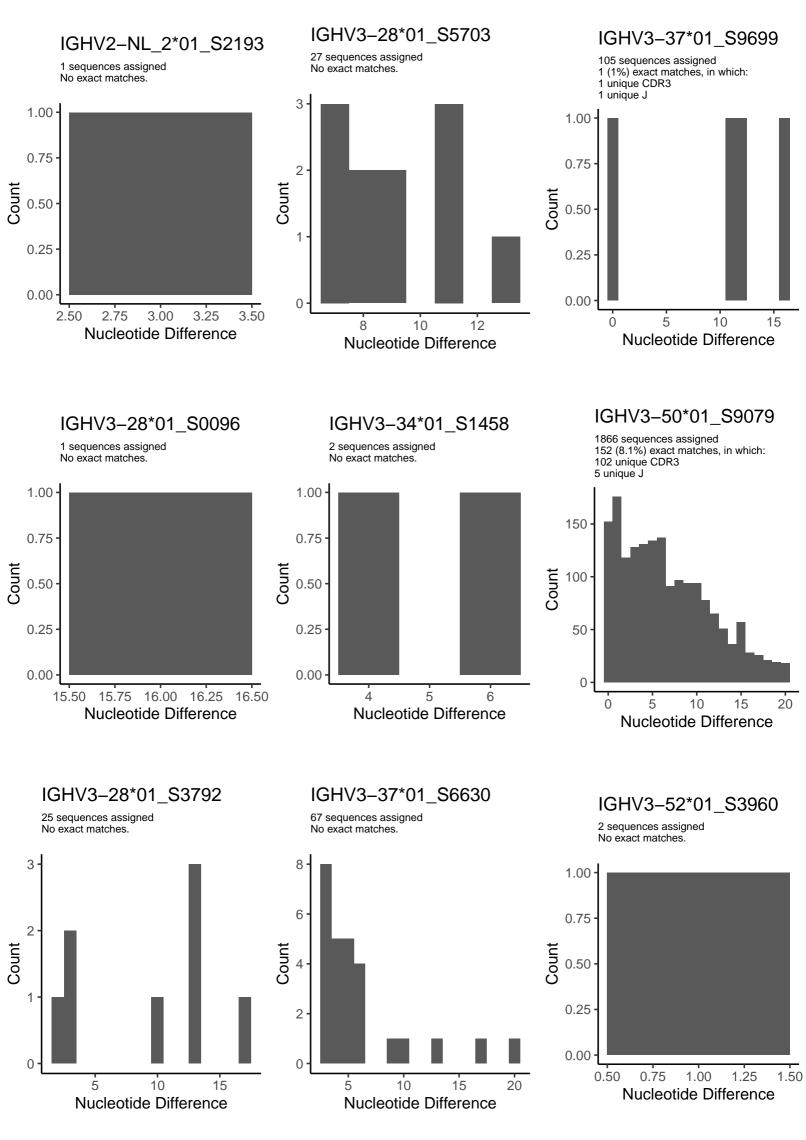
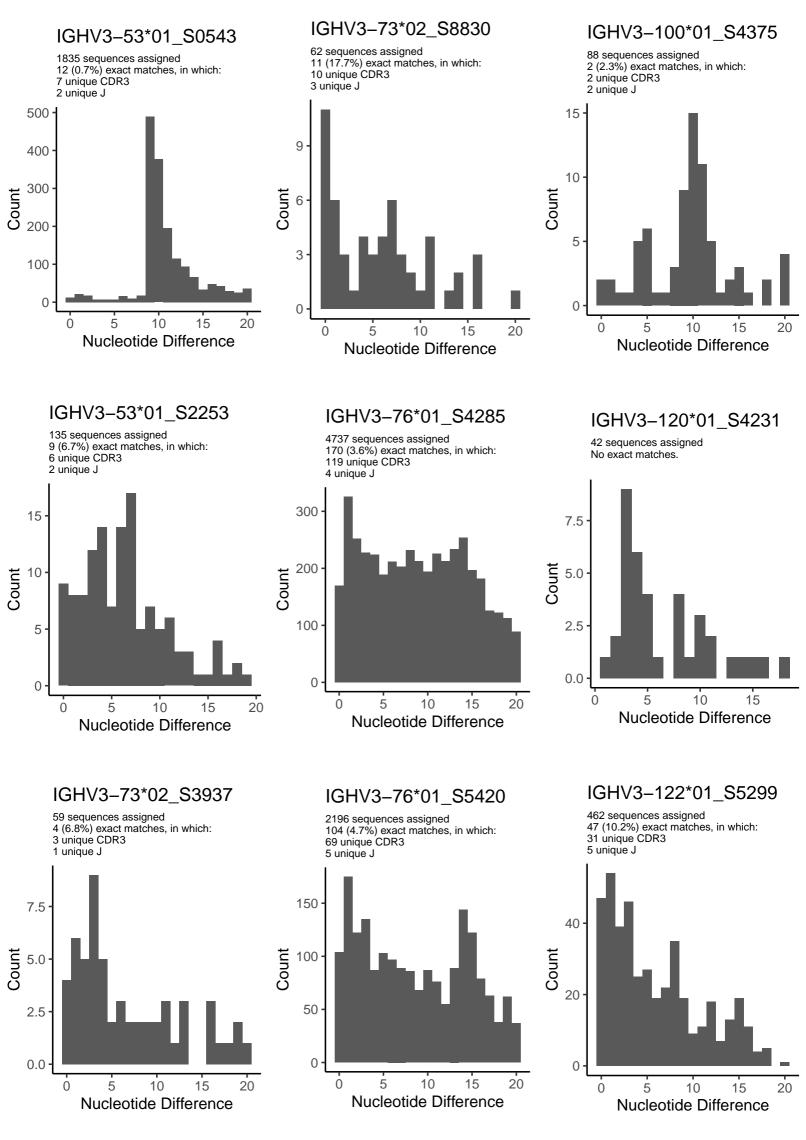


