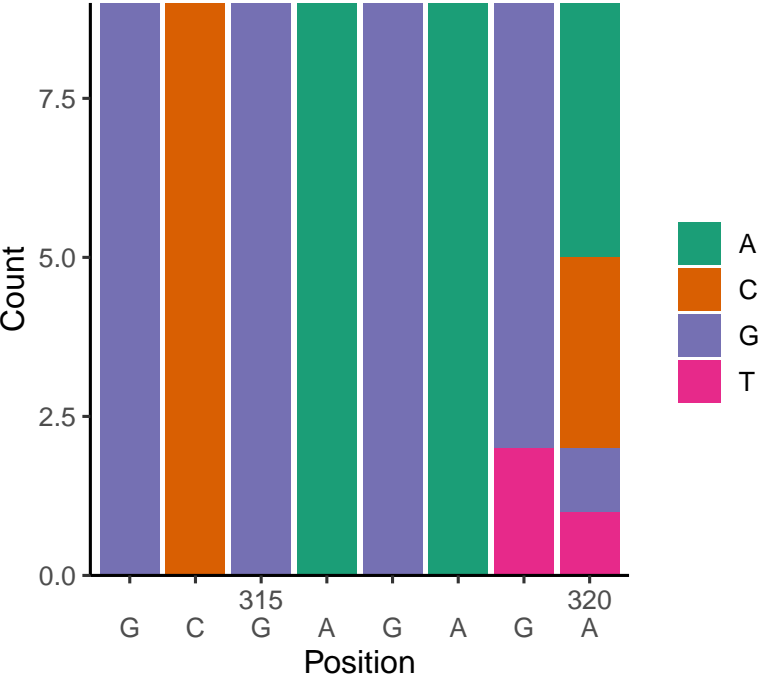


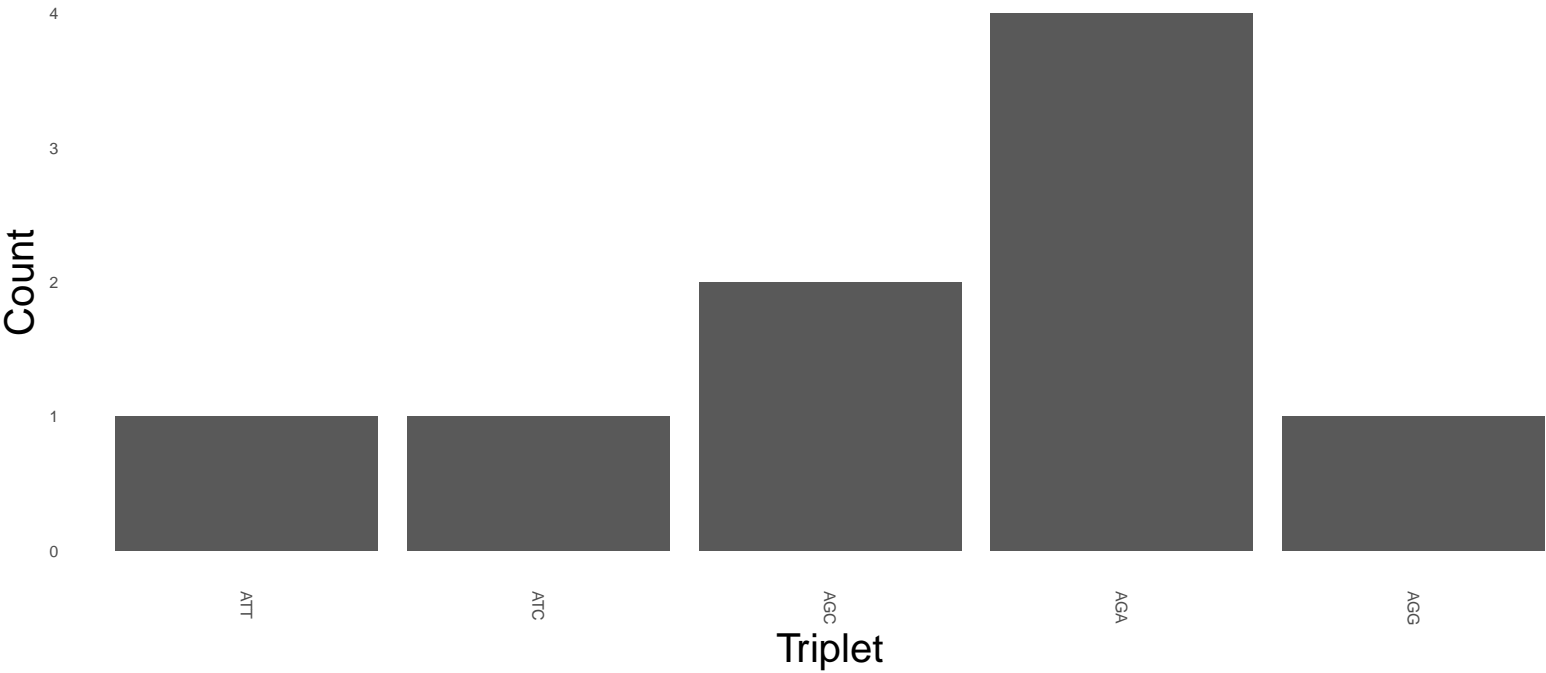
Gene IGHV3-7*03_A131C



Gene IGHV3-7*03_A131C

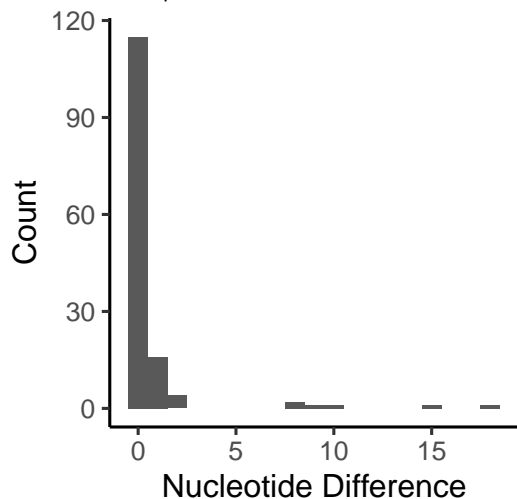


IGHV3-7*03_A131C- Final 3 nucleotides as a triplet



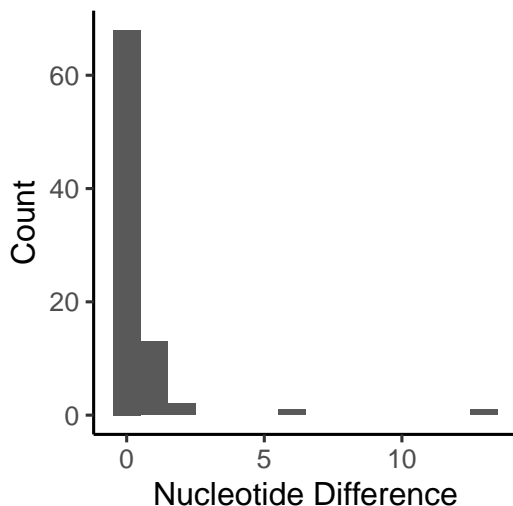
IGHV1-2*02

141 sequences assigned
115 (81.6%) exact matches, in which:
115 unique CDR3
6 unique J



