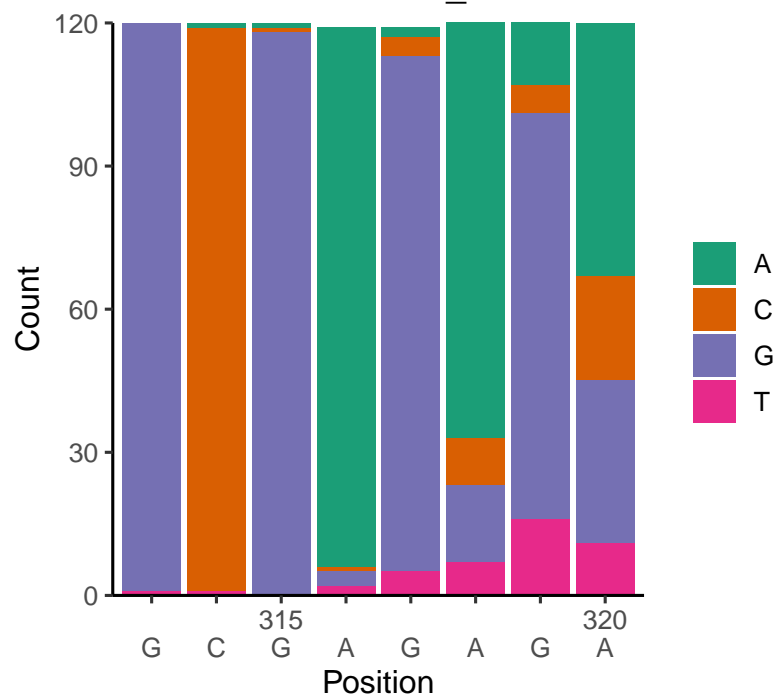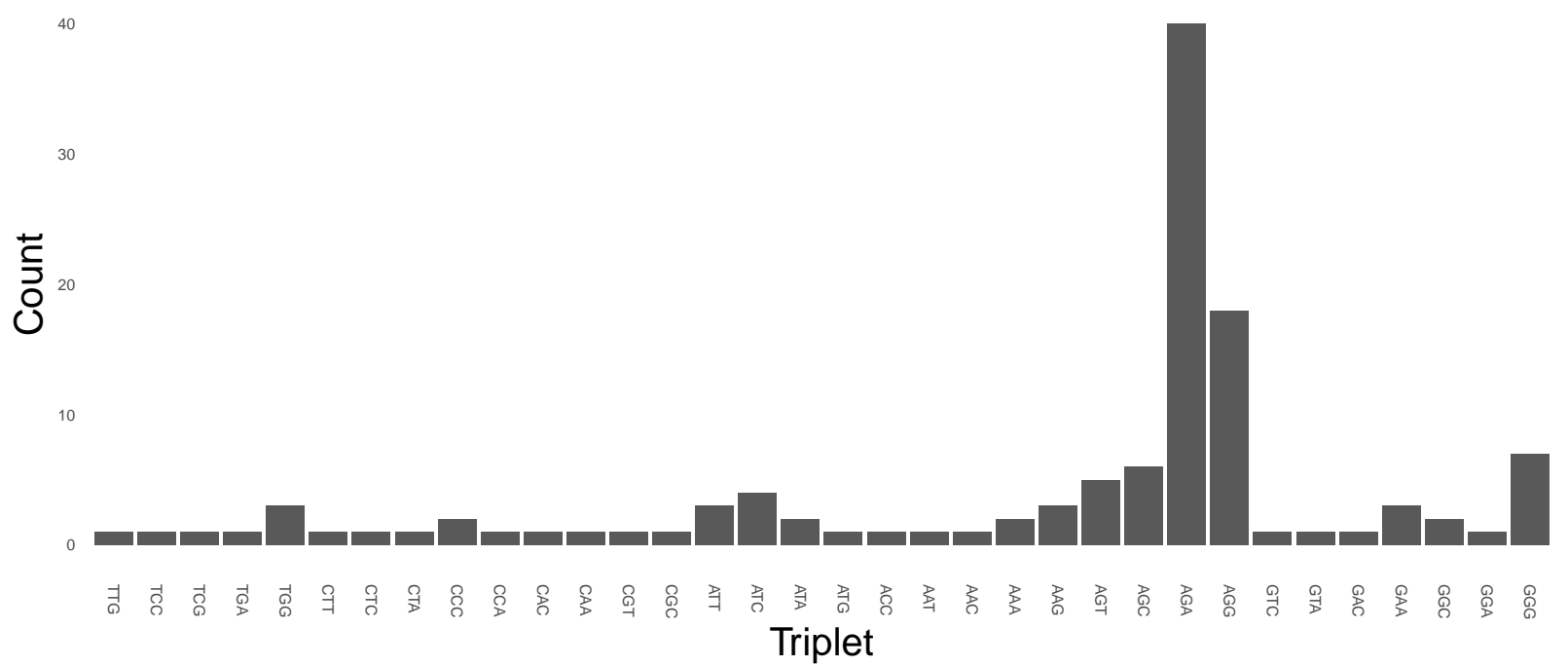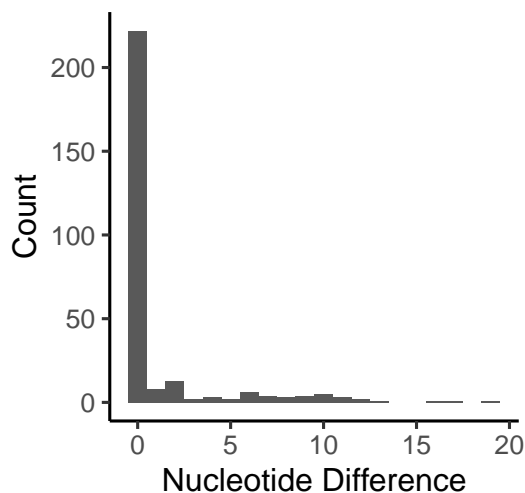Gene IGHV1−69*08_C191T

Gene IGHV1−69*08_C191T

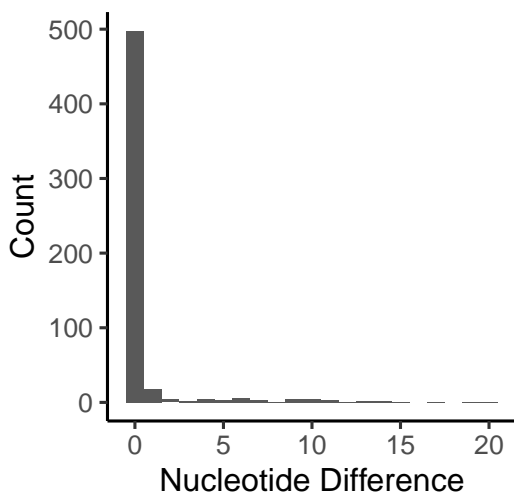IGHV1−69*08_C191T− Final 3 nucleotides as a triplet

