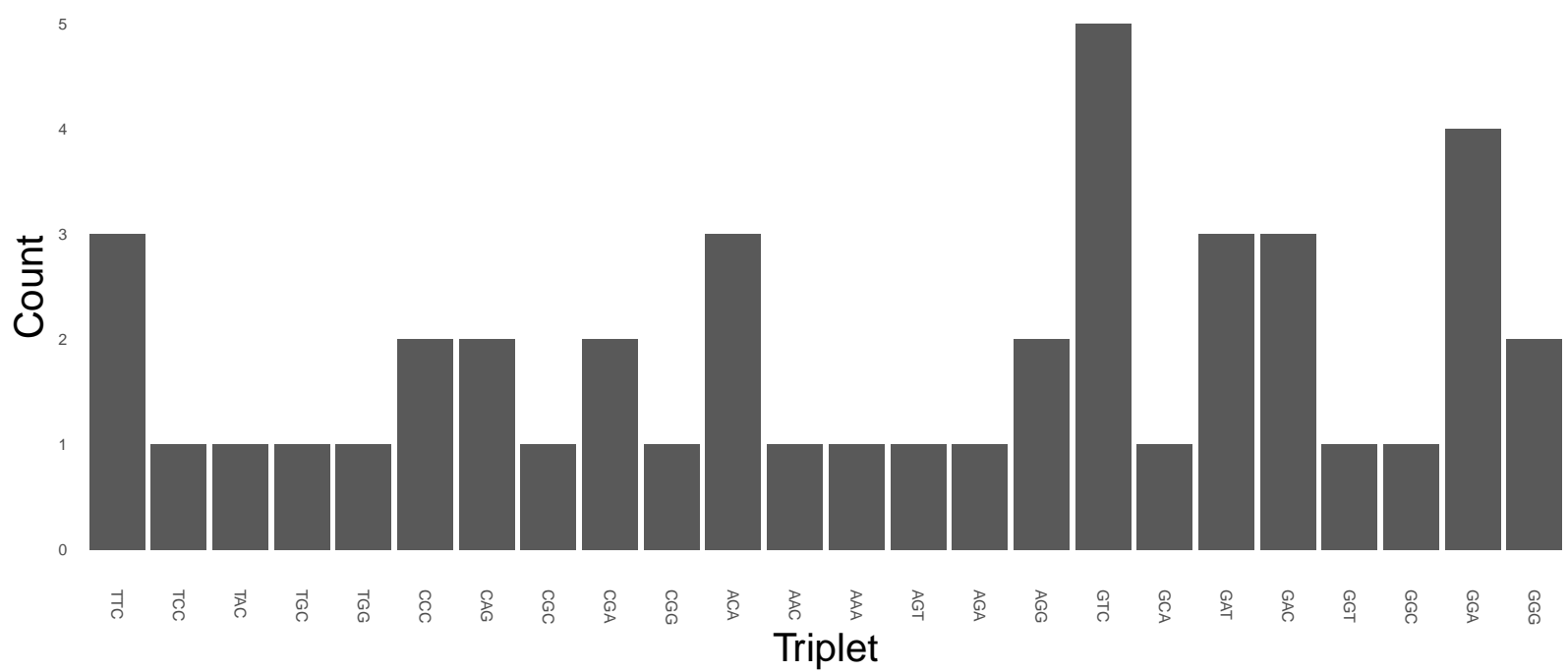
Gene TRBV5-5*02_322TTGG325



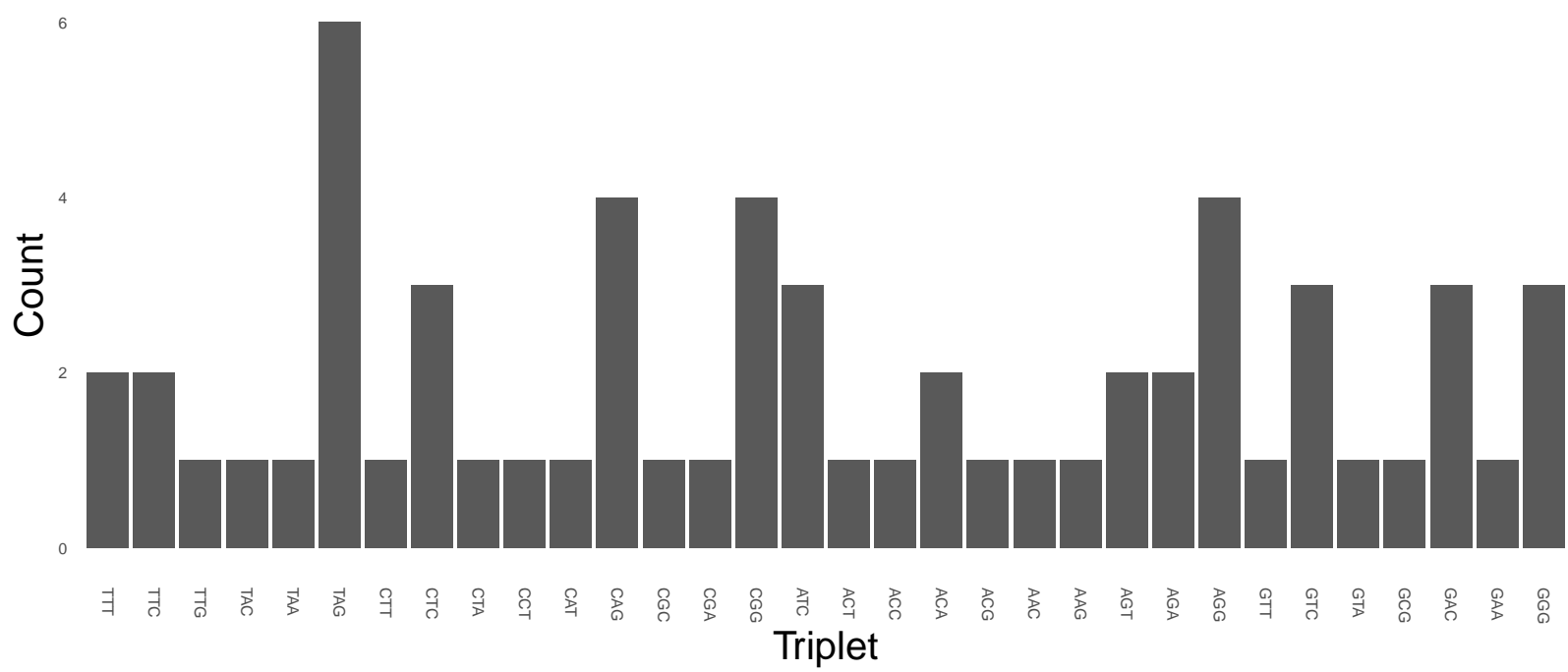
Gene TRBV7-7*01_C315T



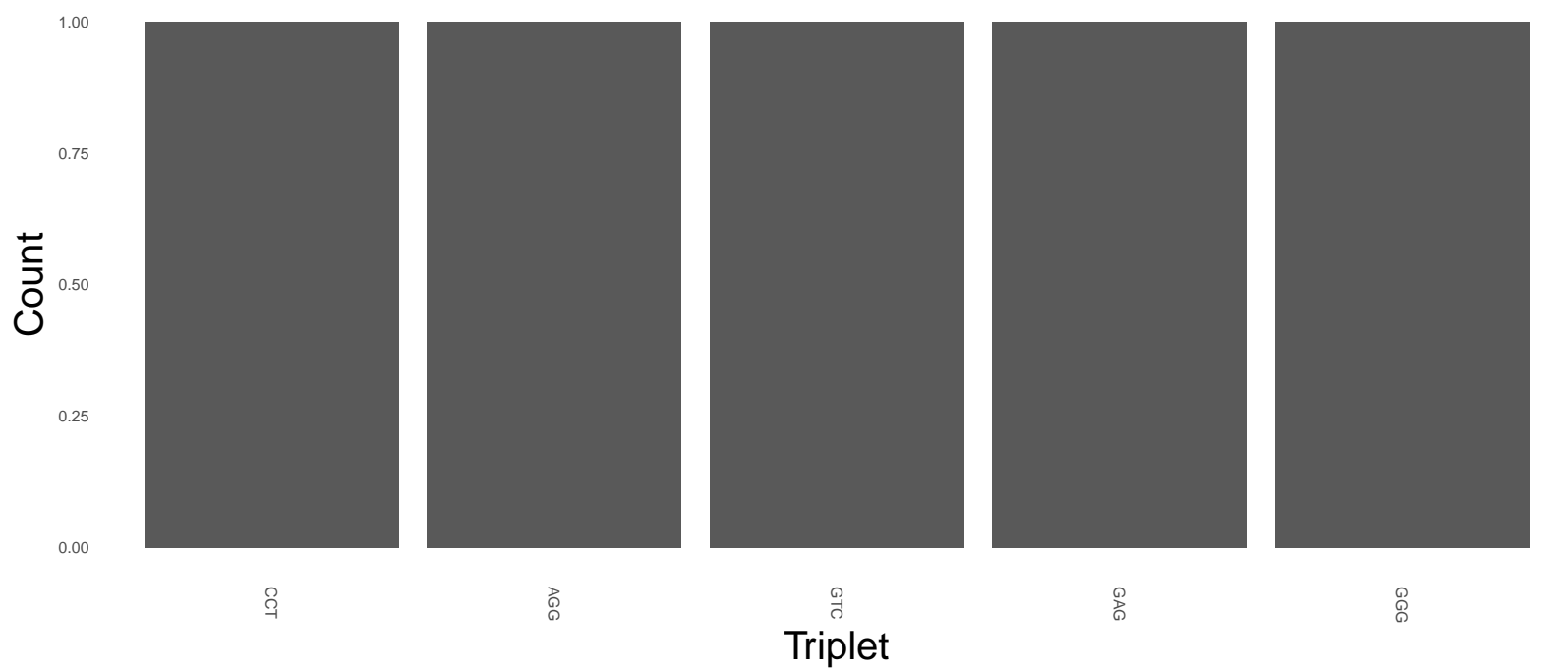
TRBV10–3*03_313GCCATCAGTGAGTC326– Final 3 nucleotides as a triplet



