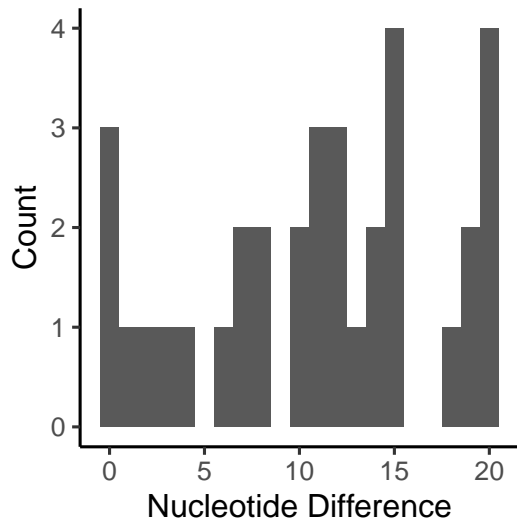


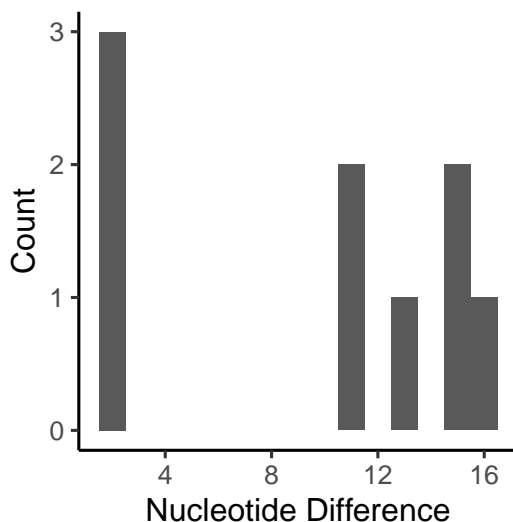
IGHV1–2*02

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



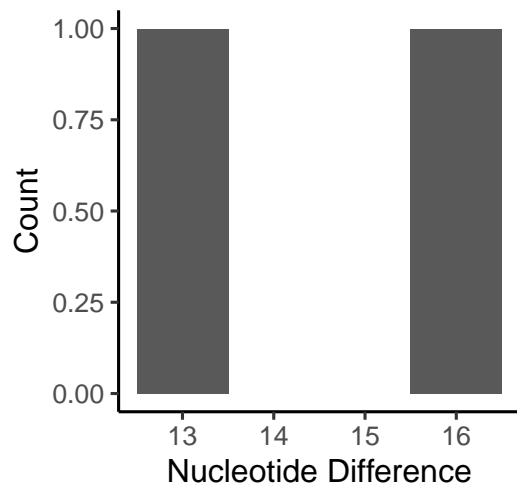
IGHV1–24*01

10 sequences assigned
No exact matches.



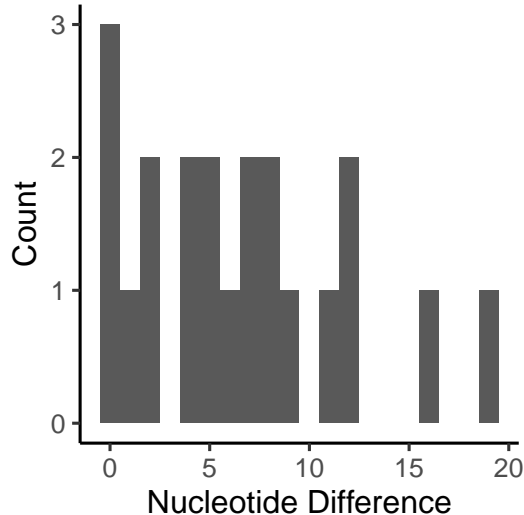
IGHV1–69–2*01

3 sequences assigned
No exact matches.



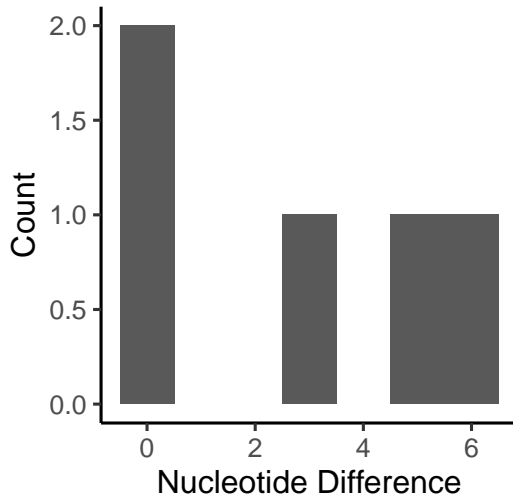
IGHV1–8*01

21 sequences assigned
3 (14.3%) exact matches, in which:
3 unique CDR3
3 unique J



