

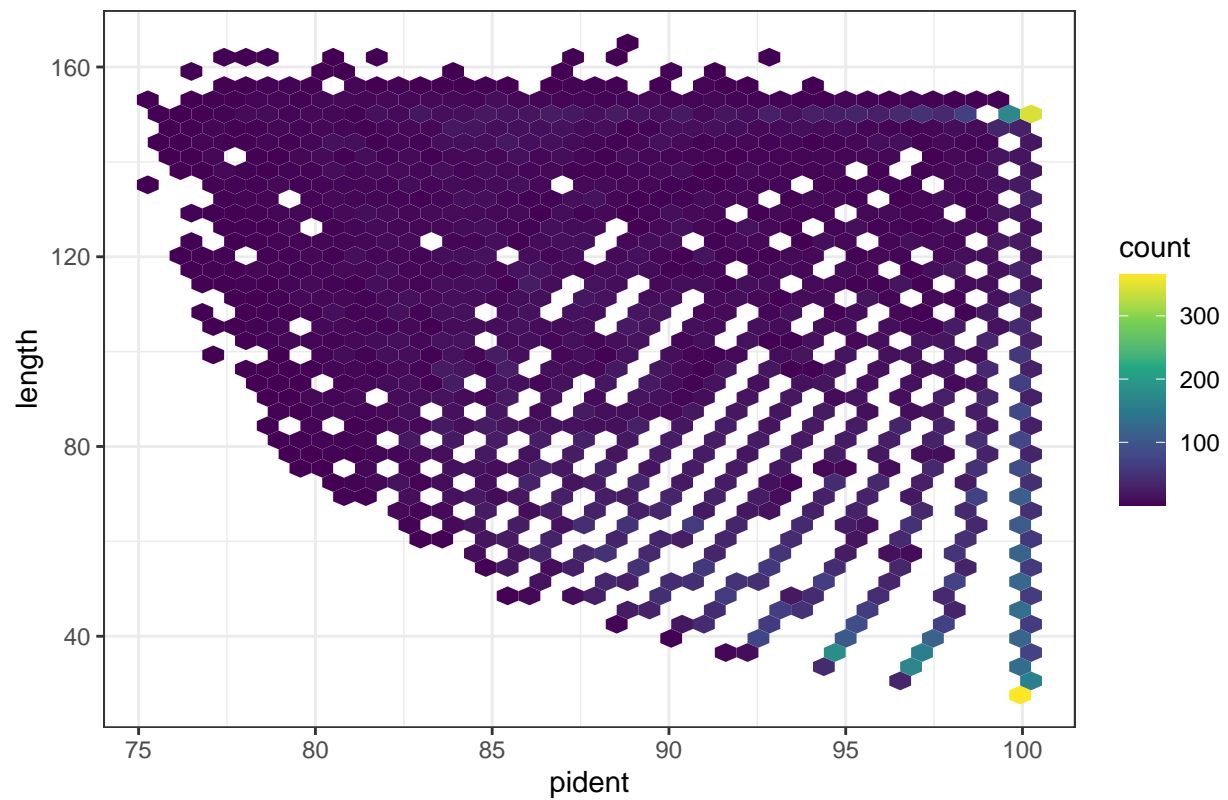
intermediate investigation on parchment data

2022-10-02

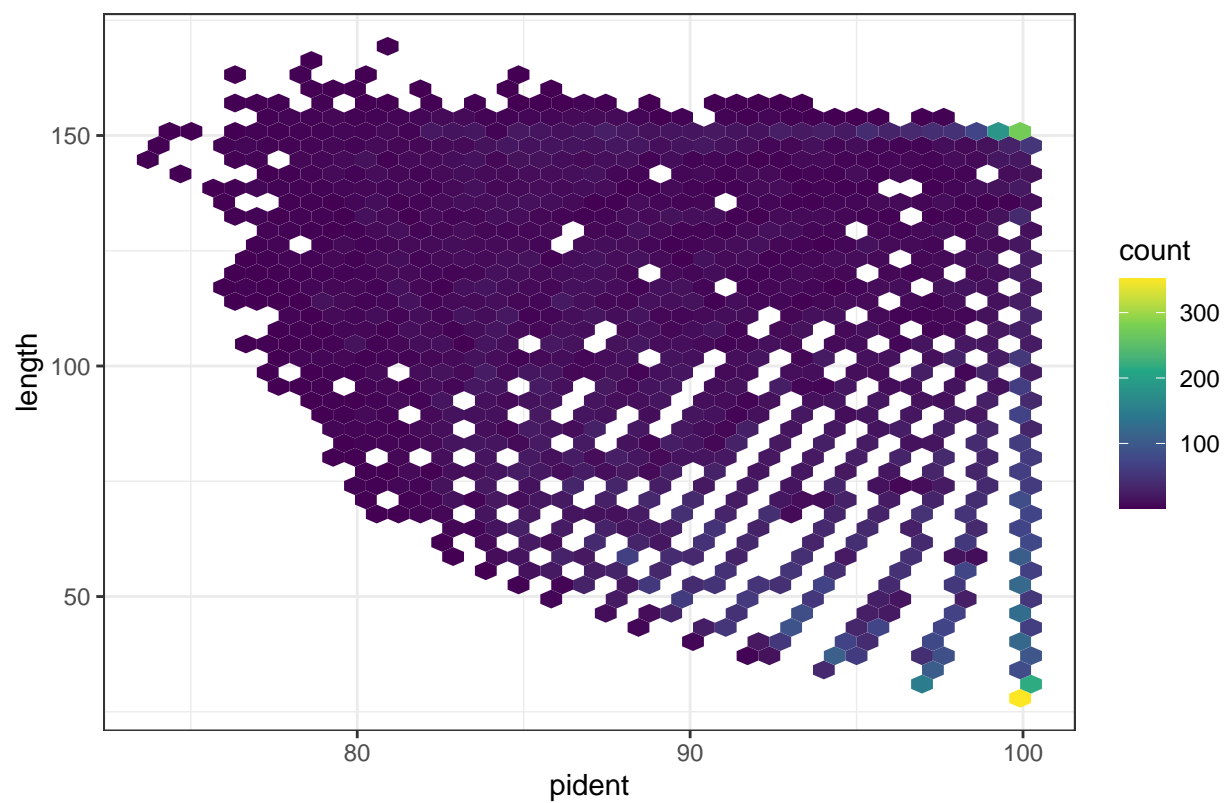
percentage of identity V.S. coverage among 22 samples.

sample SG-LibNeg_S10_L001_R1_001 and SG-LibNeg_S10_L001_R2_001 have few hits that are not informative.

