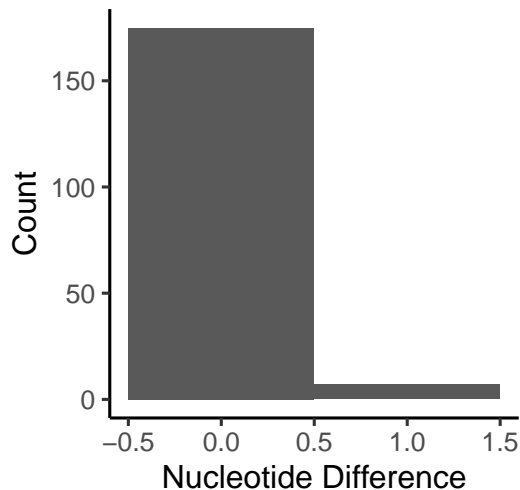


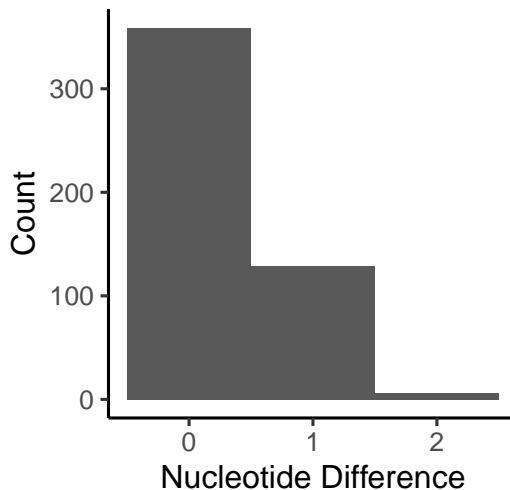
IGHV1–2*04

182 sequences assigned
175 (96.2%) exact matches, in which:
175 unique CDR3
6 unique J



IGHV1–18*01

494 sequences assigned
359 (72.7%) exact matches, in which:
318 unique CDR3
6 unique J



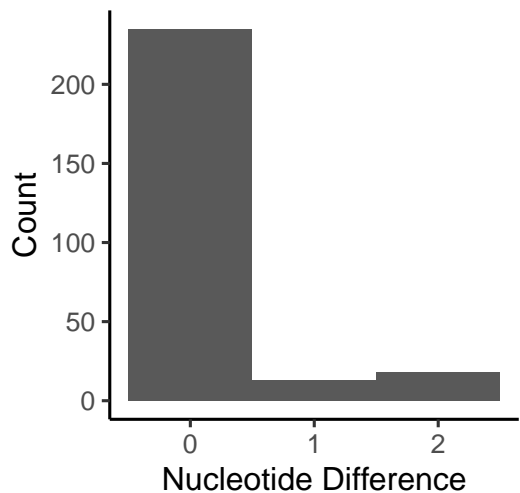
IGHV1–45*02

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



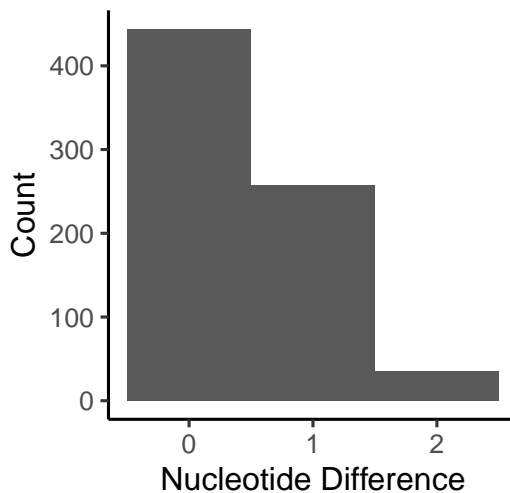
IGHV1–3*01_05

266 sequences assigned
235 (88.3%) exact matches, in which:
232 unique CDR3
6 unique J



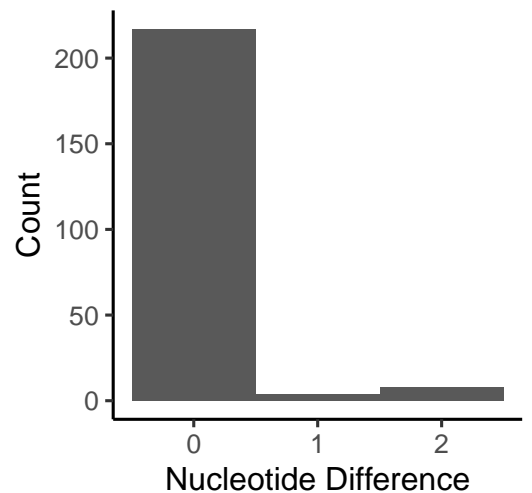
IGHV1–18*04

737 sequences assigned
444 (60.2%) exact matches, in which:
397 unique CDR3
6 unique J



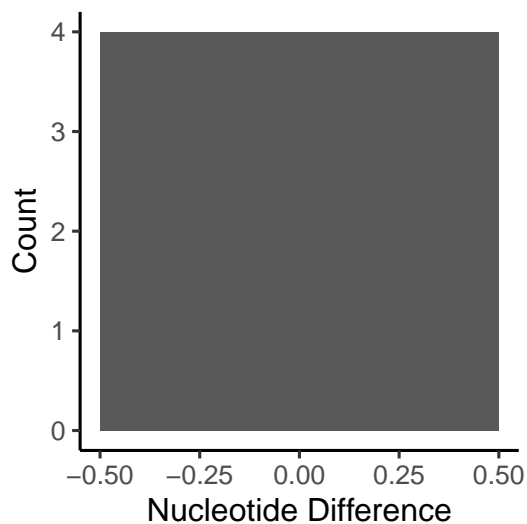
IGHV1–46*01

229 sequences assigned
217 (94.8%) exact matches, in which:
216 unique CDR3
6 unique J