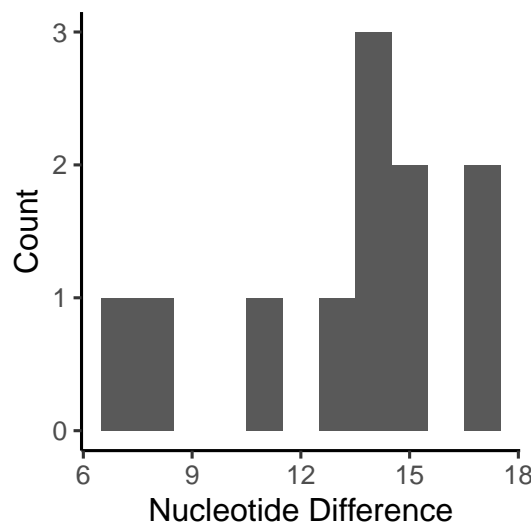
IGHV1–46*01

10 sequences assigned
2 (20%) exact matches, in which:
2 unique CDR3
2 unique J



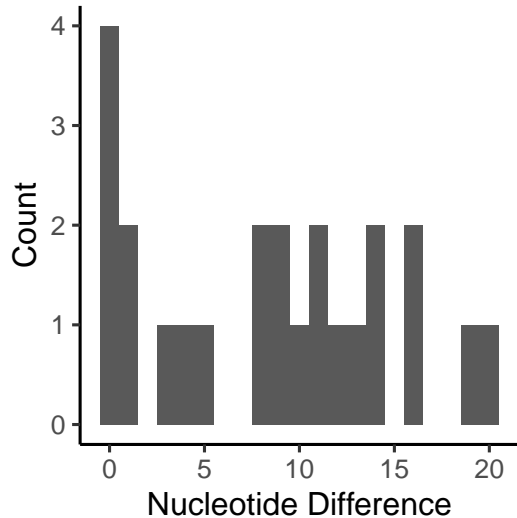
IGHV1–69*06

16 sequences assigned
No exact matches.



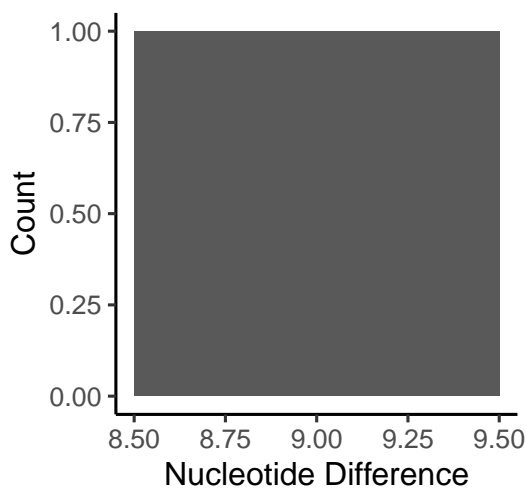
IGHV1–18*01

27 sequences assigned
4 (14.8%) exact matches, in which:
4 unique CDR3
2 unique J



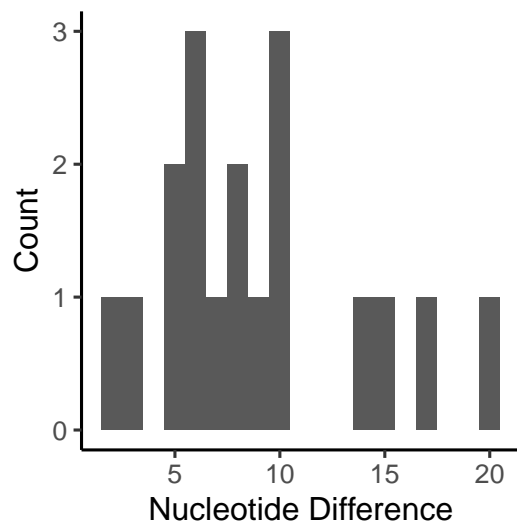
IGHV1–58*02

1 sequences assigned
No exact matches.



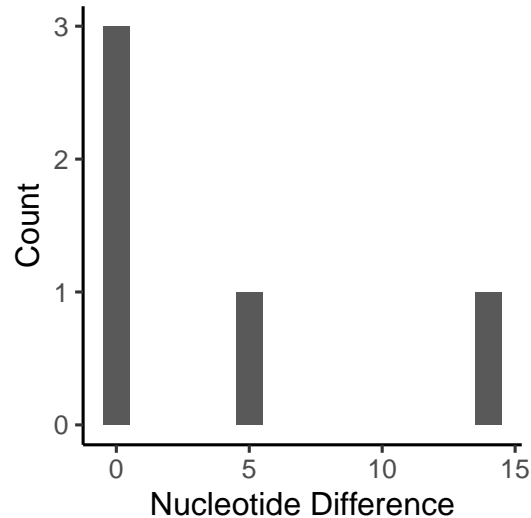
IGHV2–5*01

22 sequences assigned
No exact matches.



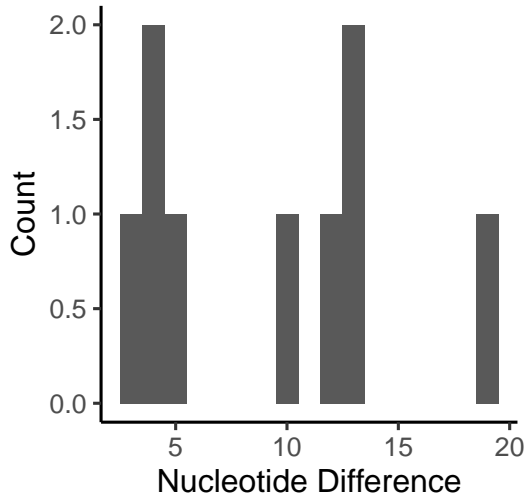
IGHV2–26*01

5 sequences assigned
3 (60%) exact matches, in which:
3 unique CDR3
2 unique J



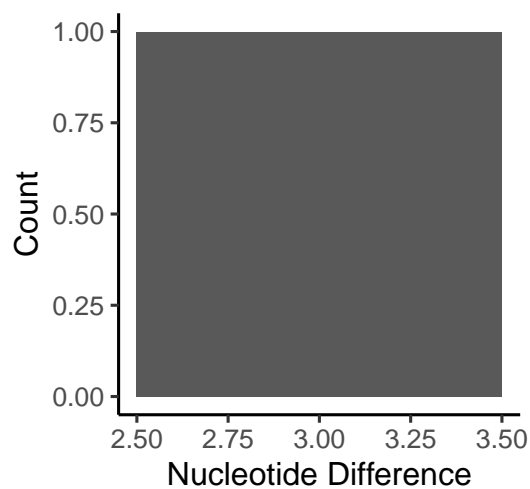
IGHV3–11*01

14 sequences assigned
No exact matches.