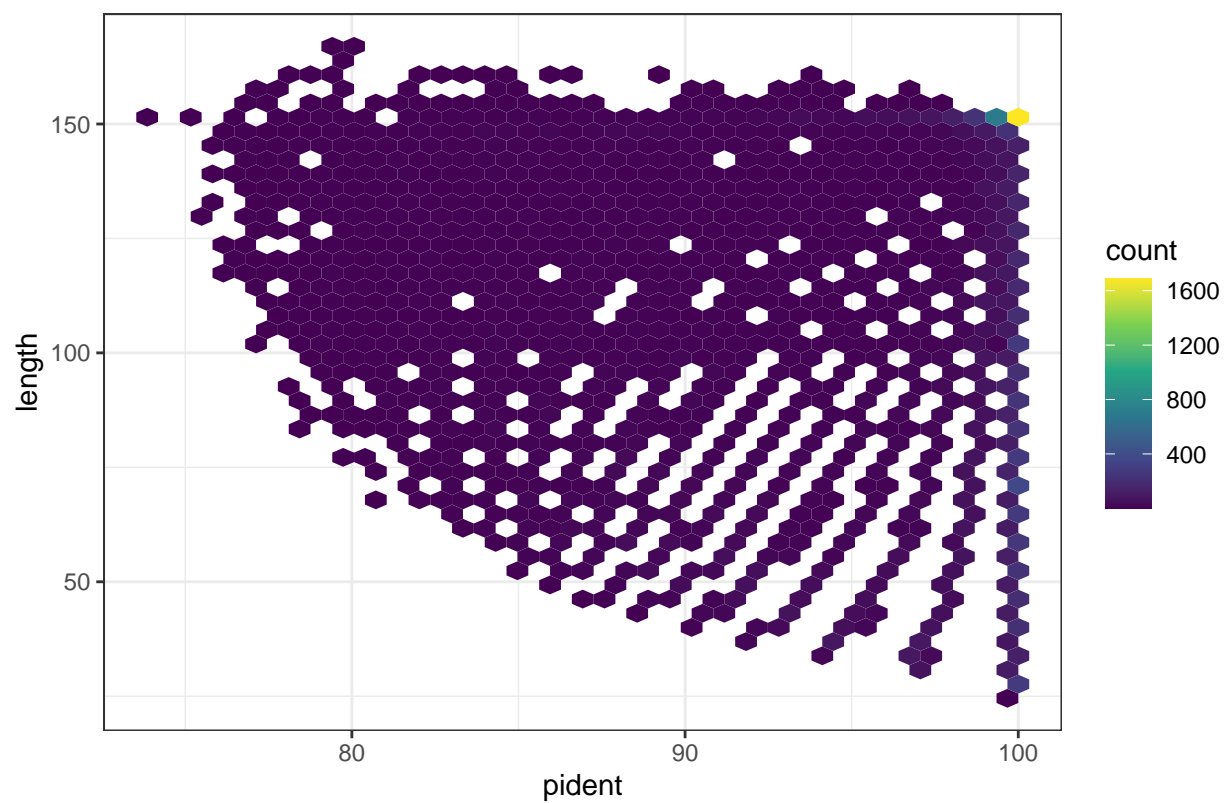
SG-002_S1_L001_R1_001

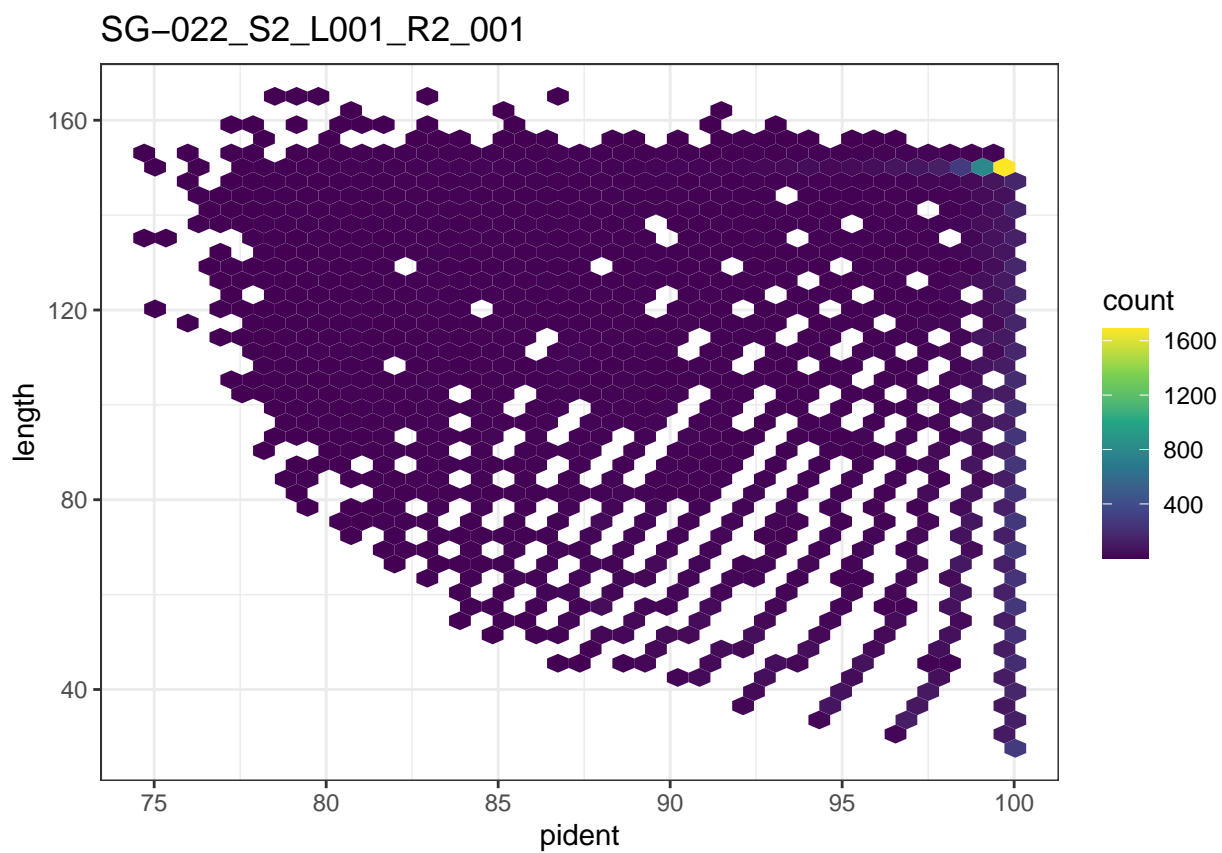


SG-002_S1_L001_R2_001

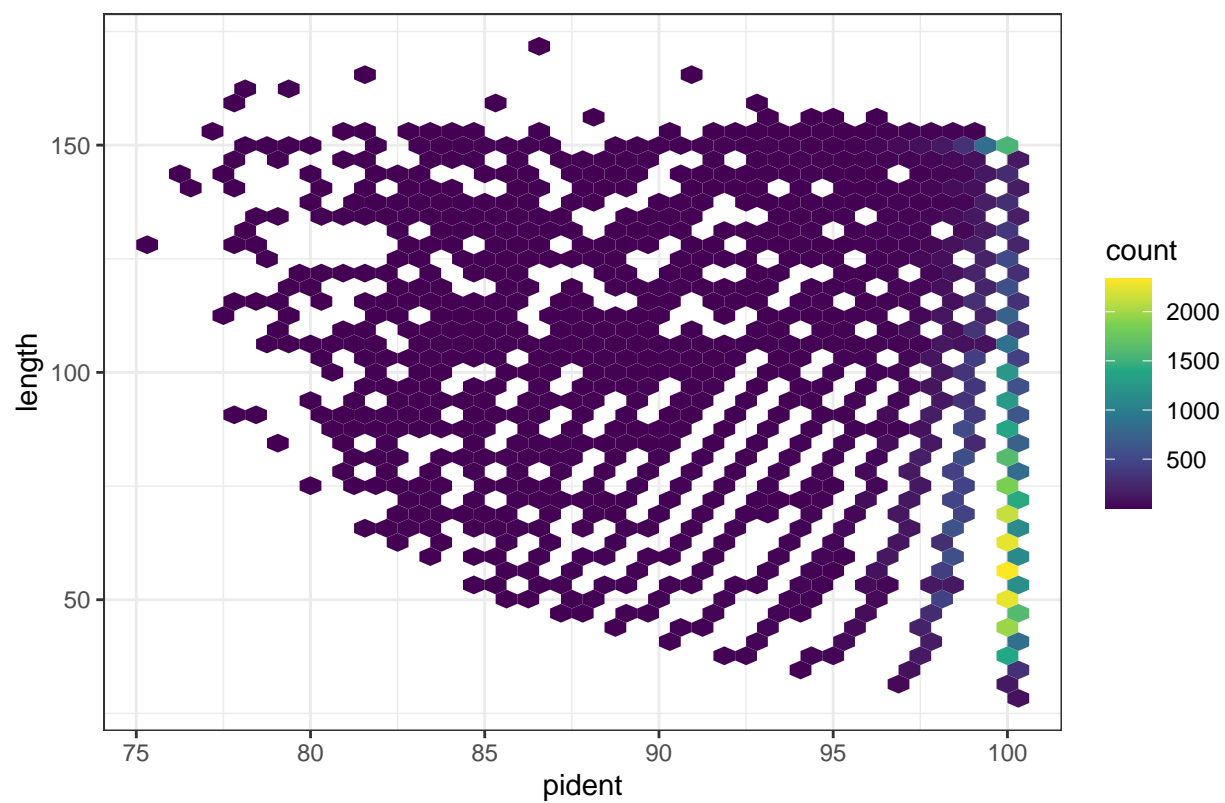


SG-022_S2_L001_R1_001

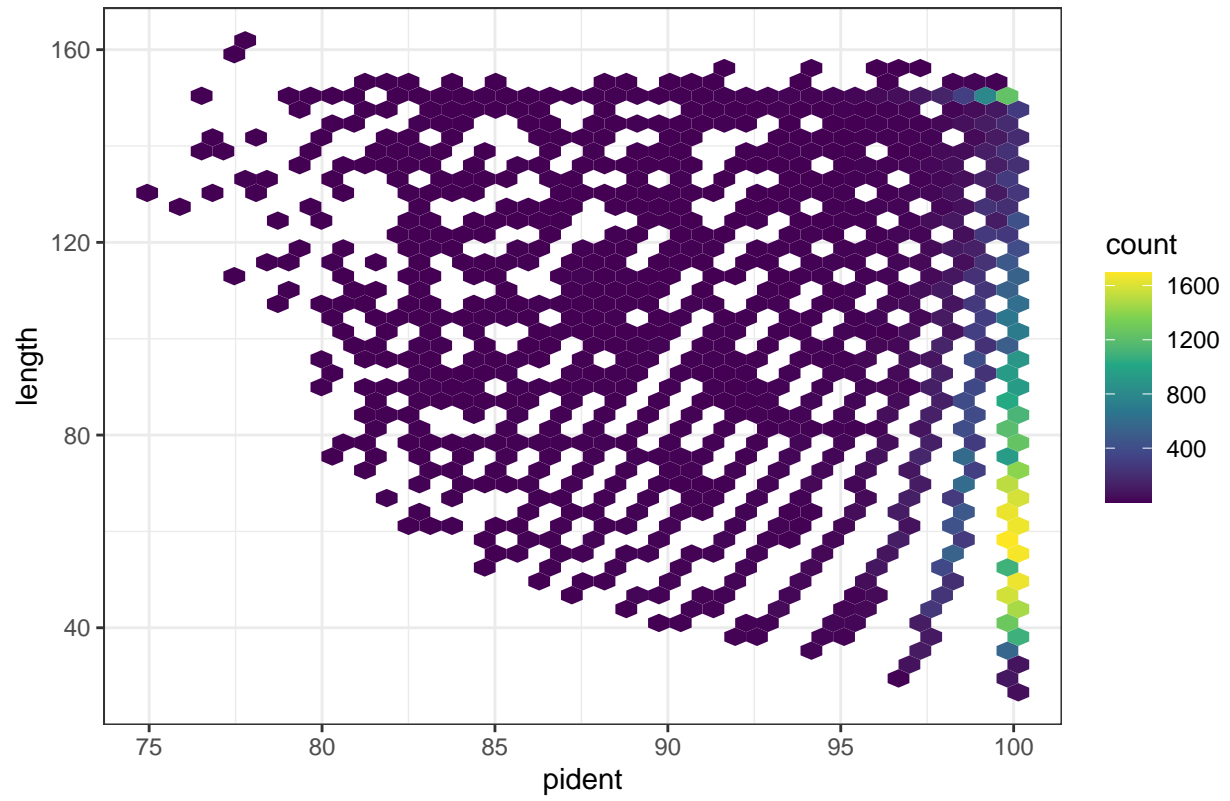




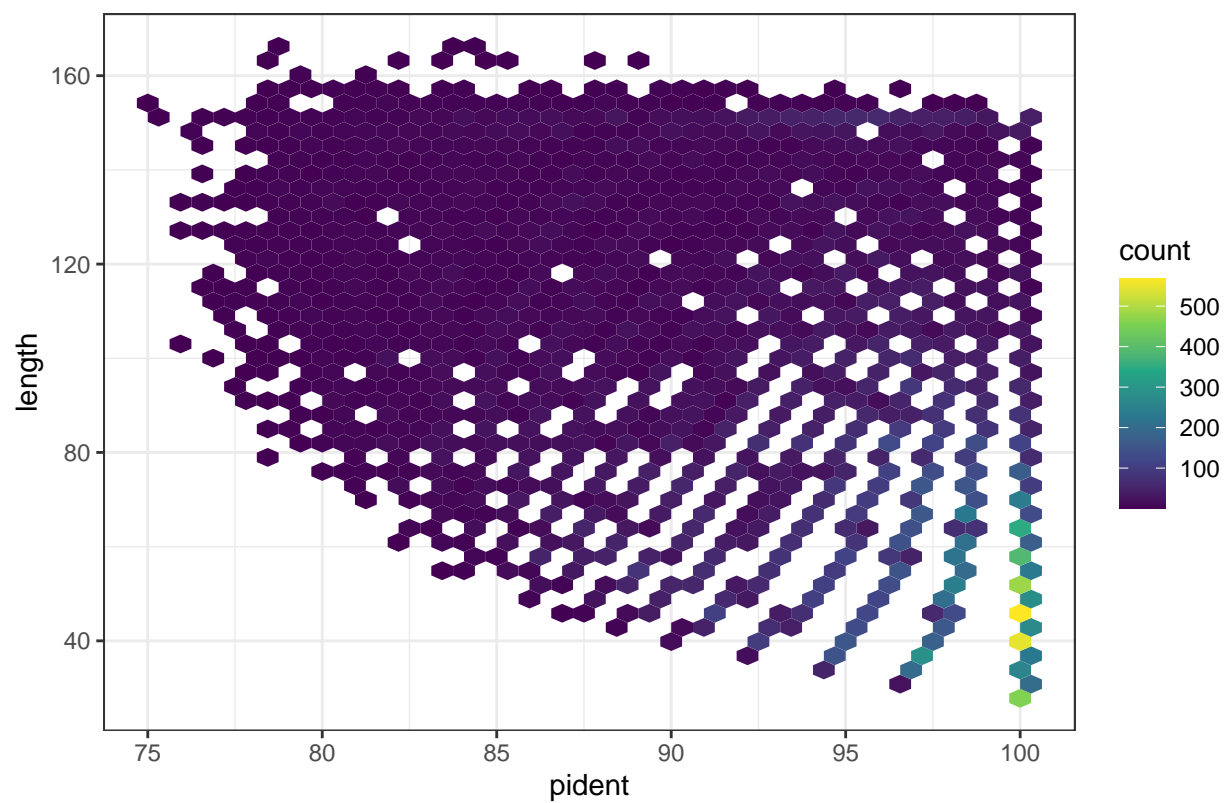
SG-047_S3_L001_R1_001