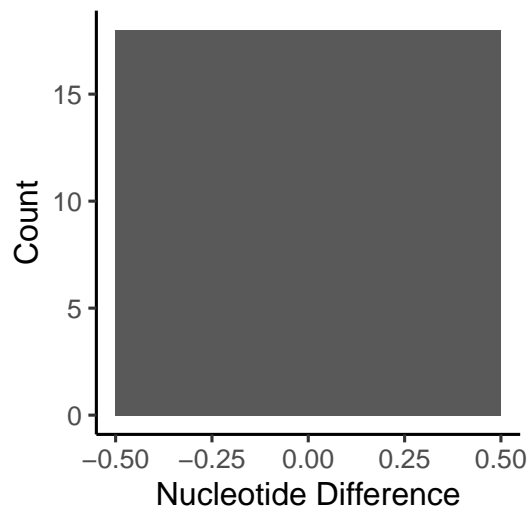
IGHV1-24*01

85 sequences assigned
68 (80%) exact matches, in which:
68 unique CDR3
5 unique J



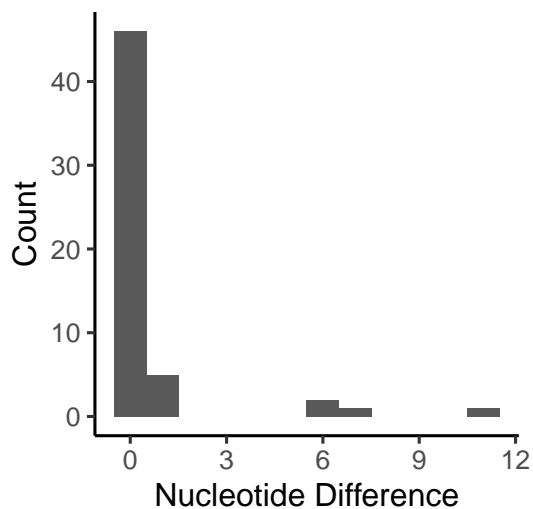
IGHV1-58*02

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



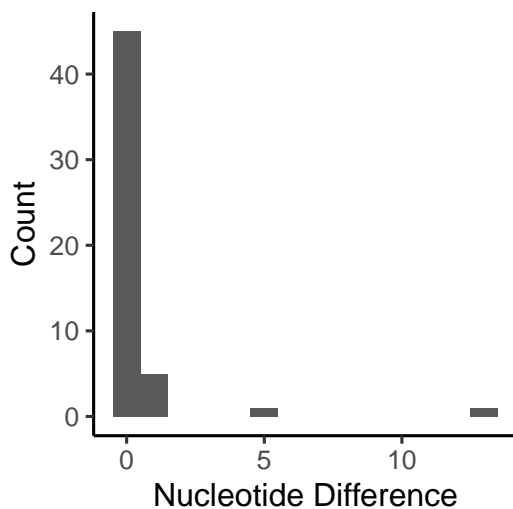
IGHV1-3*01

55 sequences assigned
46 (83.6%) exact matches, in which:
46 unique CDR3
6 unique J



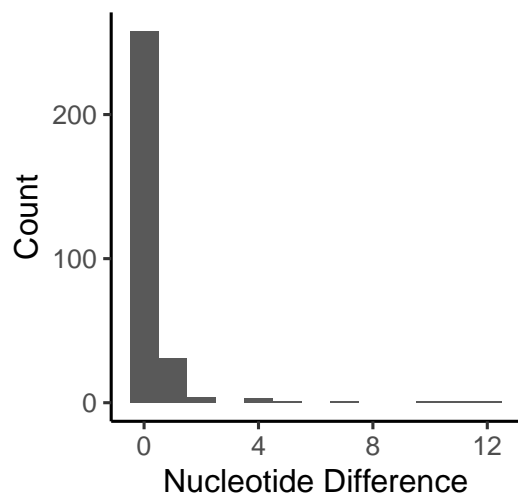
IGHV1-46*01

52 sequences assigned
45 (86.5%) exact matches, in which:
45 unique CDR3
6 unique J



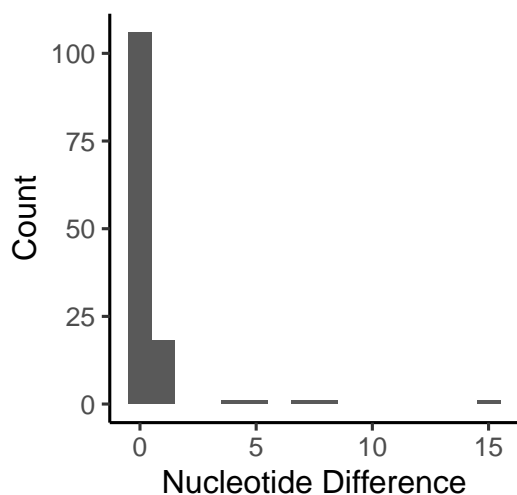
IGHV1-69*04

302 sequences assigned
258 (85.4%) exact matches, in which:
257 unique CDR3
6 unique J