IGHV1−2*02

289 sequences assigned
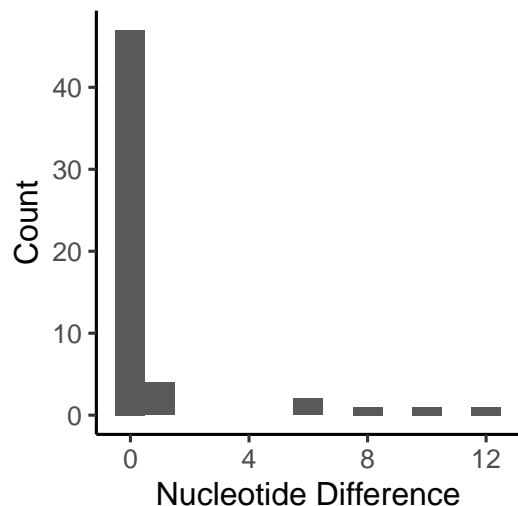222 (76.8%) exact matches, in which:
221 unique CDR3
6 unique J

IGHV1−18*01

572 sequences assigned
498 (87.1%) exact matches, in which:
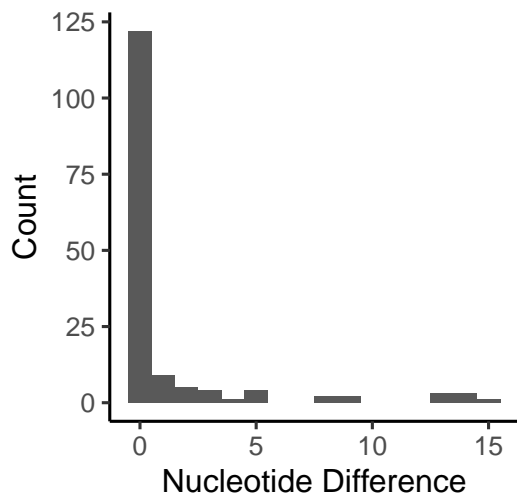498 unique CDR3
6 unique J

IGHV1−46*01

56 sequences assigned
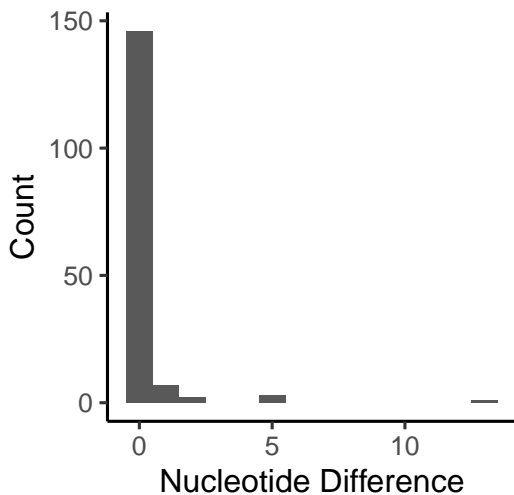47 (83.9%) exact matches, in which:
47 unique CDR3
5 unique J

IGHV1−3*01

159 sequences assigned
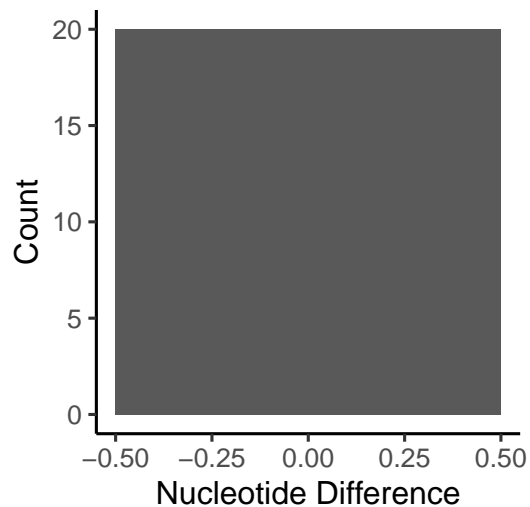122 (76.7%) exact matches, in which:
122 unique CDR3
6 unique J

IGHV1−24*01

159 sequences assigned
146 (91.8%) exact matches, in which:
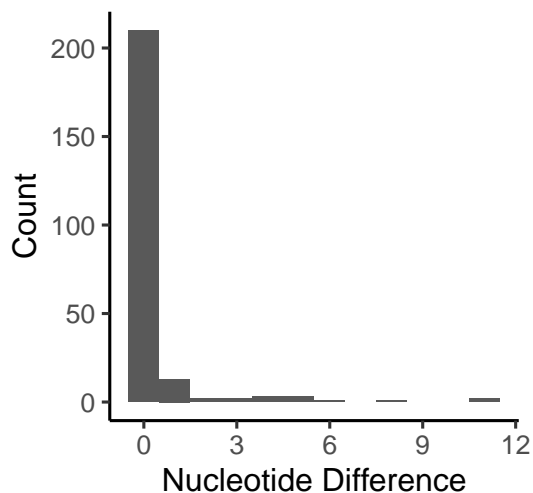146 unique CDR3
6 unique J

IGHV1−46*03

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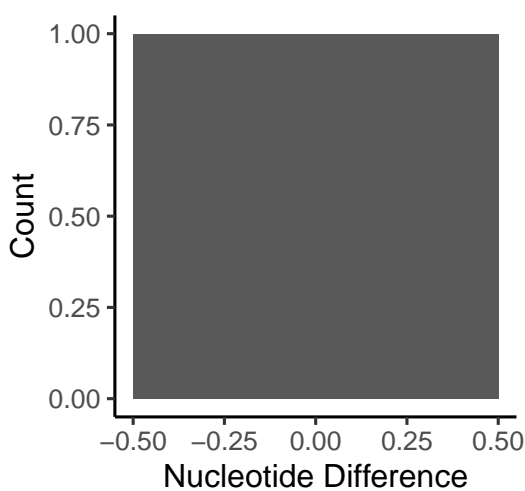20 unique CDR3
5 unique J

IGHV1−8*01

239 sequences assigned
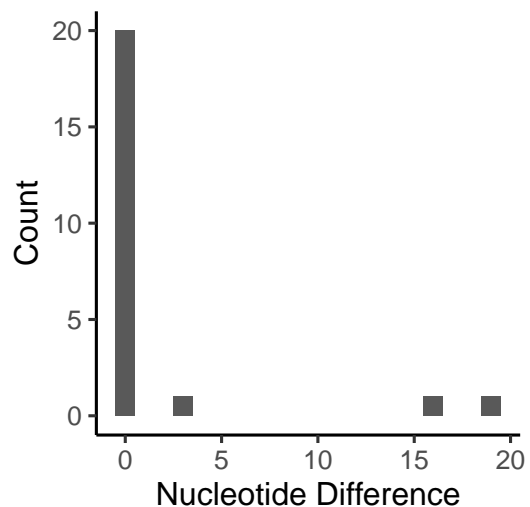210 (87.9%) exact matches, in which:
210 unique CDR3
6 unique J