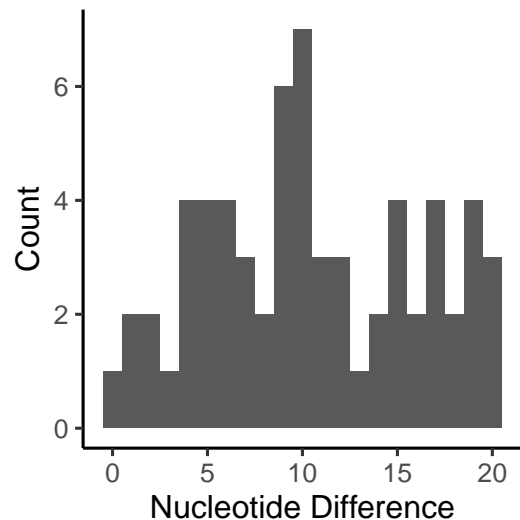
IGHV3–20*01

1 sequences assigned
No exact matches.



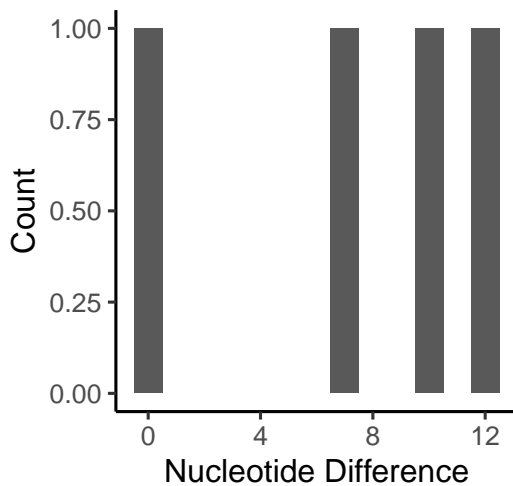
IGHV3–7*01

75 sequences assigned
1 (1.3%) exact matches, in which:
1 unique CDR3
1 unique J



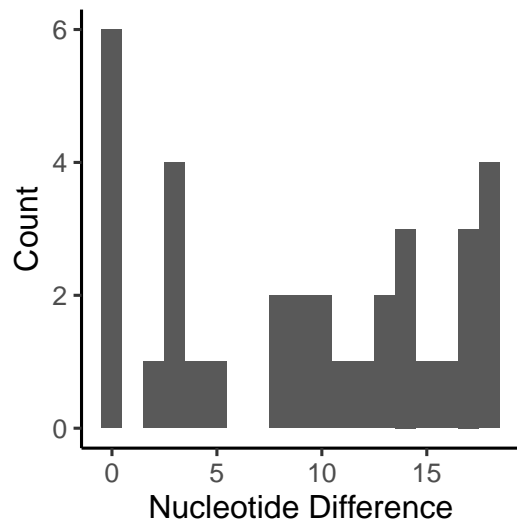
IGHV3–13*01

4 sequences assigned
1 (25%) exact matches, in which:
1 unique CDR3
1 unique J



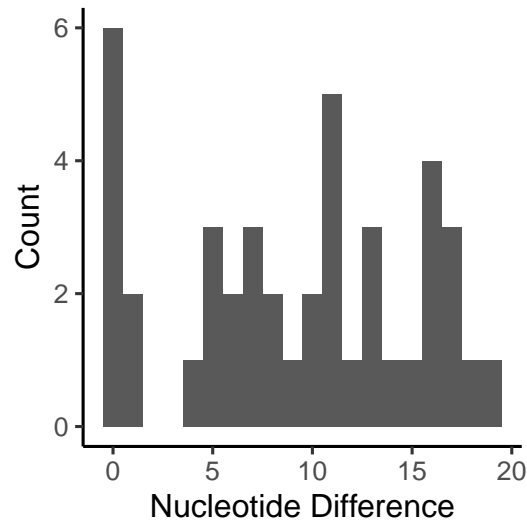
IGHV3–21*01

36 sequences assigned
6 (16.7%) exact matches, in which:
6 unique CDR3
3 unique J



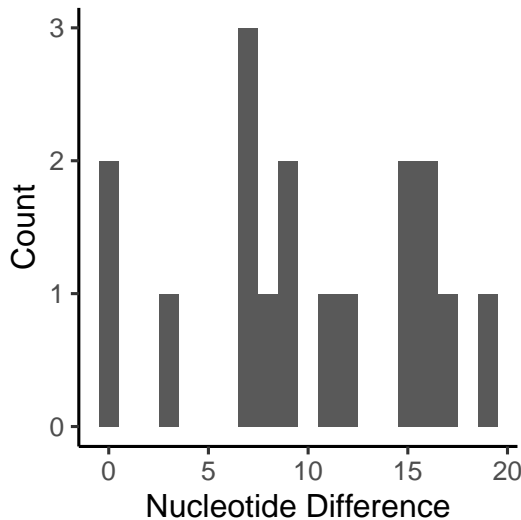
IGHV3–9*01

48 sequences assigned
6 (12.5%) exact matches, in which:
6 unique CDR3
3 unique J