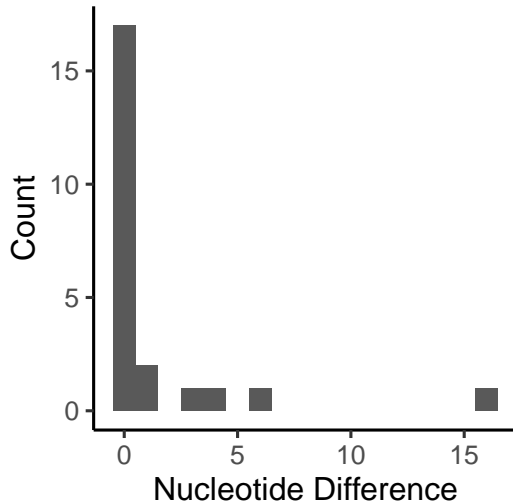
IGHV1-18*01

130 sequences assigned
106 (81.5%) exact matches, in which:
106 unique CDR3
5 unique J



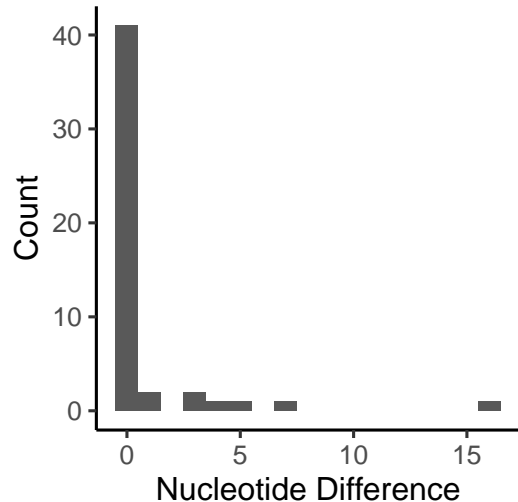
IGHV1-46*03

23 sequences assigned
17 (73.9%) exact matches, in which:
17 unique CDR3
4 unique J



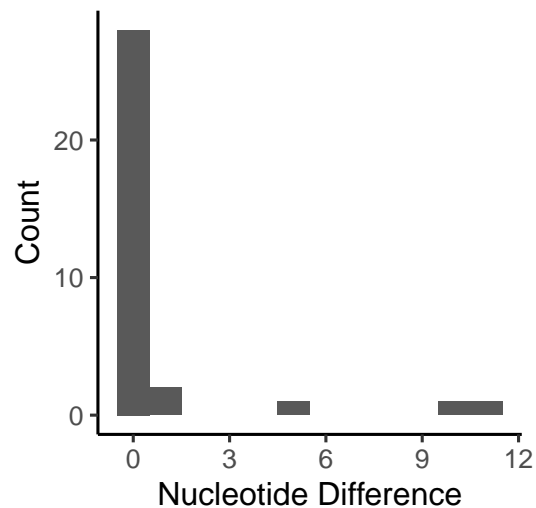
IGHV2-5*02

49 sequences assigned
41 (83.7%) exact matches, in which:
41 unique CDR3
4 unique J



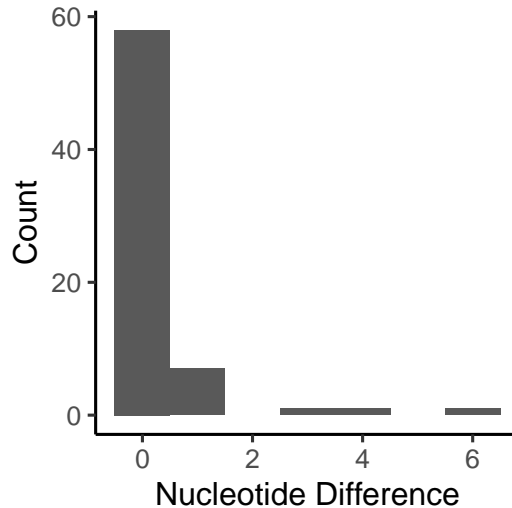
IGHV2-26*01

33 sequences assigned
28 (84.8%) exact matches, in which:
28 unique CDR3
5 unique J



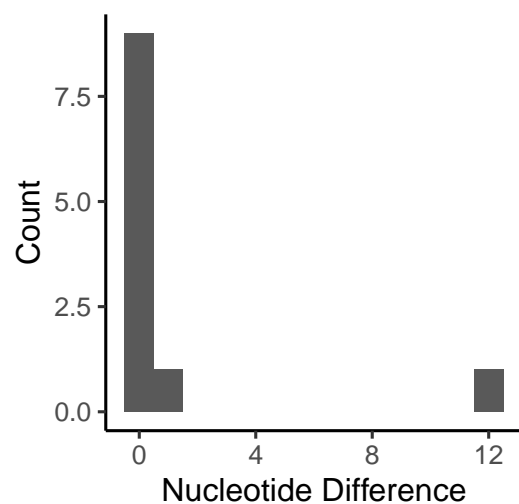
IGHV3-7*03

68 sequences assigned
58 (85.3%) exact matches, in which:
58 unique CDR3
6 unique J



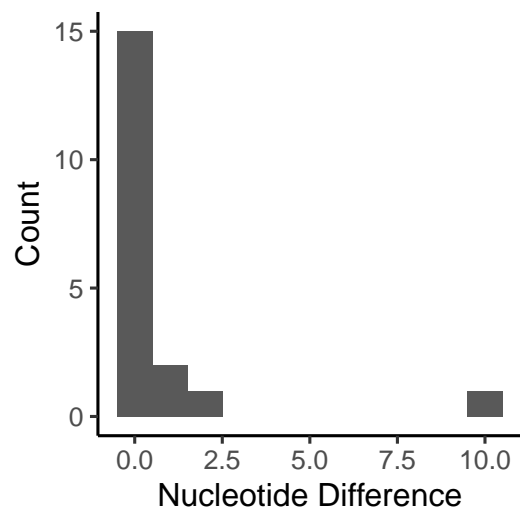
IGHV3-13*05

11 sequences assigned
9 (81.8%) exact matches, in which:
9 unique CDR3
4 unique J