IGHV1−45*02

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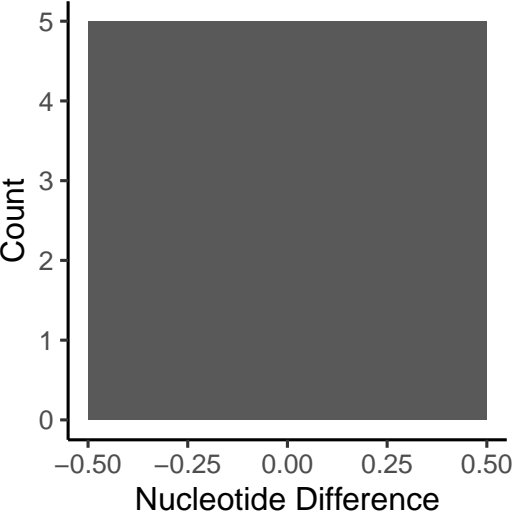1 (100%) exact matches, in which:
1 unique CDR3
1 unique J

IGHV1−58*01

29 sequences assigned
20 (69%) exact matches, in which:
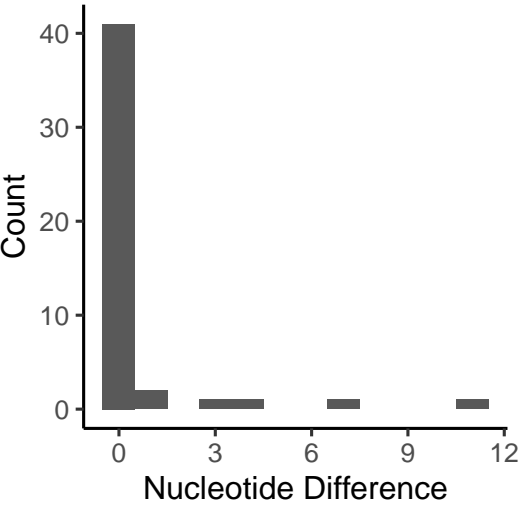20 unique CDR3
5 unique J

IGHV1–58*02

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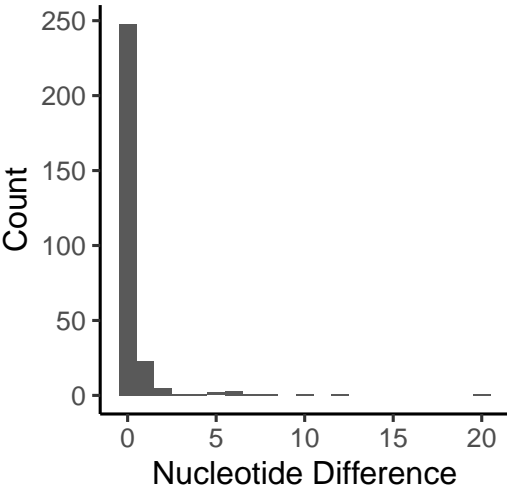5 (100%) exact matches, in which:
5 unique CDR3
2 unique J

IGHV2–26*01

47 sequences assigned
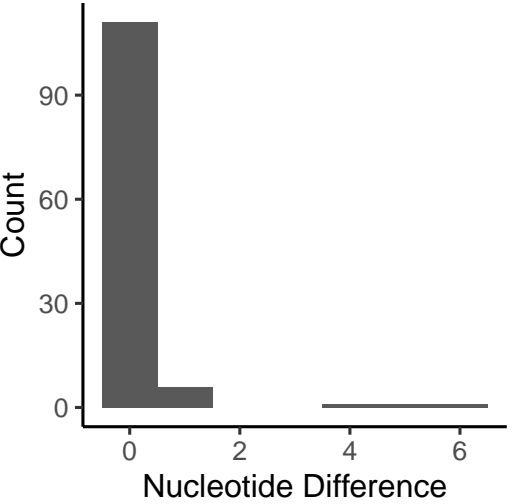41 (87.2%) exact matches, in which:
41 unique CDR3
5 unique J

IGHV3–9*01

288 sequences assigned
248 (86.1%) exact matches, in which:
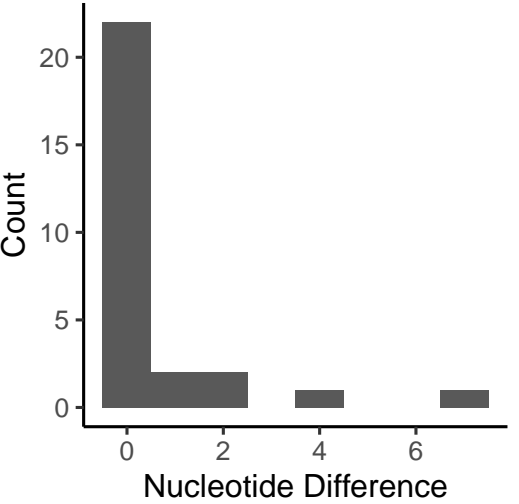248 unique CDR3
6 unique J

IGHV1–69*08_C191T

120 sequences assigned
111 (92.5%) exact matches, in which:
111 unique CDR3
6 unique J

IGHV2–70*01

28 sequences assigned
22 (78.6%) exact matches, in which:
22 unique CDR3
5 unique J

IGHV3–11*01

109 sequences assigned
82 (75.2%) exact matches, in which:
82 unique CDR3
6 unique J

IGHV2–5*01

173 sequences assigned
139 (80.3%) exact matches, in which:
139 unique CDR3
6 unique J

IGHV3–7*01

141 sequences assigned
94 (66.7%) exact matches, in which:
94 unique CDR3
6 unique J

IGHV3–13*01

19 sequences assigned
15 (78.9%) exact matches, in which:
15 unique CDR3
5 unique J

IGHV3−15*01

121 sequences assigned
91 (75.2%) exact matches, in which:
91 unique CDR3
4 unique J

IGHV3−21*01

218 sequences assigned
185 (84.9%) exact matches, in which:
185 unique CDR3
6 unique J