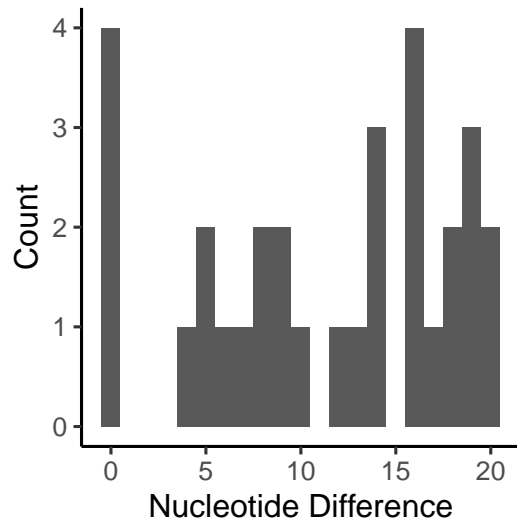
IGHV3–15*01

19 sequences assigned
2 (10.5%) exact matches, in which:
2 unique CDR3
1 unique J



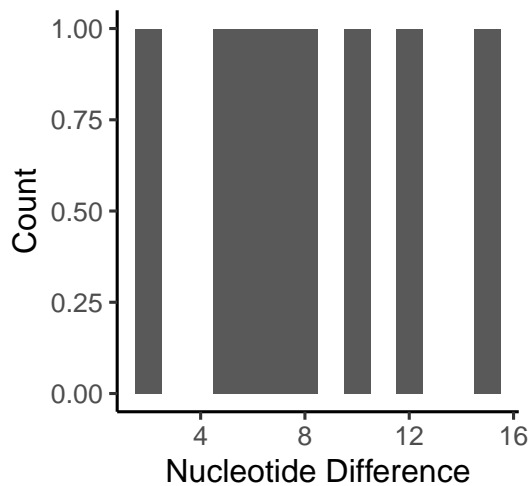
IGHV3–30–3*01

41 sequences assigned
4 (9.8%) exact matches, in which:
4 unique CDR3
3 unique J



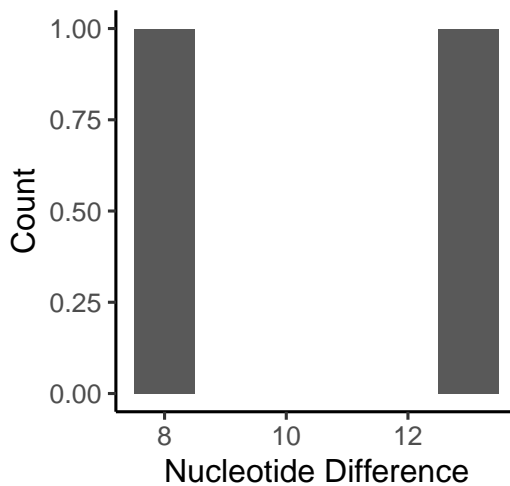
IGHV3-30*01

10 sequences assigned
No exact matches.



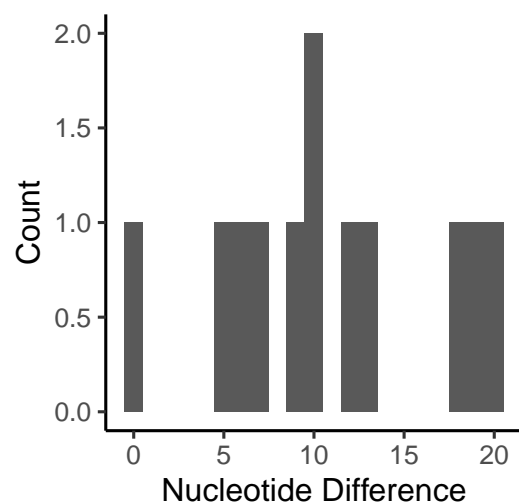
IGHV3-43*01

2 sequences assigned
No exact matches.



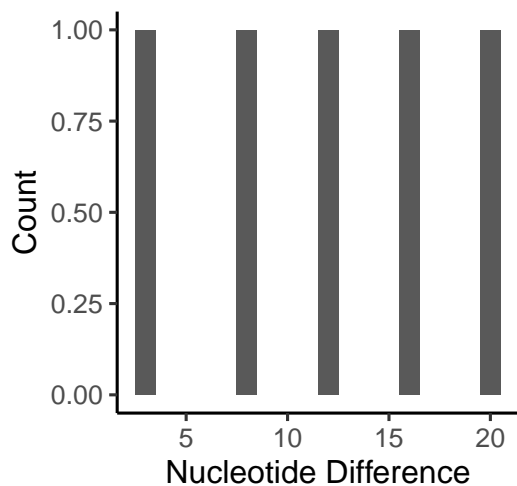
IGHV3-49*03

14 sequences assigned
1 (7.1%) exact matches, in which:
1 unique CDR3
1 unique J



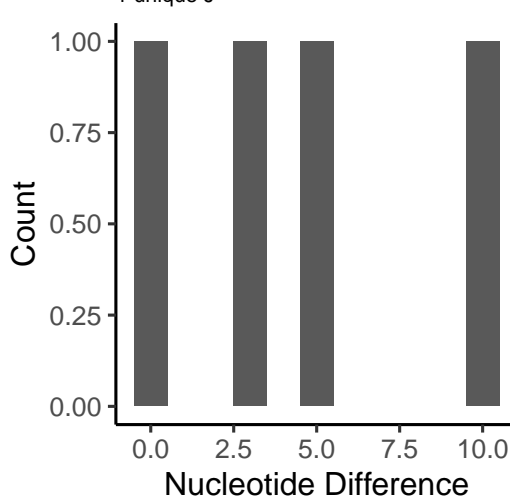
IGHV3-30*03

8 sequences assigned
No exact matches.