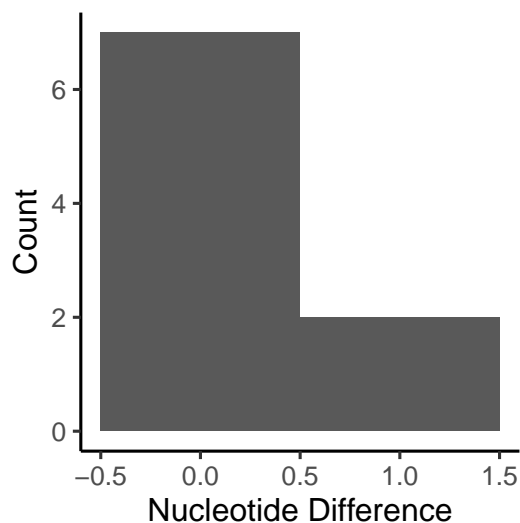
IGHV2-70*15

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



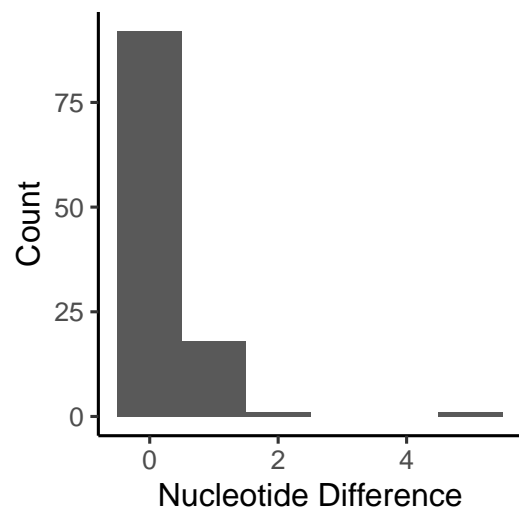
IGHV3-7*03_A131C

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



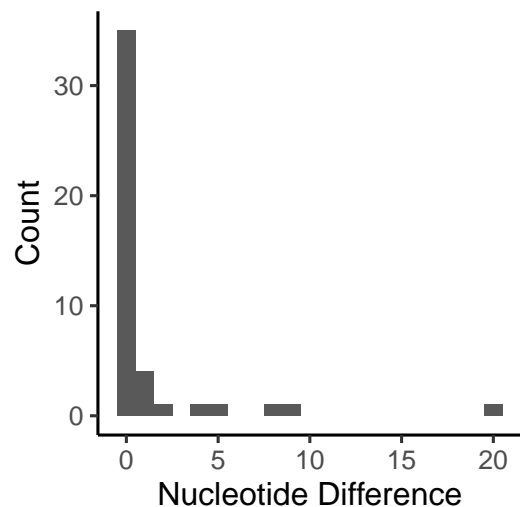
IGHV3-15*01

112 sequences assigned
92 (82.1%) exact matches, in which:
92 unique CDR3
6 unique J



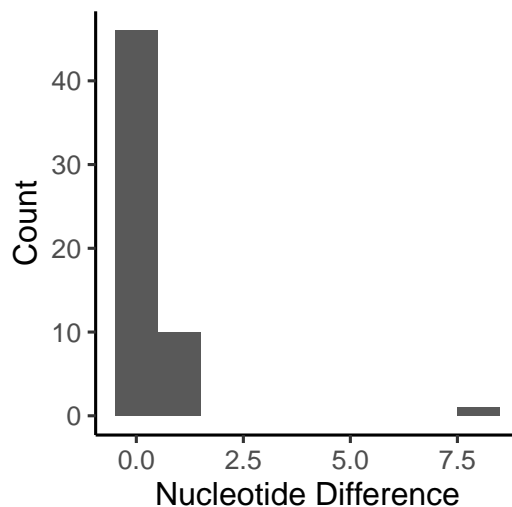
IGHV3-7*01

46 sequences assigned
35 (76.1%) exact matches, in which:
35 unique CDR3
5 unique J



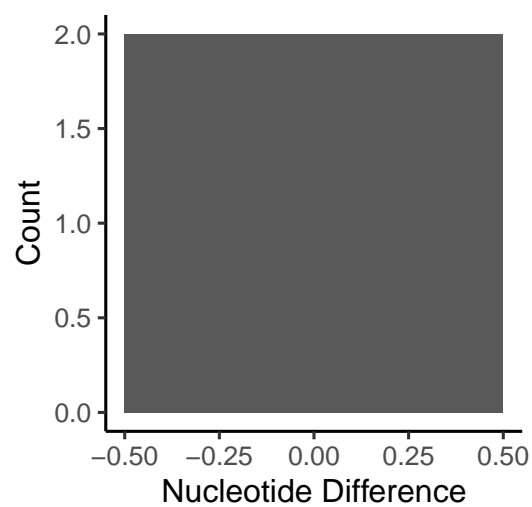
IGHV3-11*06

57 sequences assigned
46 (80.7%) exact matches, in which:
46 unique CDR3
5 unique J



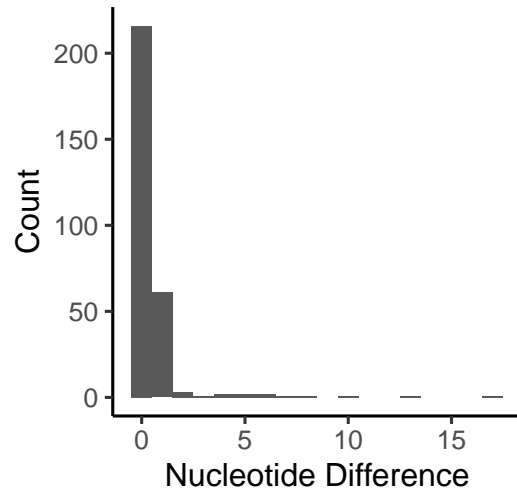
IGHV3-20*01

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



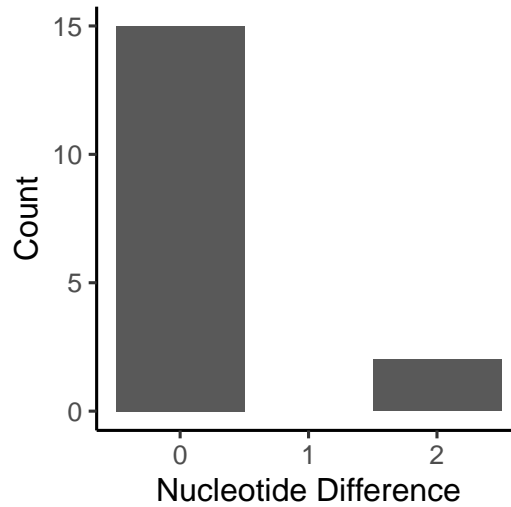
IGHV3-21*01

293 sequences assigned
216 (73.7%) exact matches, in which:
216 unique CDR3
6 unique J