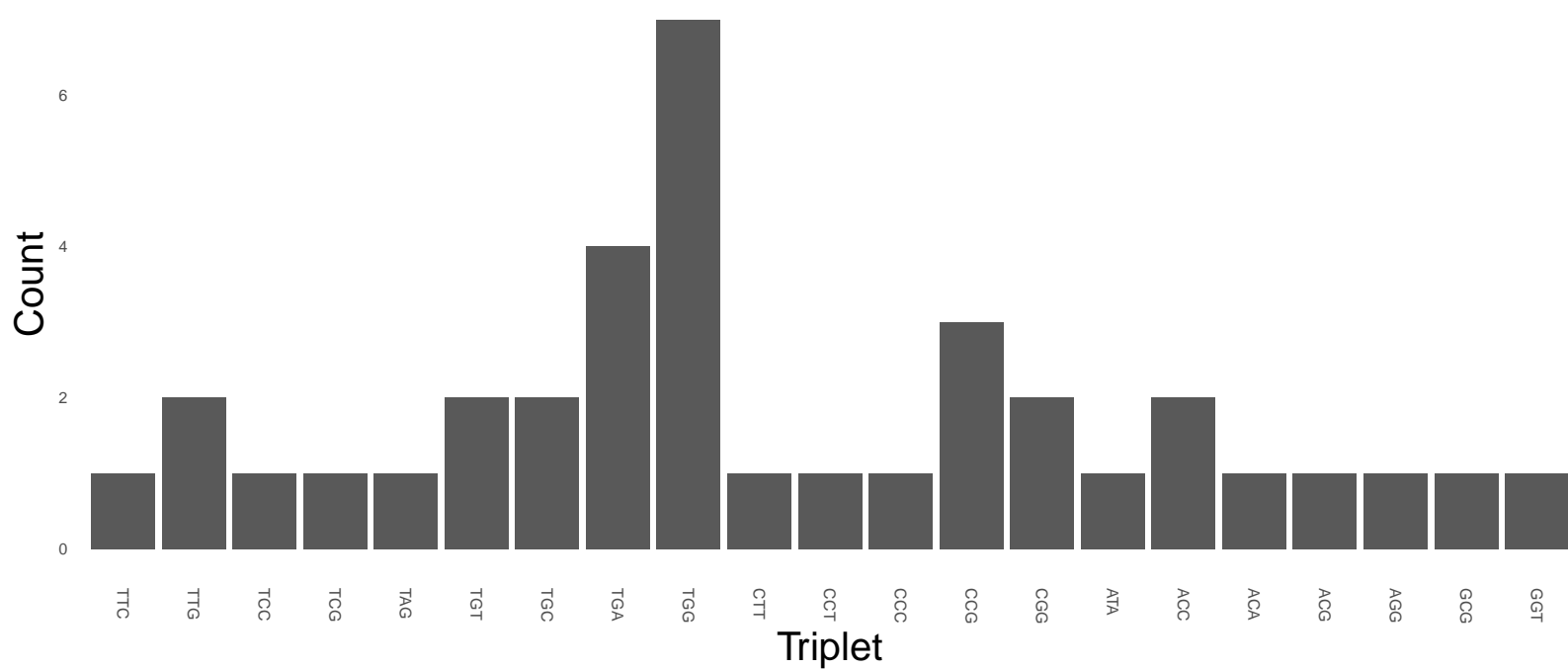
TRBV12–4*01_C87T– Final 3 nucleotides as a triplet



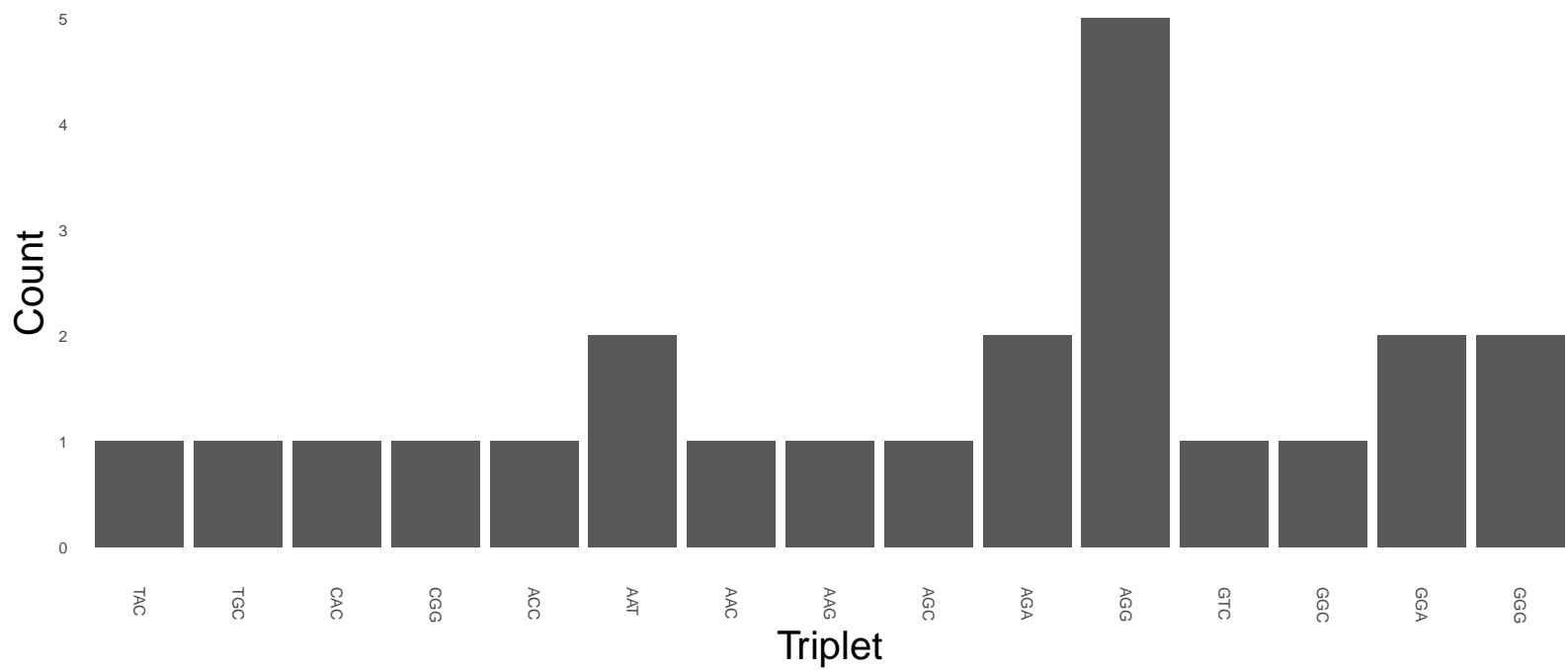
TRBV12–5*01_C28G– Final 3 nucleotides as a triplet