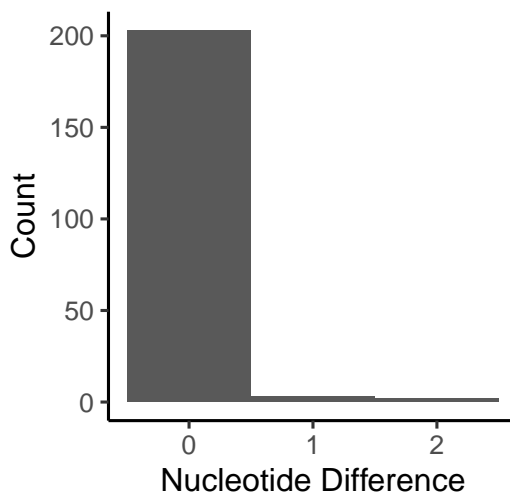
IGHV1–8*01

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



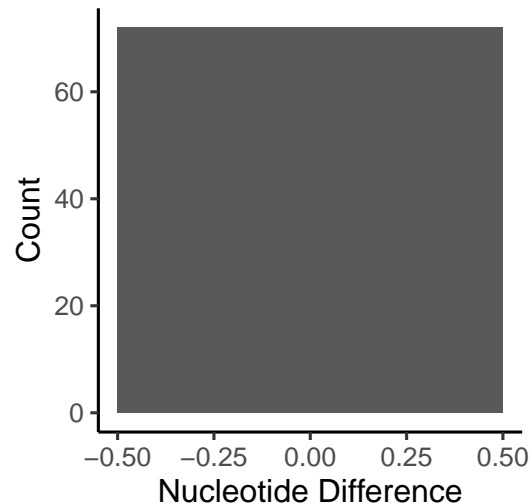
IGHV1–24*01

208 sequences assigned
203 (97.6%) exact matches, in which:
202 unique CDR3
6 unique J



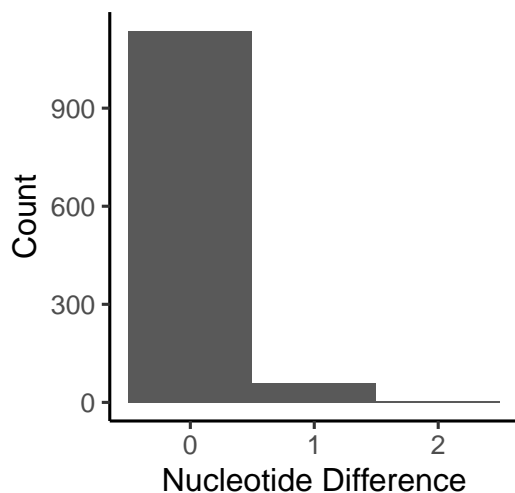
IGHV1–58*01_03

72 sequences assigned
72 (100%) exact matches, in which:
72 unique CDR3
6 unique J



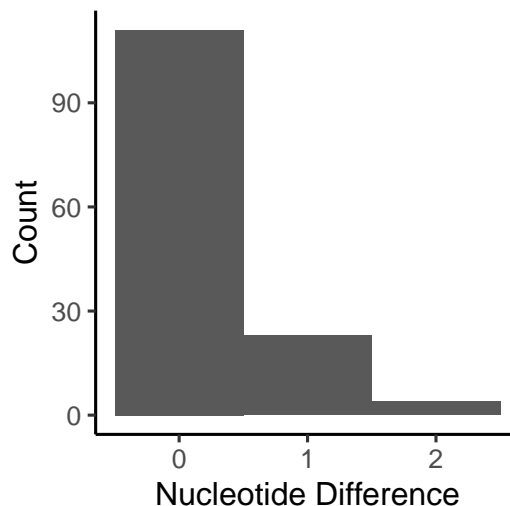
IGHV1–69*01_12_13

1200 sequences assigned
1137 (94.8%) exact matches, in which:
1129 unique CDR3
6 unique J



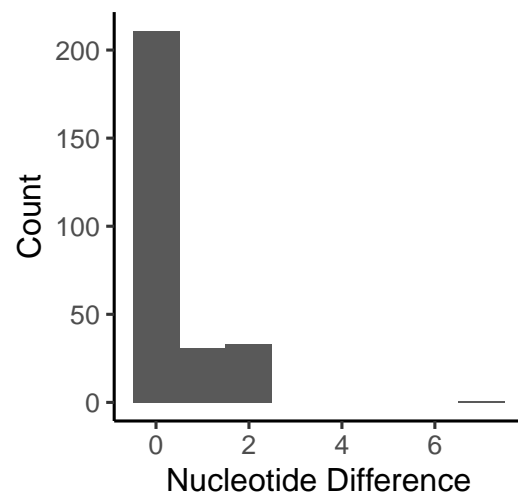
IGHV2–70*01

138 sequences assigned
111 (80.4%) exact matches, in which:
94 unique CDR3
5 unique J



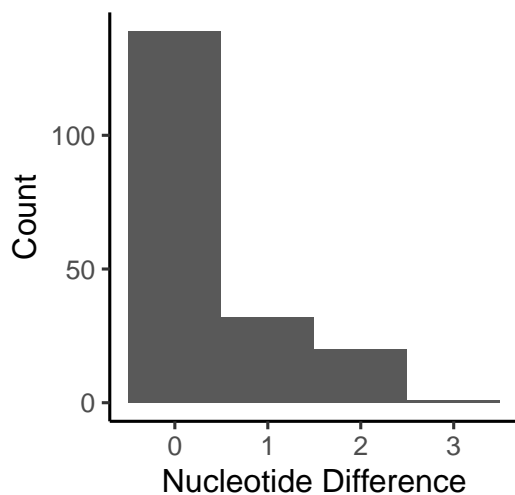
IGHV3–11*06

276 sequences assigned
211 (76.4%) exact matches, in which:
181 unique CDR3
6 unique J