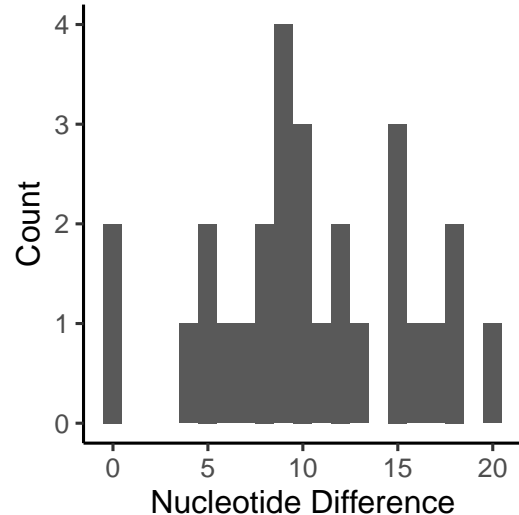
IGHV3-48*01

5 sequences assigned
1 (20%) exact matches, in which:
1 unique CDR3
1 unique J



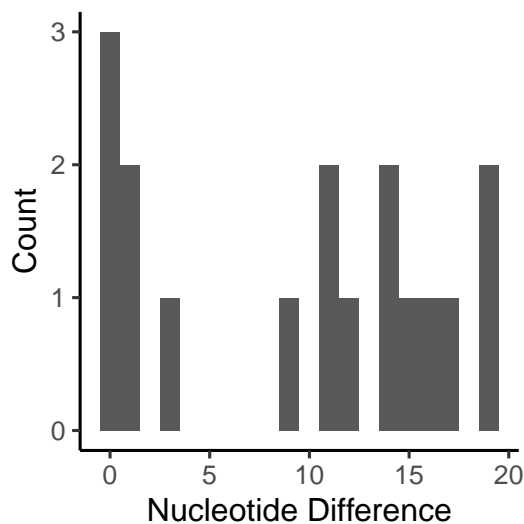
IGHV3-53*01

33 sequences assigned
2 (6.1%) exact matches, in which:
2 unique CDR3
1 unique J



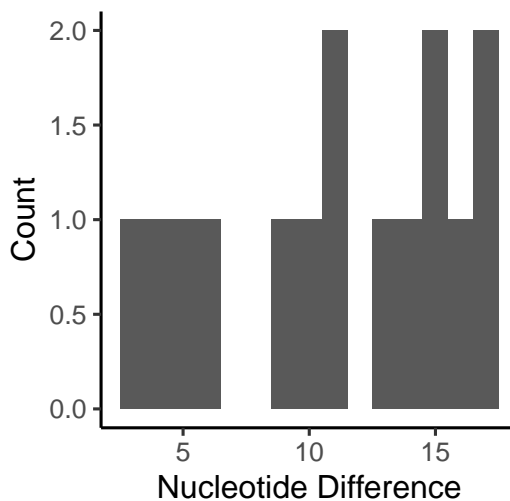
IGHV3-33*01

21 sequences assigned
3 (14.3%) exact matches, in which:
3 unique CDR3
2 unique J



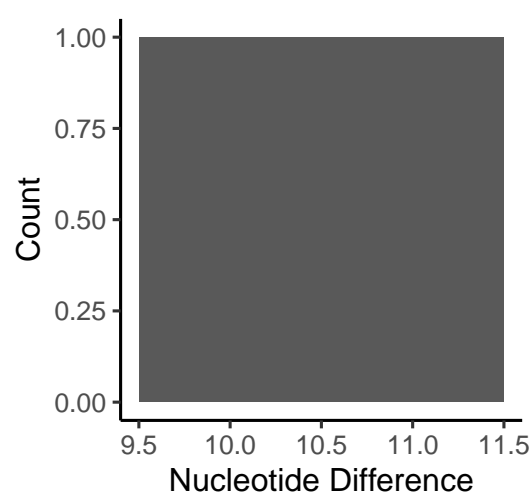
IGHV3-48*02

15 sequences assigned
No exact matches.



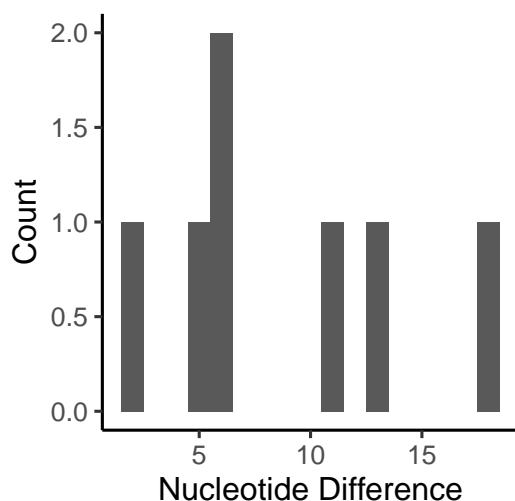
IGHV3-64*07

2 sequences assigned
No exact matches.



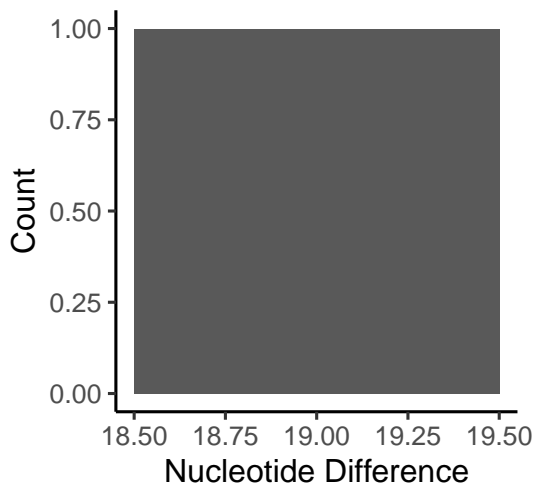
IGHV3-72*01

9 sequences assigned
No exact matches.