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

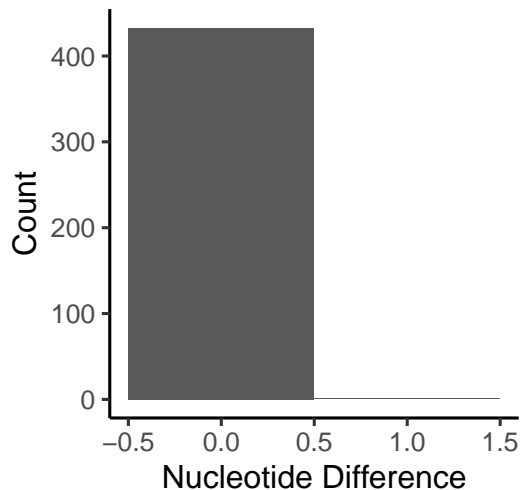


TRBV7–7*01_C315T– Final 3 nucleotides as a triplet



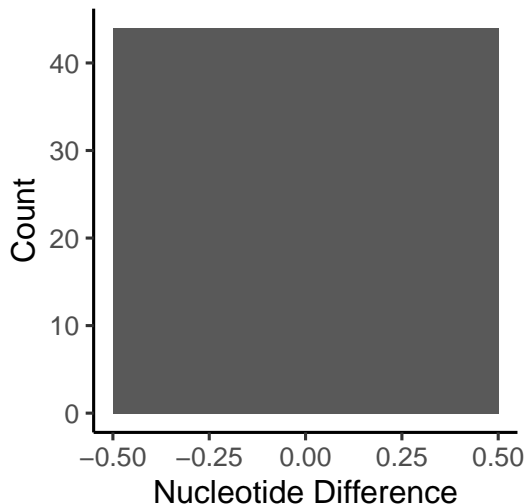
TRBV20–1*01

434 sequences assigned
433 (99.8%) exact matches, in which:
433 unique CDR3
13 unique J



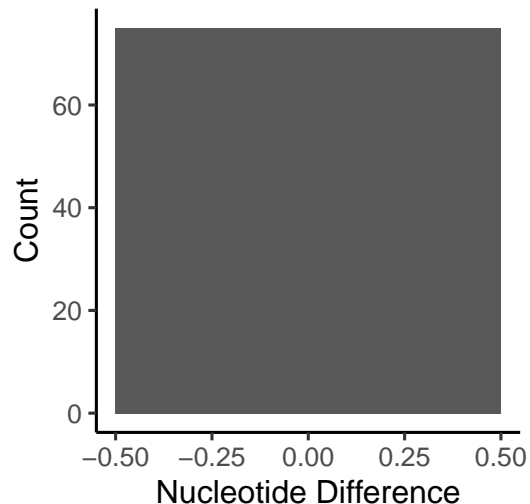
TRBV10–3*02

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



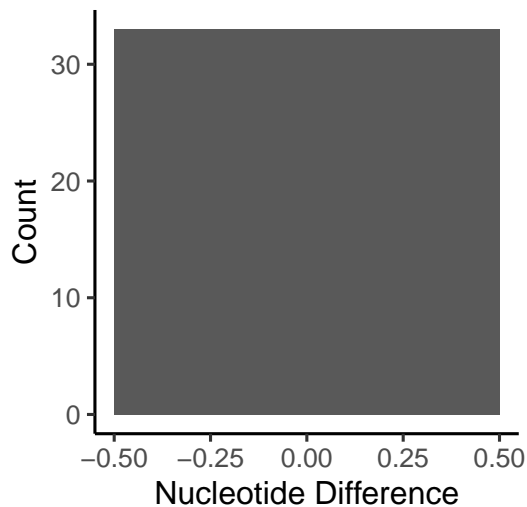
TRBV14*01

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



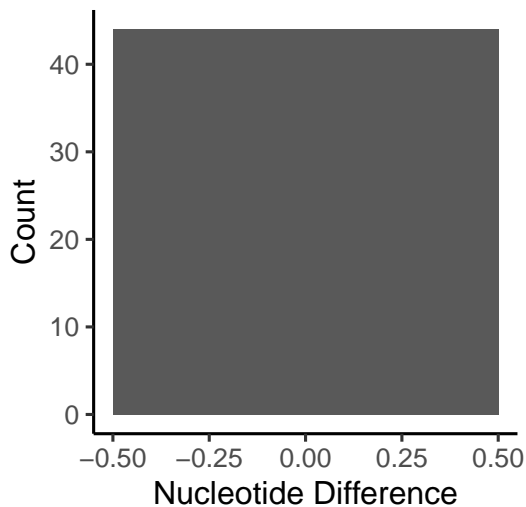
TRBV10–1*01

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



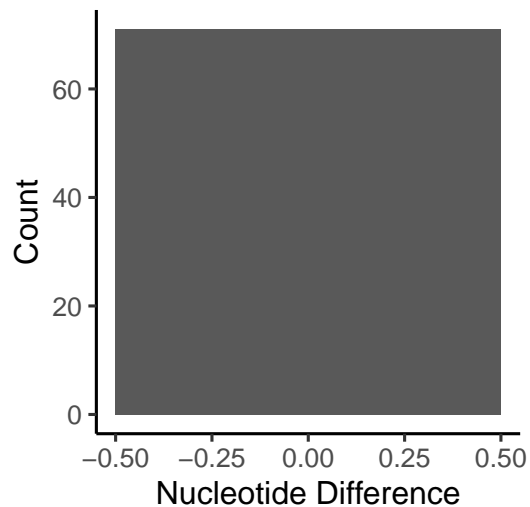
TRBV10–3*03_313GCCA1

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



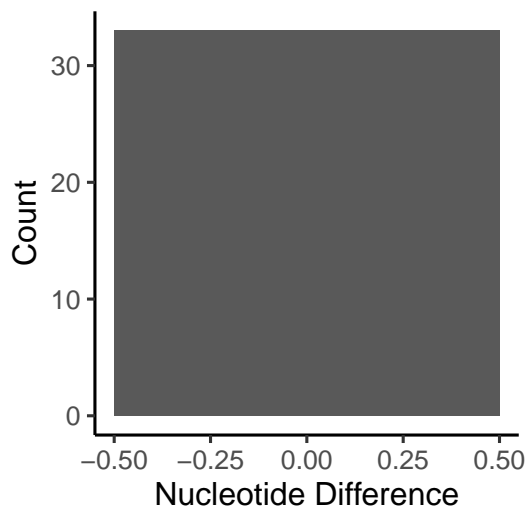
TRBV15*02

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



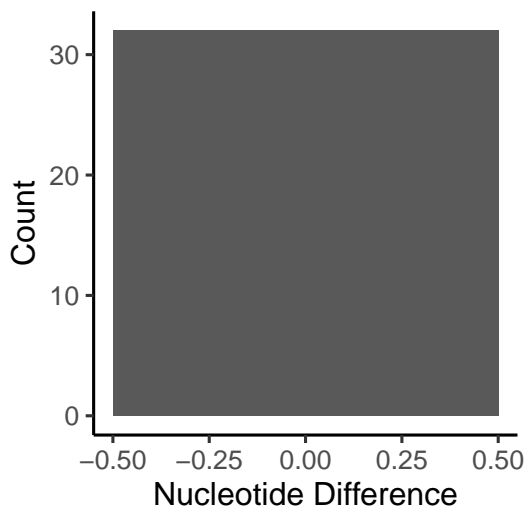
TRBV10–2*01

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



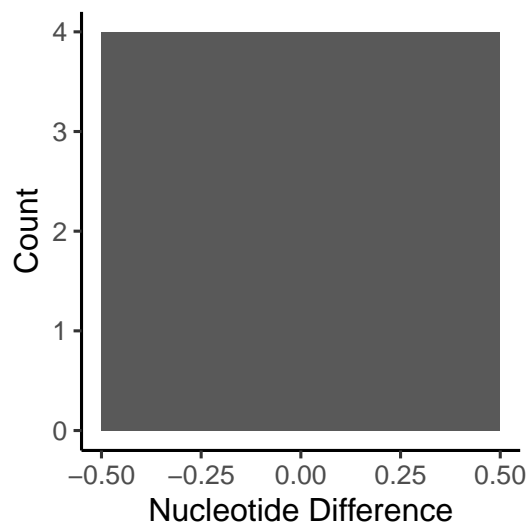
TRBV13*01

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



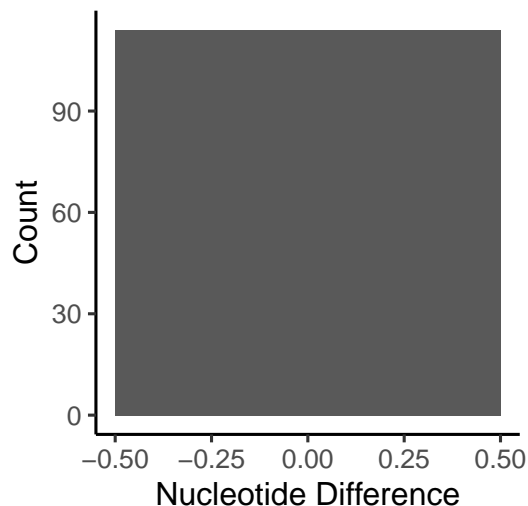
TRBV16*01

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



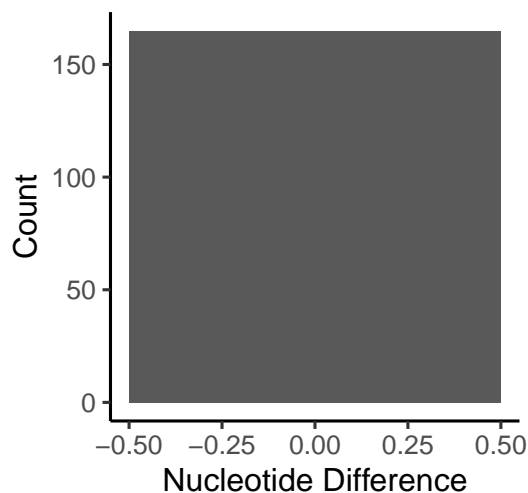
TRBV18*01

114 sequences assigned
114 (100%) exact matches, in which:
114 unique CDR3
13 unique J