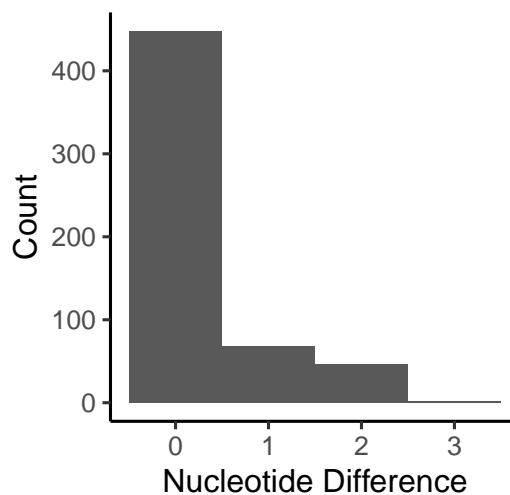
IGHV2–5*02

192 sequences assigned
139 (72.4%) exact matches, in which:
135 unique CDR3
5 unique J



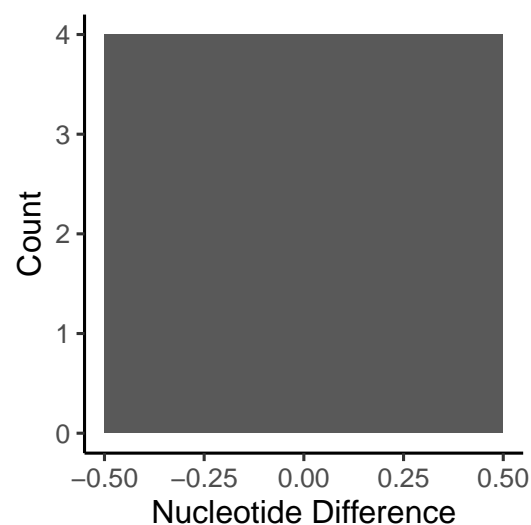
IGHV3–7*03

565 sequences assigned
448 (79.3%) exact matches, in which:
387 unique CDR3
6 unique J



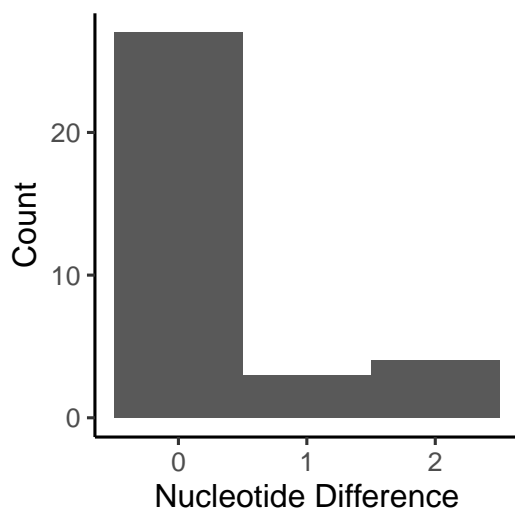
IGHV3–13*05

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



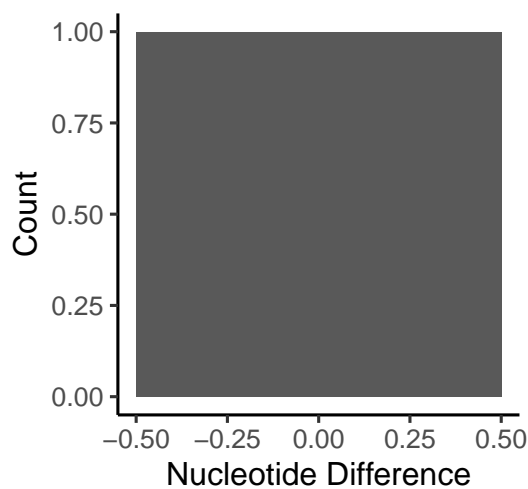
IGHV2–26*01

34 sequences assigned
27 (79.4%) exact matches, in which:
27 unique CDR3
4 unique J