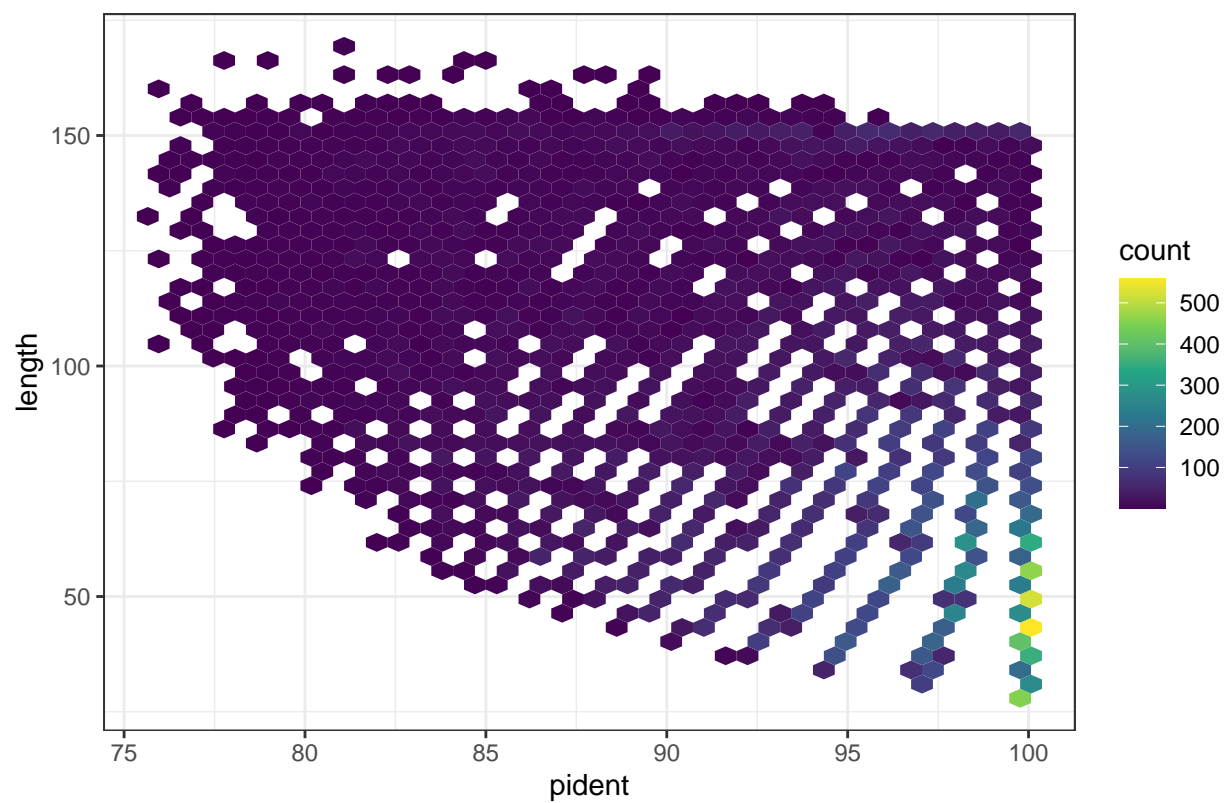
SG-047_S3_L001_R2_001



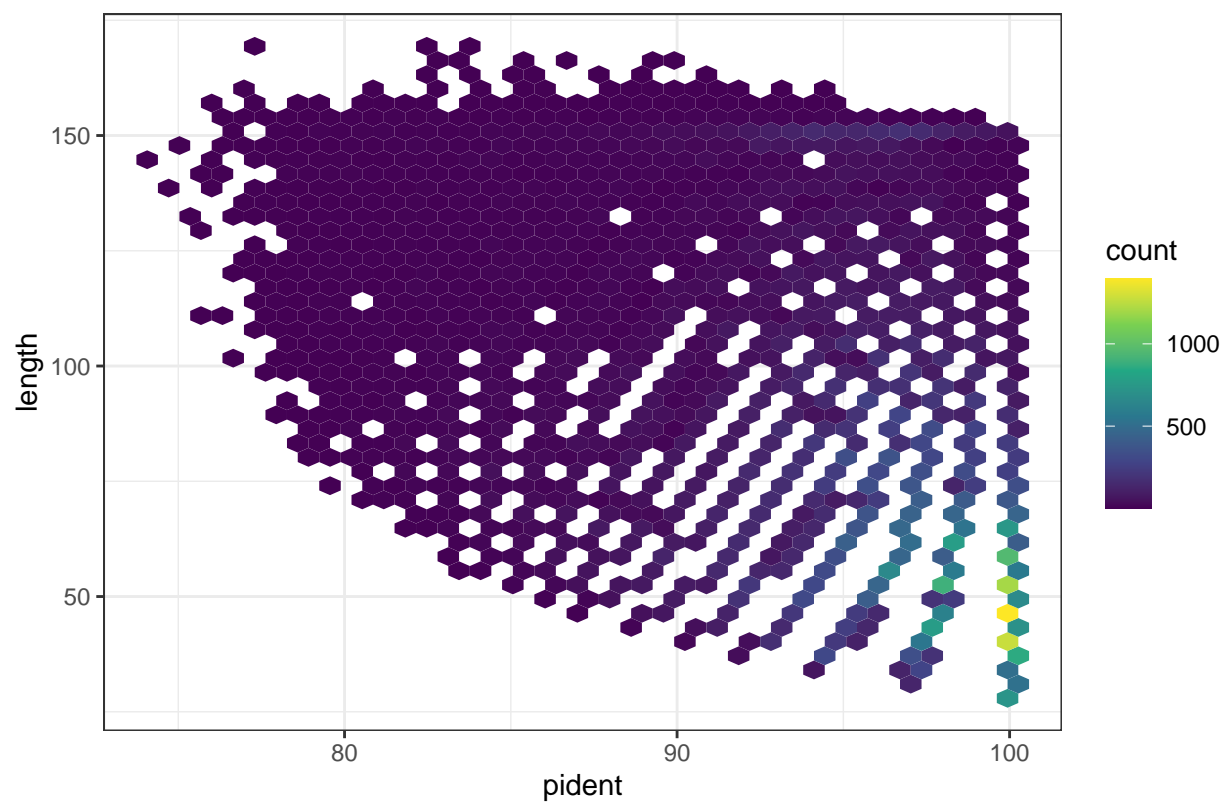
SG-087_S4_L001_R1_001



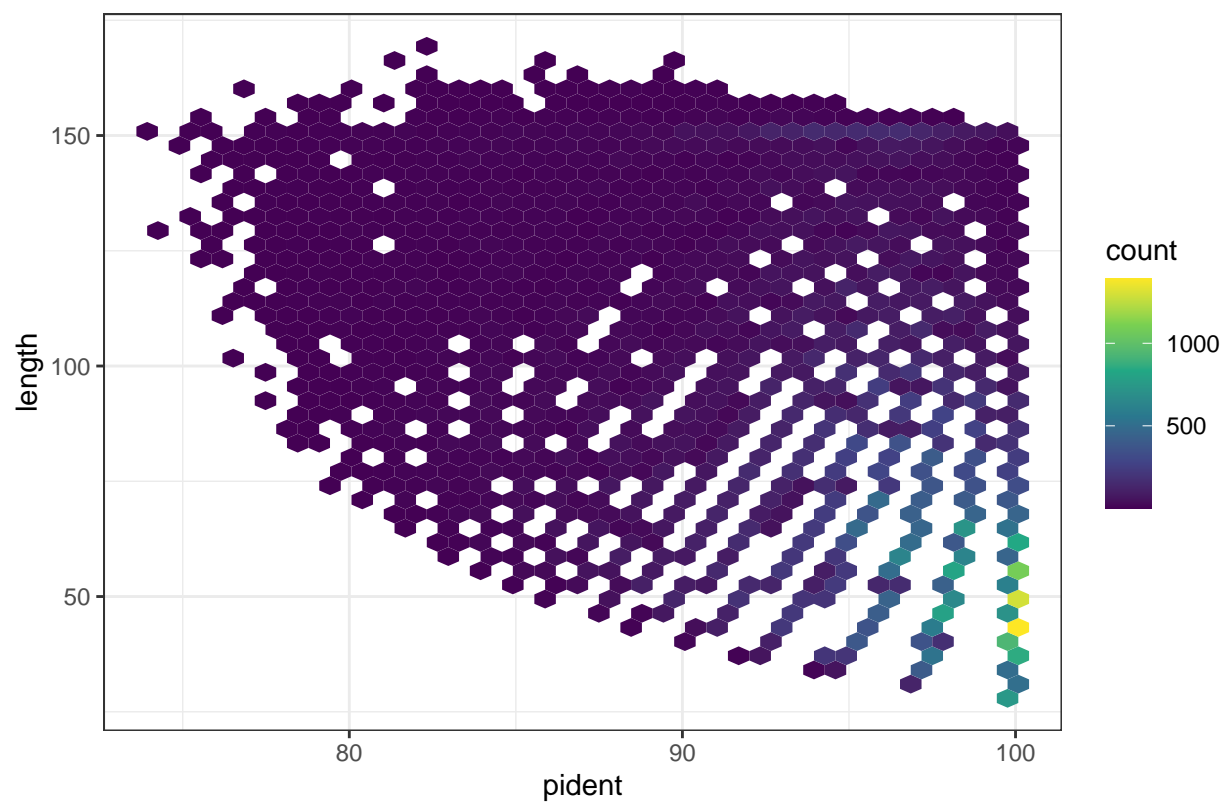
SG-087_S4_L001_R2_001



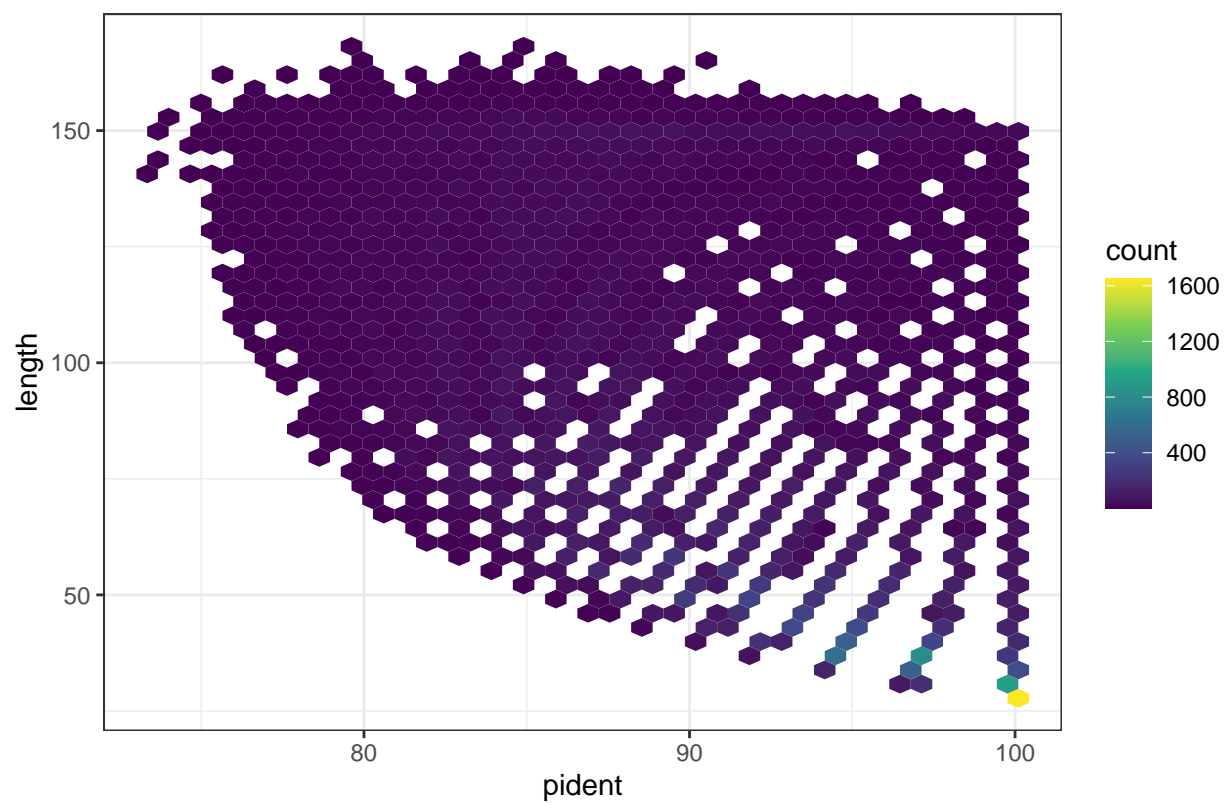
SG-112_S5_L001_R1_001



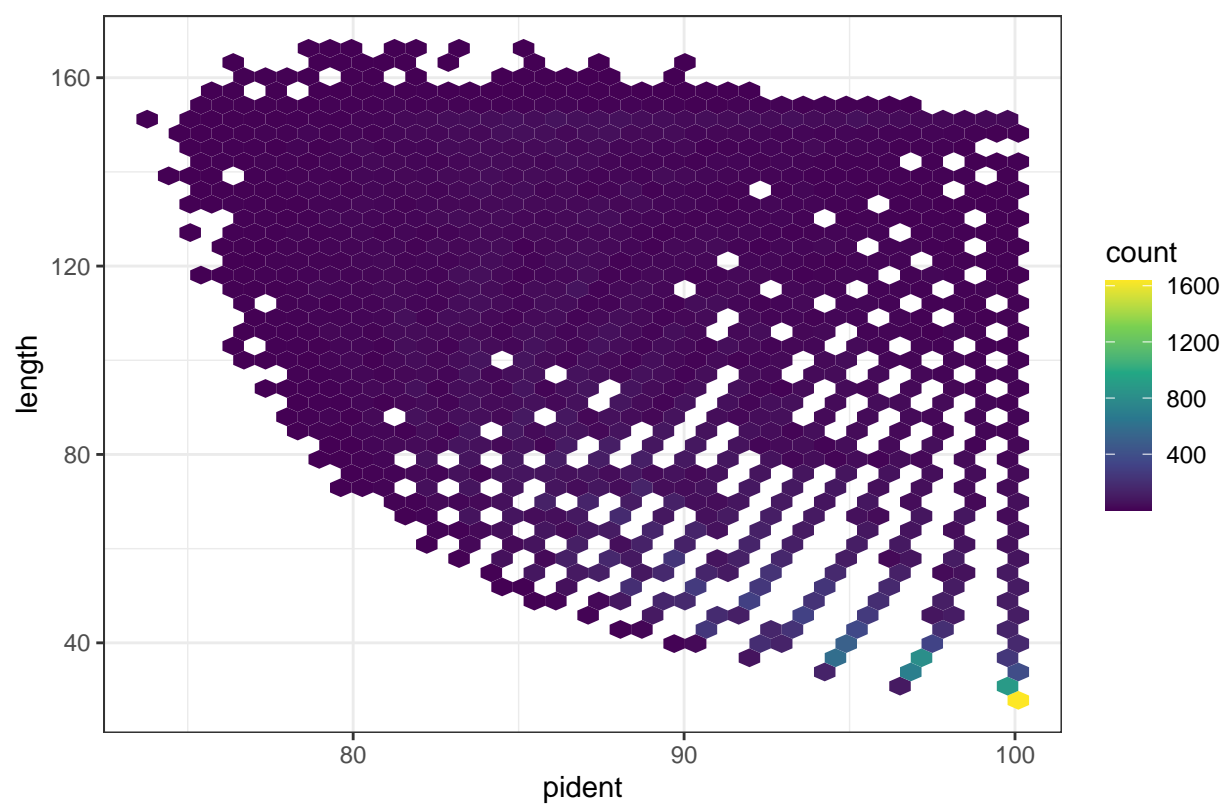
SG-112_S5_L001_R2_001