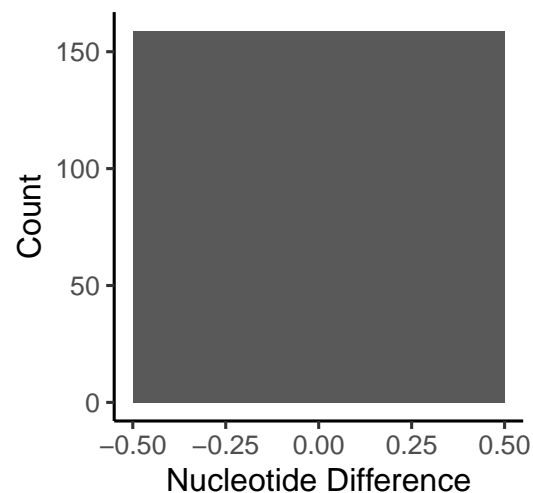
TRBV27*01

165 sequences assigned
165 (100%) exact matches, in which:
165 unique CDR3
12 unique J



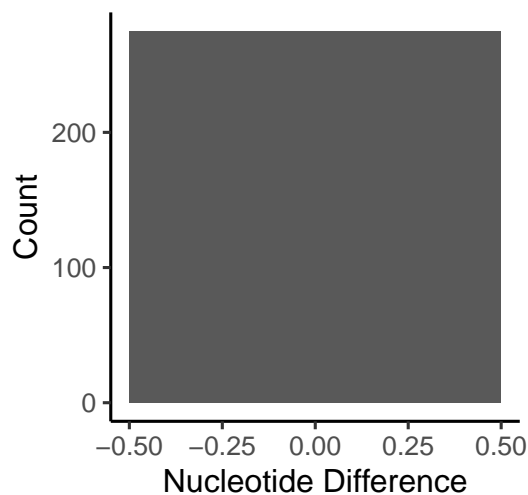
TRBV9*01

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



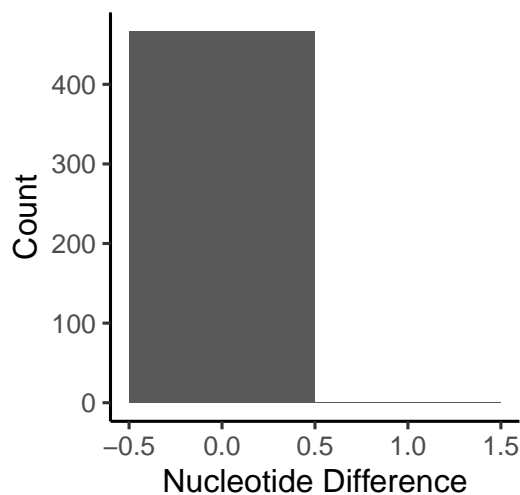
TRBV19*01

275 sequences assigned
275 (100%) exact matches, in which:
275 unique CDR3
13 unique J



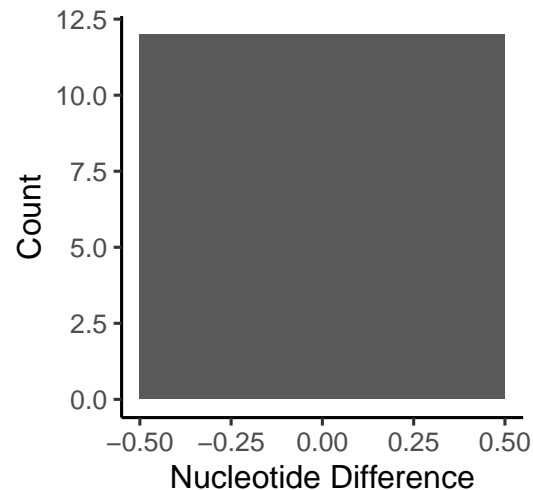
TRBV28*01

468 sequences assigned
467 (99.8%) exact matches, in which:
467 unique CDR3
13 unique J



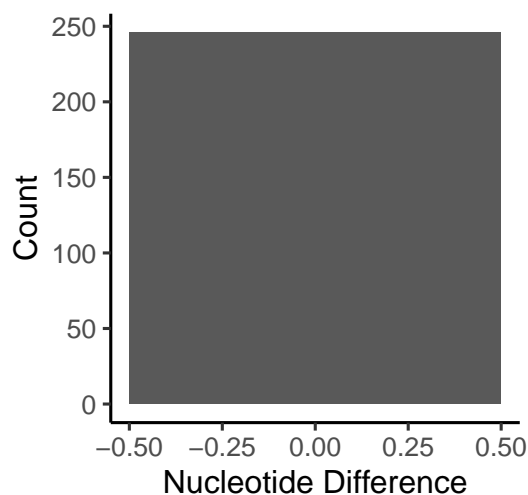
TRBV11-1*01

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



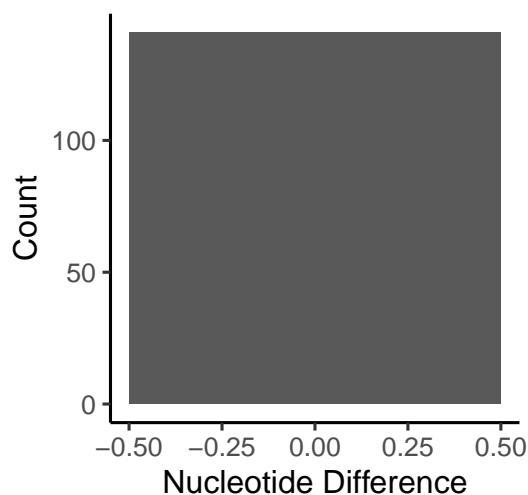
TRBV2*01

246 sequences assigned
246 (100%) exact matches, in which:
246 unique CDR3
13 unique J



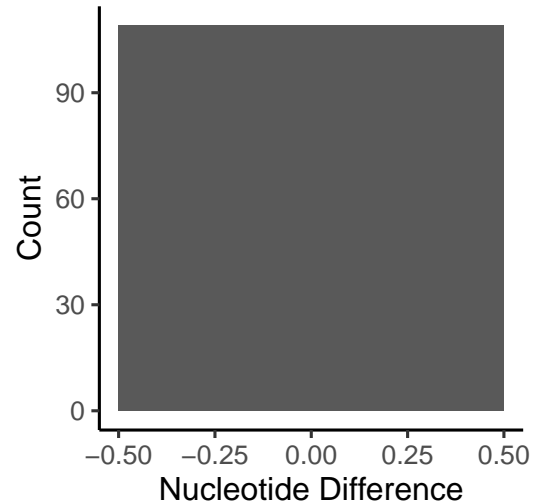
TRBV30*01

141 sequences assigned
141 (100%) exact matches, in which:
141 unique CDR3
13 unique J



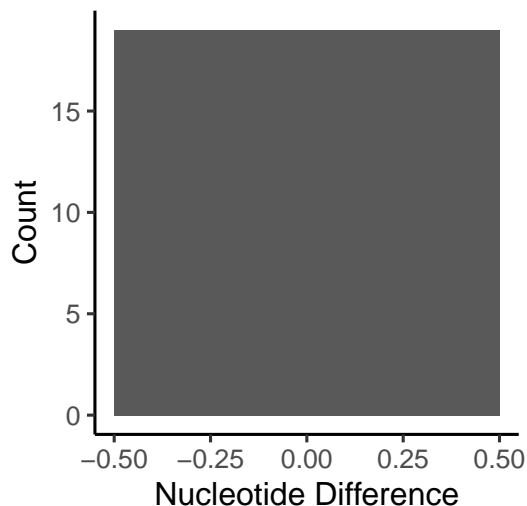
TRBV11-2*01

109 sequences assigned
109 (100%) exact matches, in which:
109 unique CDR3
12 unique J