## IGHV2-69\*01\_S2408 IGHV2-7\*02\_S1732 IGHV2-173\*01\_S2781 364 sequences assigned 36 (9.9%) exact matches, in which: 2134 sequences assigned 137 (6.4%) exact matches, in which: 86 unique CDR3 13 sequences assigned No exact matches. 18 unique CDR3 3 unique J 4 unique J 3 200 40 Count 100 20 0 0 5 10 15 20 0 10 0 15 Nucleotide Difference Nucleotide Difference **Nucleotide Difference** IGHV2-7\*02\_S4285 IGHV2-69\*01\_S6153 IGHV2-NL\_1\*01\_S5779 639 sequences assigned 52 (8.1%) exact matches, in which: 326 sequences assigned 1020 sequences assigned 115 (11.3%) exact matches, in which: 37 (11.3%) exact matches, in which: 27 unique CDR3 30 unique CDR3 68 unique CDR3 3 unique J 4 unique J 5 unique J 125 50 100 40 40 75 Count 30 Count 50 20 20 25 10 0 0 0 10 20 0 5 15 10 20 10 15 20 0 5 15 0 5 Nucleotide Difference Nucleotide Difference Nucleotide Difference IGHV2-69\*01\_C6G\_C36G IGHV2-NL\_1\*01\_S5779\_0 IGHV2-118\*01\_S4527 314 sequences assigned 183 sequences assigned 3322 sequences assigned 17 (9.3%) exact matches, in which: 45 (14.3%) exact matches, in which: 216 (6.5%) exact matches, in which: 17 unique CDR3 39 unique CDR3 134 unique CDR3 3 unique J 4 unique J 5 unique J 40 300 30 Count 200 20 5 100 10 0 0 0 20 5 10 15 0 5 10 15 20 5 10 15 20 **Nucleotide Difference** Nucleotide Difference Nucleotide Difference