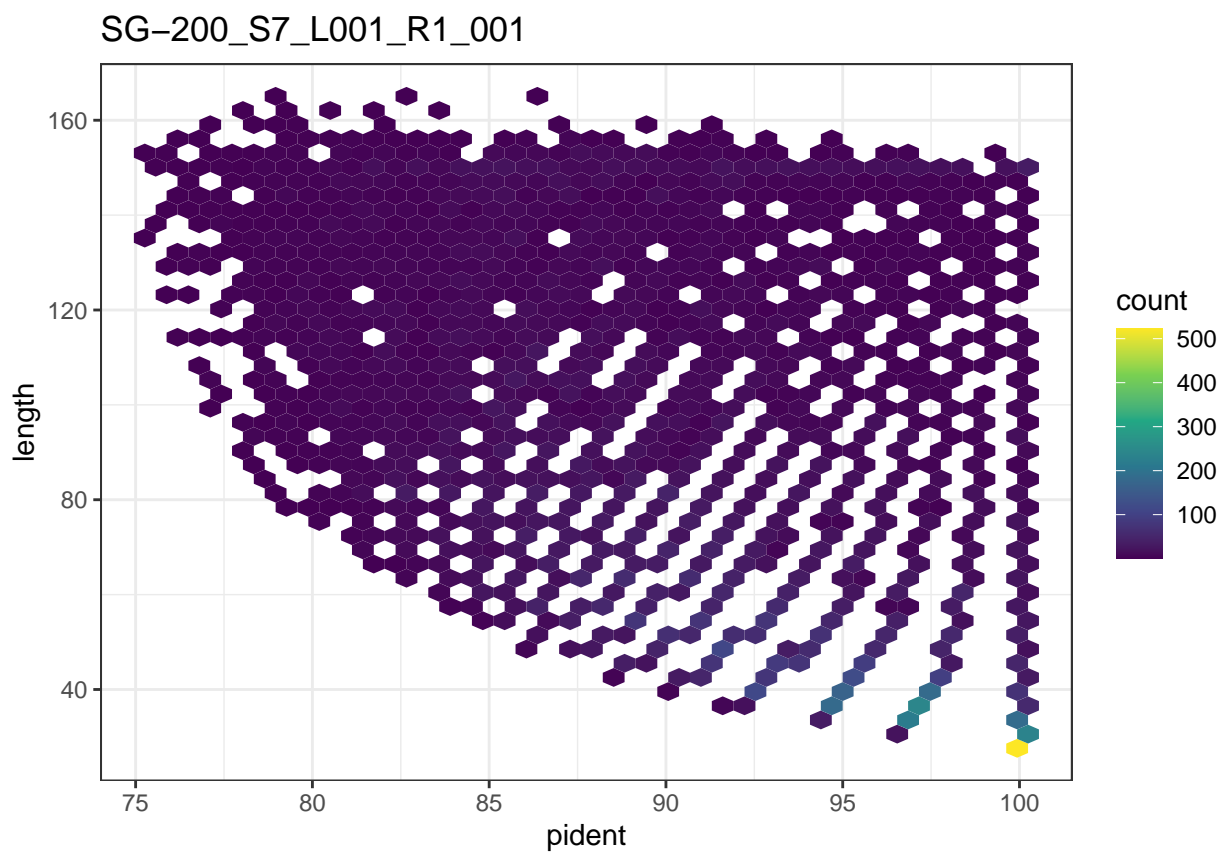


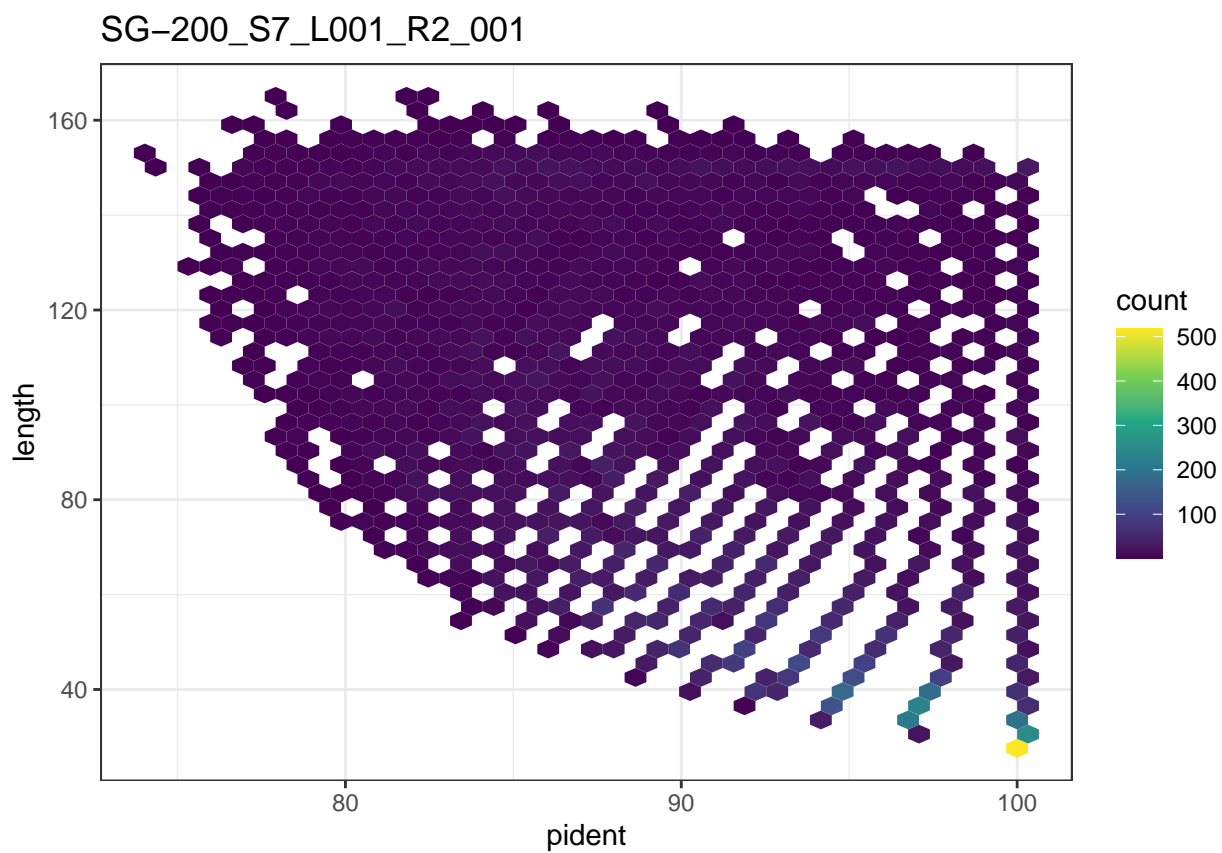
SG-193_S6_L001_R1_001



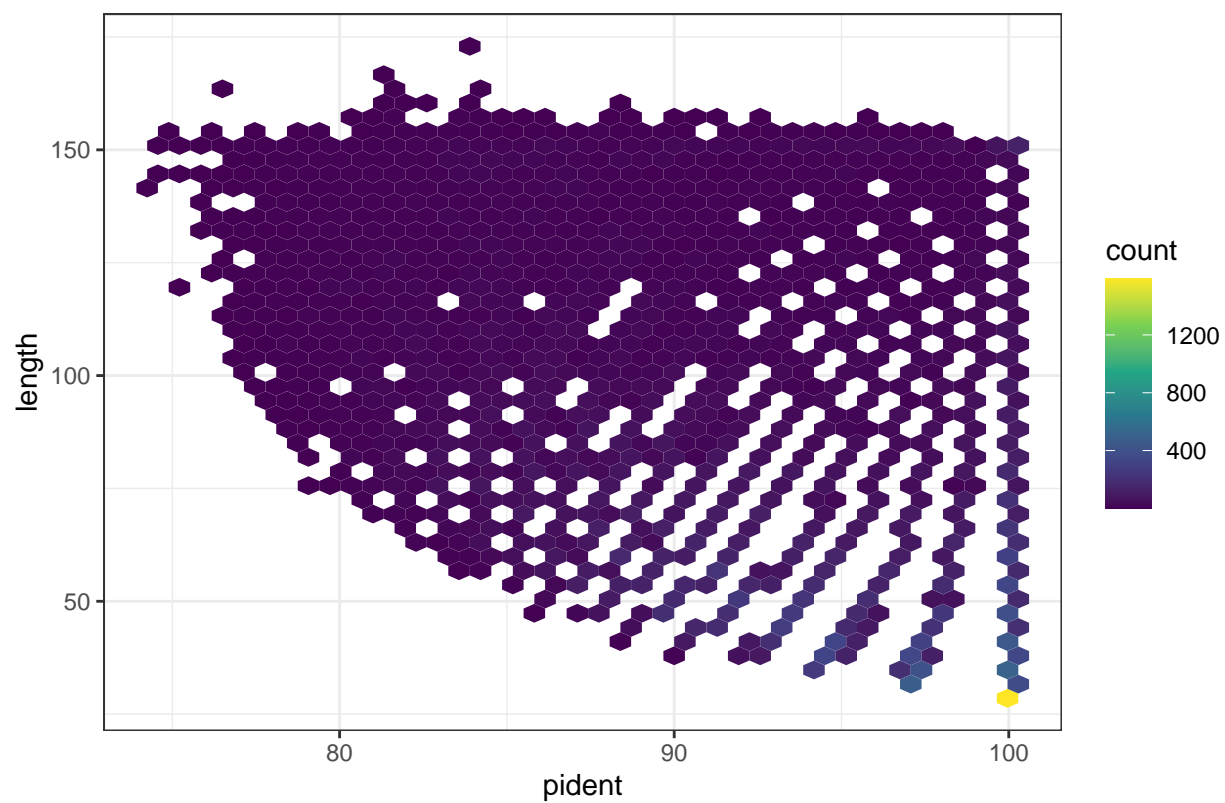
SG-193_S6_L001_R2_001



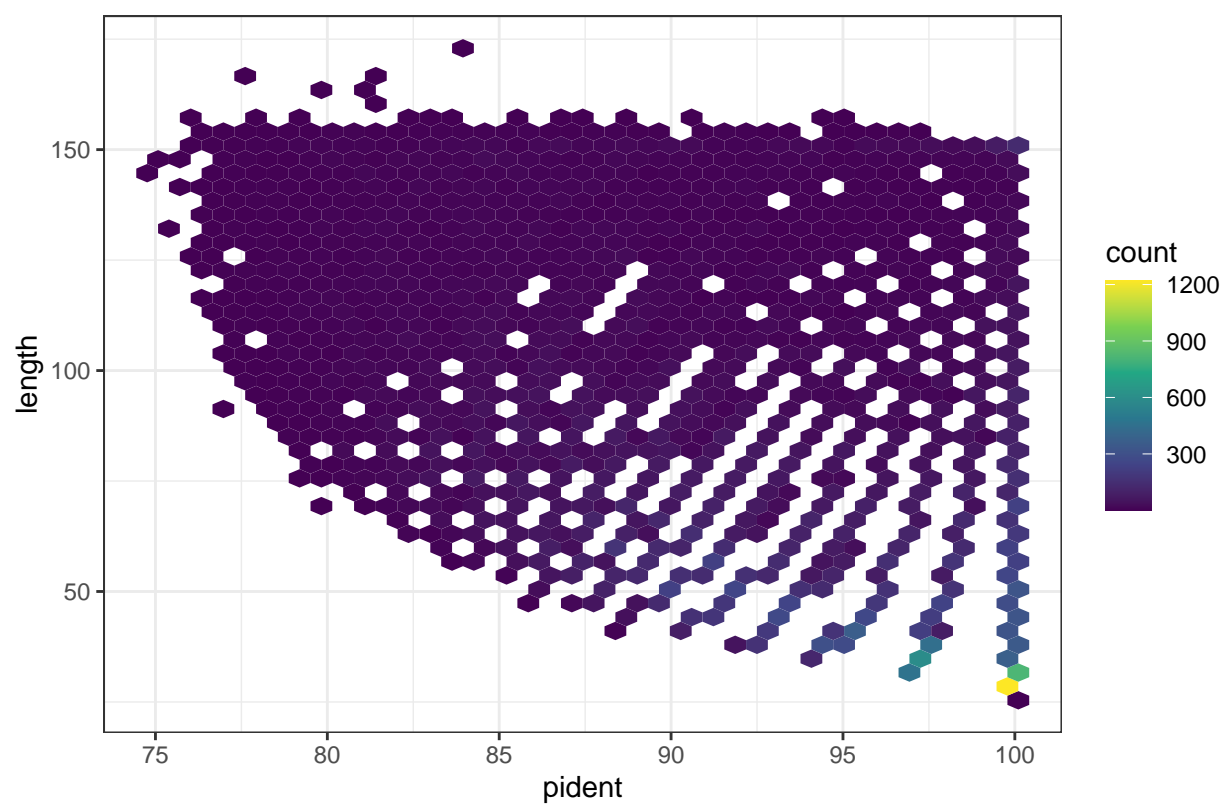


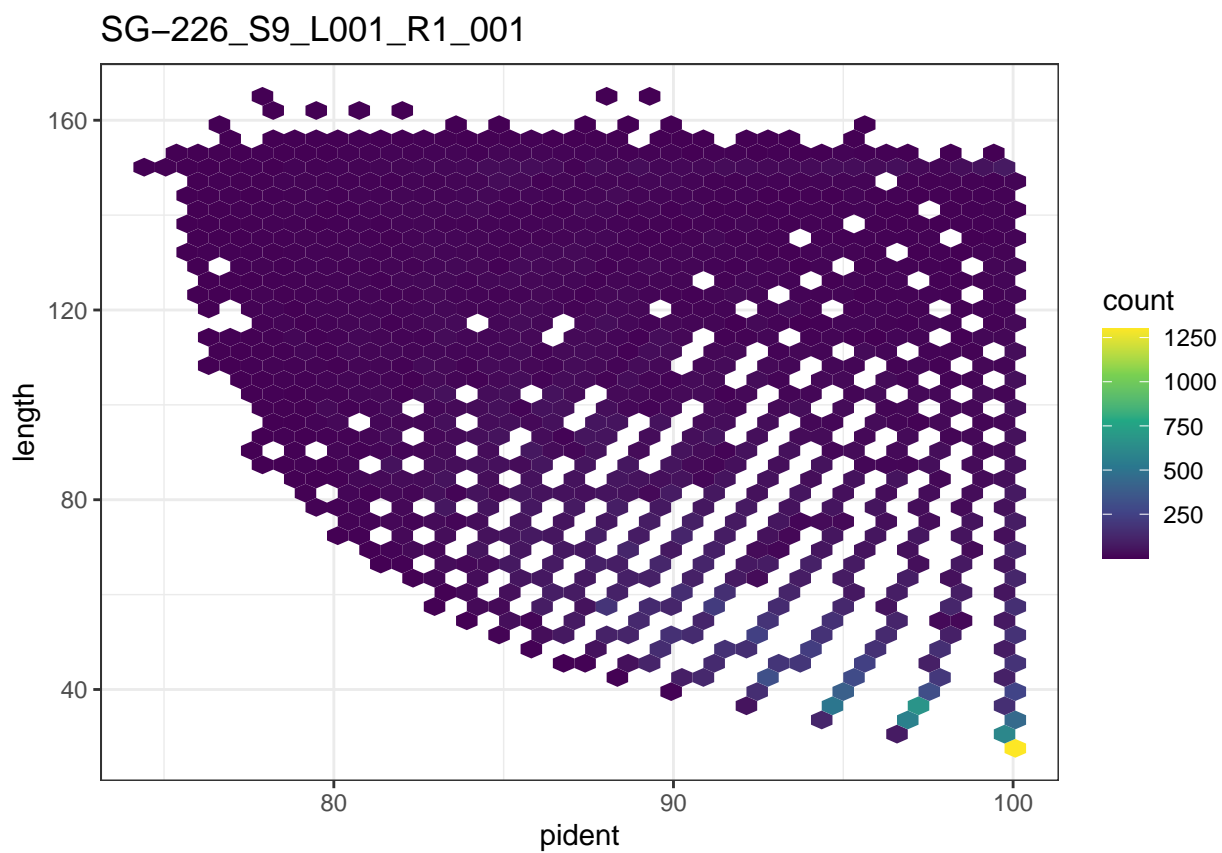


SG-224_S8_L001_R1_001

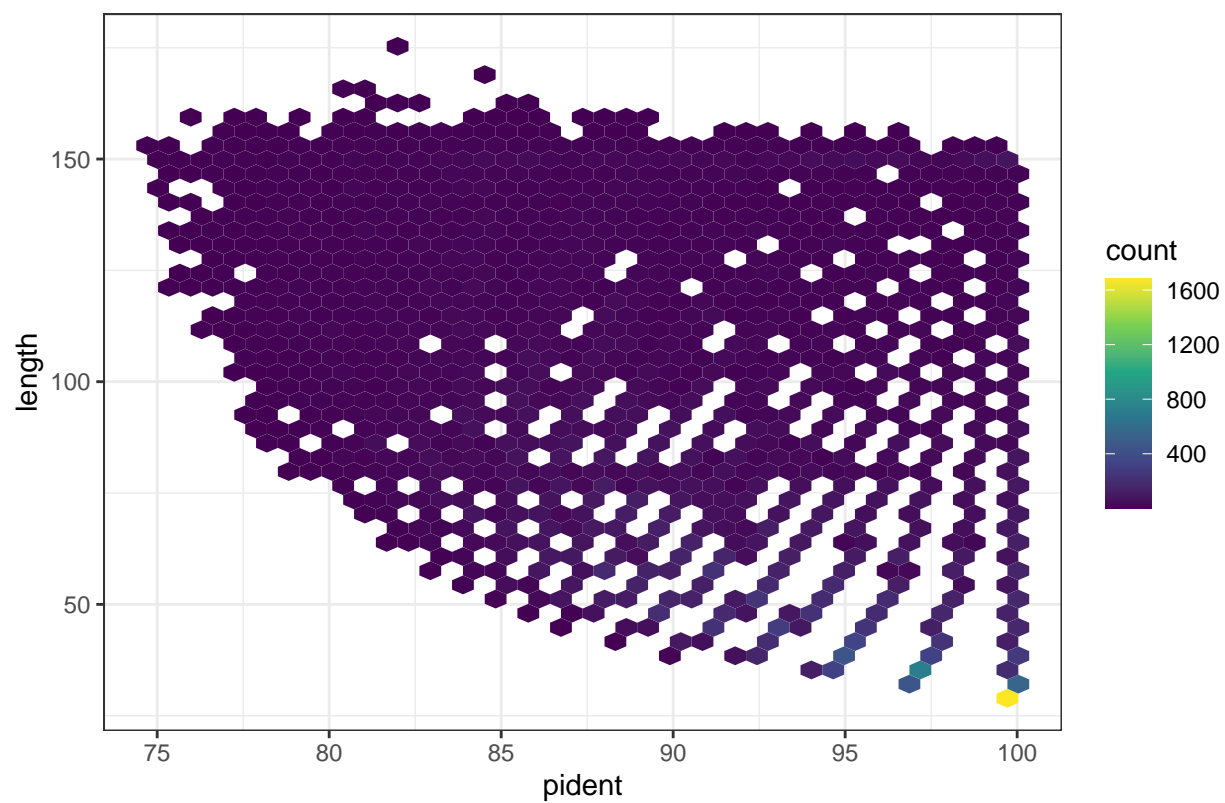