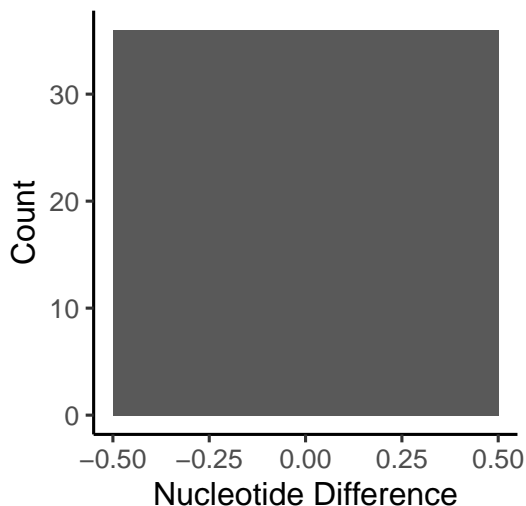
TRBV11–3*01

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



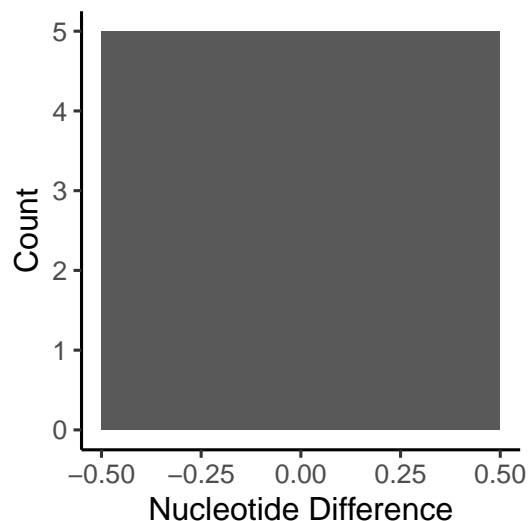
TRBV12–4*01

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



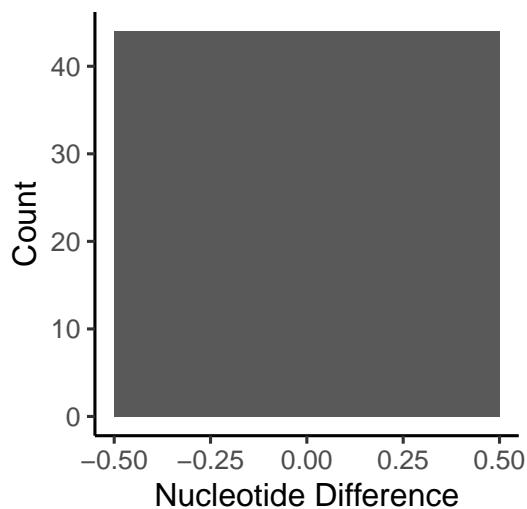
TRBV12–5*01_C28G

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



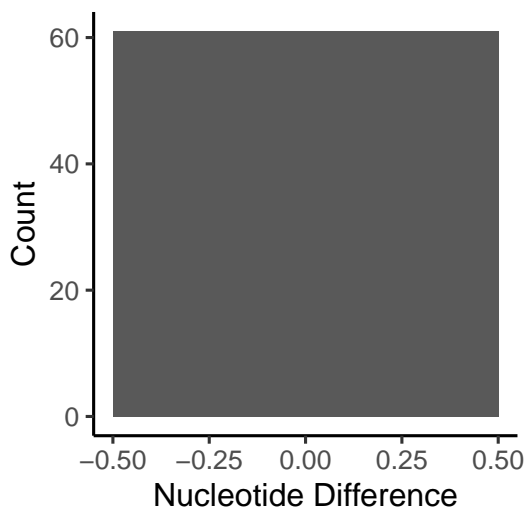
TRBV11–3*04

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



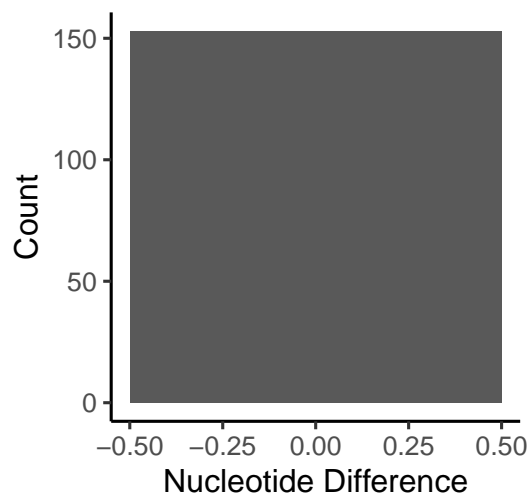
TRBV12–4*01_C87T

61 sequences assigned
61 (100%) exact matches, in which:
61 unique CDR3
11 unique J



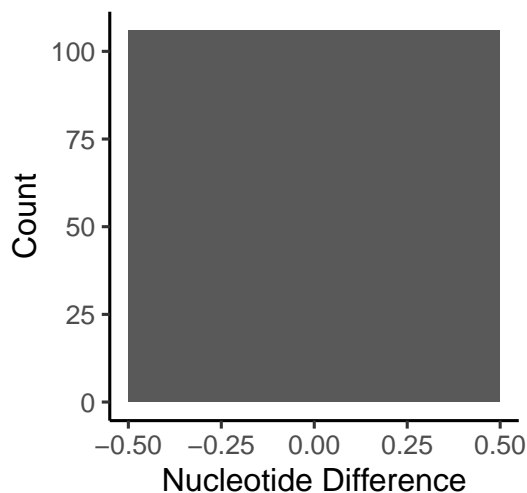
TRBV3–1*01

153 sequences assigned
153 (100%) exact matches, in which:
153 unique CDR3
13 unique J



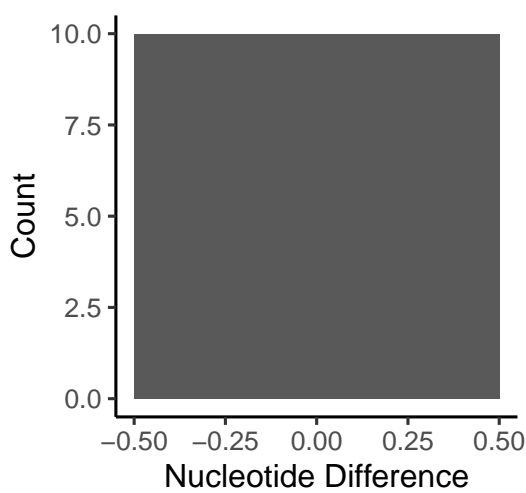
TRBV12–3*01

106 sequences assigned
106 (100%) exact matches, in which:
106 unique CDR3
13 unique J