IGHV3−43*01

16 sequences assigned
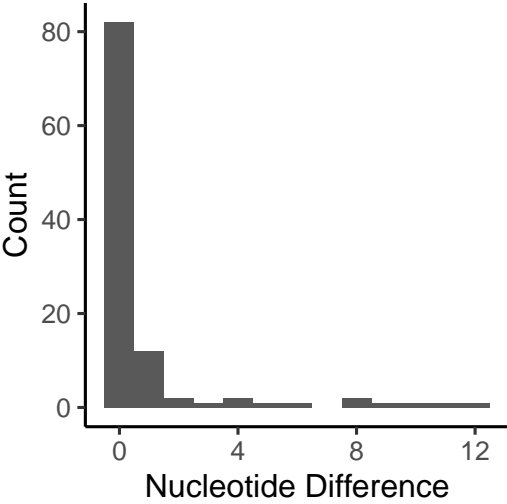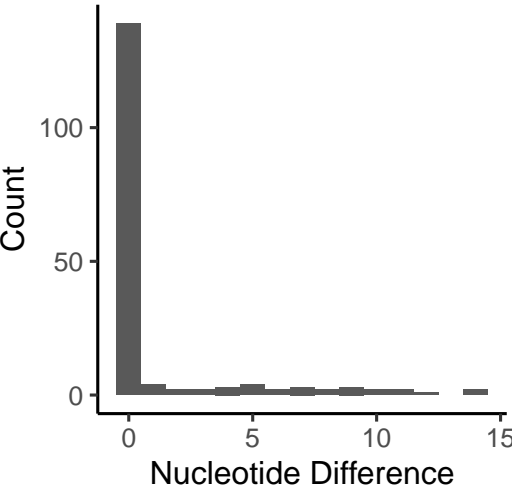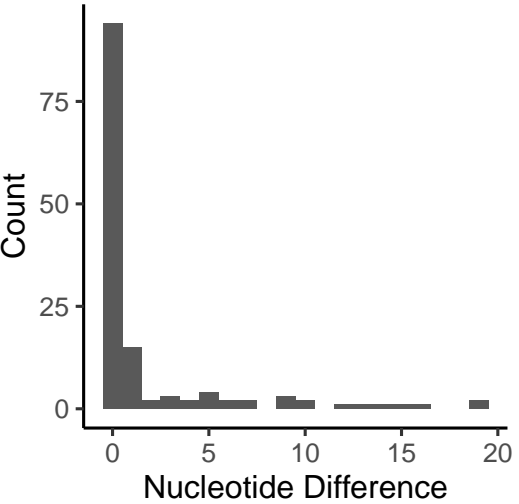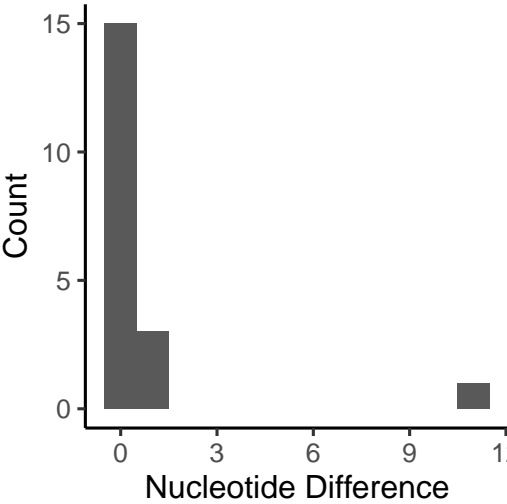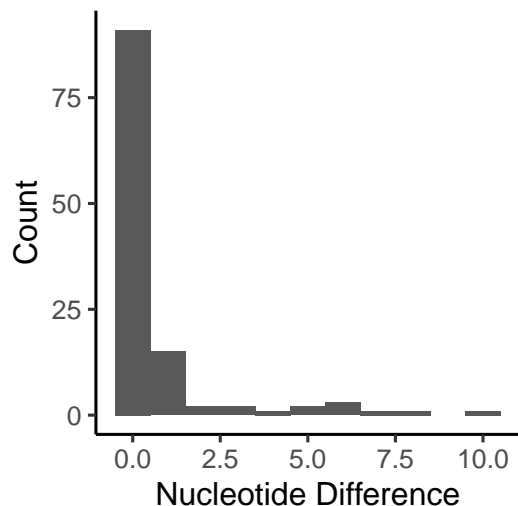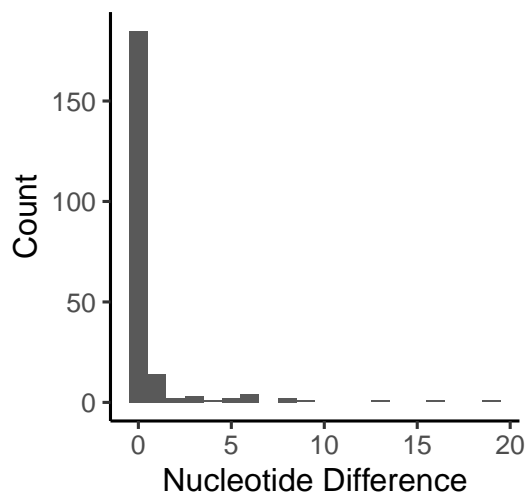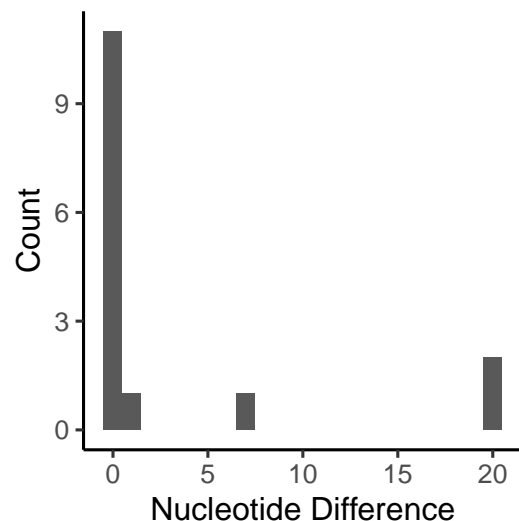11 (68.8%) exact matches, in which:
11 unique CDR3
4 unique J

IGHV3−20*01

9 sequences assigned
7 (77.8%) exact matches, in which:
7 unique CDR3
4 unique J

IGHV3−30−3*01

114 sequences assigned
87 (76.3%) exact matches, in which:
87 unique CDR3
6 unique J

IGHV3−48*01

40 sequences assigned
32 (80%) exact matches, in which:
32 unique CDR3
5 unique J

IGHV3−20*04

3 sequences assigned
No exact matches.

IGHV3−33*01

170 sequences assigned
129 (75.9%) exact matches, in which:
129 unique CDR3
5 unique J

IGHV3−48*02

72 sequences assigned
59 (81.9%) exact matches, in which:
59 unique CDR3
5 unique J

IGHV3-49*03
24 sequences assigned
16 (66.7%) exact matches, in which:
16 unique CDR3
4 unique J

IGHV3-53*02
21 sequences assigned
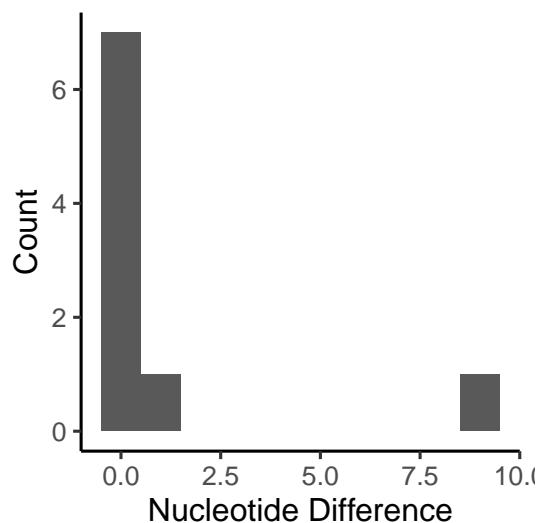15 (71.4%) exact matches, in which:
15 unique CDR3
4 unique J