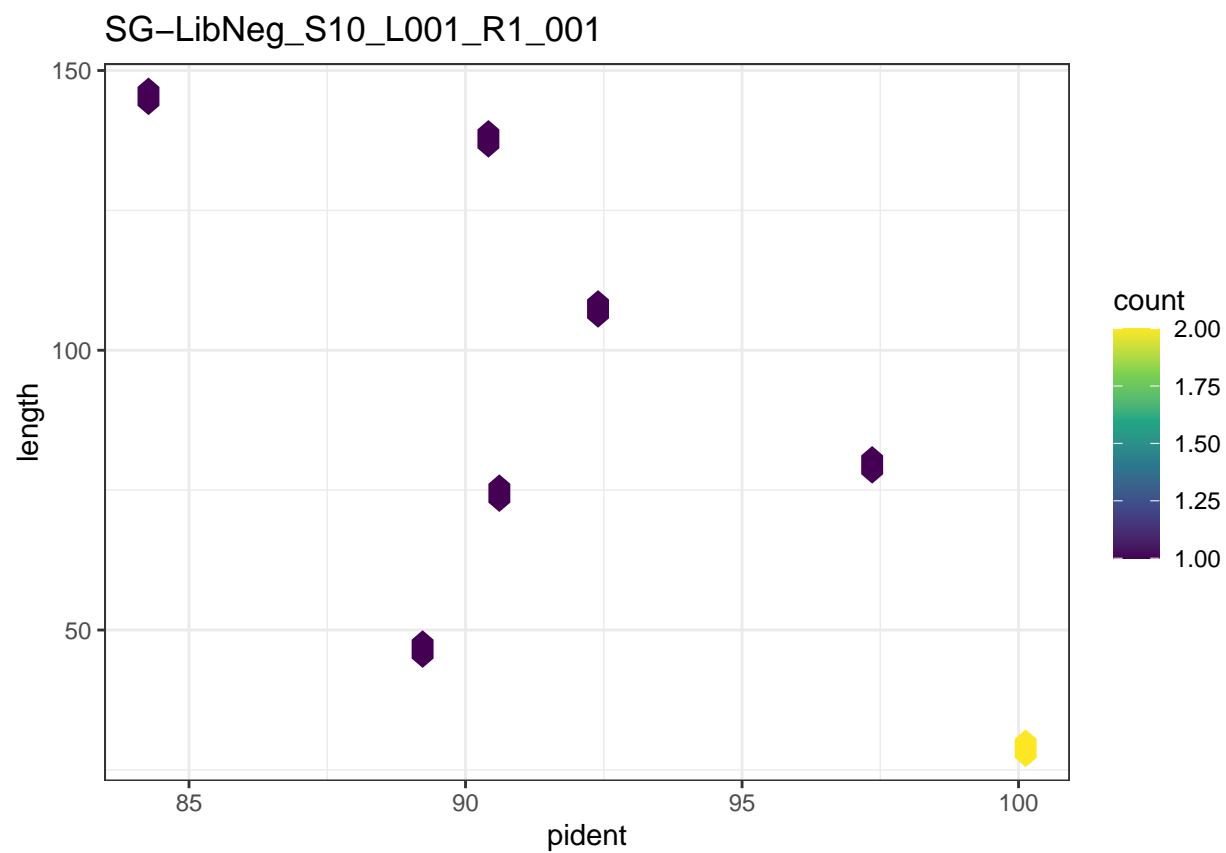
SG-224_S8_L001_R2_001



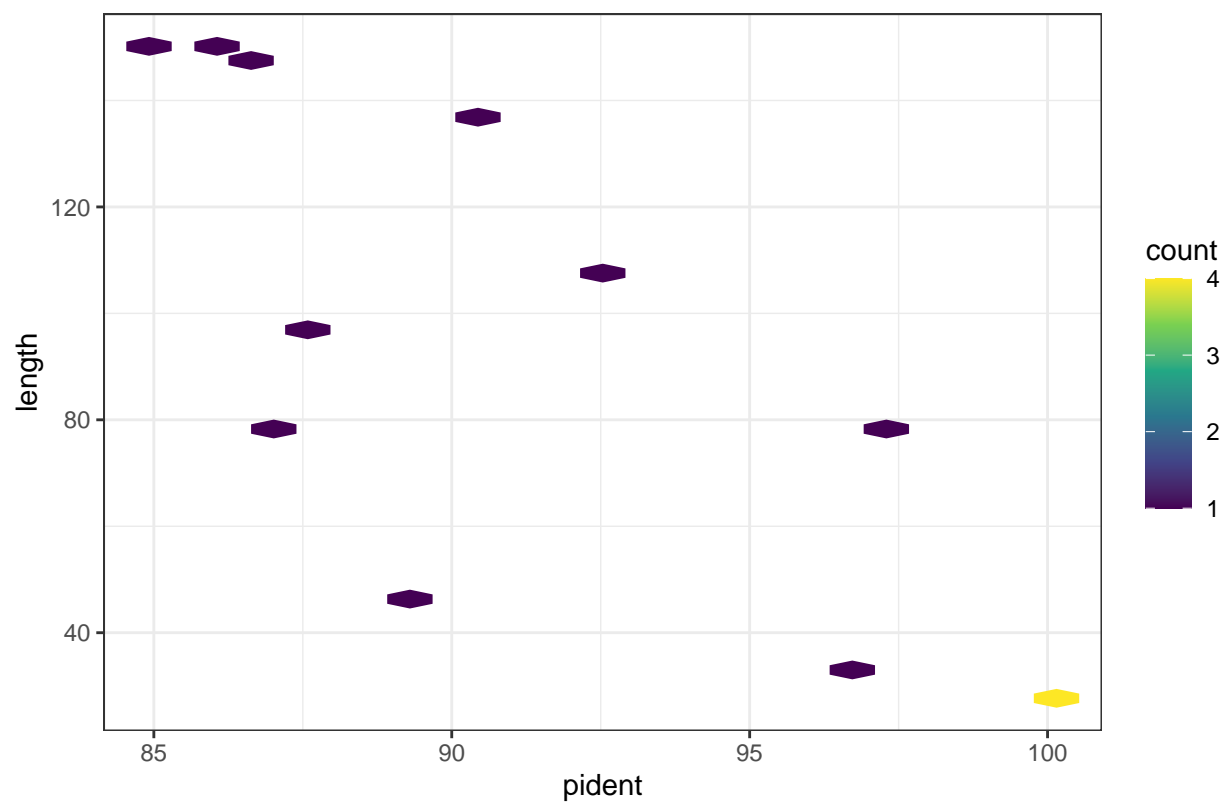


SG-226_S9_L001_R2_001

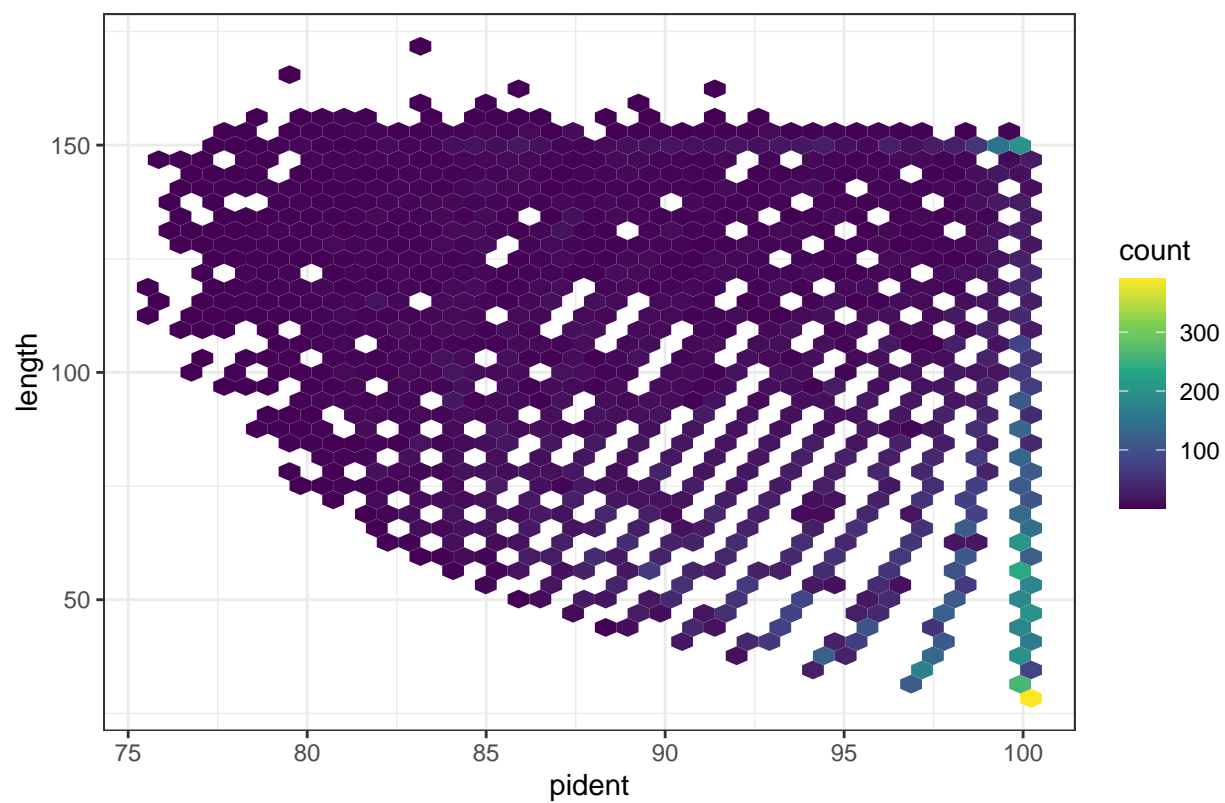


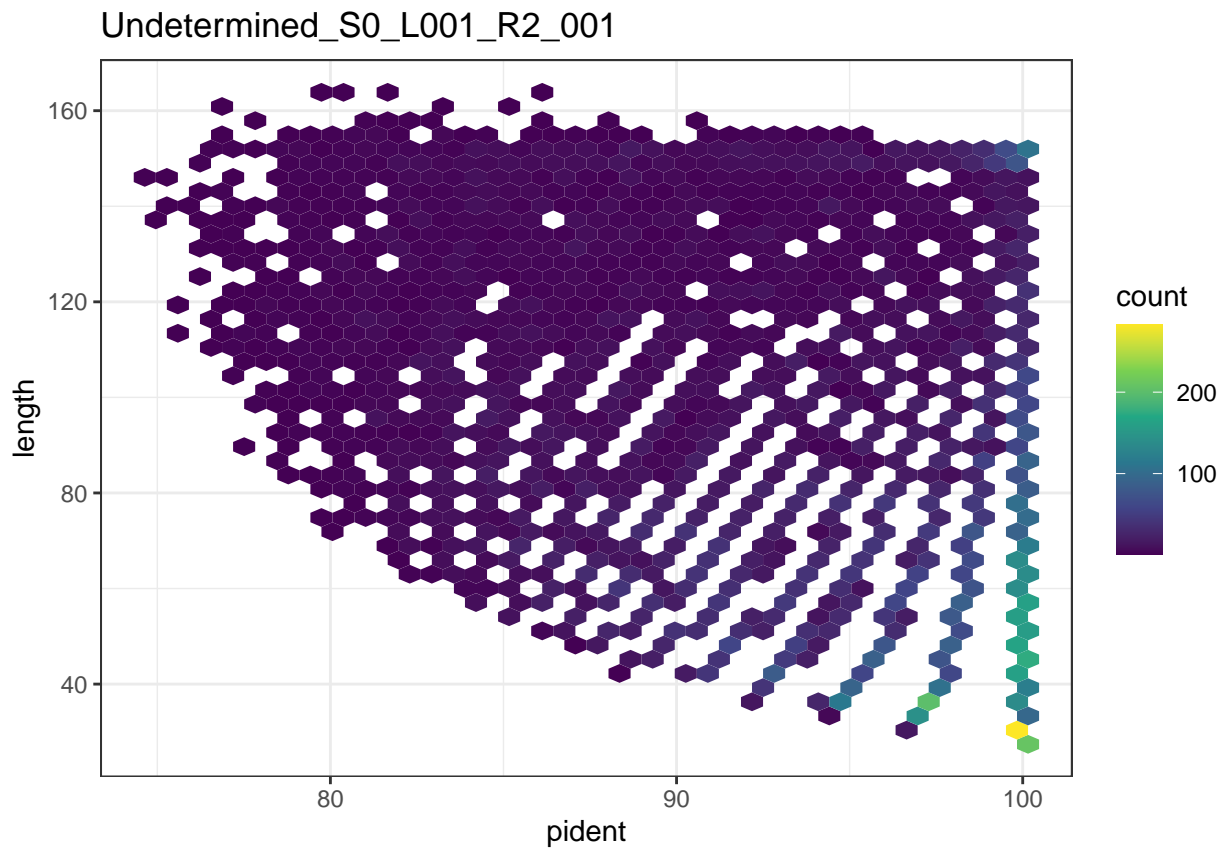


SG-LibNeg_S10_L001_R2_001

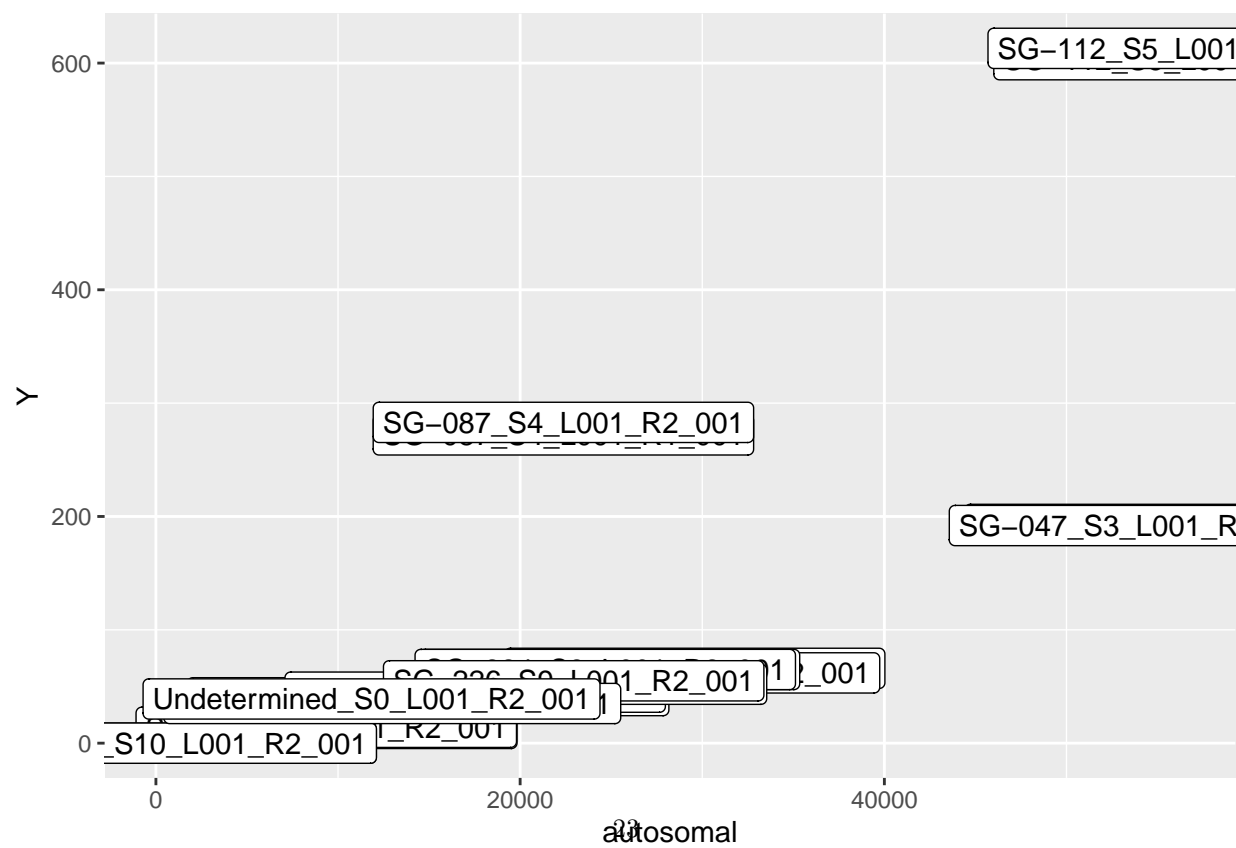
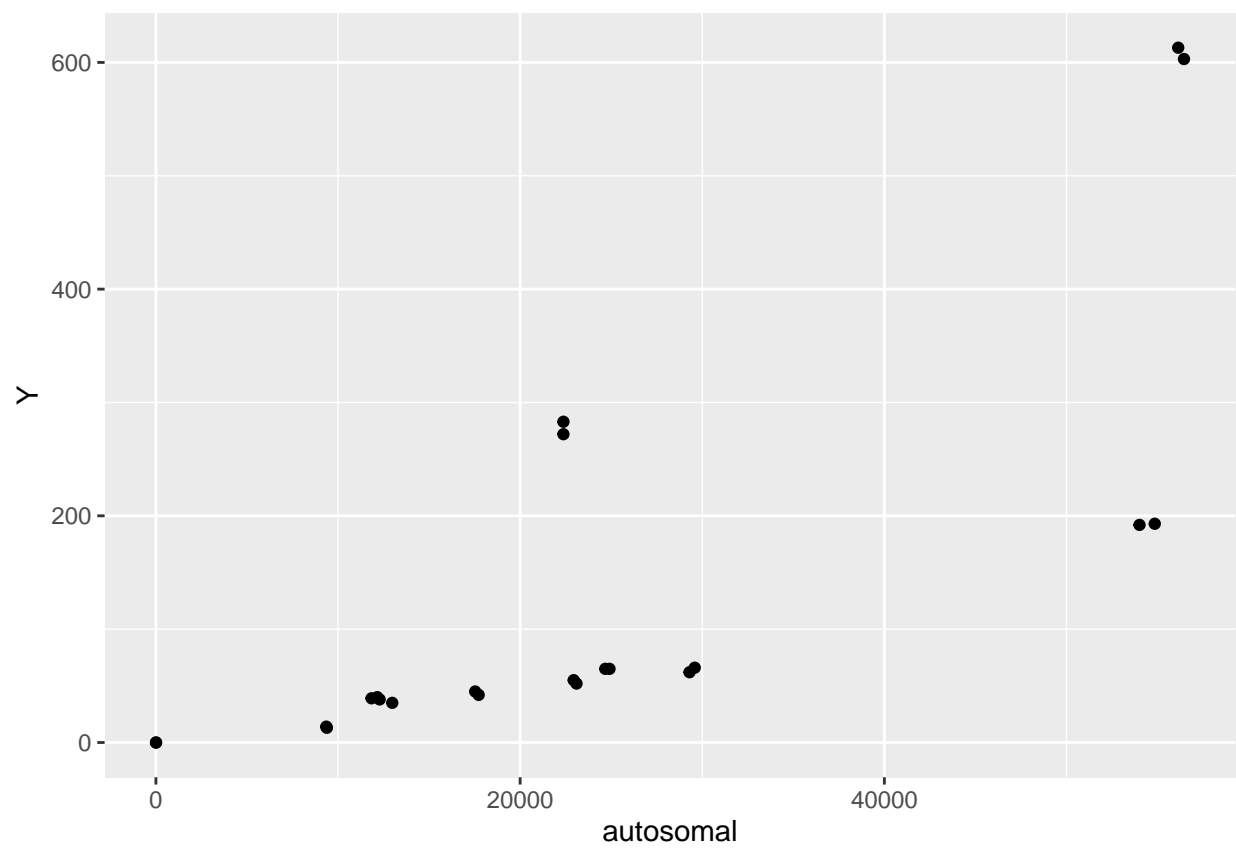