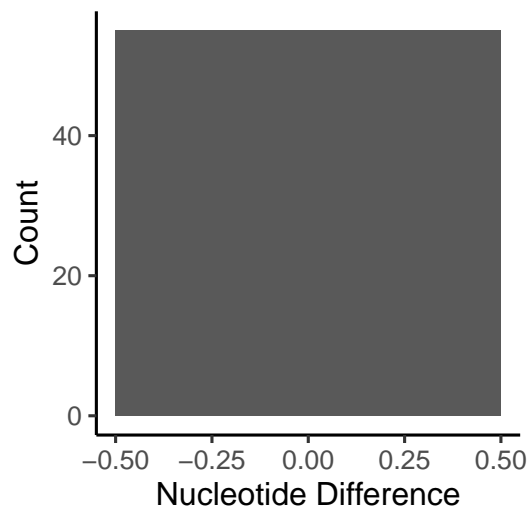
TRBV12–5*01

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



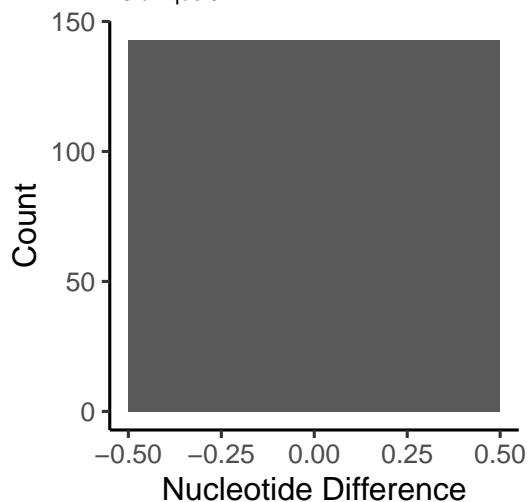
TRBV24–1*01

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



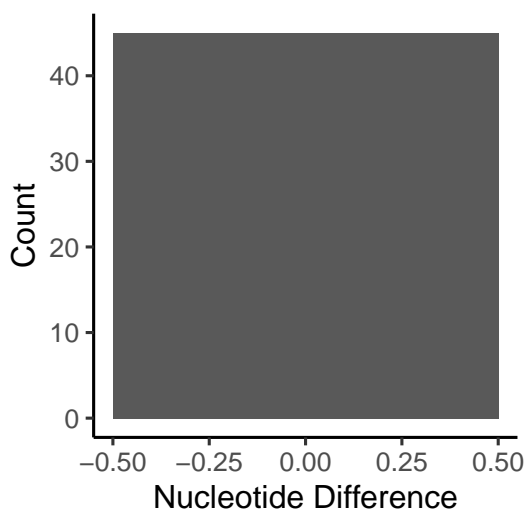
TRBV4–1*01

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



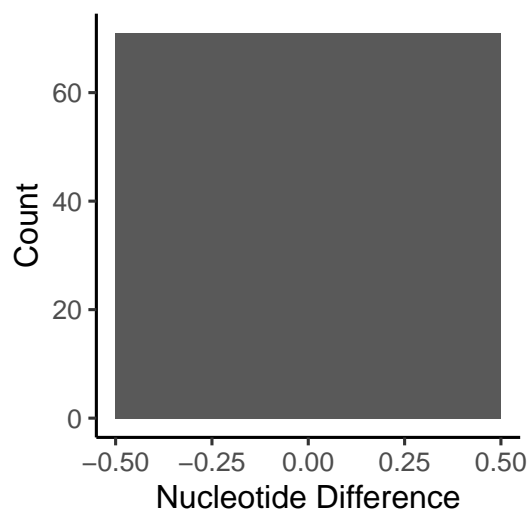
TRBV25–1*01

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



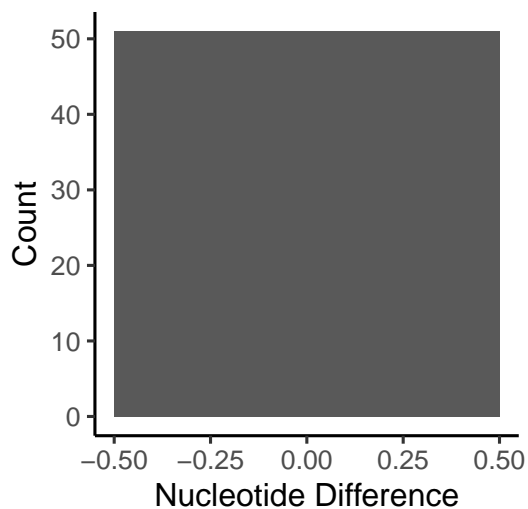
TRBV5–6*01

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



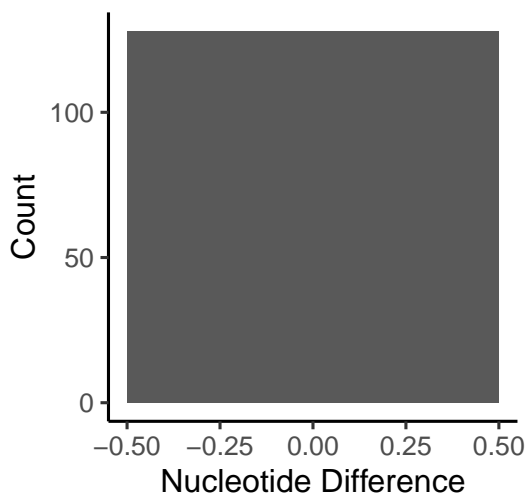
TRBV4–2*01

51 sequences assigned
51 (100%) exact matches, in which:
51 unique CDR3
12 unique J



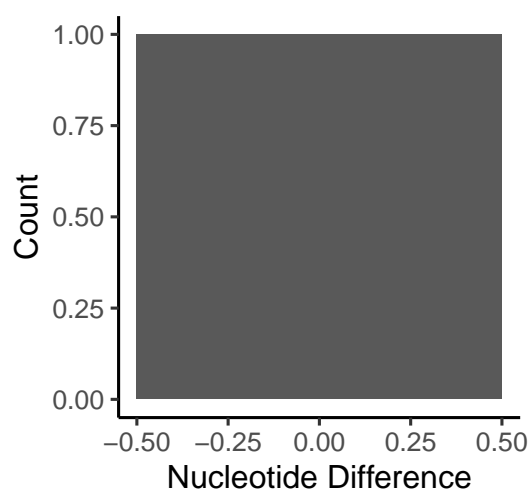
TRBV5–4*01

128 sequences assigned
128 (100%) exact matches, in which:
128 unique CDR3
11 unique J



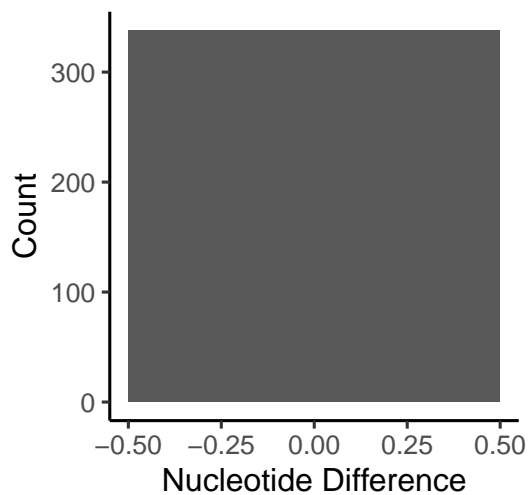
TRBV5–7*01

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



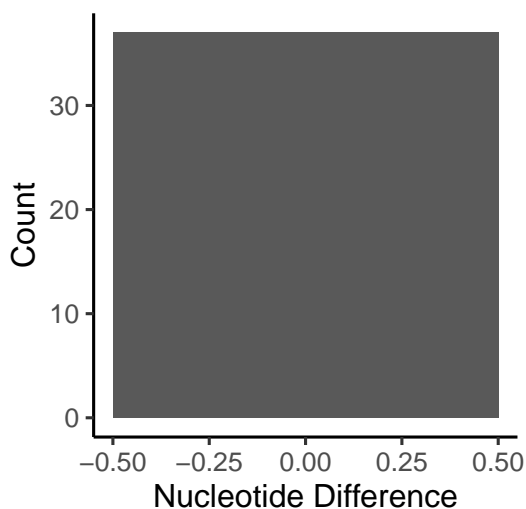
TRBV5–1*01

338 sequences assigned
338 (100%) exact matches, in which:
338 unique CDR3
13 unique J



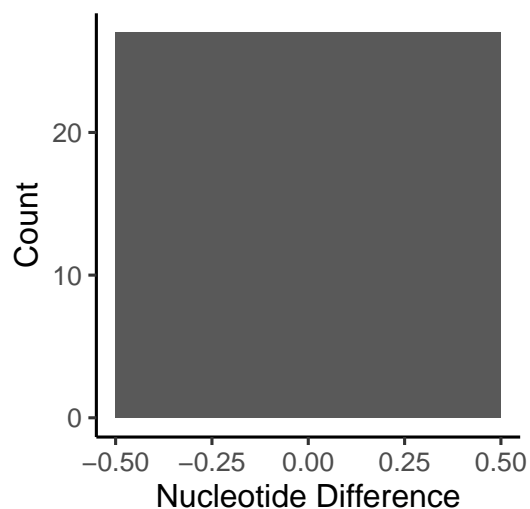
TRBV5–5*02_322TTGG32

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



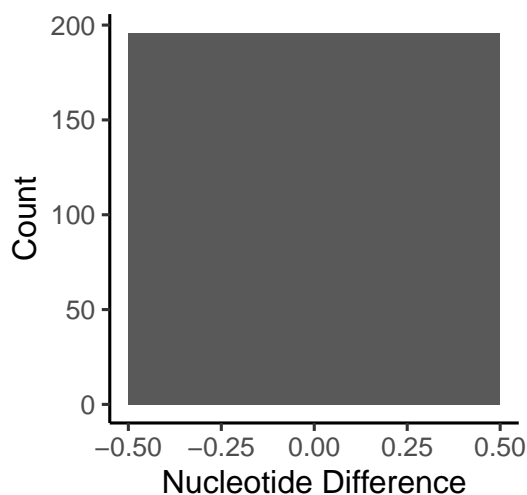
TRBV5–8*01

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



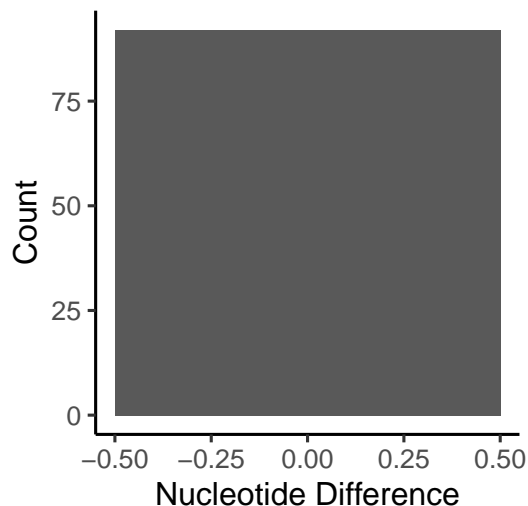
TRBV6-1*01

196 sequences assigned
196 (100%) exact matches, in which:
196 unique CDR3
13 unique J