IGHV3-66*02
25 sequences assigned
17 (68%) exact matches, in which:
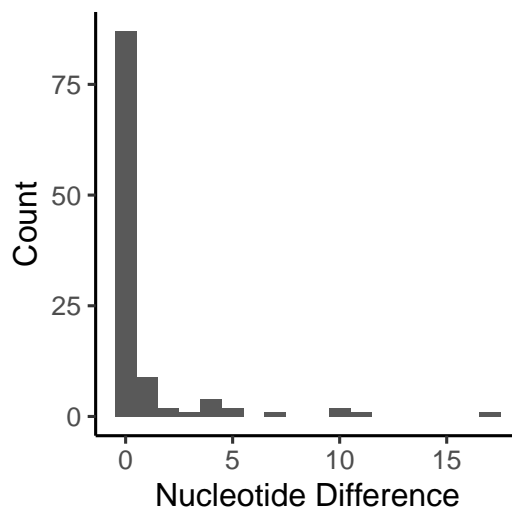17 unique CDR3
4 unique J

IGHV3-49*05
24 sequences assigned
20 (83.3%) exact matches, in which:
20 unique CDR3
4 unique J

IGHV3-64*01
4 sequences assigned
2 (50%) exact matches, in which:
2 unique CDR3
2 unique J

IGHV3-72*01
12 sequences assigned
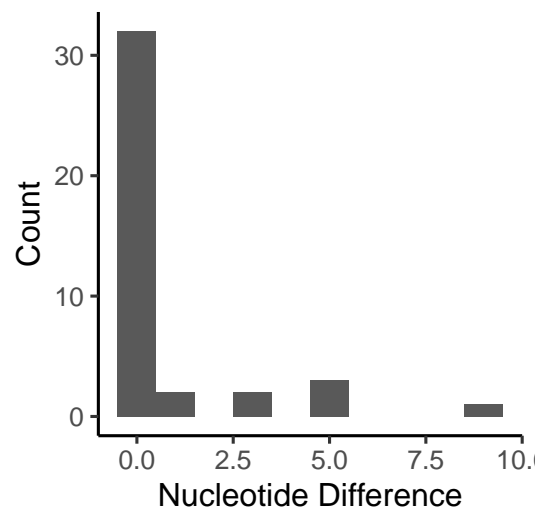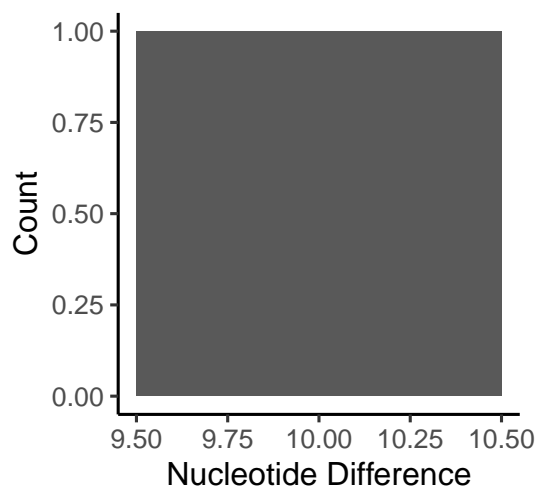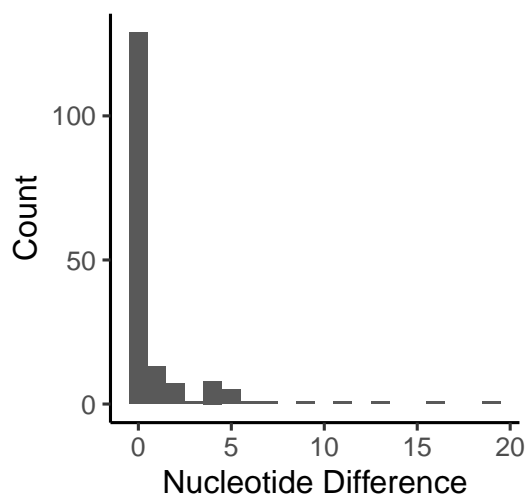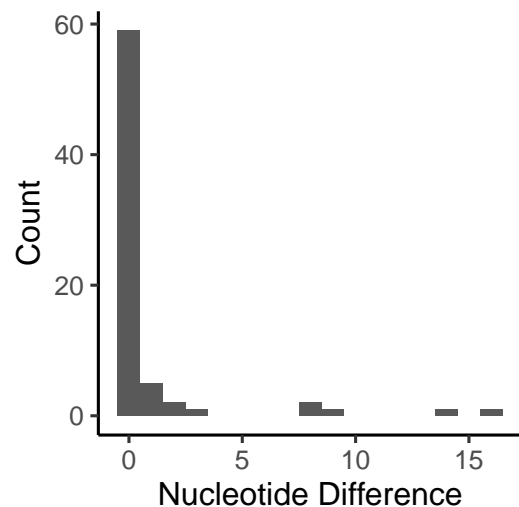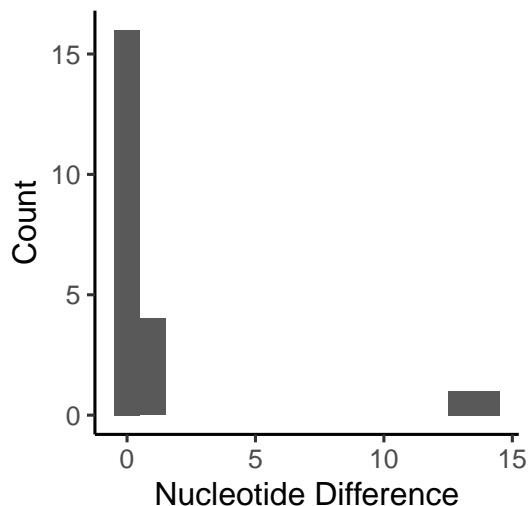1 (8.3%) exact matches, in which:
1 unique CDR3
0 unique J

IGHV3-53*01
79 sequences assigned
65 (82.3%) exact matches, in which:
65 unique CDR3
6 unique J

IGHV3-64*02
1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J

IGHV3-73*01
15 sequences assigned
11 (73.3%) exact matches, in which:
11 unique CDR3
4 unique J

IGHV3−73*02

15 sequences assigned
9 (60%) exact matches, in which:
9 unique CDR3
4 unique J

IGHV4−30−4*01

37 sequences assigned
34 (91.9%) exact matches, in which:
34 unique CDR3
6 unique J

IGHV4−34*01

776 sequences assigned
705 (90.9%) exact matches, in which:
704 unique CDR3
6 unique J
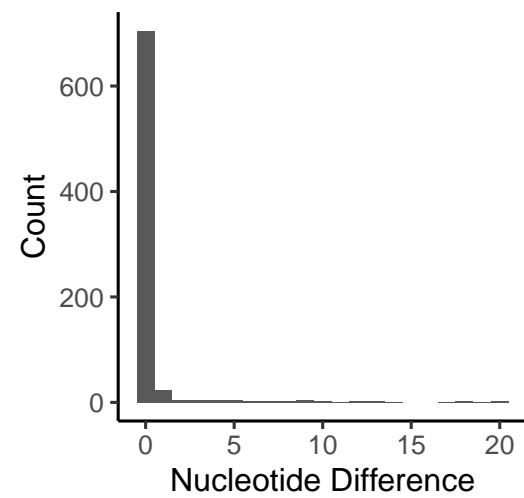
IGHV3−74*01