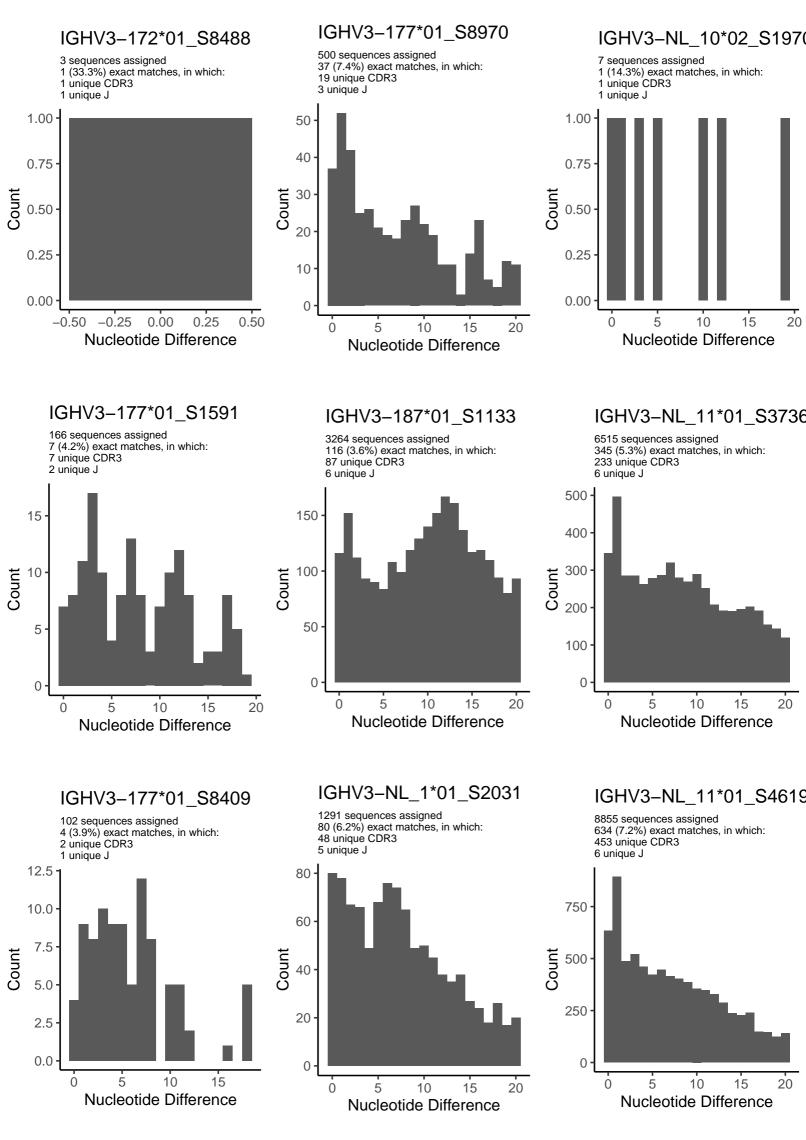


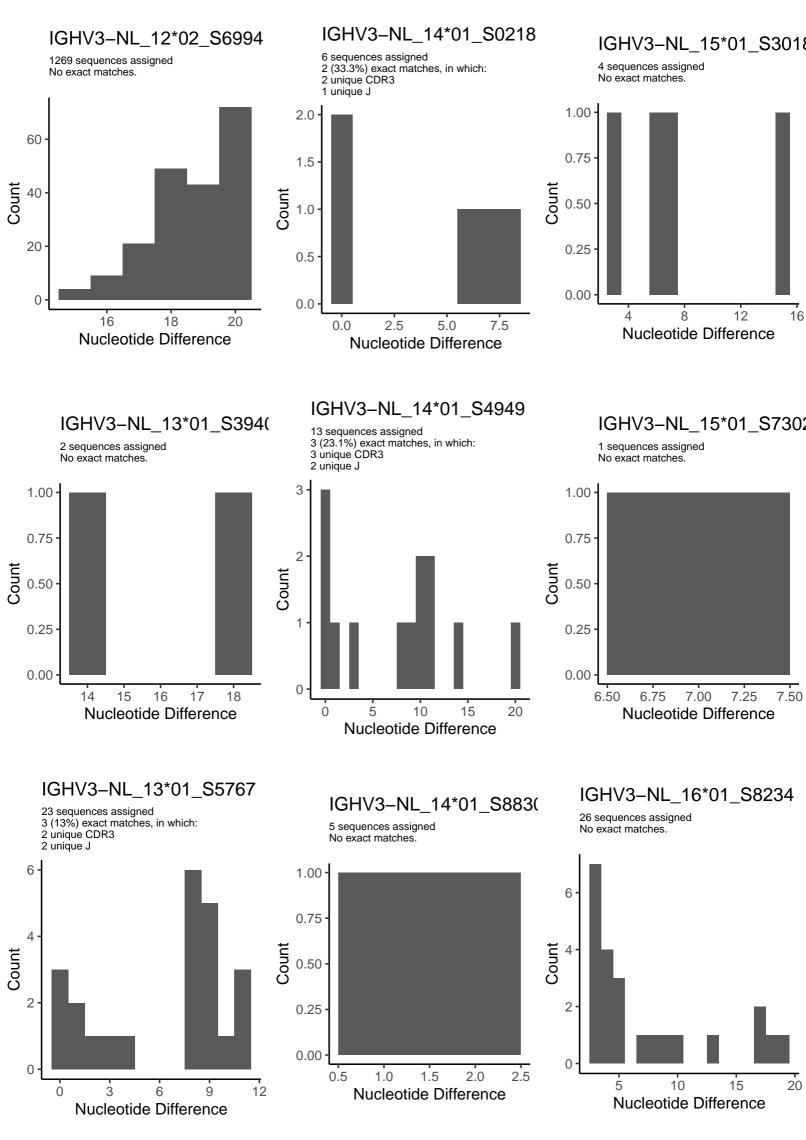
## IGHV3-122\*01\_S8388 IGHV3-128\*01\_S1128 IGHV3-141\*01\_S8690 880 sequences assigned 5634 sequences assigned 86 sequences assigned 68 (7.7%) exact matches, in which: 38 unique CDR3 40 (0.7%) exact matches, in which: 28 unique CDR3 6 (7%) exact matches, in which: 4 unique CDR3 4 unique J 4 unique J 2 unique J 12.5 500 10.0 400 7.5 300 Count Count 5.0 200 20 2.5 100 0.0 0 0 20 5 10 10 15 20 0 5 10 15 20 0 5 **Nucleotide Difference** Nucleotide Difference Nucleotide Difference IGHV3-127\*01\_S5171 IGHV3-128\*02 S1570 IGHV3-141\*02 S4530 1142 sequences assigned 124 sequences assigned 611 sequences assigned 33 (5.4%) exact matches, in which: 22 unique CDR3 56 (4.9%) exact matches, in which: 19 (15.3%) exact matches, in which: 13 unique CDR3 35 unique CDR3 4 unique J 4 unique J 3 unique J 50 20 60 40 15 30 Count Count Count 40 10 20 20 5 10 0 0 0 10 15 20 10 15 20 15 0 5 0 5 0 10 Nucleotide Difference Nucleotide Difference Nucleotide Difference IGHV3-127\*01\_S6769 IGHV3-141\*01 S5017 IGHV3-152\*01\_S5891 1048 sequences assigned 116 sequences assigned 1198 sequences assigned 20 (17.2%) exact matches, in which: 20 (1.7%) exact matches, in which: 16 (1.5%) exact matches, in which: 8 unique CDR3 15 unique CDR3 14 unique CDR3 2 unique J 4 unique J 3 unique J 20 80 90 15 60 Count 10 40 30 5 20 0 0 0 0 10 15 20 0 10 15 20 0 5 10 15 20 5