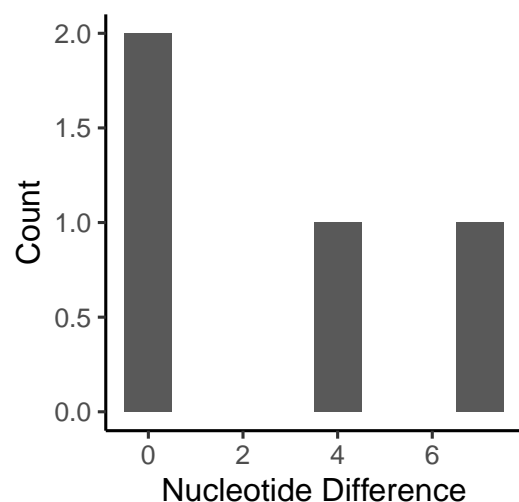
IGHV3-74*03

1 sequences assigned
No exact matches.



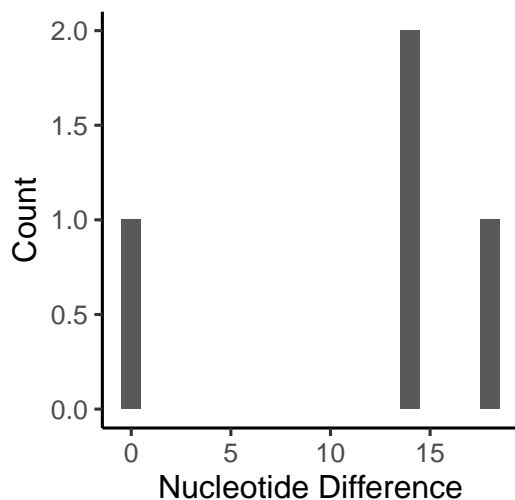
IGHV4-30-4*01

5 sequences assigned
2 (40%) exact matches, in which:
2 unique CDR3
2 unique J



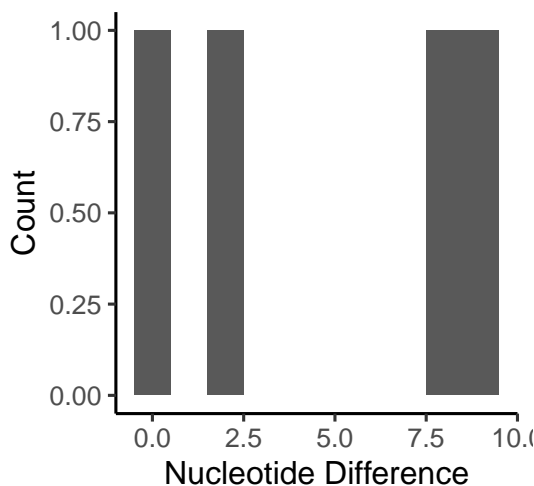
IGHV3-73*02

9 sequences assigned
1 (11.1%) exact matches, in which:
1 unique CDR3
1 unique J



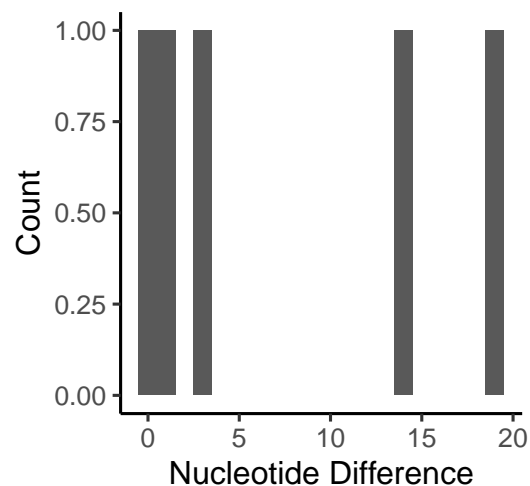
IGHV4-4*07

5 sequences assigned
1 (20%) exact matches, in which:
1 unique CDR3
1 unique J



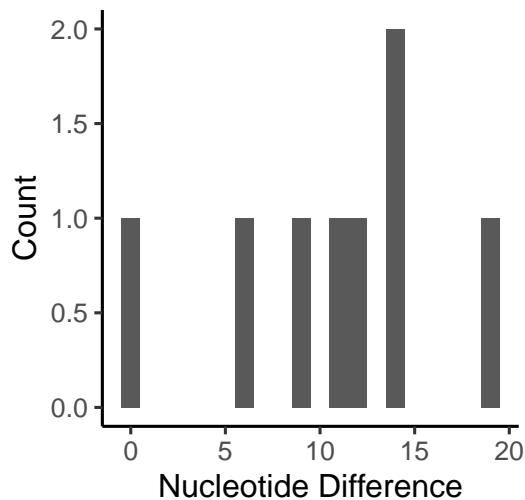
IGHV4-30-2*01

6 sequences assigned