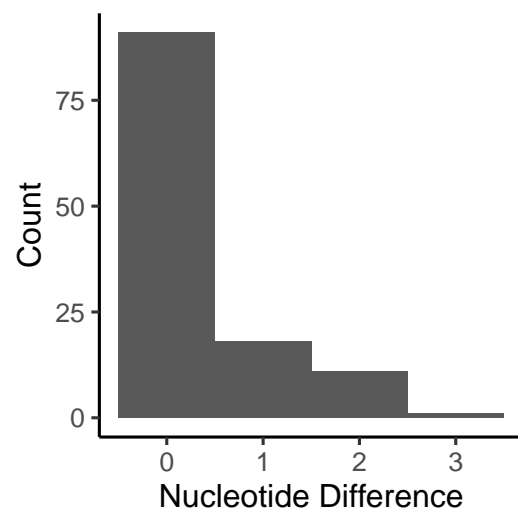
IGHV3–9*01

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



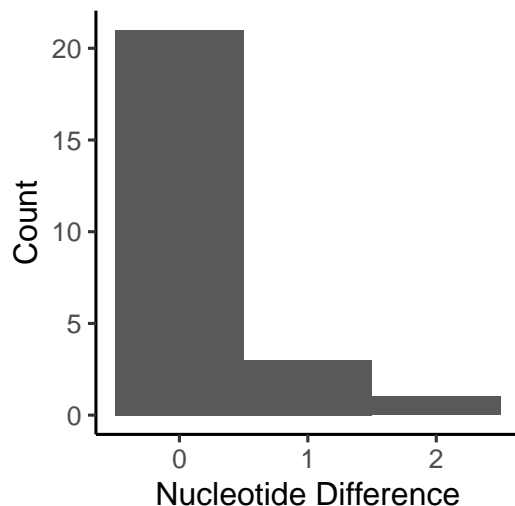
IGHV3–15*01_02

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



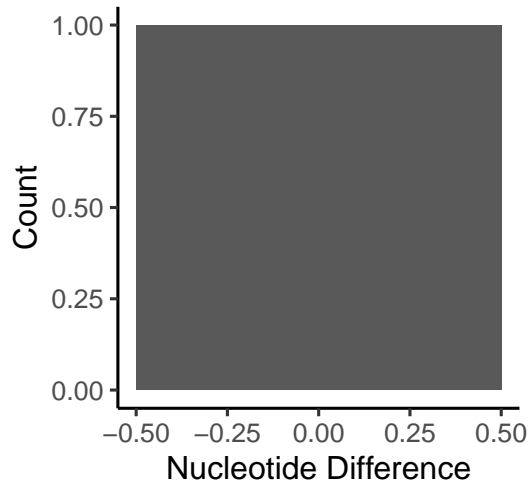
IGHV3-20*01_02

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



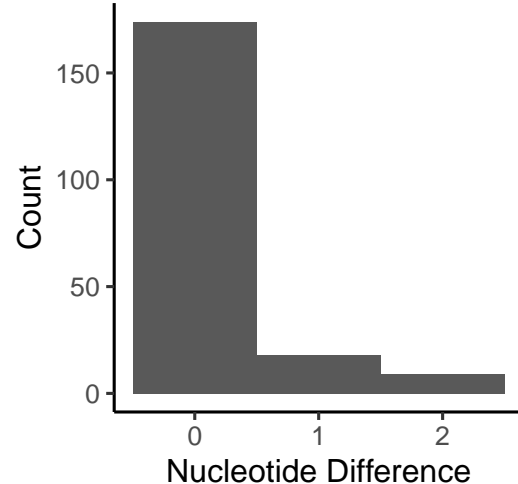
IGHV3-30-3*01

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



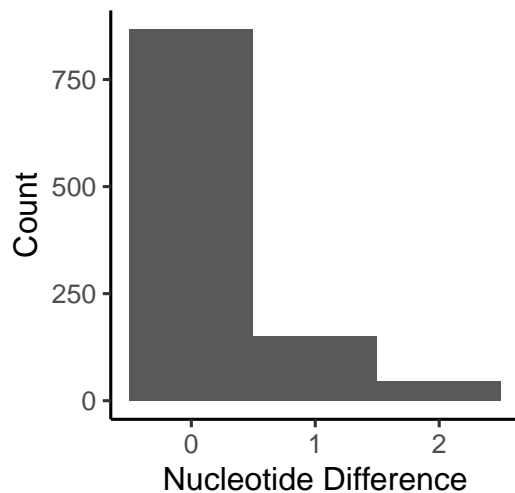
IGHV3-48*02

201 sequences assigned
174 (86.6%) exact matches, in which:
131 unique CDR3
6 unique J



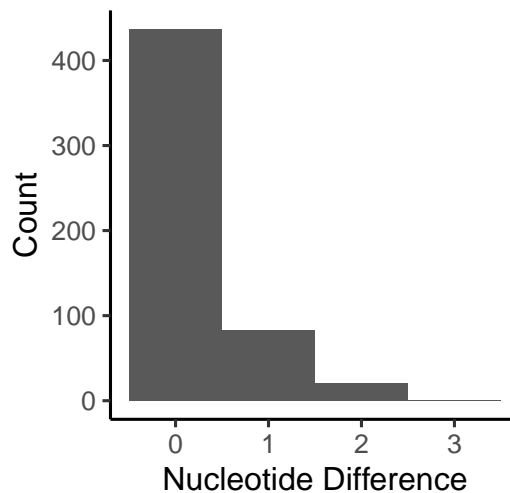
IGHV3-21*01_02

1065 sequences assigned
868 (81.5%) exact matches, in which:
685 unique CDR3
6 unique J



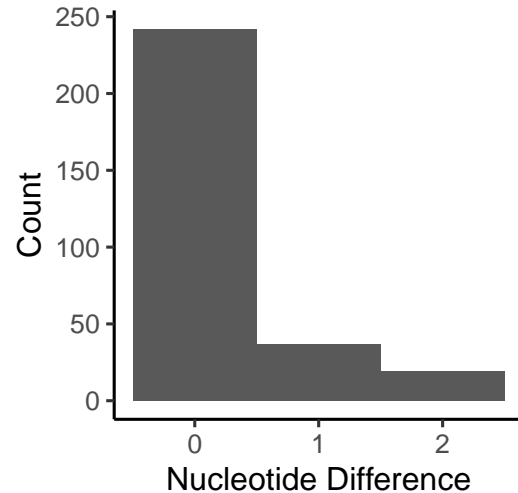
IGHV3-33*01

542 sequences assigned
437 (80.6%) exact matches, in which:
361 unique CDR3
6 unique J



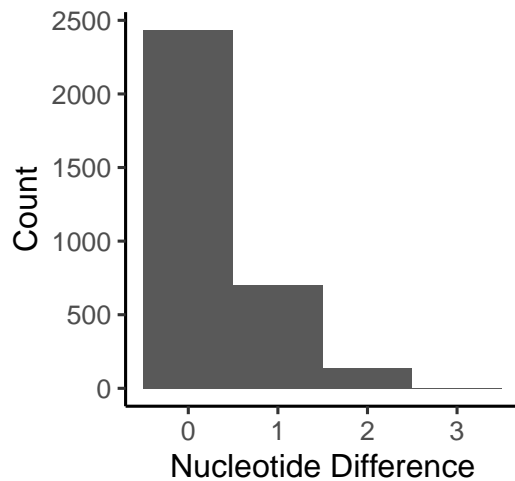
IGHV3-48*04

298 sequences assigned
242 (81.2%) exact matches, in which:
203 unique CDR3
5 unique J