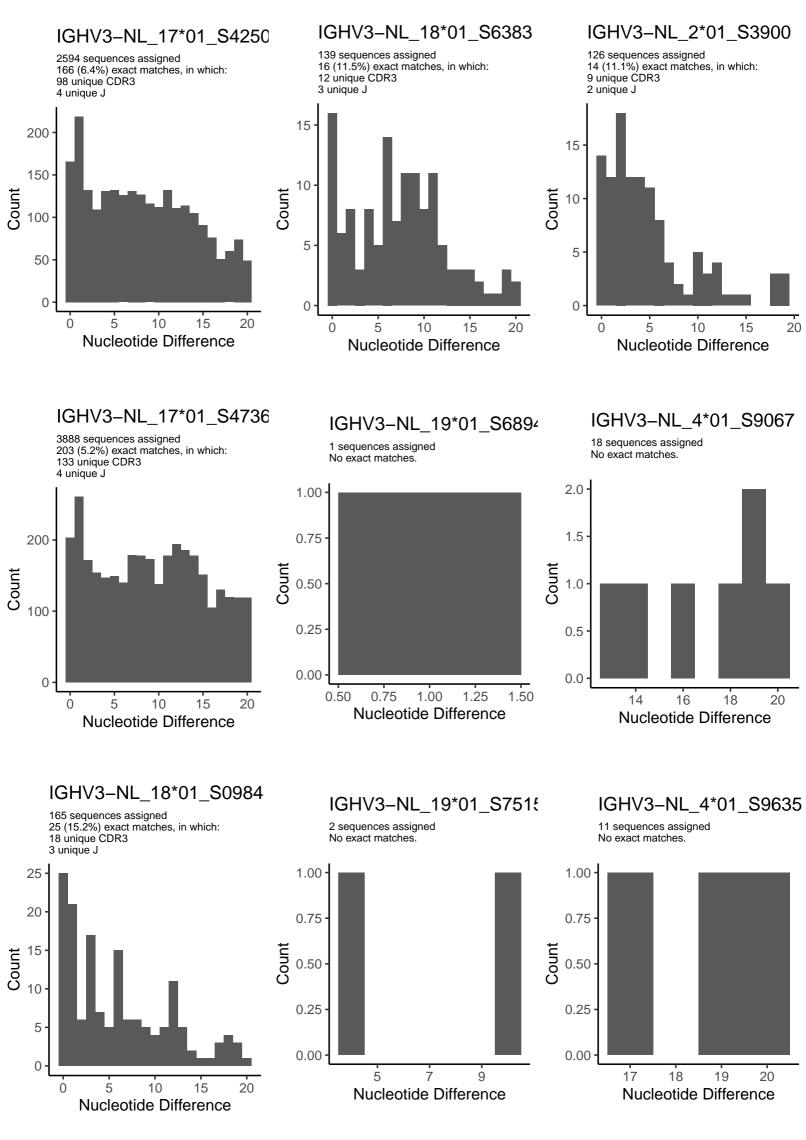
**Nucleotide Difference** 

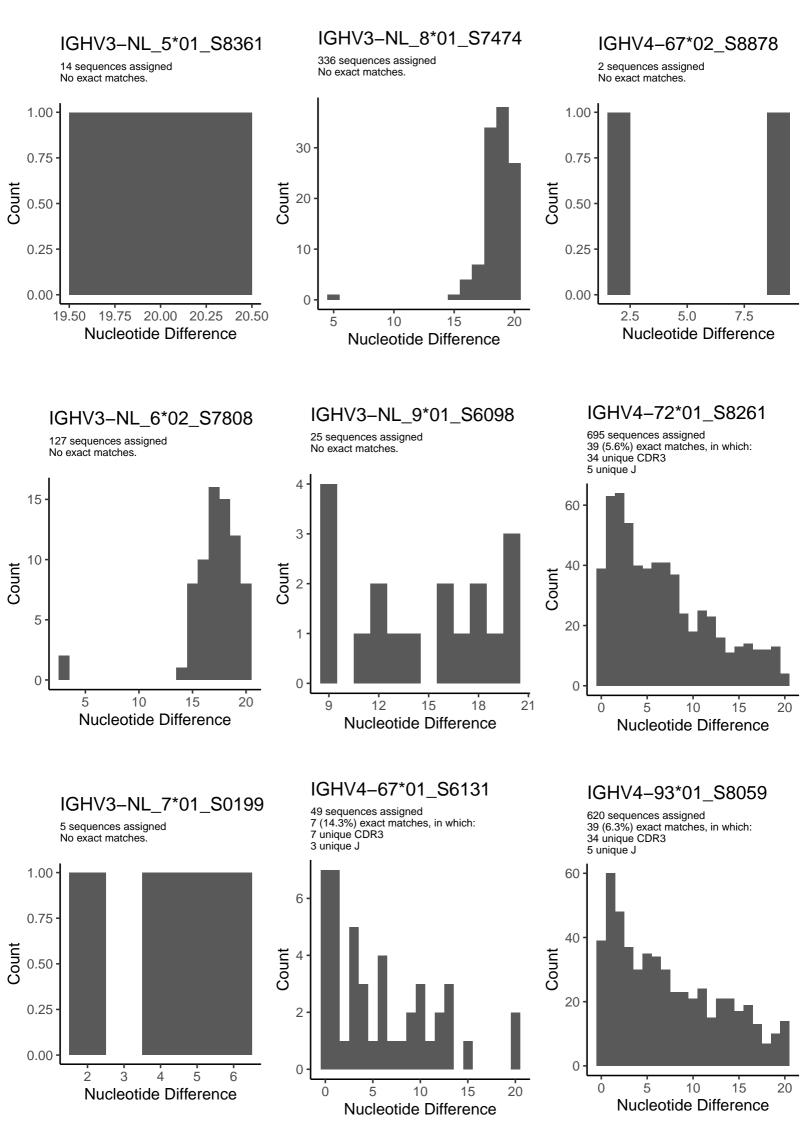
Nucleotide Difference

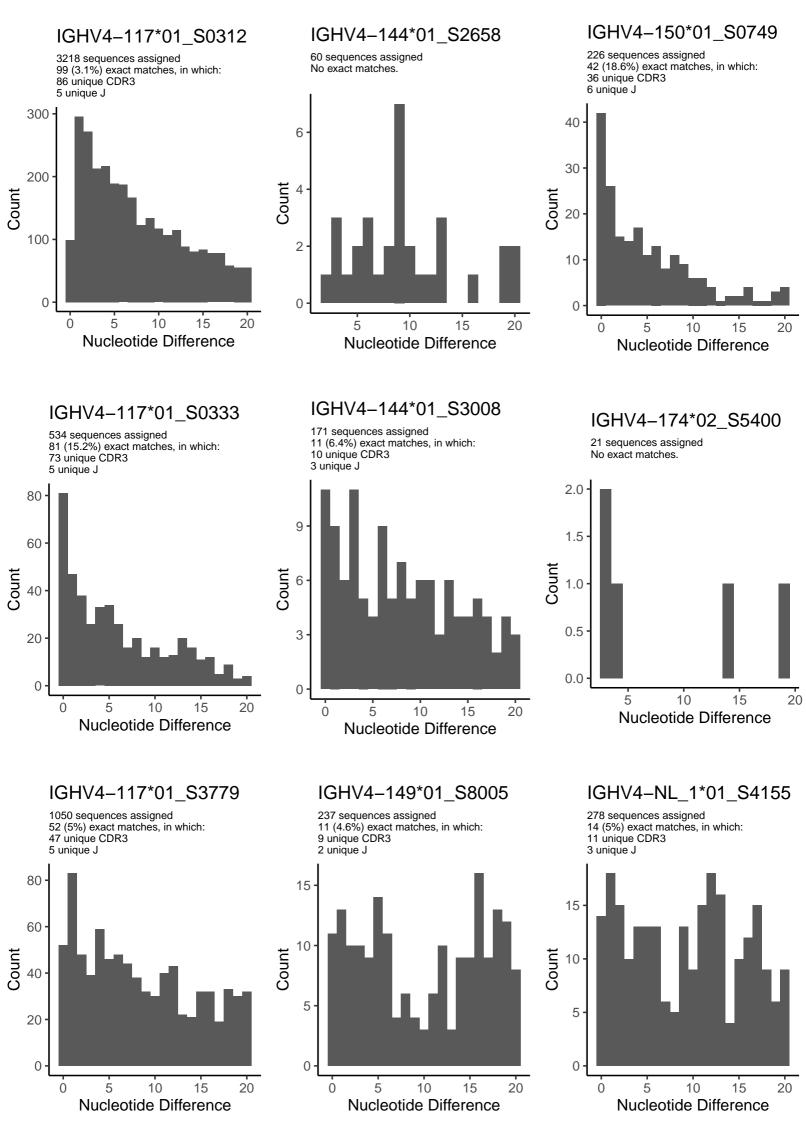
Nucleotide Difference

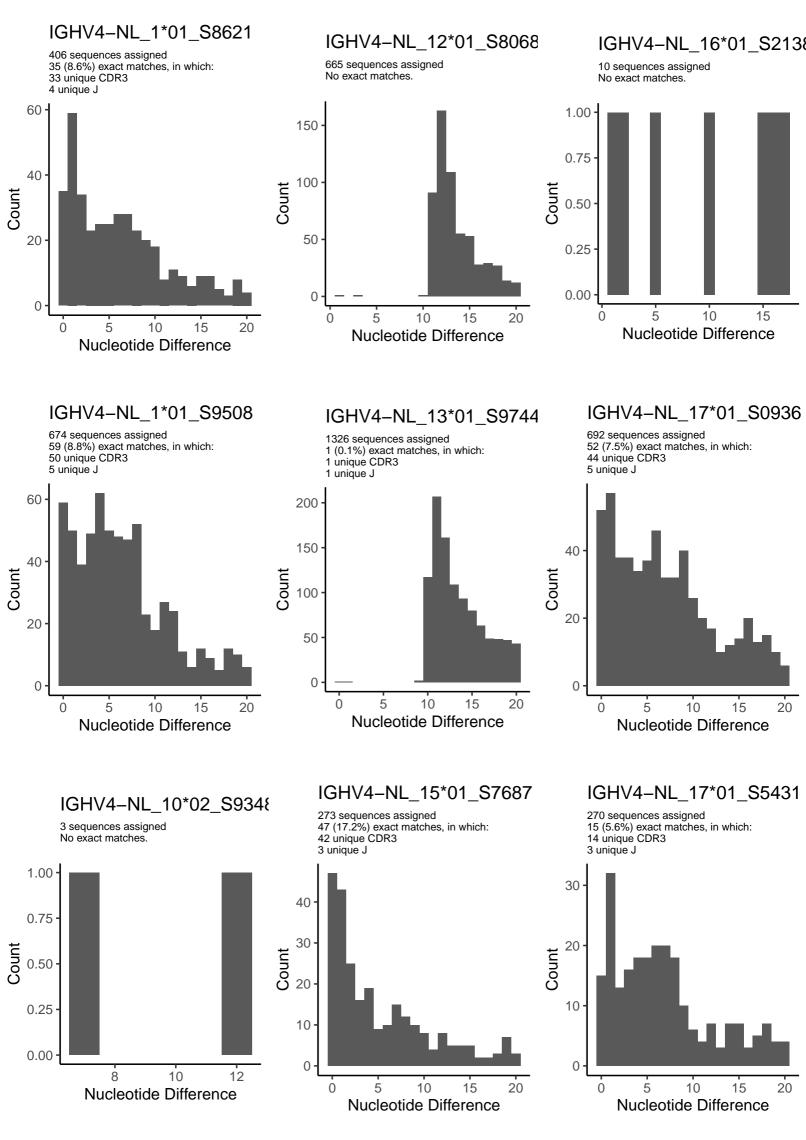


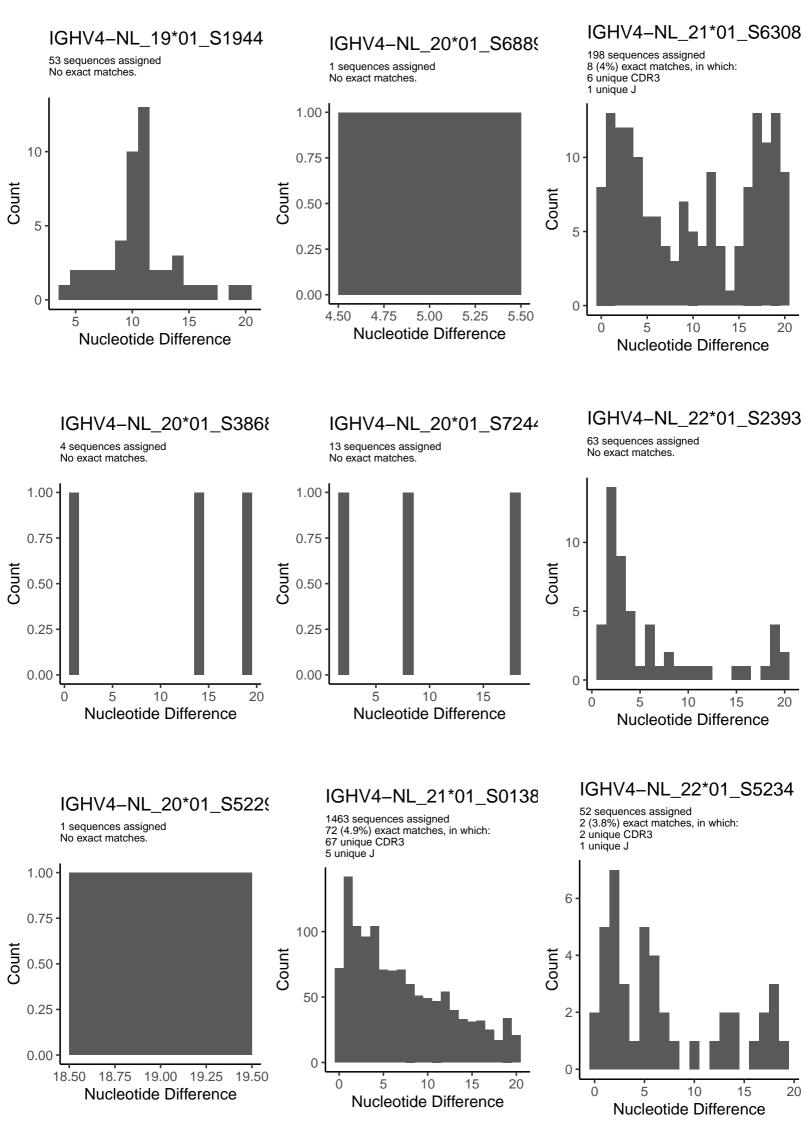