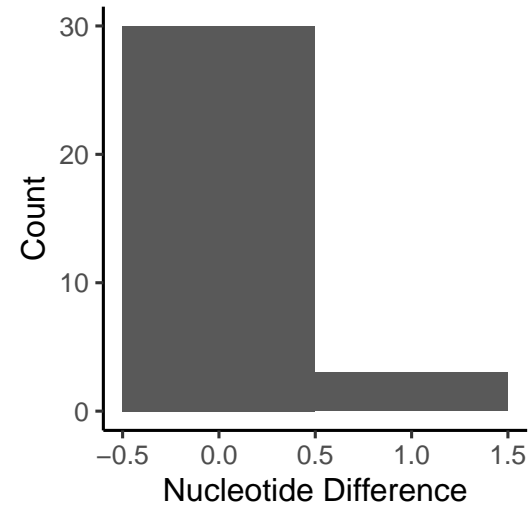
IGHV3-43*01

17 sequences assigned
15 (88.2%) exact matches, in which:
15 unique CDR3
2 unique J



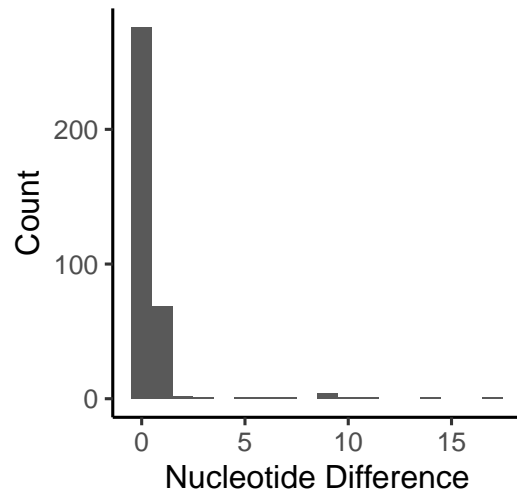
IGHV3-53*02

33 sequences assigned
30 (90.9%) exact matches, in which:
30 unique CDR3
4 unique J



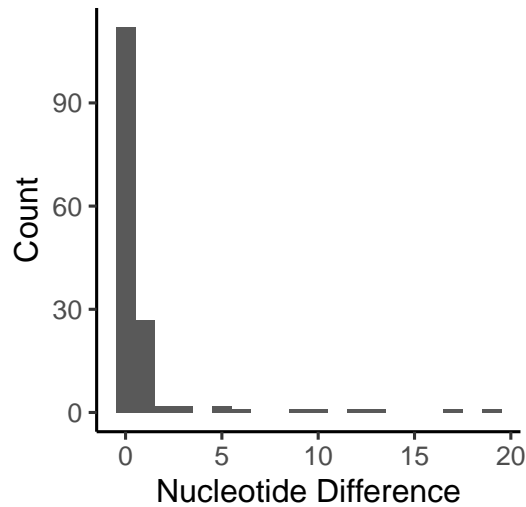
IGHV3-30-3*01

361 sequences assigned
276 (76.5%) exact matches, in which:
276 unique CDR3
6 unique J



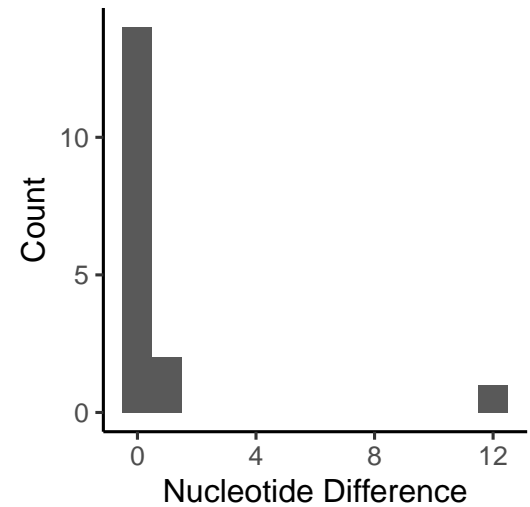
IGHV3-48*01

152 sequences assigned
112 (73.7%) exact matches, in which:
112 unique CDR3
6 unique J



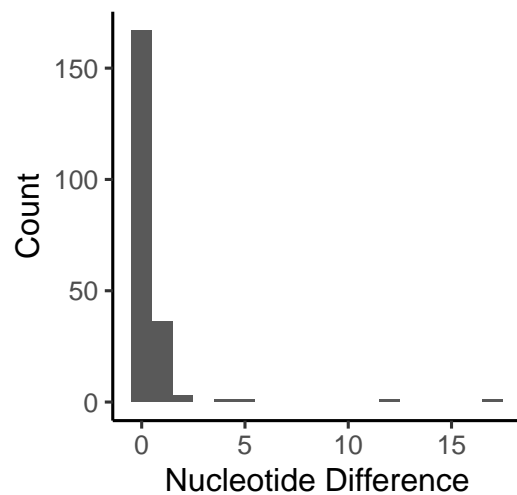
IGHV3-64*01

17 sequences assigned
14 (82.4%) exact matches, in which:
14 unique CDR3
3 unique J



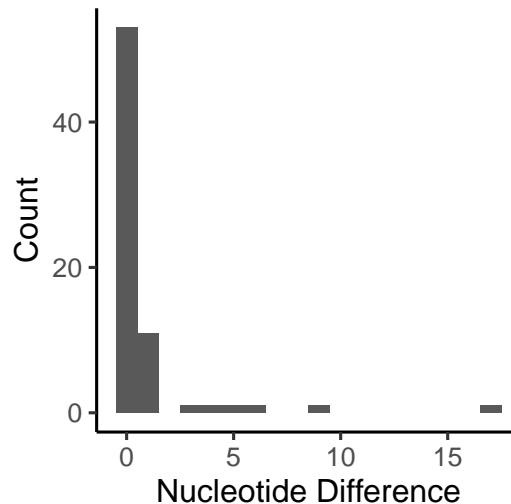
IGHV3-33*01

210 sequences assigned
167 (79.5%) exact matches, in which:
167 unique CDR3
6 unique J