Undetermined_S0_L001_R1_001



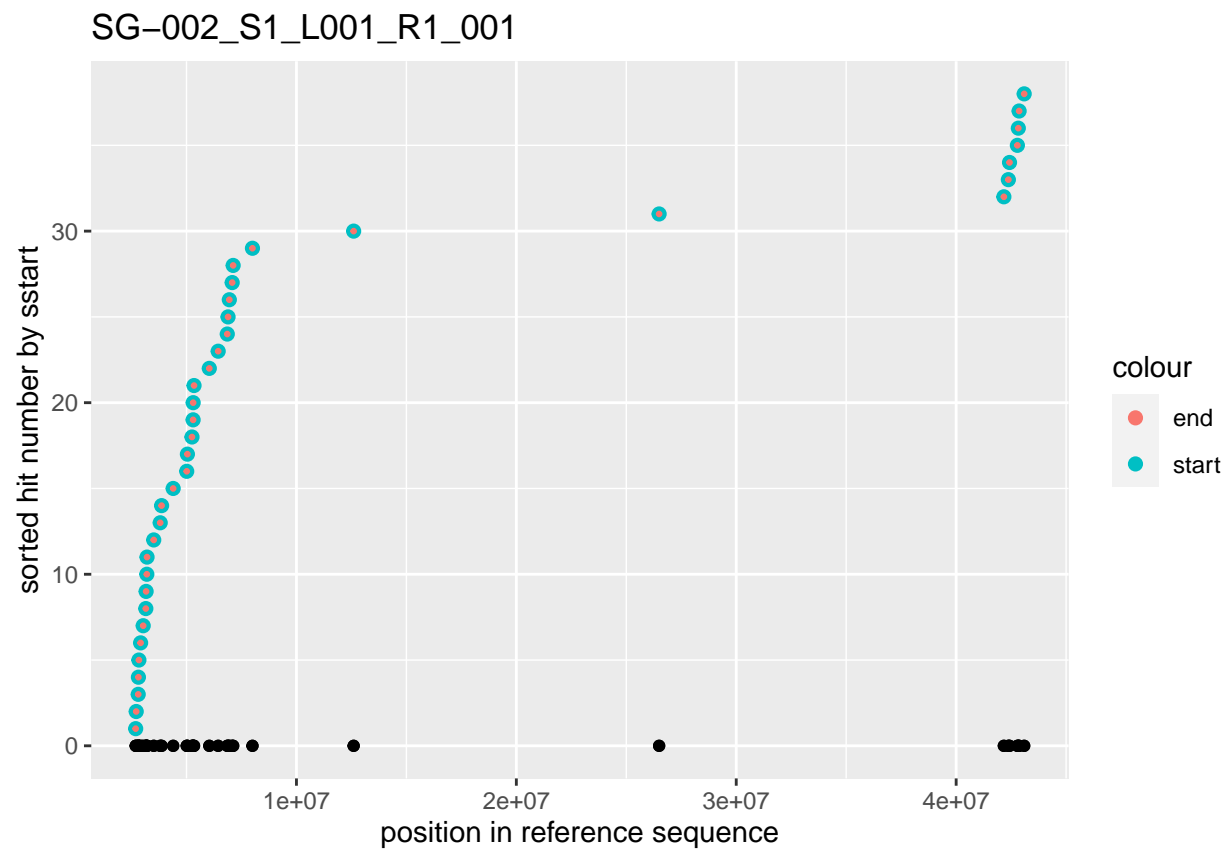


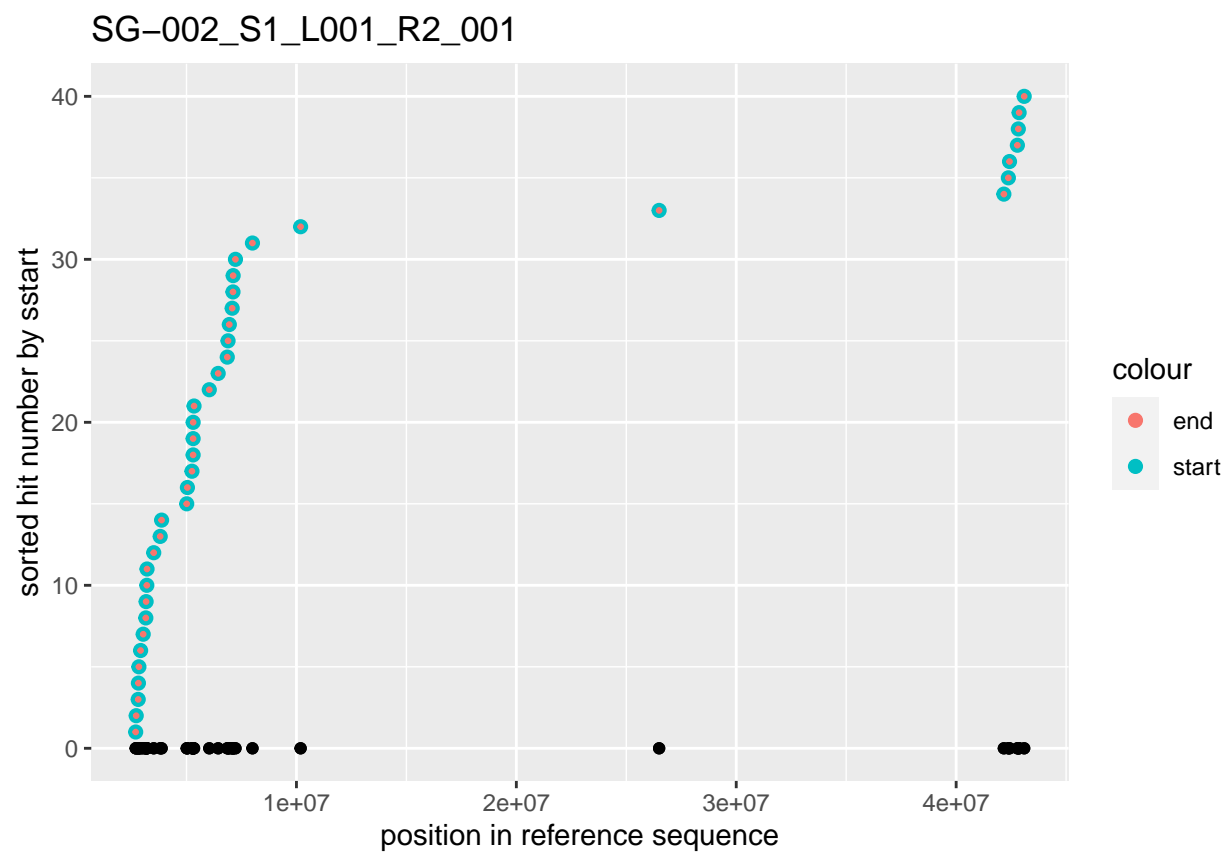
Number of autosomal chromosome hits V.S. number of Y chromosome hits

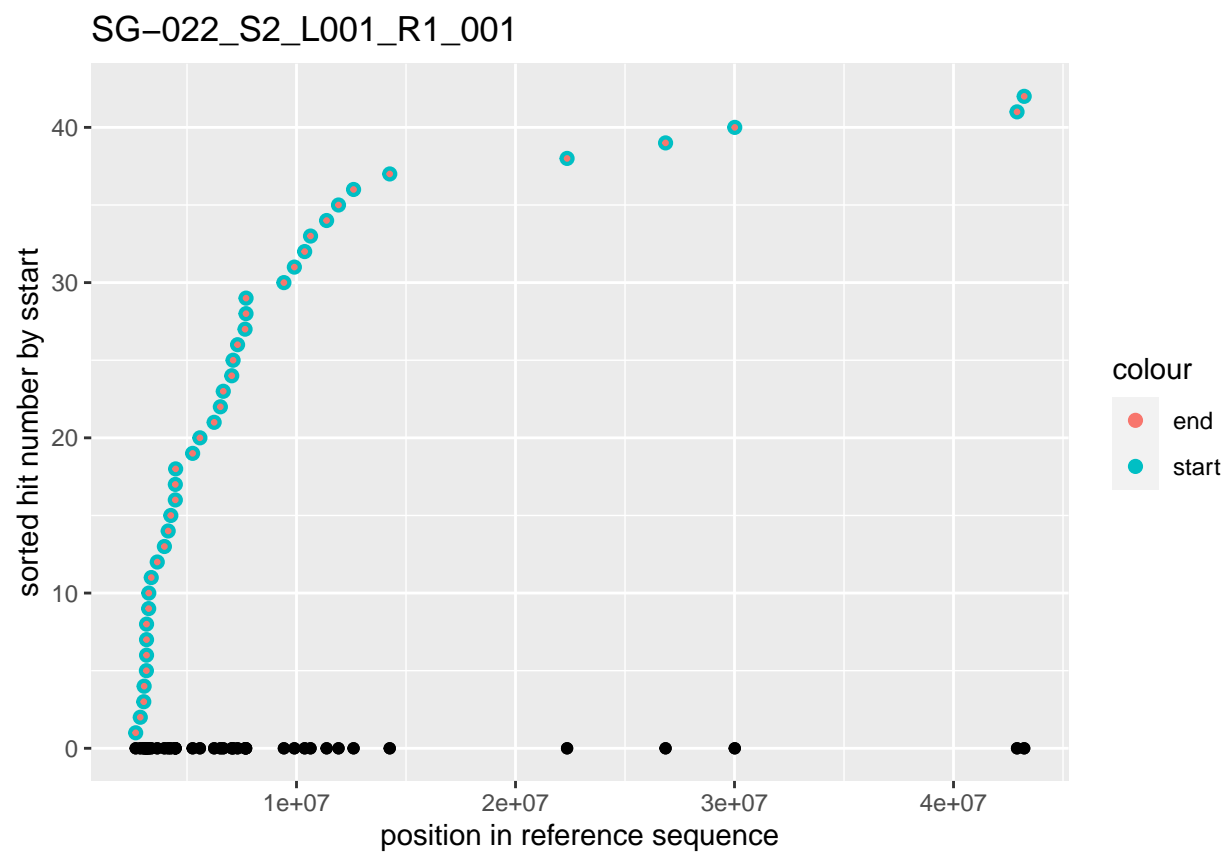


map of hits in Y chromosome reference sequence

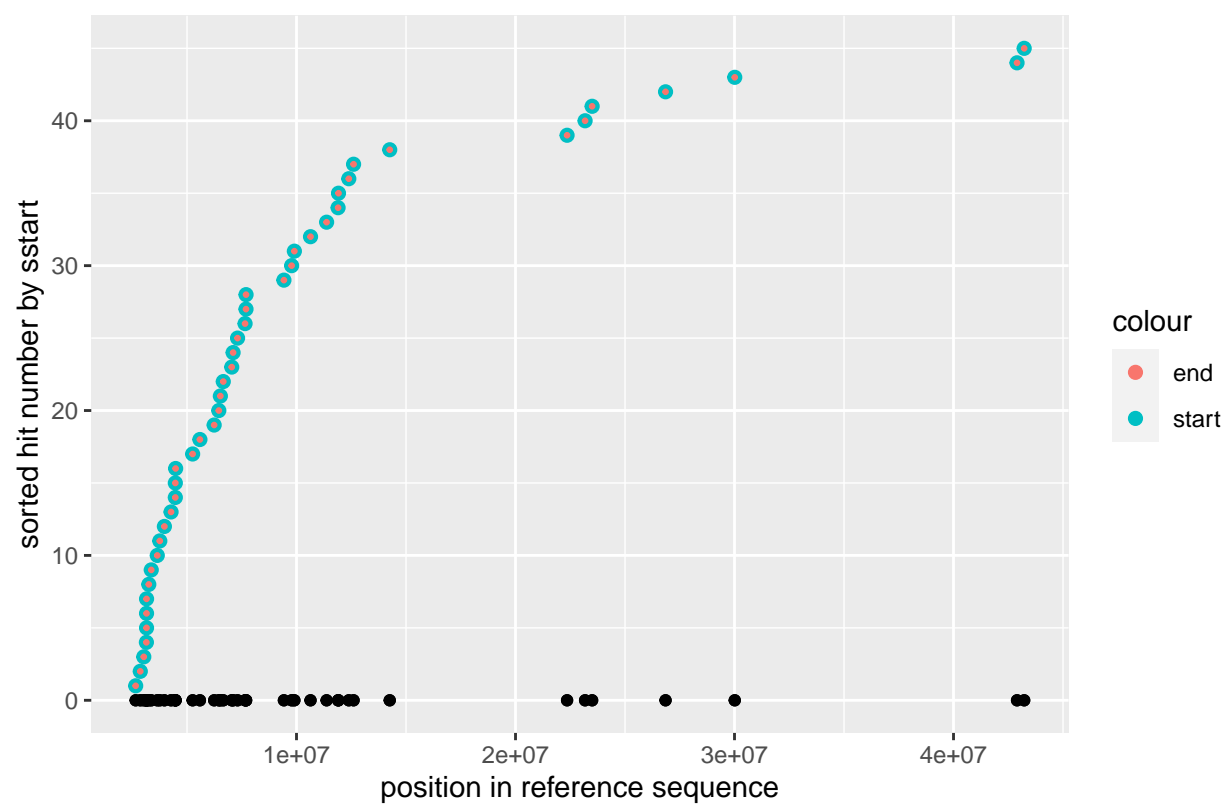
x dots are projections of the curve on x axis