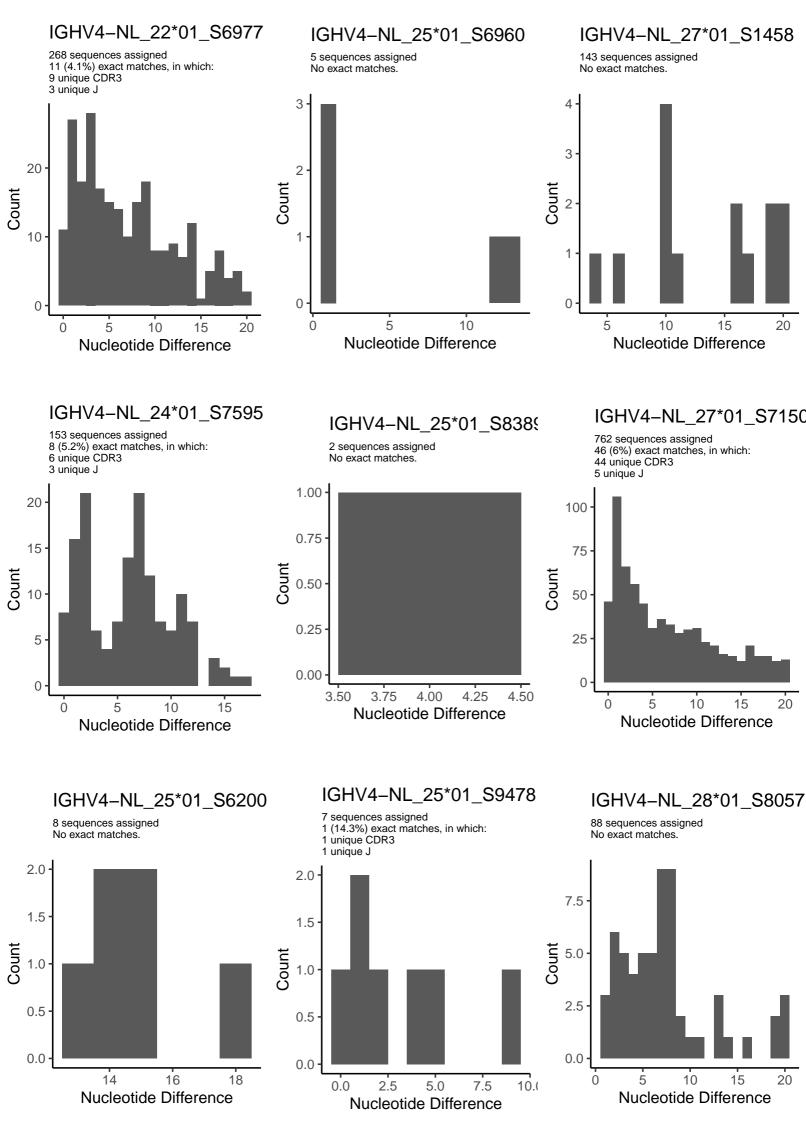


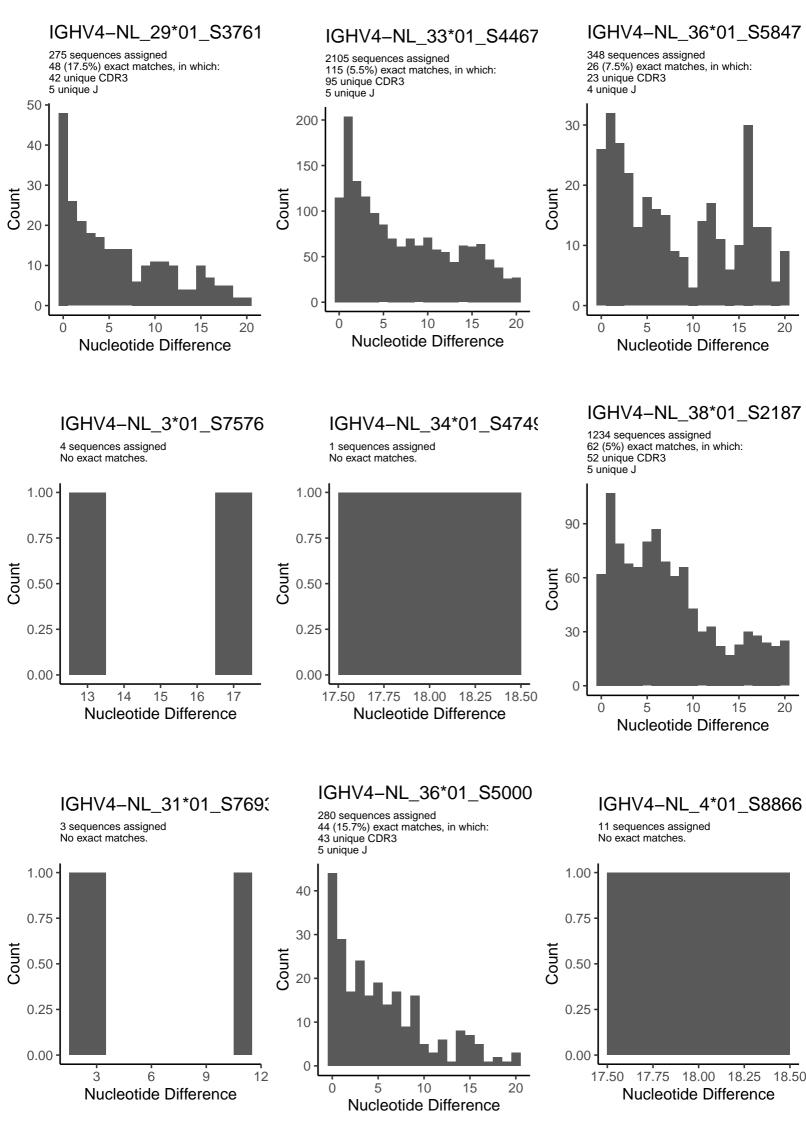


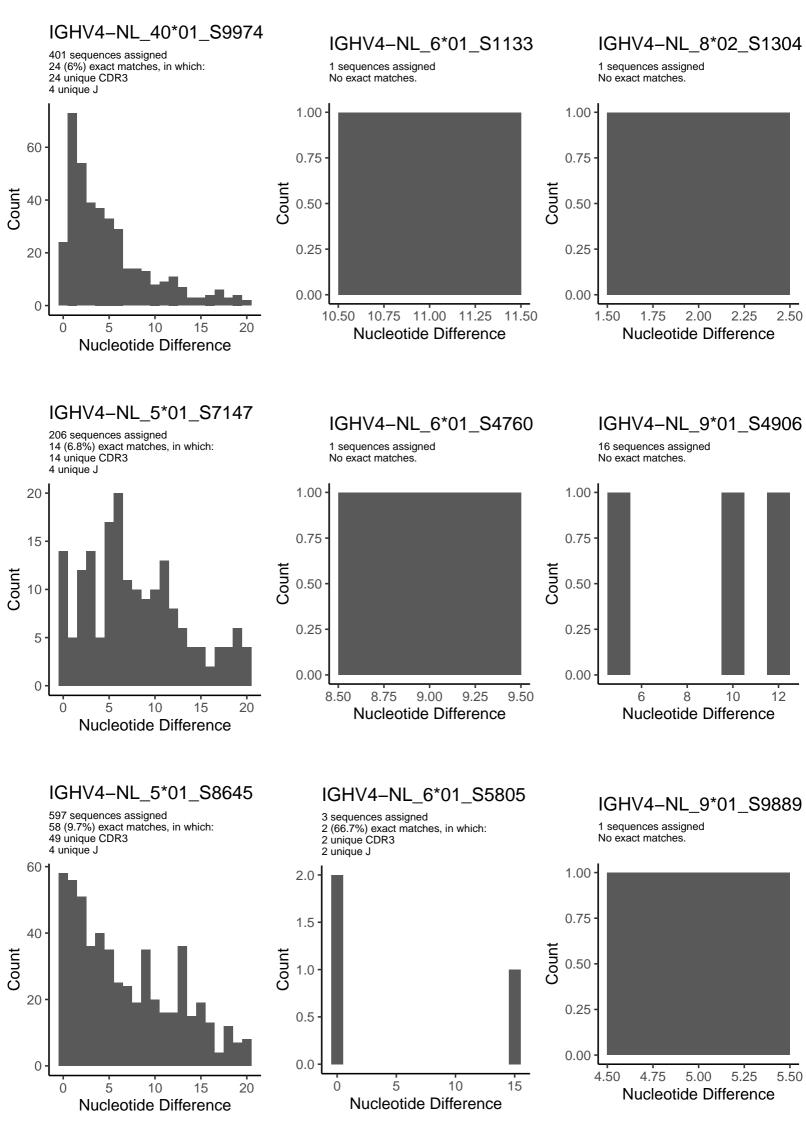












## IGHV7-86\*01\_S0200 IGHV5-15\*01\_S0308 751 sequences assigned 127 (16.9%) exact matches, in which: 68 unique CDR3 487 sequences assigned 63 (12.9%) exact matches, in which: 35 unique CDR3 4 unique J 5 unique J 150 60 100 40 Count Count 50 20 0 0 5 10 15 20 0 0 5 10 15 **Nucleotide Difference** Nucleotide Difference IGHV7-135\*01\_S4281 IGHV5-15\*01\_S4790 11 sequences assigned 3 (27.3%) exact matches, in which: 1 unique CDR3 1312 sequences assigned 180 (13.7%) exact matches, in which: 101 unique CDR3 1 unique J 5 unique J 3 250 -200 2 150 100 50 0 0 5 10 15 20 0 10 15 0 5 **Nucleotide Difference** Nucleotide Difference IGHV5-157\*02\_S8705 IGHV7-135\*02\_S1424 11 sequences assigned 3560 sequences assigned No exact matches. 414 (11.6%) exact matches, in which: 254 unique CDR3 5 unique J 3 600 400 Count 200 0

10

**Nucleotide Difference** 

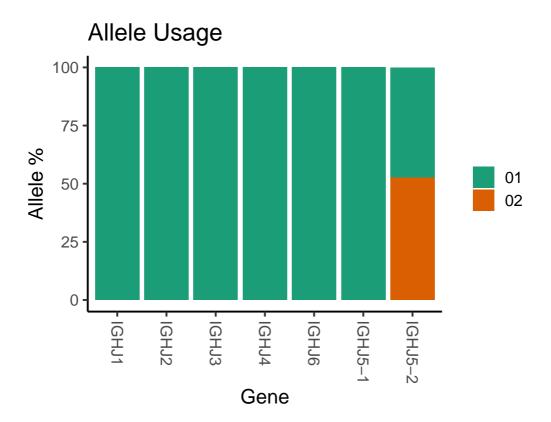
15

20

5.0

**Nucleotide Difference** 

7.5



## Sequence Count by IGHJ5-2 allele usage