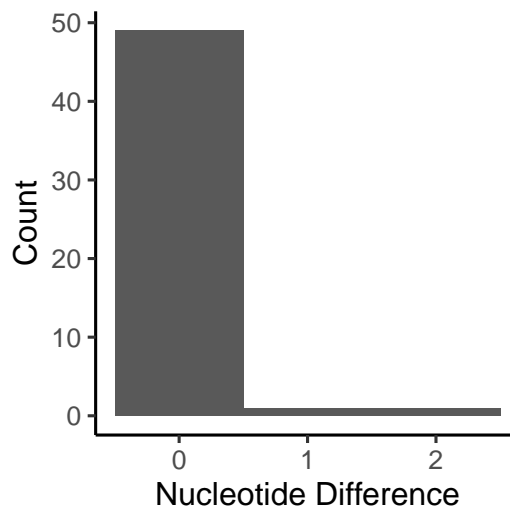
IGHV3-23*01_04

3276 sequences assigned
2434 (74.3%) exact matches, in which:
1975 unique CDR3
6 unique J



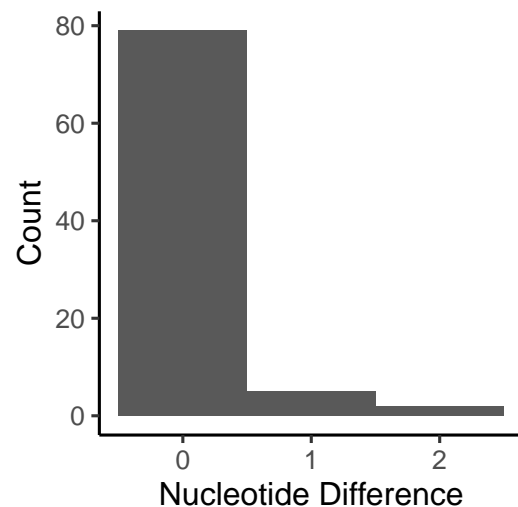
IGHV3-43*01

51 sequences assigned
49 (96.1%) exact matches, in which:
44 unique CDR3
5 unique J



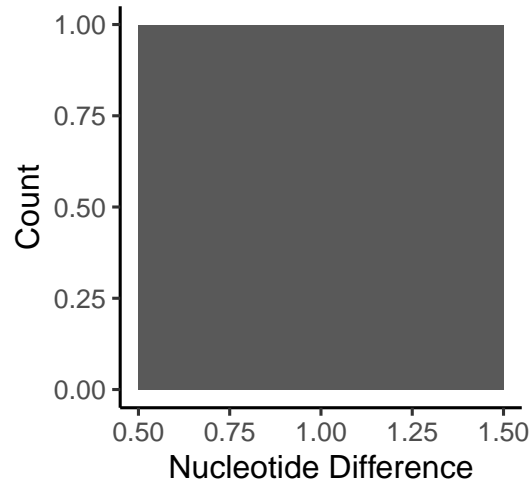
IGHV3-49*03_05

86 sequences assigned
79 (91.9%) exact matches, in which:
76 unique CDR3
5 unique J



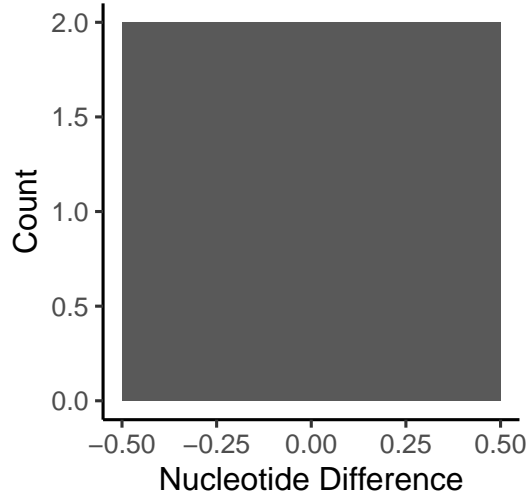
IGHV3-52*01_03

1 sequences assigned
No exact matches.



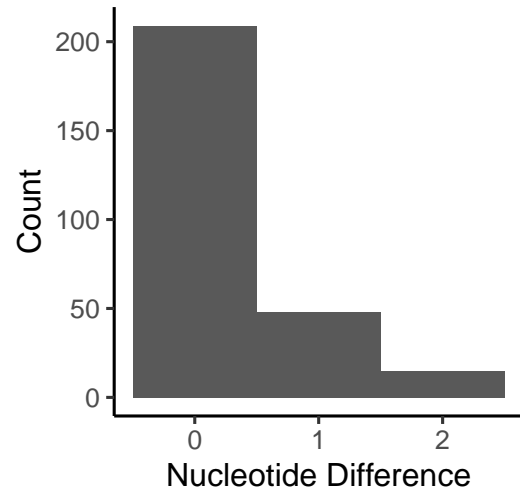
IGHV3-66*03

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



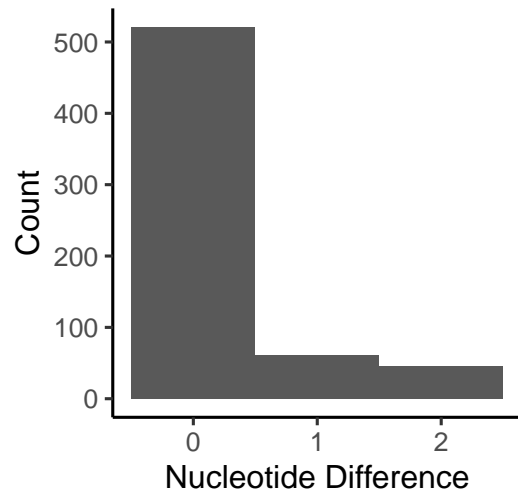
IGHV3-74*01_02

272 sequences assigned
209 (76.8%) exact matches, in which:
167 unique CDR3
6 unique J