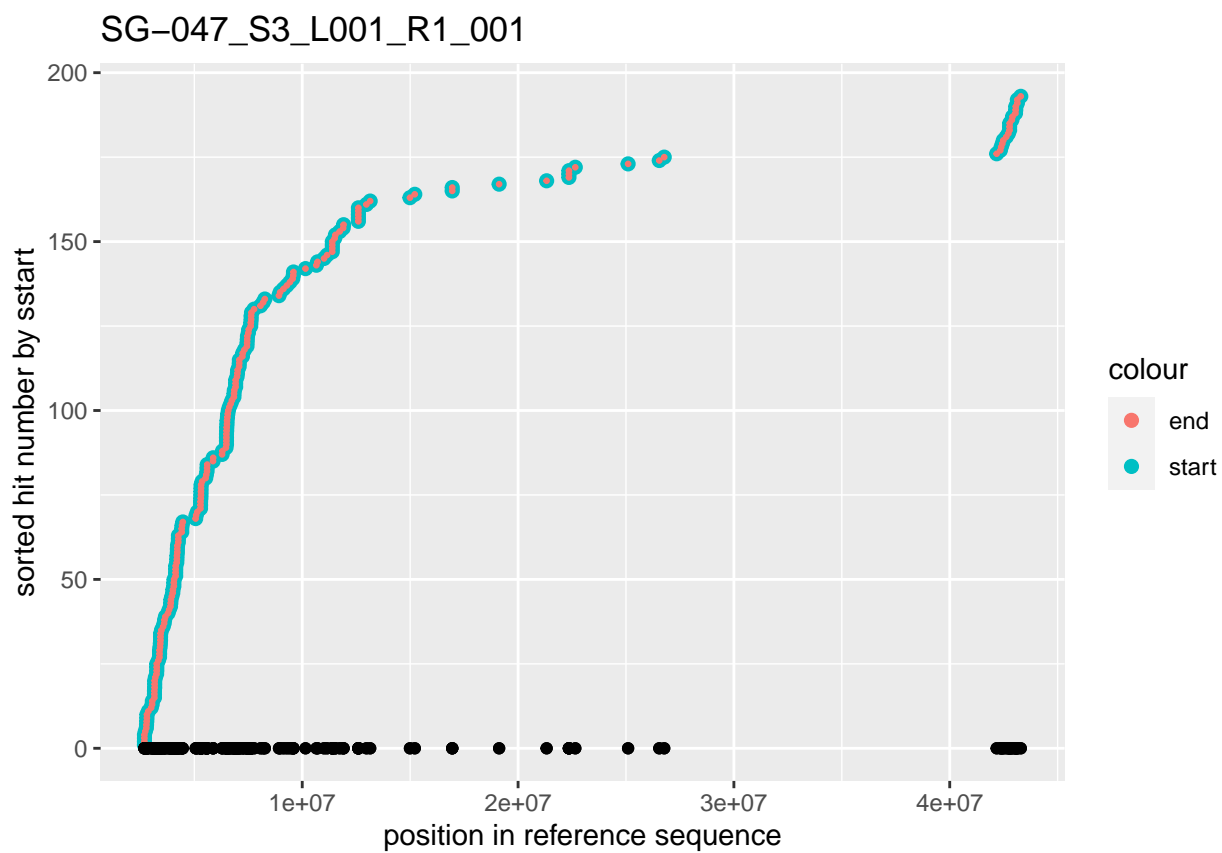


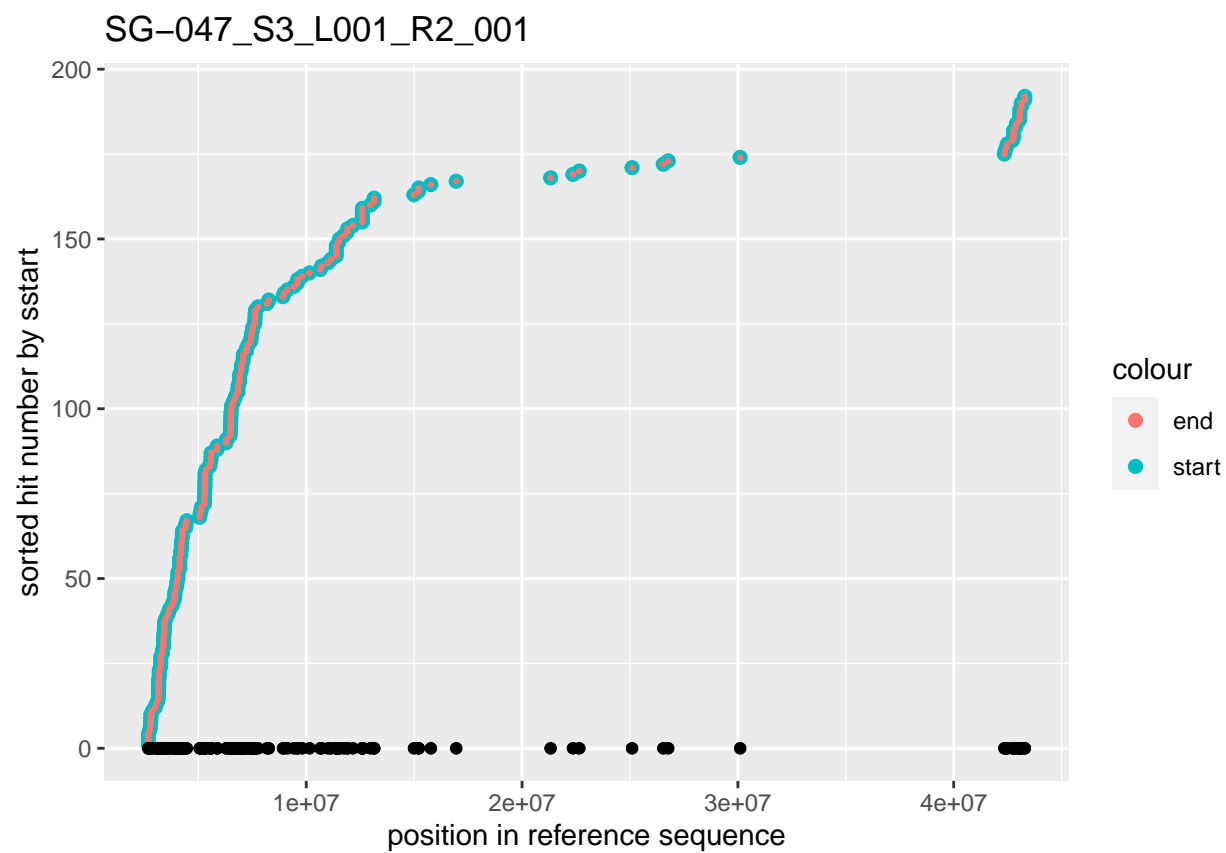




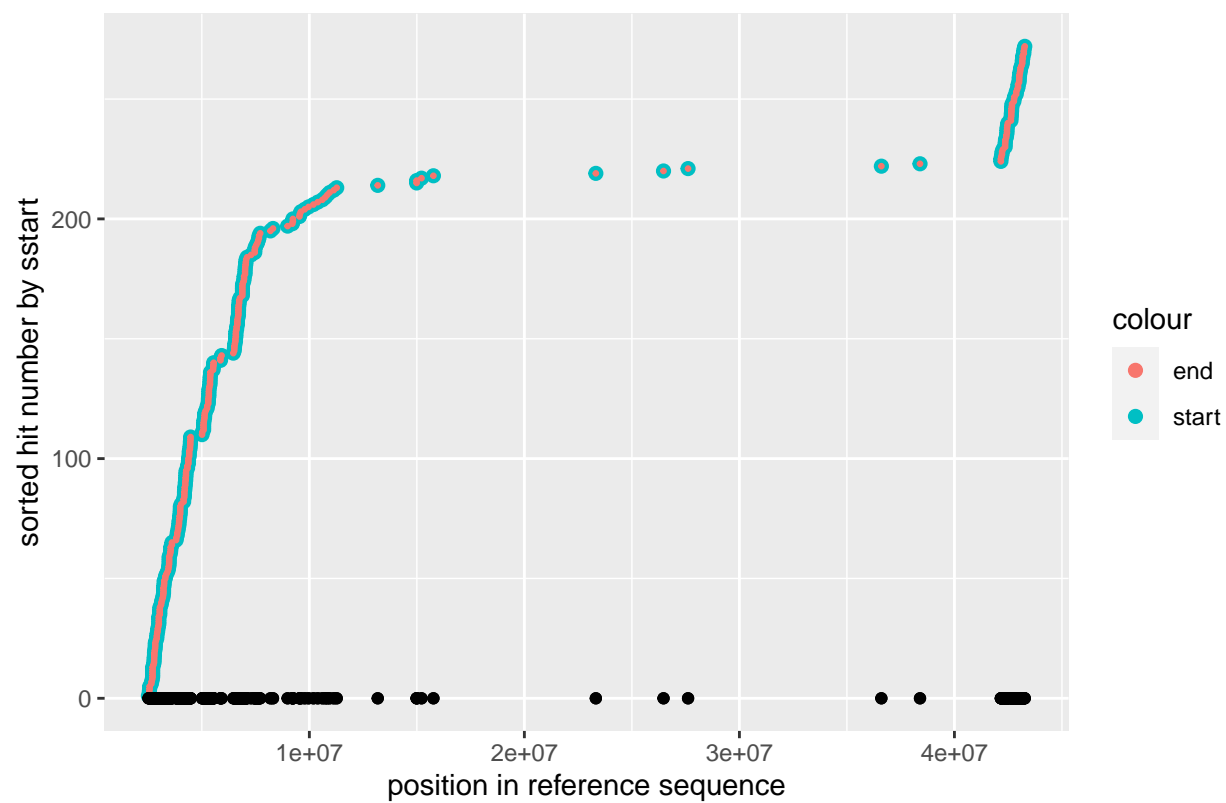
SG-022_S2_L001_R2_001



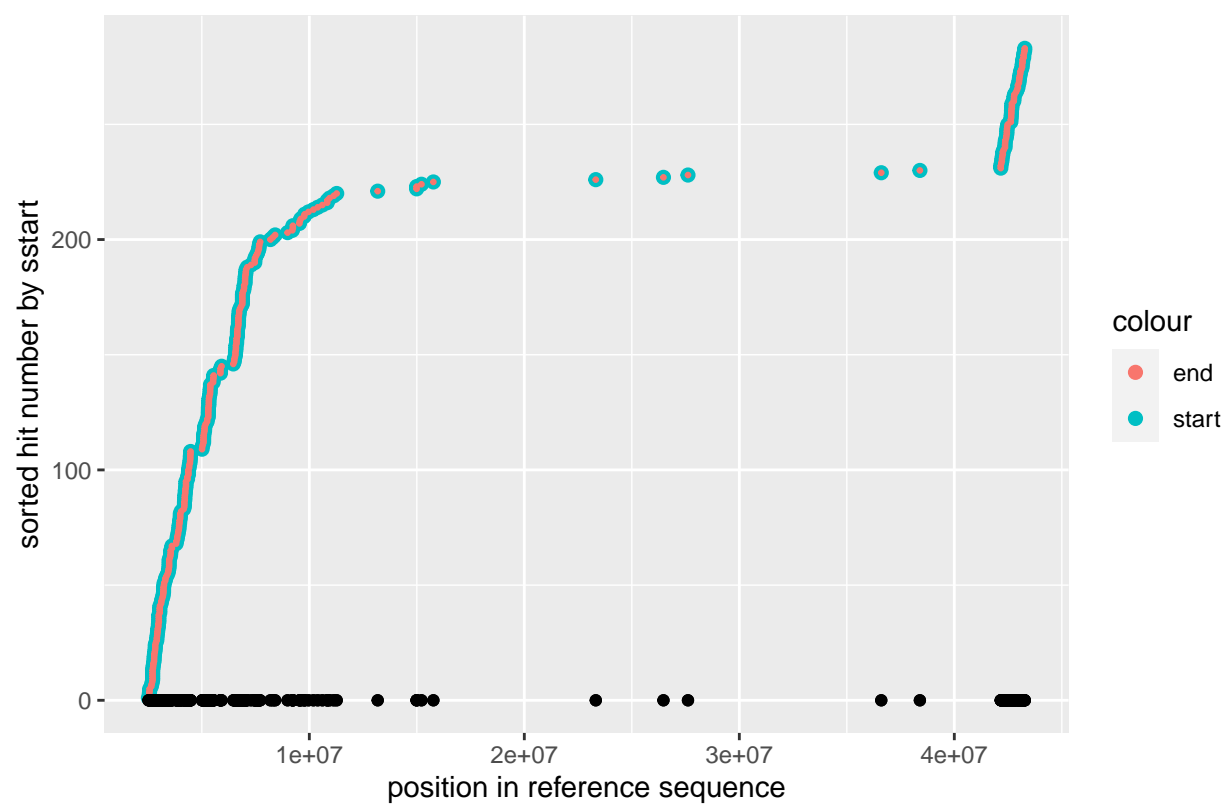


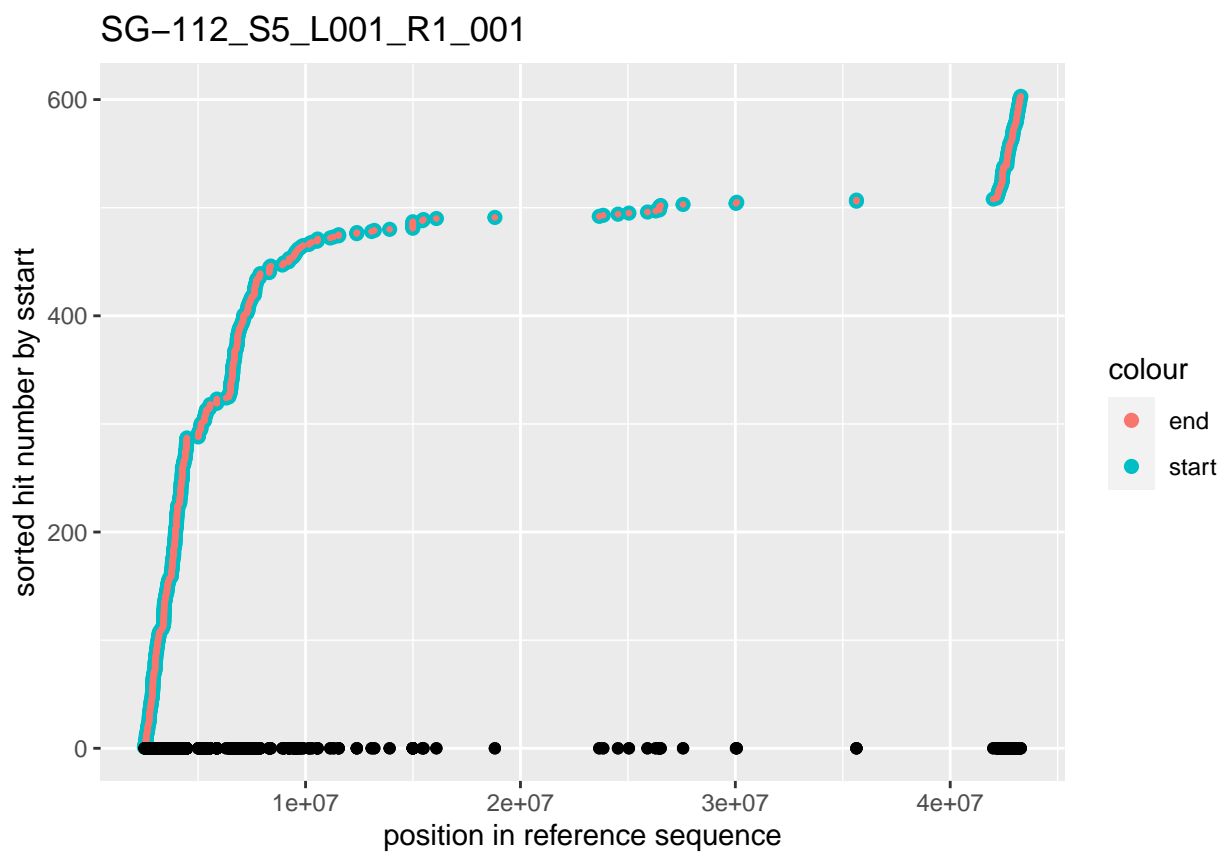