1 (16.7%) exact matches, in which:
1 unique CDR3
1 unique J



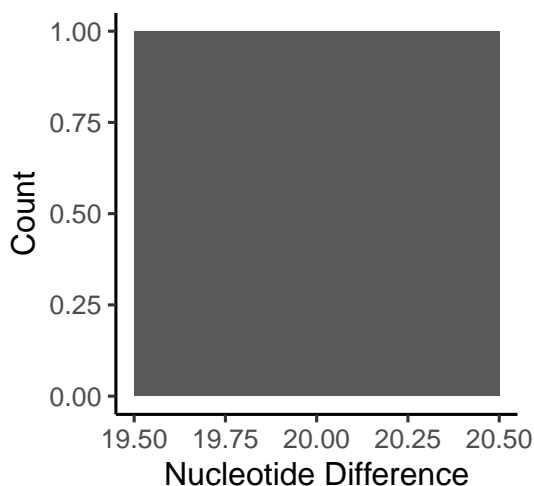
IGHV3-74*01

9 sequences assigned
1 (11.1%) exact matches, in which:
1 unique CDR3
1 unique J



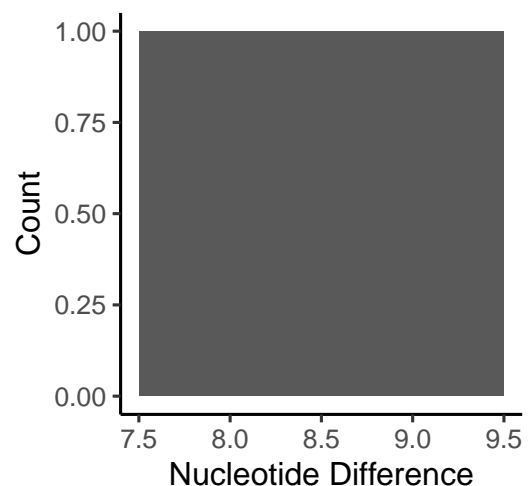
IGHV4-28*03

1 sequences assigned
No exact matches.



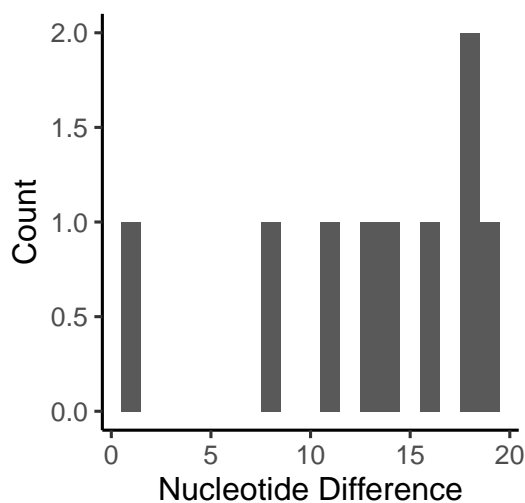
IGHV4-31*01

2 sequences assigned
No exact matches.



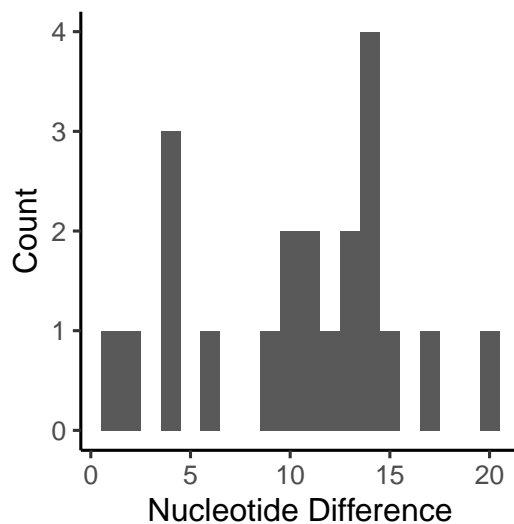
IGHV4-31*03

11 sequences assigned
No exact matches.



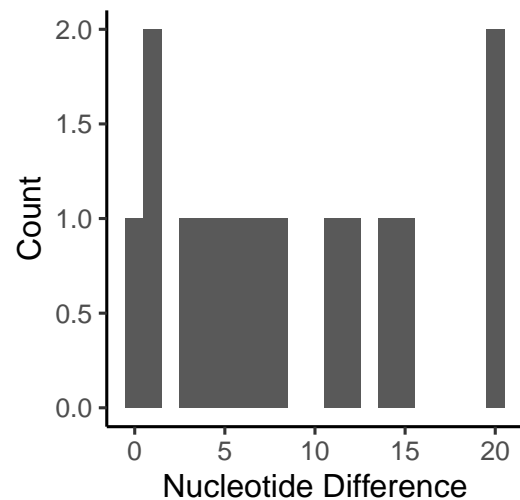
IGHV4-59*01

27 sequences assigned
No exact matches.