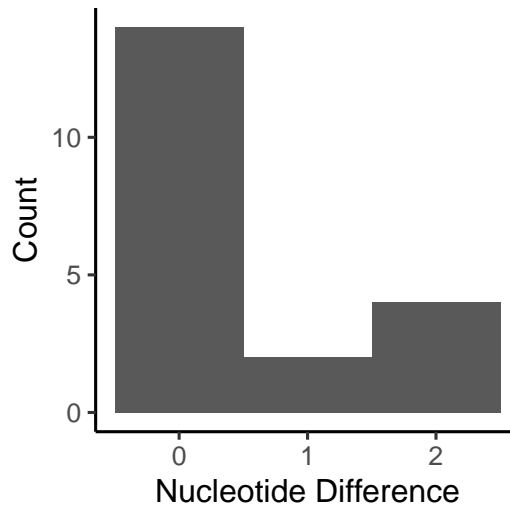
IGHV3-53*01_02

628 sequences assigned
521 (83%) exact matches, in which:
460 unique CDR3
6 unique J



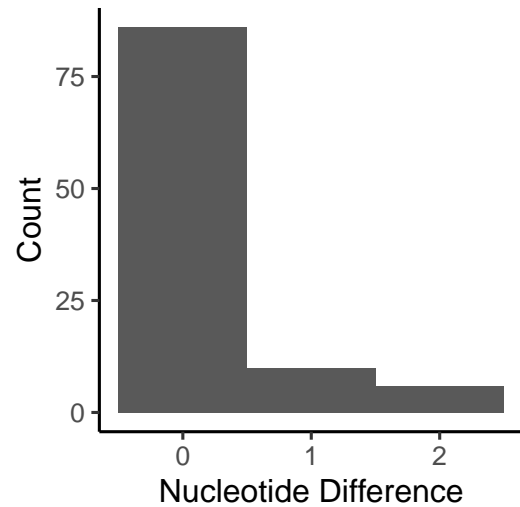
IGHV3-72*01

20 sequences assigned
14 (70%) exact matches, in which:
13 unique CDR3
4 unique J



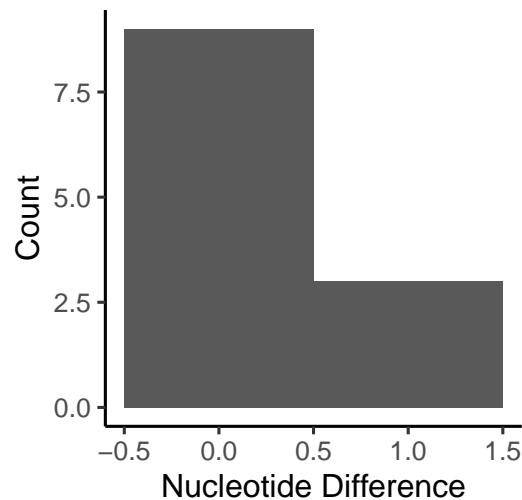
IGHV3-64D*06

102 sequences assigned
86 (84.3%) exact matches, in which:
73 unique CDR3
5 unique J



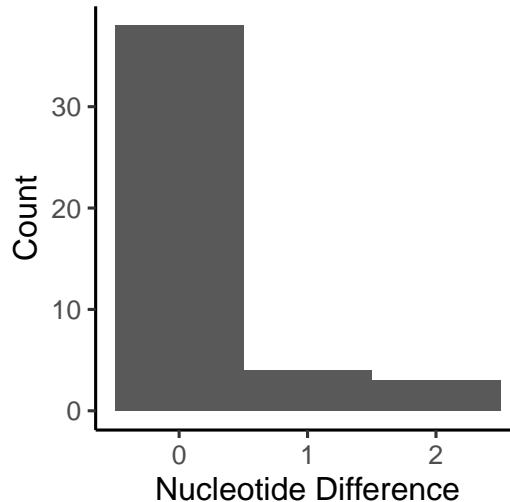
IGHV3-64*02_07

12 sequences assigned
9 (75%) exact matches, in which:
6 unique CDR3
2 unique J



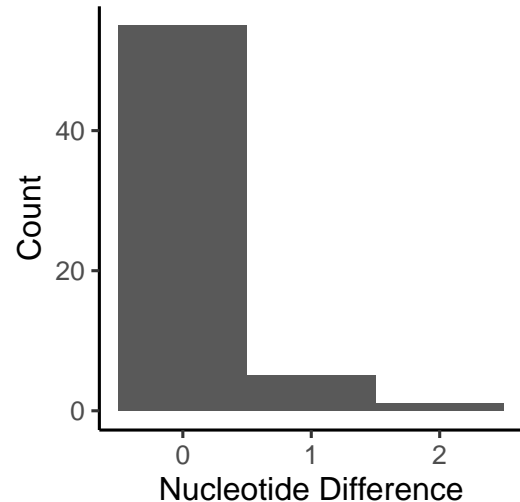
IGHV3-73*01_02

45 sequences assigned
38 (84.4%) exact matches, in which:
28 unique CDR3
3 unique J