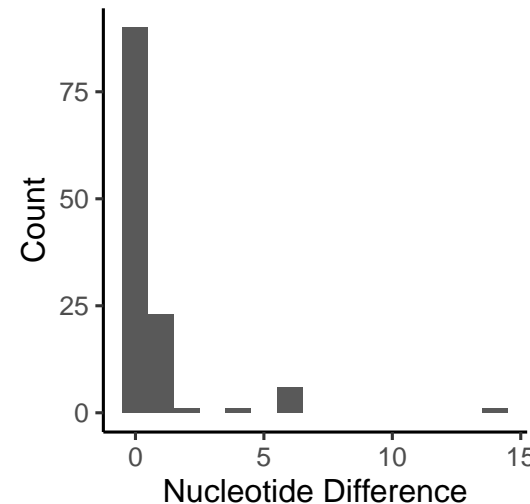
IGHV3-49*03

71 sequences assigned
53 (74.6%) exact matches, in which:
53 unique CDR3
5 unique J



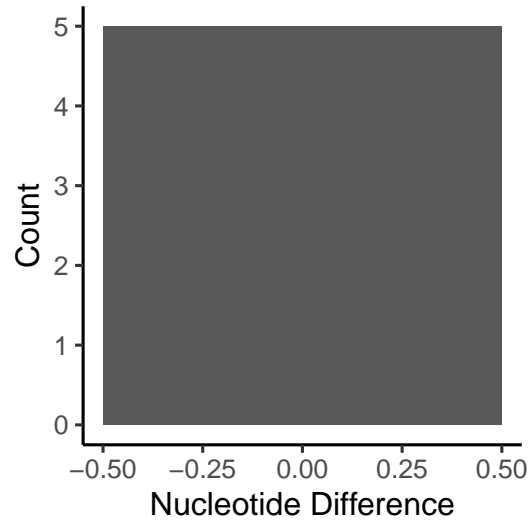
IGHV3-66*01

123 sequences assigned
90 (73.2%) exact matches, in which:
90 unique CDR3
6 unique J



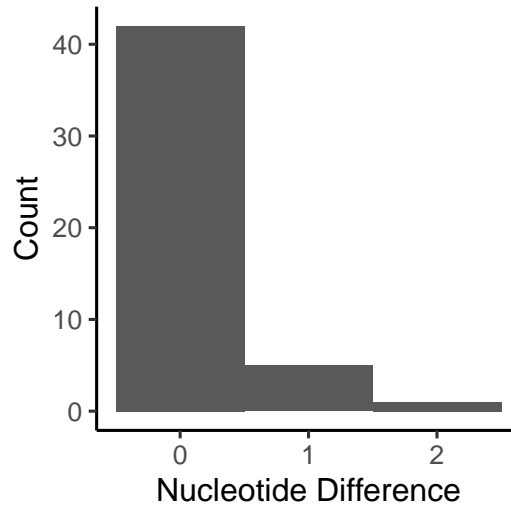
IGHV3-72*01

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



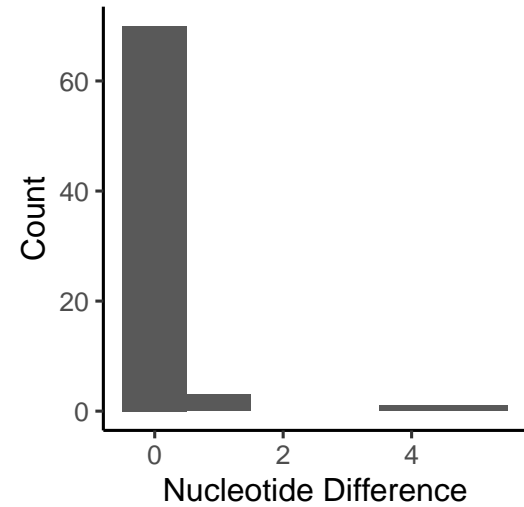
IGHV3-64D*06

48 sequences assigned
42 (87.5%) exact matches, in which:
42 unique CDR3
5 unique J



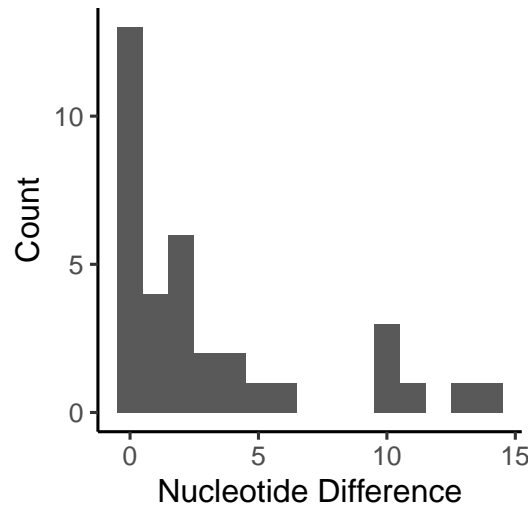
IGHV4-30-4*01

75 sequences assigned
70 (93.3%) exact matches, in which:
70 unique CDR3
4 unique J



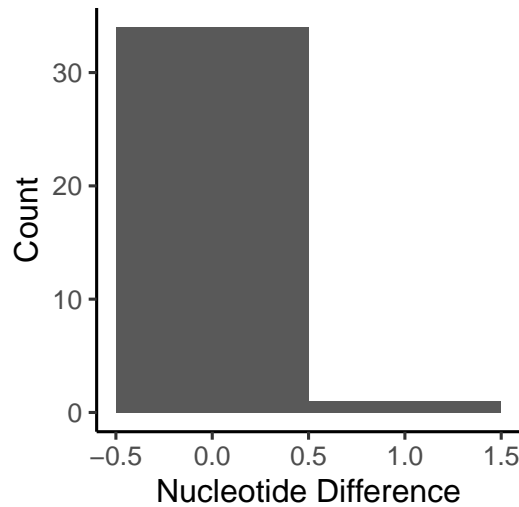
IGHV3-73*01

35 sequences assigned
13 (37.1%) exact matches, in which:
13 unique CDR3
3 unique J