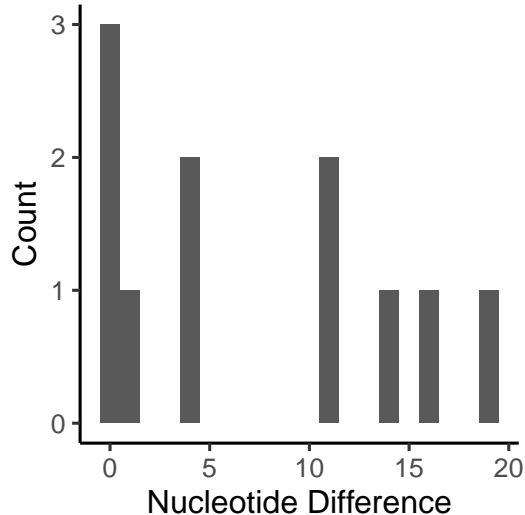
IGHV6-1*01

16 sequences assigned
1 (6.2%) exact matches, in which:
1 unique CDR3
1 unique J



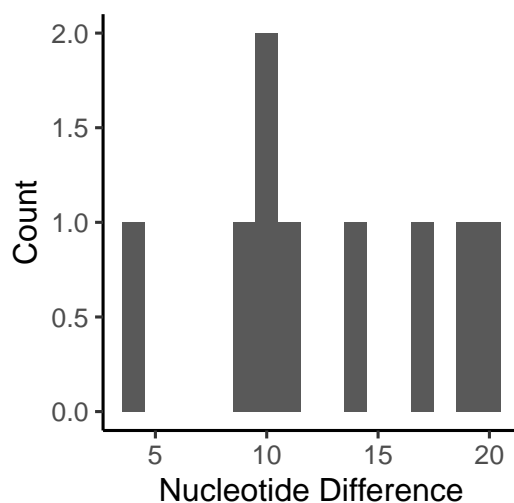
IGHV4-34*01

12 sequences assigned
3 (25%) exact matches, in which:
3 unique CDR3
2 unique J



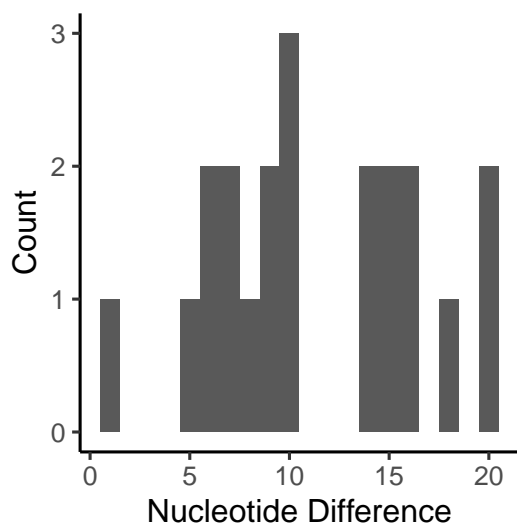
IGHV4-61*02

13 sequences assigned
No exact matches.



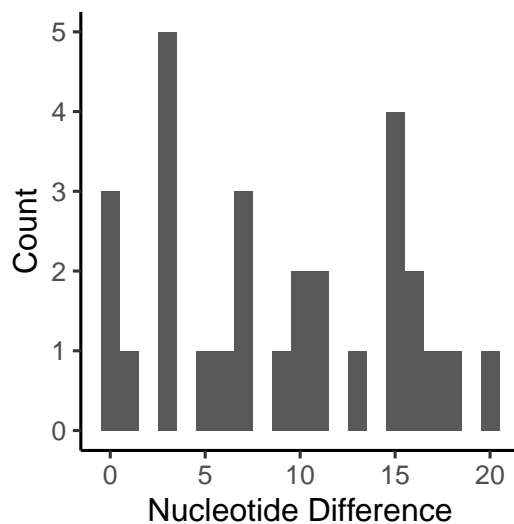
IGHV4-39*01

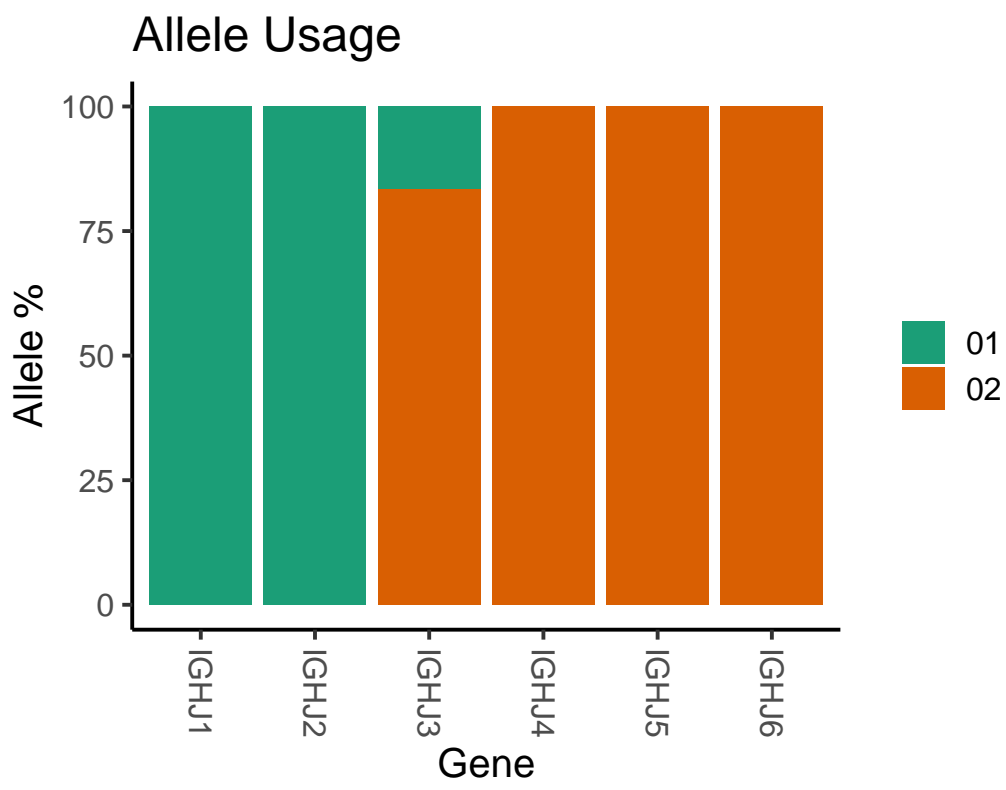
32 sequences assigned
No exact matches.



IGHV5-51*01

36 sequences assigned
3 (8.3%) exact matches, in which:
3 unique CDR3
3 unique J





Warning – no inferred sequences found.

Warning – no inferred sequences found.