100 sequences assigned
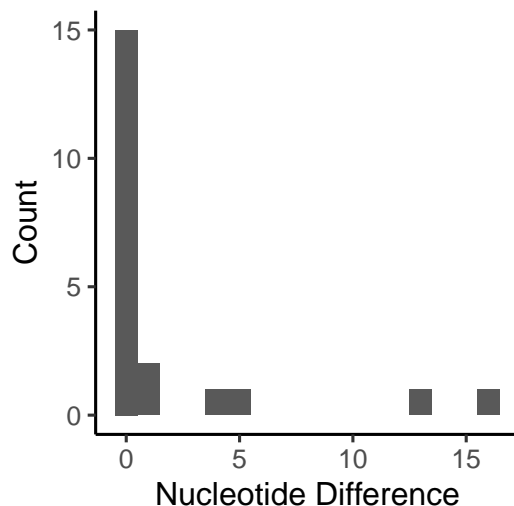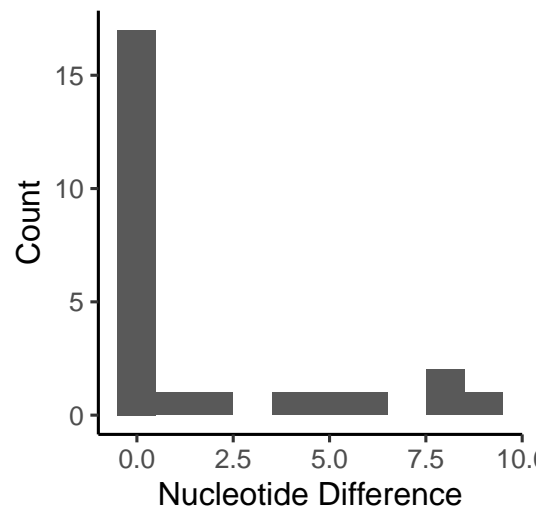69 (69%) exact matches, in which:
69 unique CDR3
5 unique J
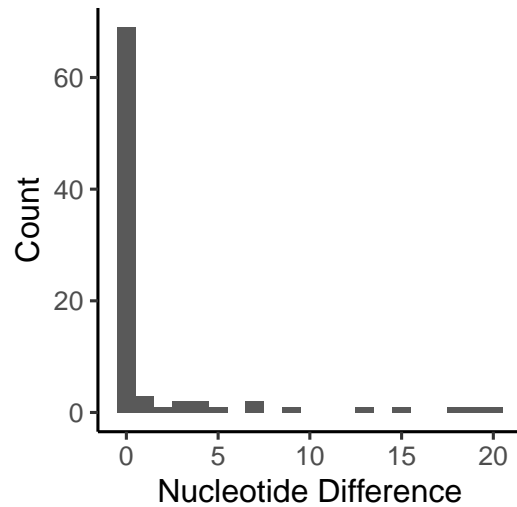
IGHV4−30−2*01

25 sequences assigned
21 (84%) exact matches, in which:
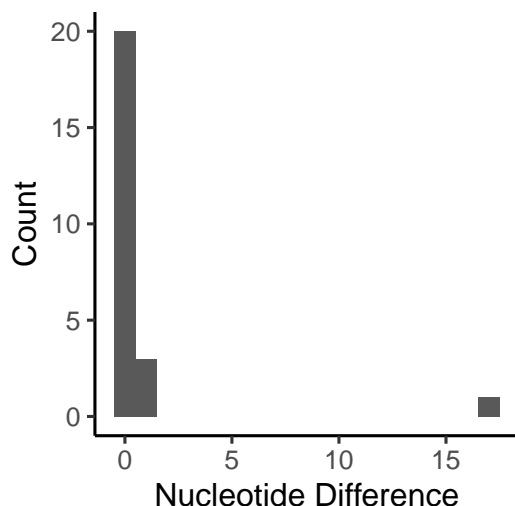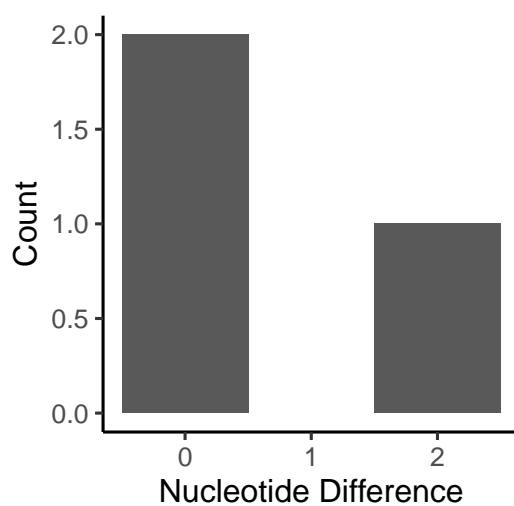21 unique CDR3
5 unique J

IGHV4−39*01

694 sequences assigned
589 (84.9%) exact matches, in which:
589 unique CDR3
6 unique J

IGHV4−4*07

29 sequences assigned
22 (75.9%) exact matches, in which:
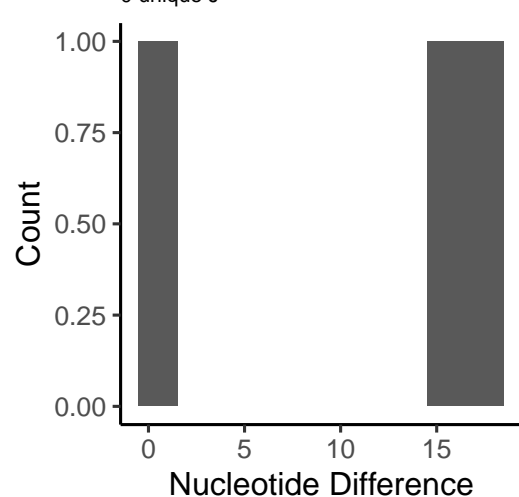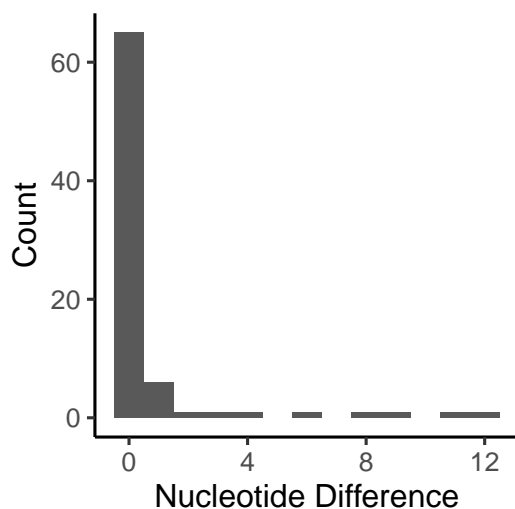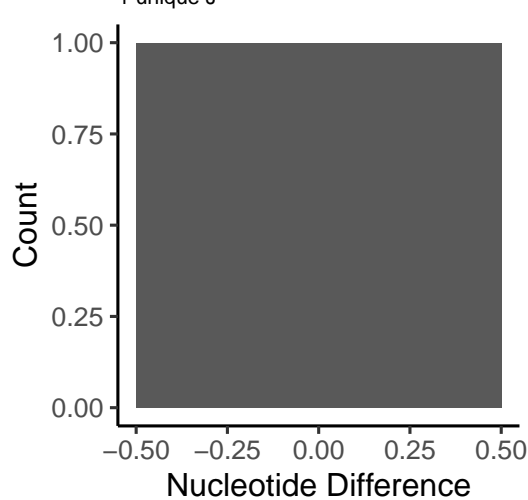22 unique CDR3
5 unique J