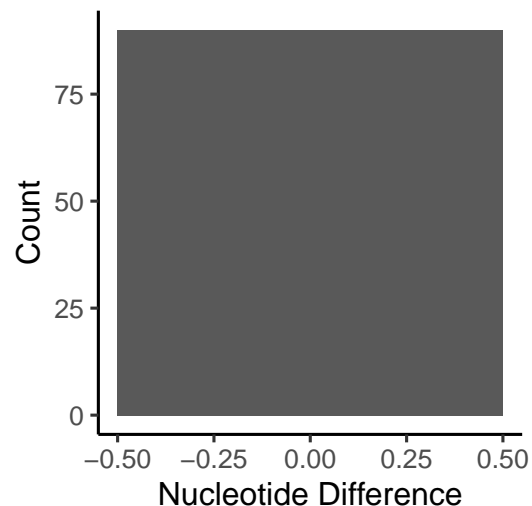
TRBV6-6*01

92 sequences assigned
92 (100%) exact matches, in which:
92 unique CDR3
13 unique J



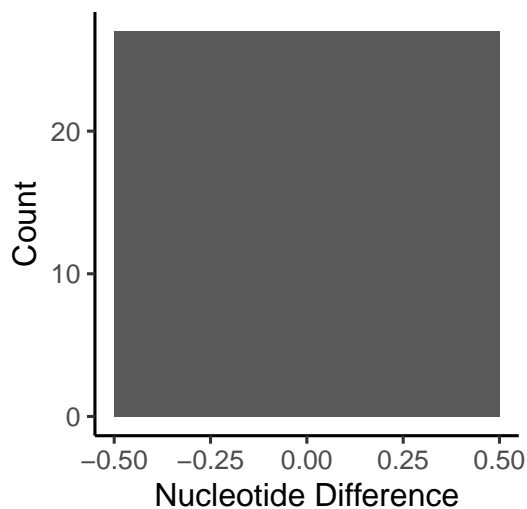
TRBV7-2*02

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



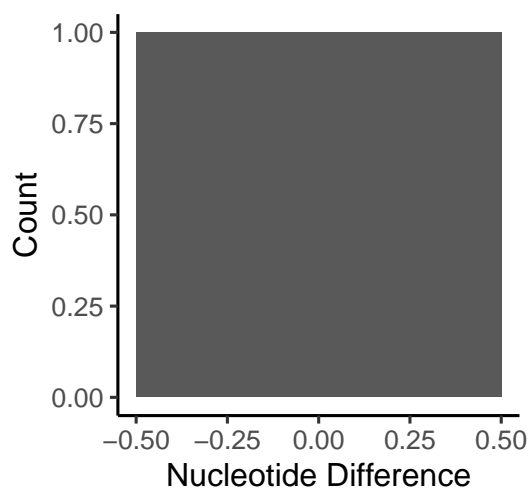
TRBV6-4*01

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



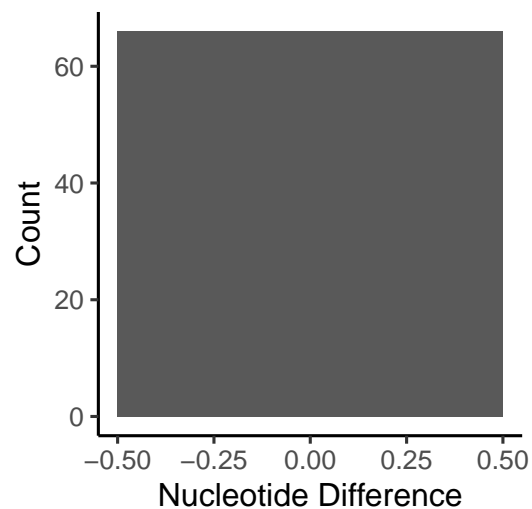
TRBV6-9*01

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



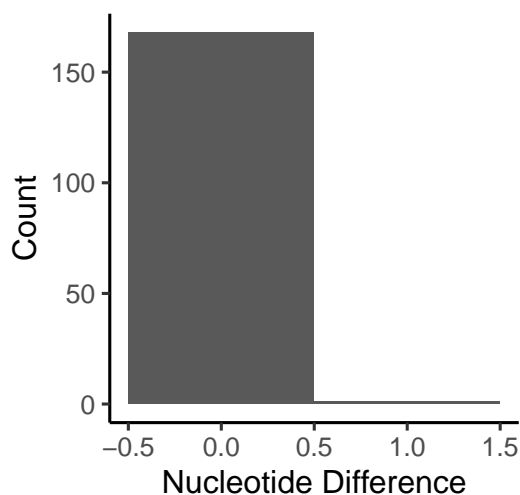
TRBV7-3*01

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



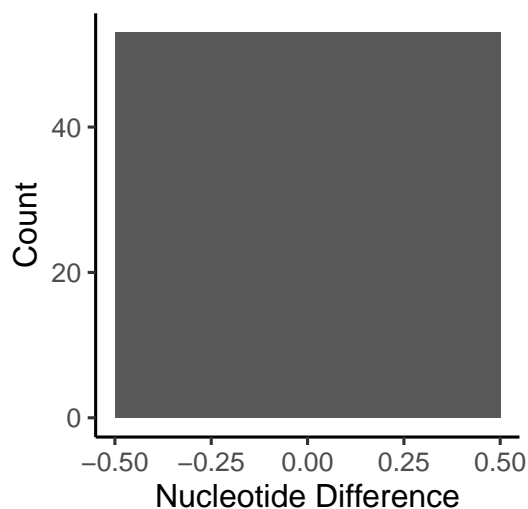
TRBV6-5*01

169 sequences assigned
168 (99.4%) exact matches, in which:
168 unique CDR3
13 unique J



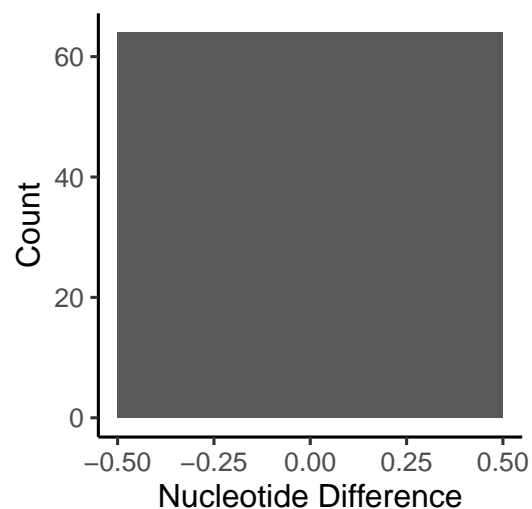
TRBV6-23*01

53 sequences assigned
53 (100%) exact matches, in which:
53 unique CDR3
11 unique J



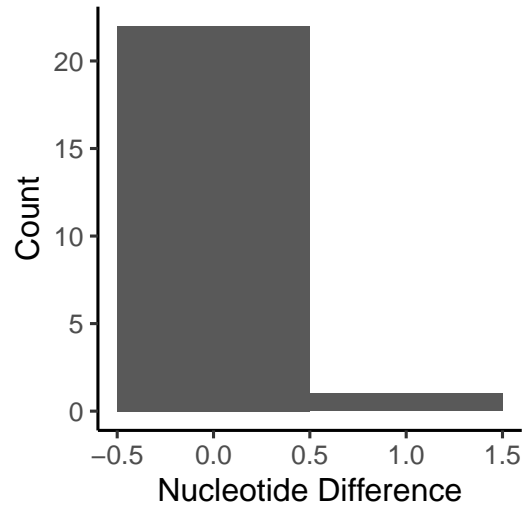
TRBV7-6*01

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



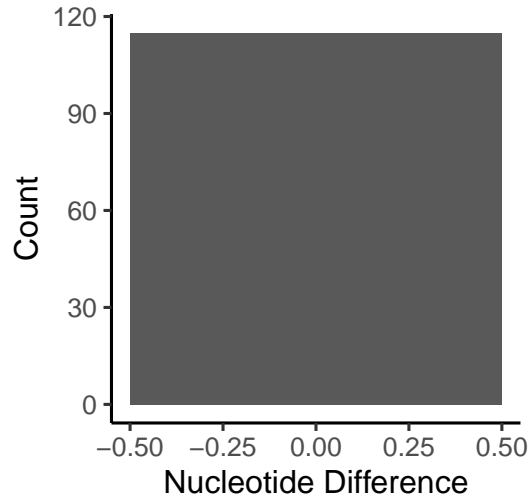
TRBV7-7*01_C315T

23 sequences assigned
22 (95.7%) exact matches, in which:
22 unique CDR3
5 unique J



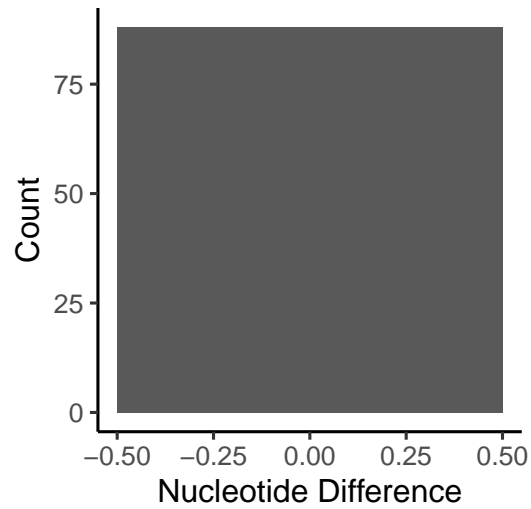
TRBV29-1*01

115 sequences assigned
115 (100%) exact matches, in which:
115 unique CDR3
13 unique J