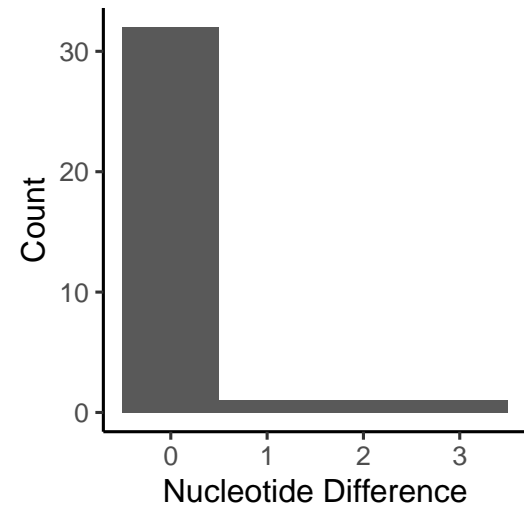
IGHV4-4*02

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



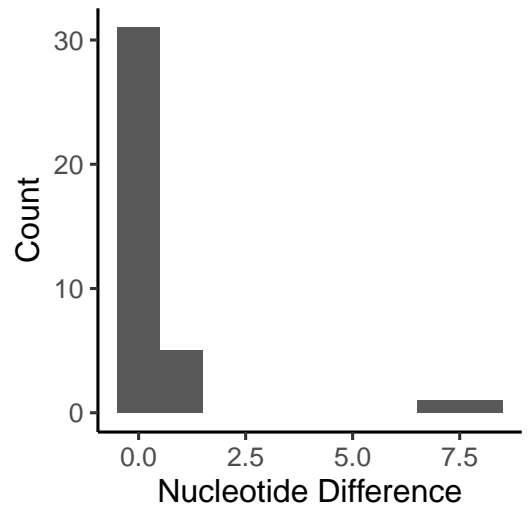
IGHV4-30-2*01

35 sequences assigned
32 (91.4%) exact matches, in which:
32 unique CDR3
6 unique J



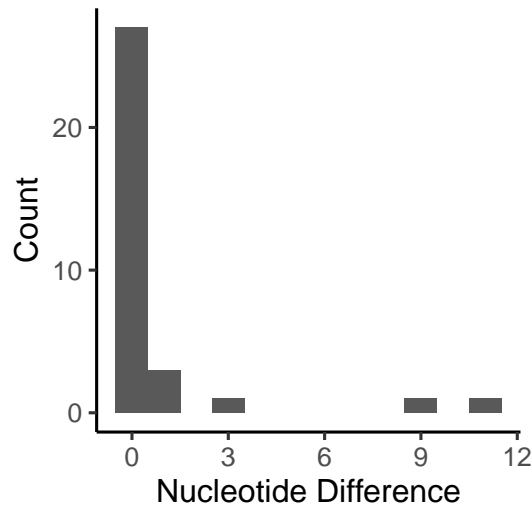
IGHV3-74*01

38 sequences assigned
31 (81.6%) exact matches, in which:
31 unique CDR3
5 unique J



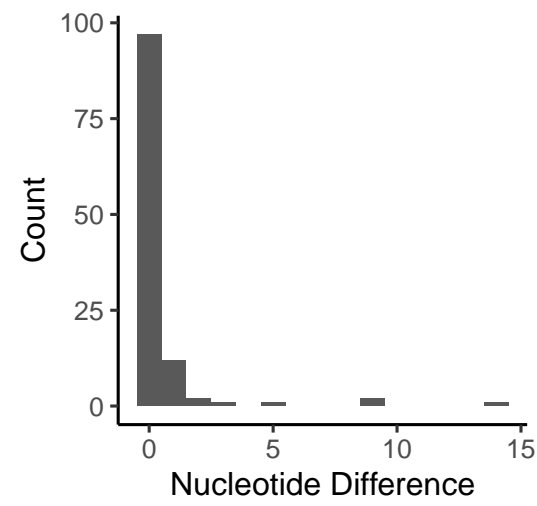
IGHV4-4*07

33 sequences assigned
27 (81.8%) exact matches, in which:
27 unique CDR3
4 unique J



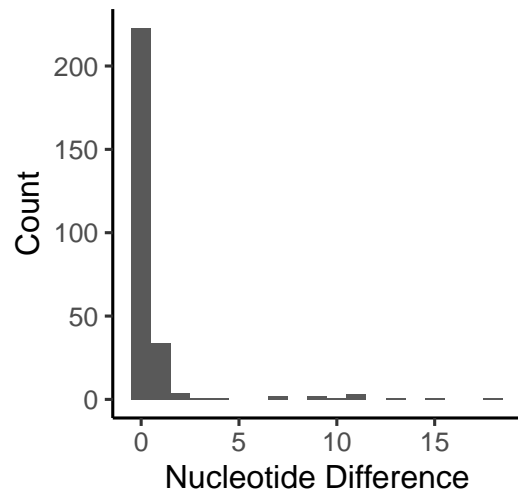
IGHV4-31*03

116 sequences assigned
97 (83.6%) exact matches, in which:
97 unique CDR3
5 unique J



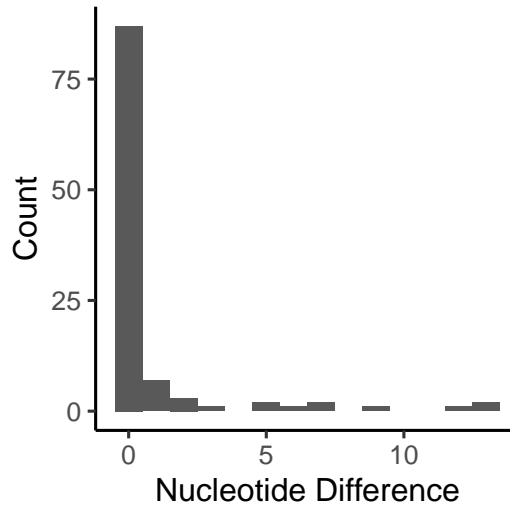
IGHV4–34*01

274 sequences assigned
223 (81.4%) exact matches, in which:
222 unique CDR3
6 unique J



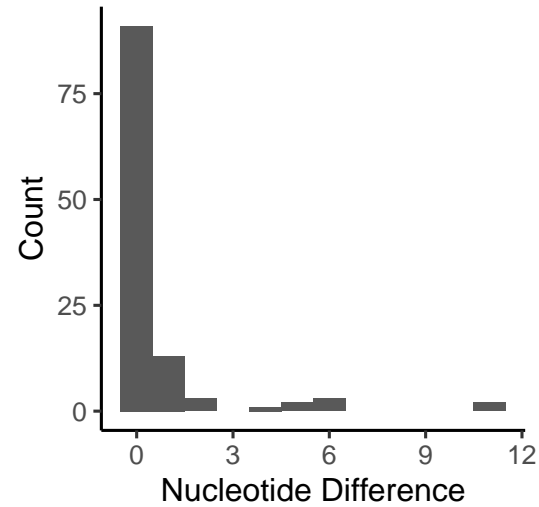
IGHV4–59*08

108 sequences assigned
87 (80.6%) exact matches, in which:
87 unique CDR3
6 unique J