IGHV4−31*03

159 sequences assigned
134 (84.3%) exact matches, in which:
134 unique CDR3
6 unique J

IGHV4−59*01

156 sequences assigned
120 (76.9%) exact matches, in which:
120 unique CDR3
5 unique J

IGHV4−61*02

143 sequences assigned
129 (90.2%) exact matches, in which:
129 unique CDR3
6 unique J

IGHV6−1*01

112 sequences assigned
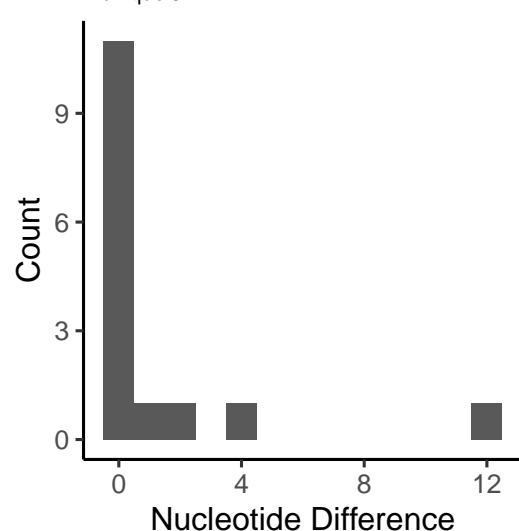89 (79.5%) exact matches, in which:
89 unique CDR3
5 unique J

IGHV5−51*01

376 sequences assigned
278 (73.9%) exact matches, in which:
278 unique CDR3
6 unique J

IGHV7−4−1*02

251 sequences assigned
220 (87.6%) exact matches, in which:
220 unique CDR3
6 unique J