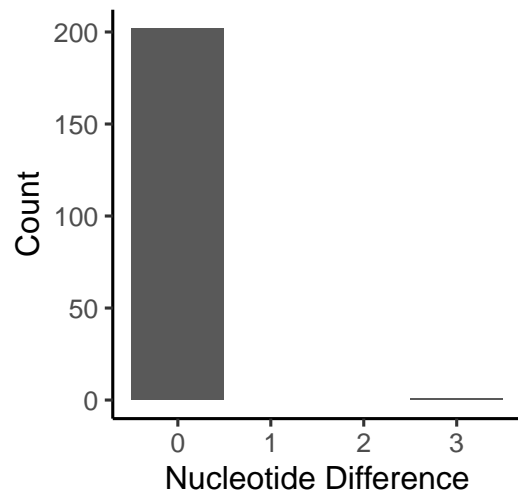
TRBV7-8*01

88 sequences assigned
88 (100%) exact matches, in which:
88 unique CDR3
13 unique J

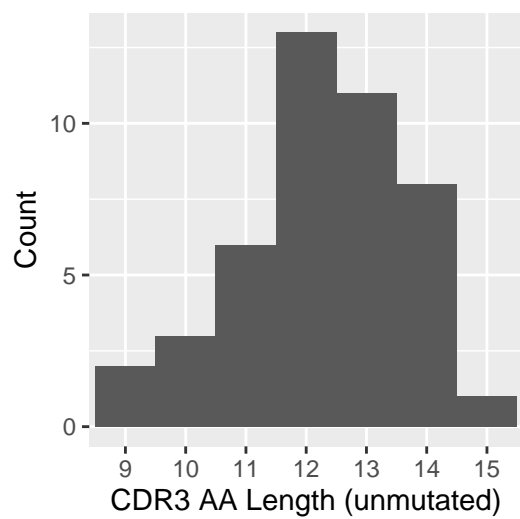


TRBV7-9*03

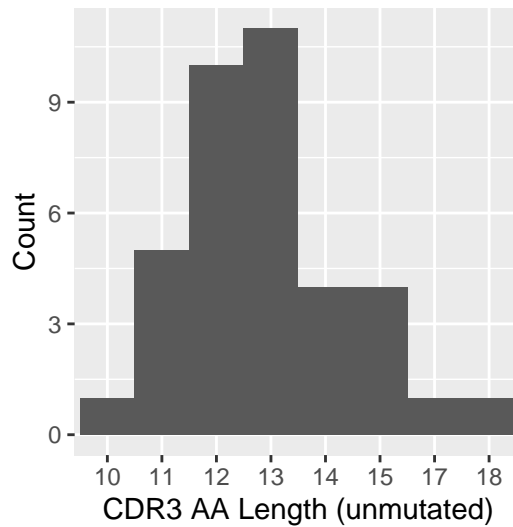
203 sequences assigned
202 (99.5%) exact matches, in which:
202 unique CDR3
13 unique J



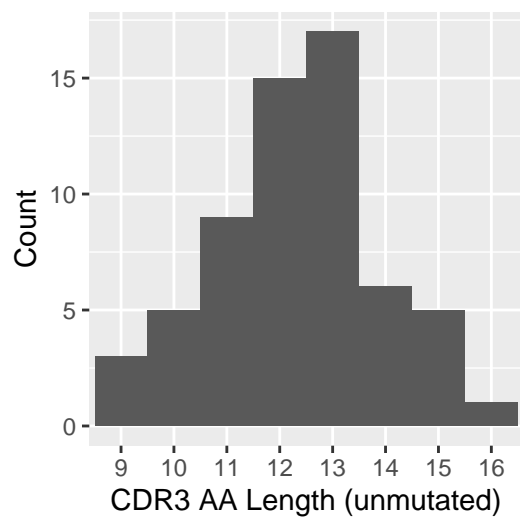
TRBV10-3*03_313GCCATC,



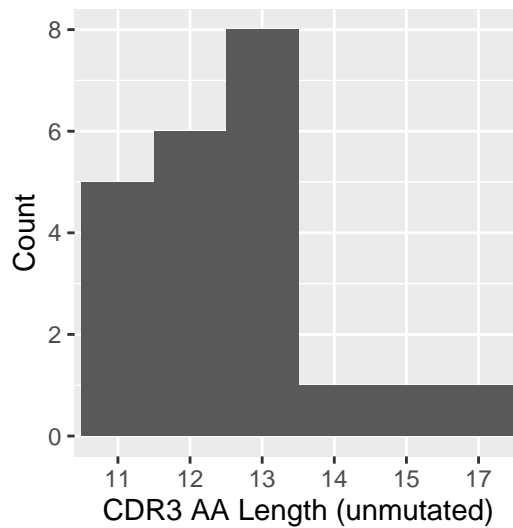
TRBV5-5*02_322TTGG325



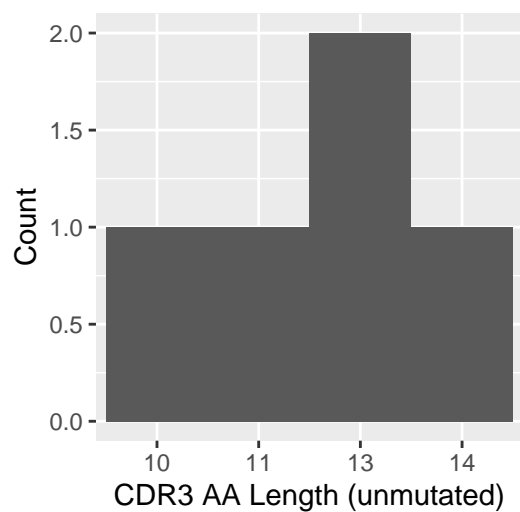
TRBV12-4*01_C87T

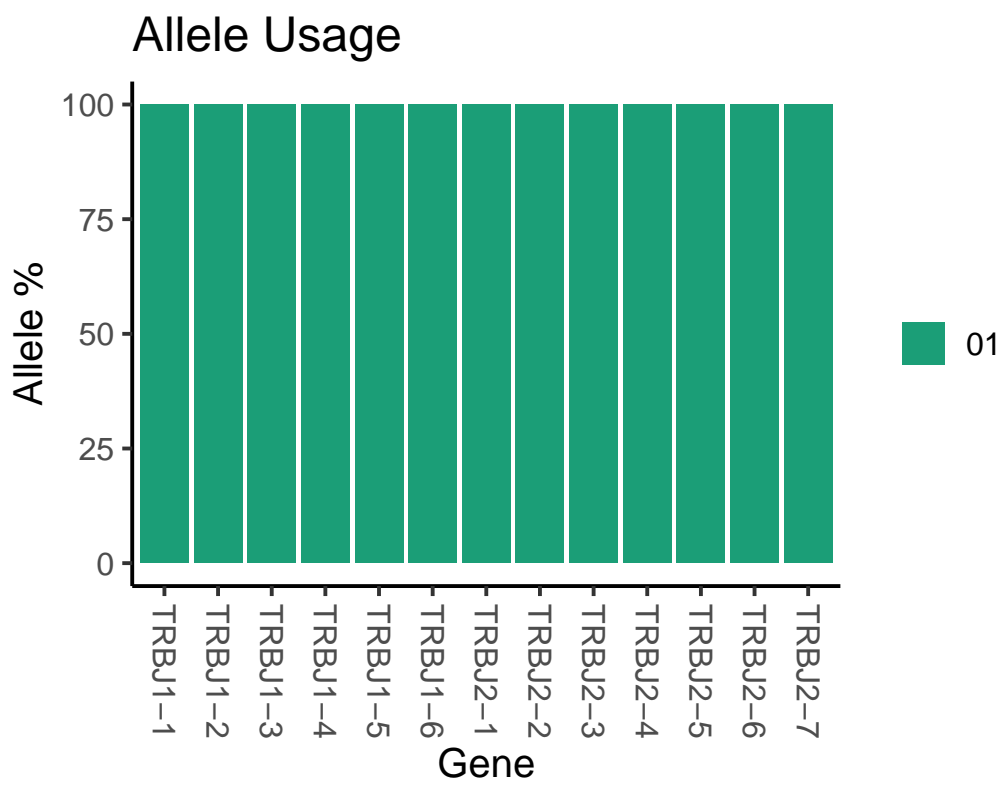


TRBV7-7*01_C315T



TRBV12-5*01_C28G





Warning – no inferred sequences found.

Warning – no inferred sequences found.