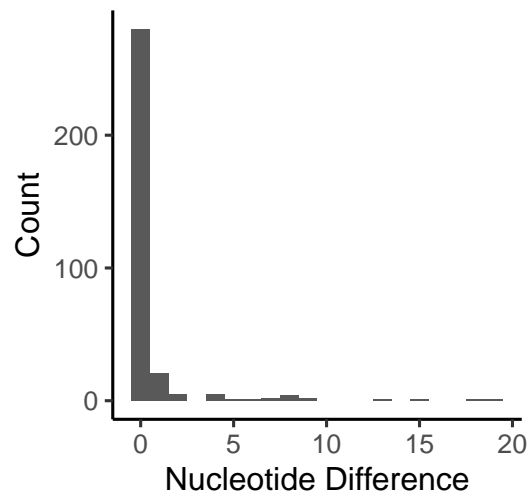
IGHV5–51*03

115 sequences assigned
91 (79.1%) exact matches, in which:
91 unique CDR3
6 unique J



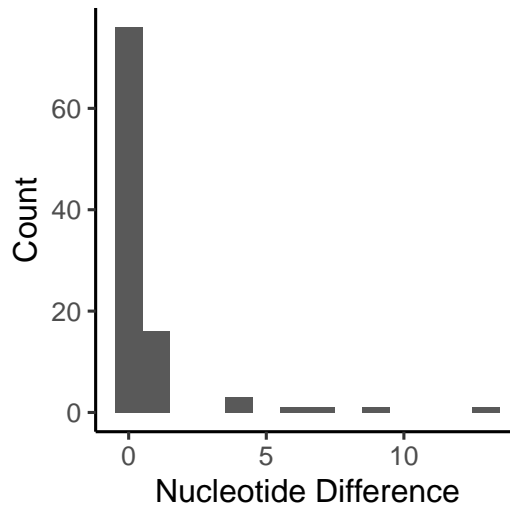
IGHV4–39*01

325 sequences assigned
280 (86.2%) exact matches, in which:
279 unique CDR3
6 unique J



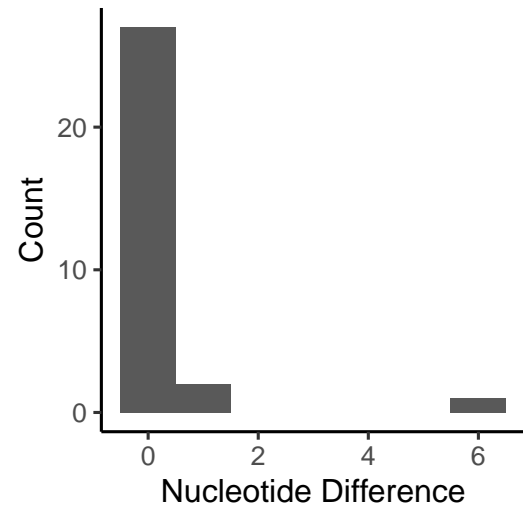
IGHV5–10–1*03

99 sequences assigned
76 (76.8%) exact matches, in which:
76 unique CDR3
6 unique J



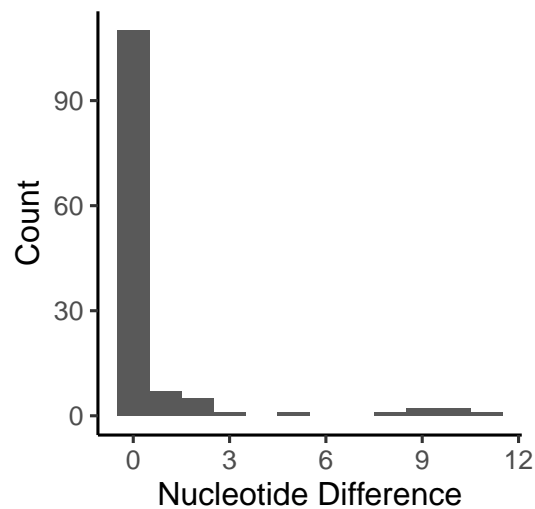
IGHV6–1*01

30 sequences assigned
27 (90%) exact matches, in which:
27 unique CDR3
5 unique J



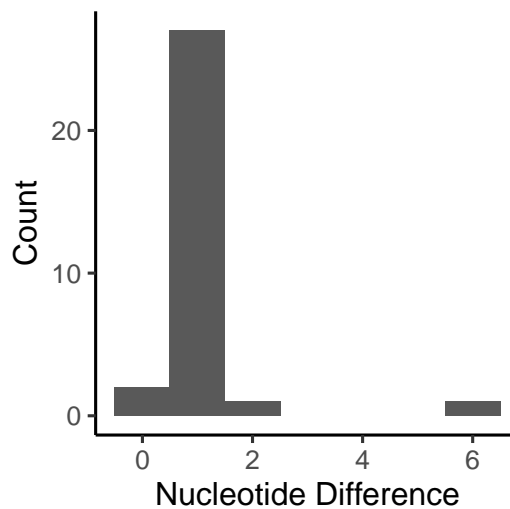
IGHV4–59*01

131 sequences assigned
110 (84%) exact matches, in which:
110 unique CDR3
6 unique J



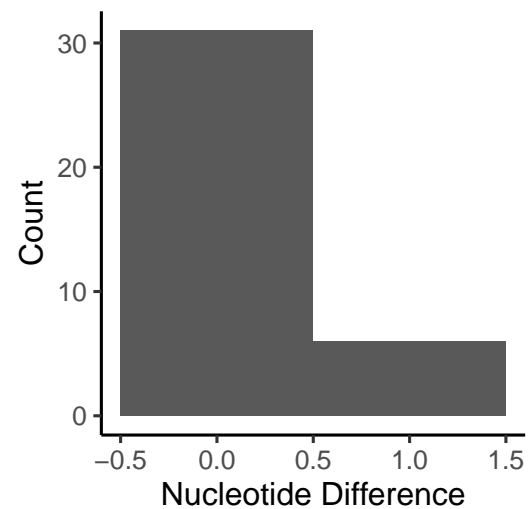
IGHV5–51*01

31 sequences assigned
2 (6.5%) exact matches, in which:
2 unique CDR3
2 unique J



IGHV7–4–1*02

37 sequences assigned
31 (83.8%) exact matches, in which:
31 unique CDR3
5 unique J



IGHV3-7*03_A131C