IGHV5−51*03

133 sequences assigned
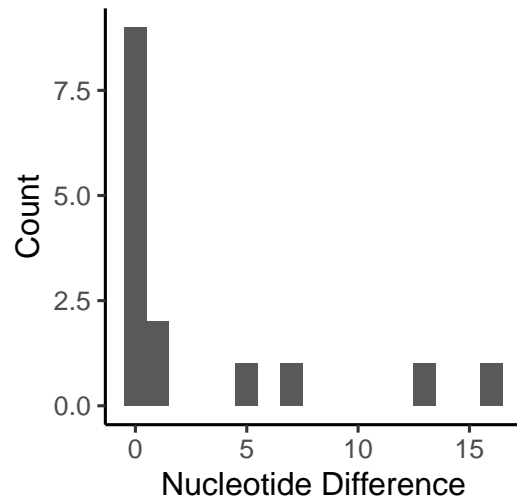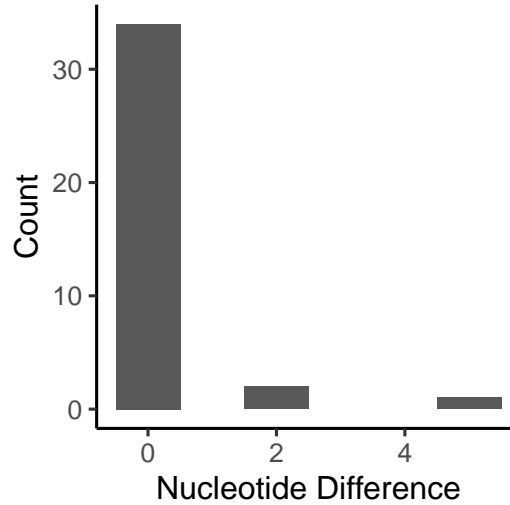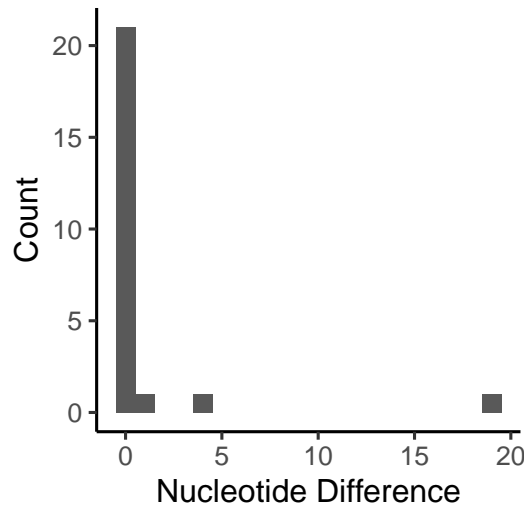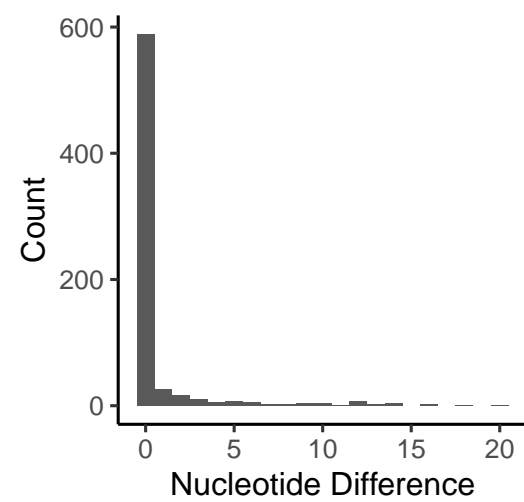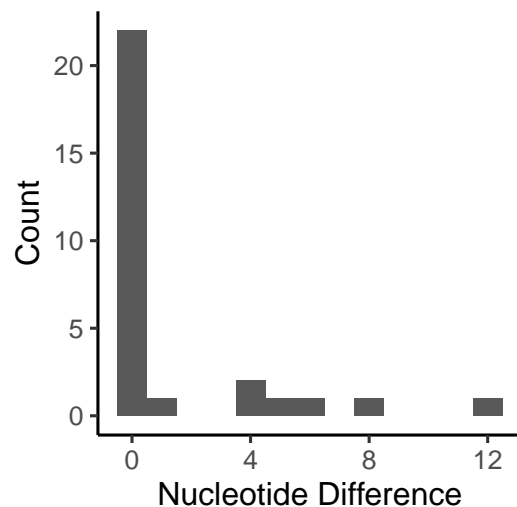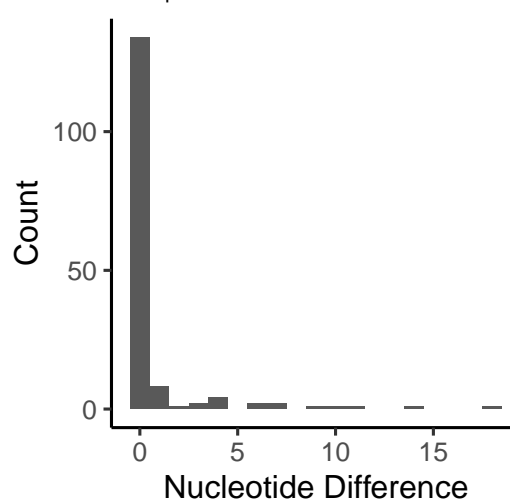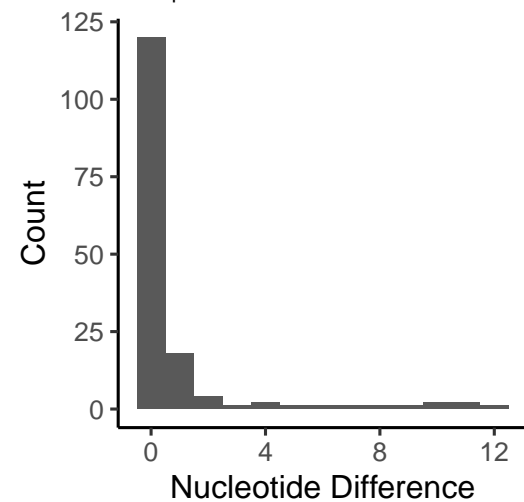119 (89.5%) exact matches, in which:
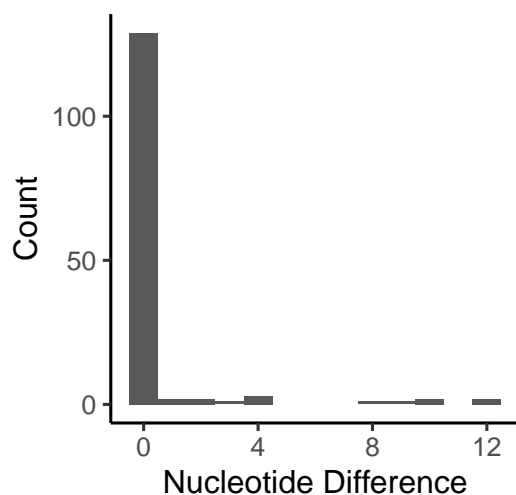119 unique CDR3
5 unique J
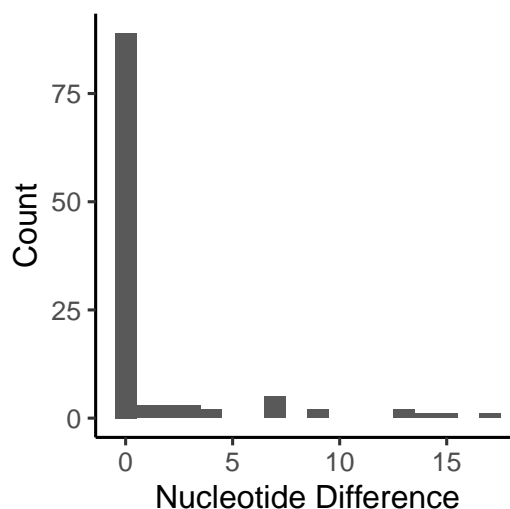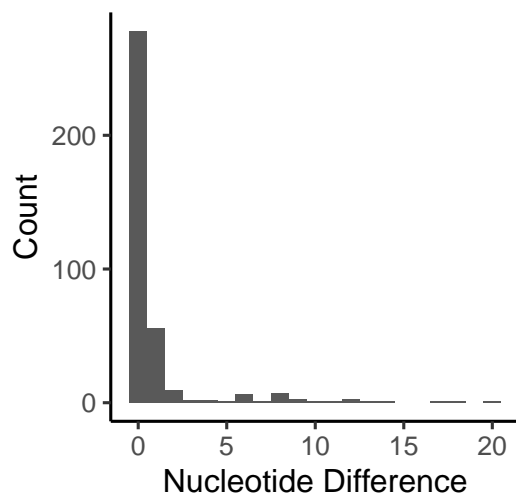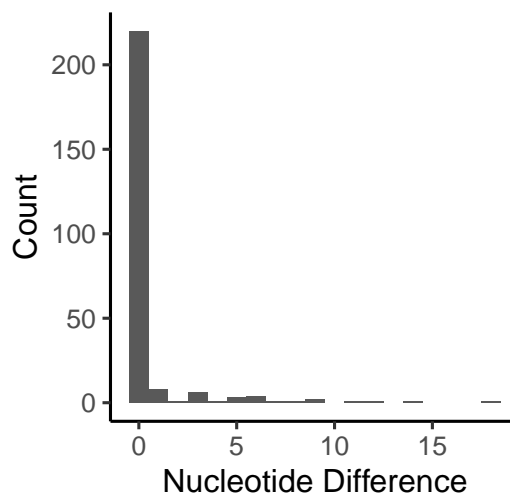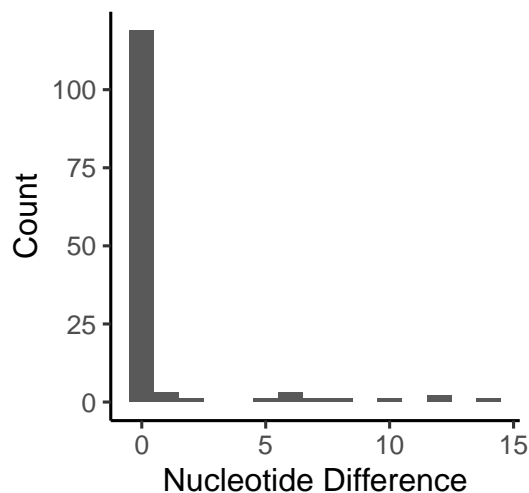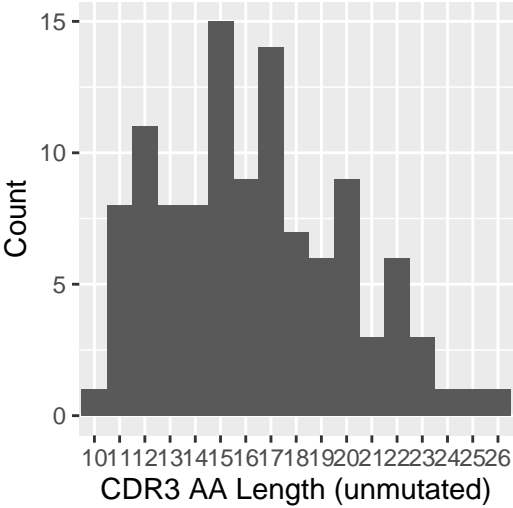
IGHV1−69*08_C191T

Allele Usage

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.

Warning – no inferred sequences found.