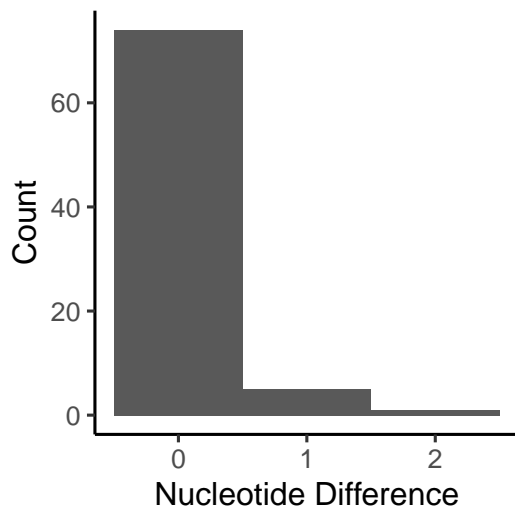
IGHV3-64D*08

61 sequences assigned
55 (90.2%) exact matches, in which:
51 unique CDR3
5 unique J



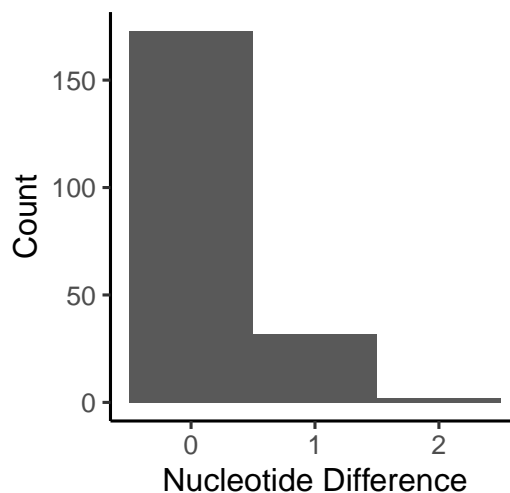
IGHV4-4*02_03

80 sequences assigned
74 (92.5%) exact matches, in which:
70 unique CDR3
5 unique J



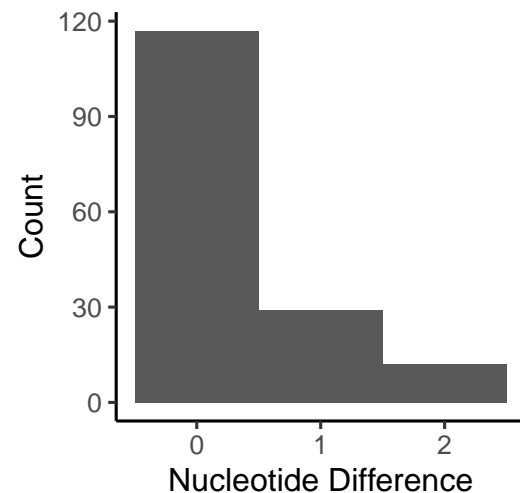
IGHV4-31*02

207 sequences assigned
173 (83.6%) exact matches, in which:
152 unique CDR3
6 unique J



IGHV4-39*02_C258G

158 sequences assigned
117 (74.1%) exact matches, in which:
107 unique CDR3
5 unique J



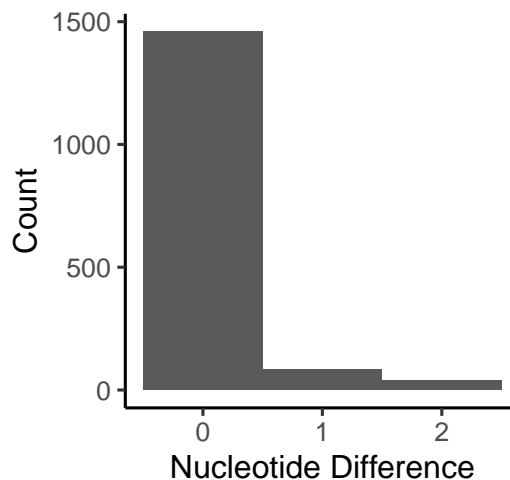
IGHV4-28*01_07

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



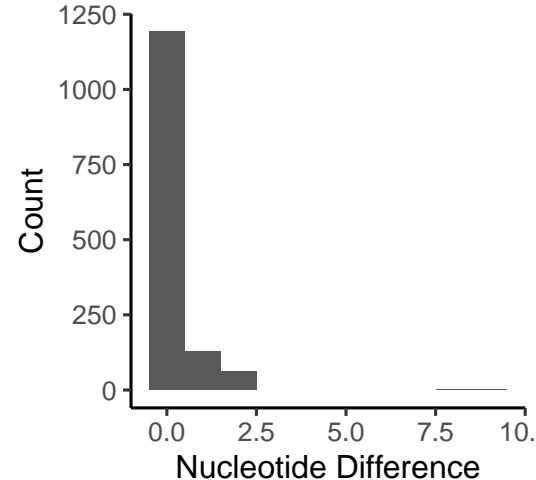
IGHV4-34*01_02

1583 sequences assigned
1459 (92.2%) exact matches, in which:
1405 unique CDR3
6 unique J



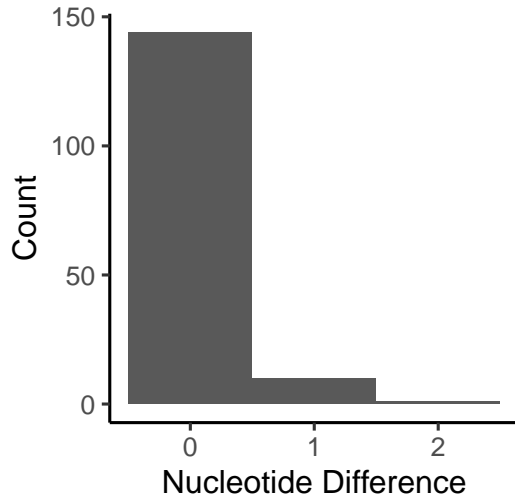
IGHV4-59*02_G88A

1384 sequences assigned
1192 (86.1%) exact matches, in which:
1127 unique CDR3
6 unique J



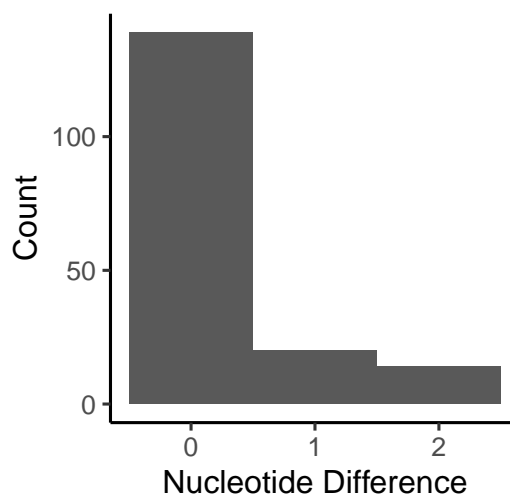
IGHV4-30-2*01

155 sequences assigned
144 (92.9%) exact matches, in which:
128 unique CDR3
5 unique J



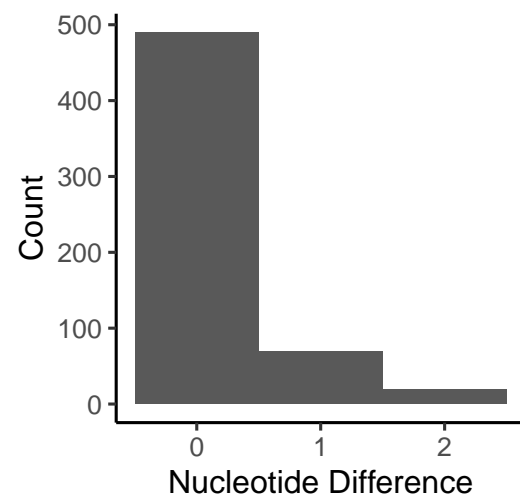
IGHV4-39*07

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



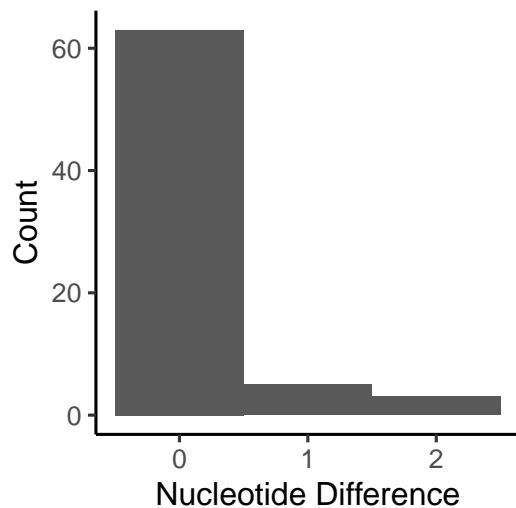
IGHV4-61*01

578 sequences assigned
490 (84.8%) exact matches, in which:
447 unique CDR3
6 unique J



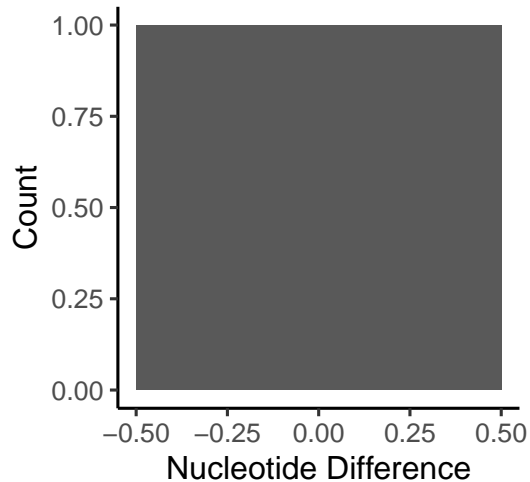
IGHV5-10-1*01_03

71 sequences assigned
63 (88.7%) exact matches, in which:
63 unique CDR3
6 unique J



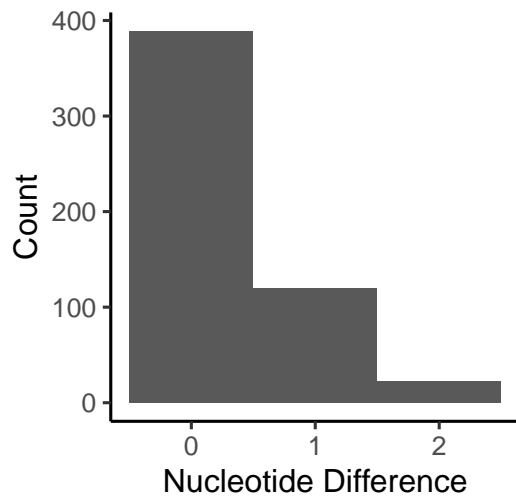
IGHV7-4-1*02

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



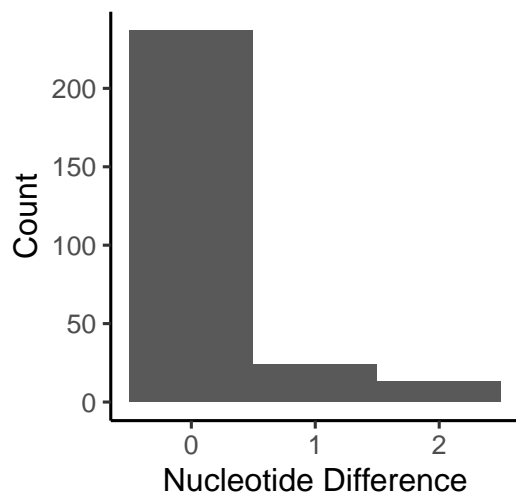
IGHV5-51*07_A128G

531 sequences assigned
389 (73.3%) exact matches, in which:
367 unique CDR3
6 unique J



IGHV6-1*01_02

274 sequences assigned
237 (86.5%) exact matches, in which:
234 unique CDR3
6 unique J





Warning – no inferred sequences found.