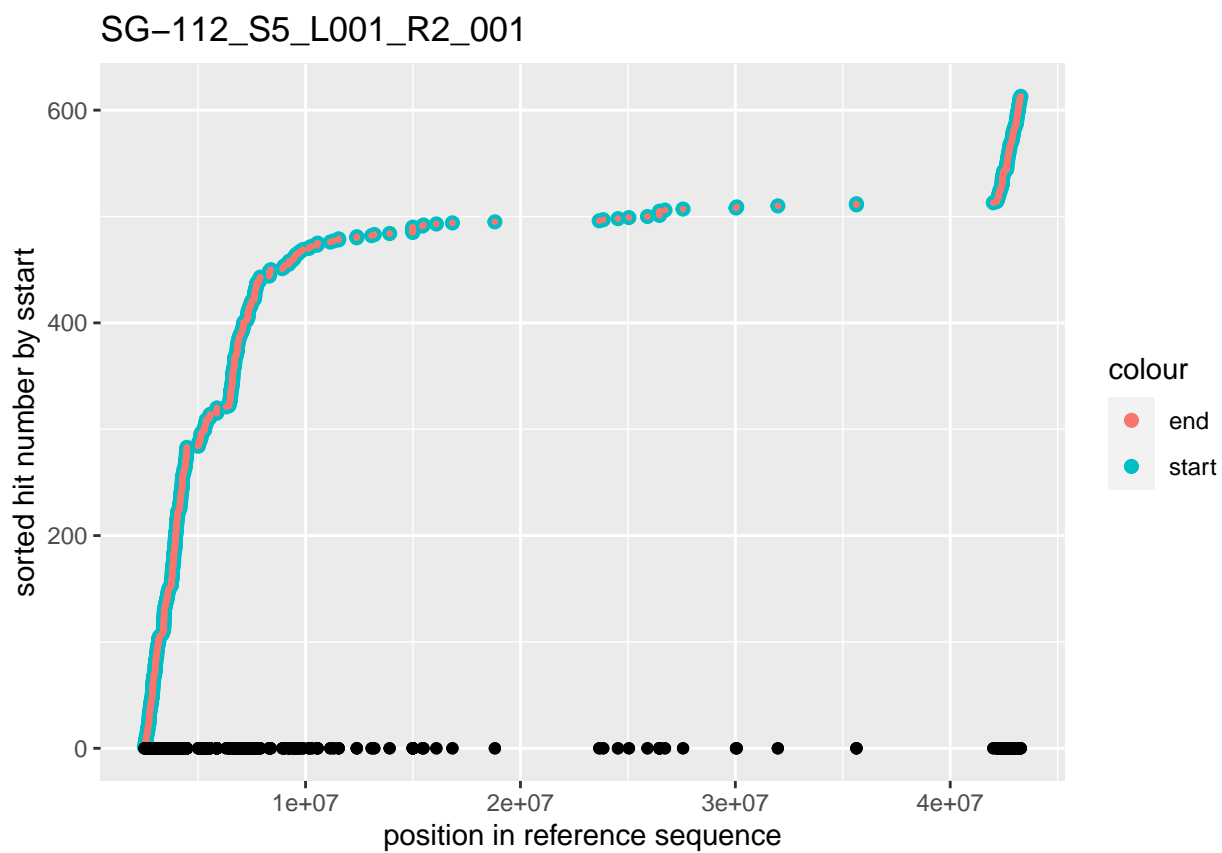
SG-087_S4_L001_R1_001

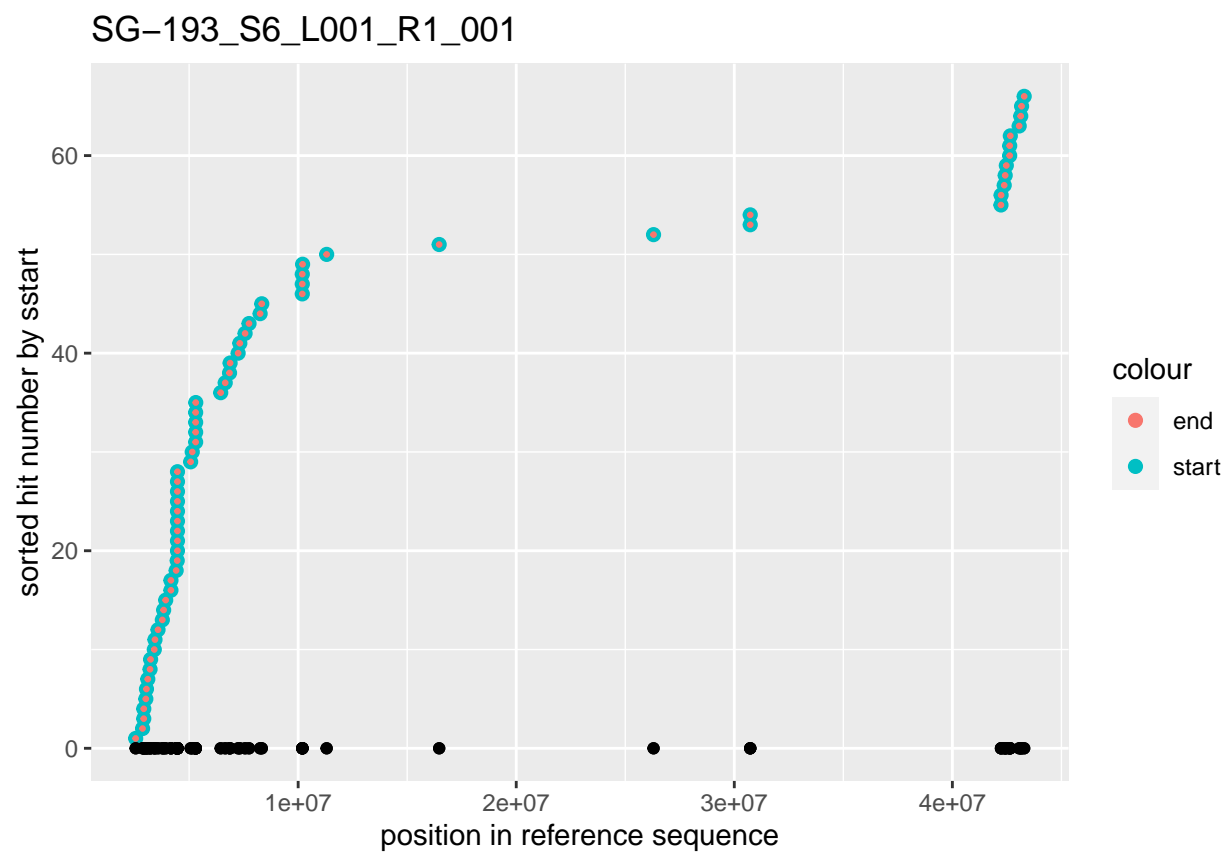


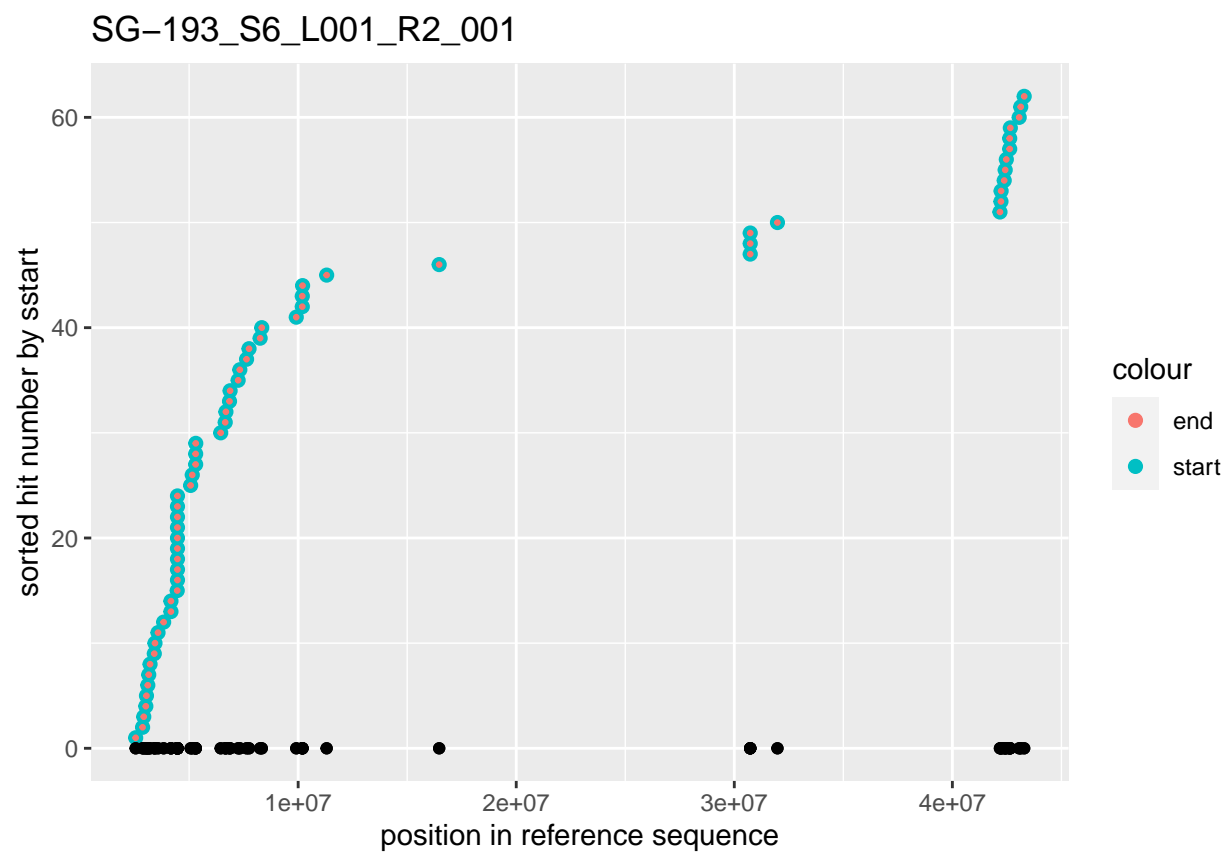
SG-087_S4_L001_R2_001



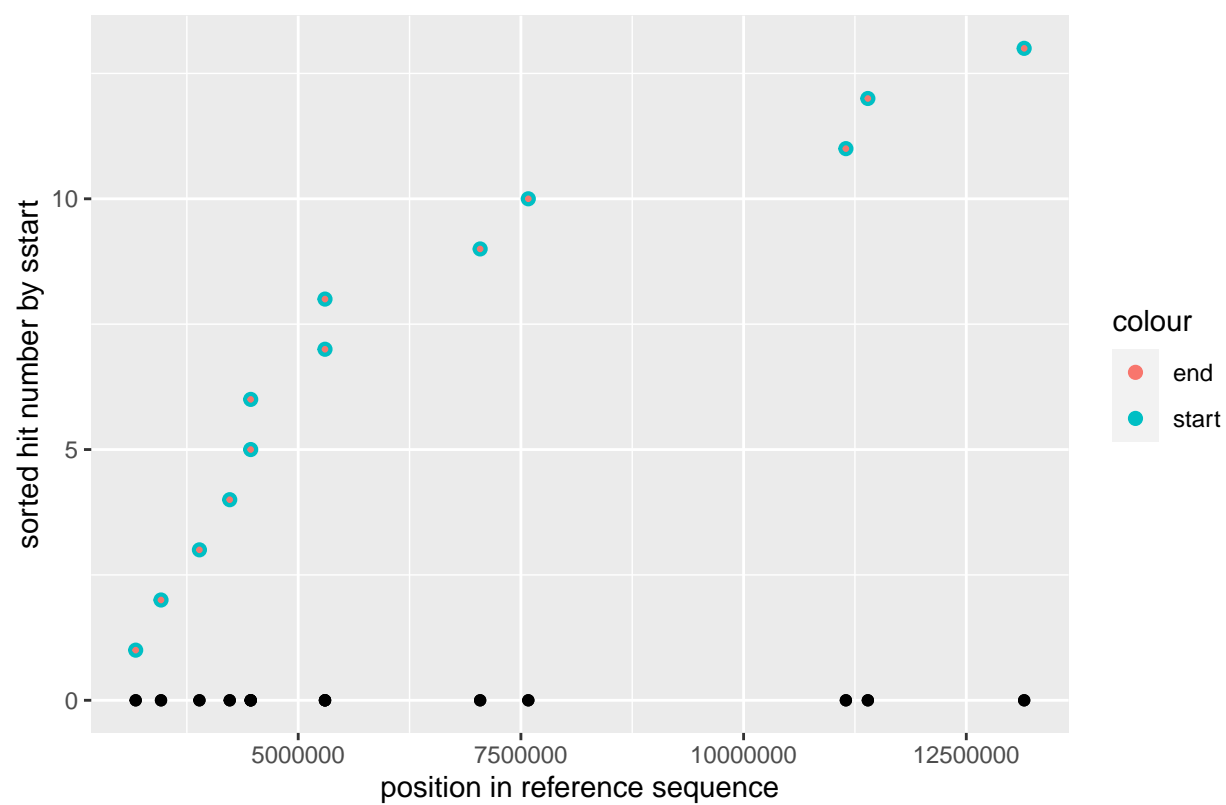


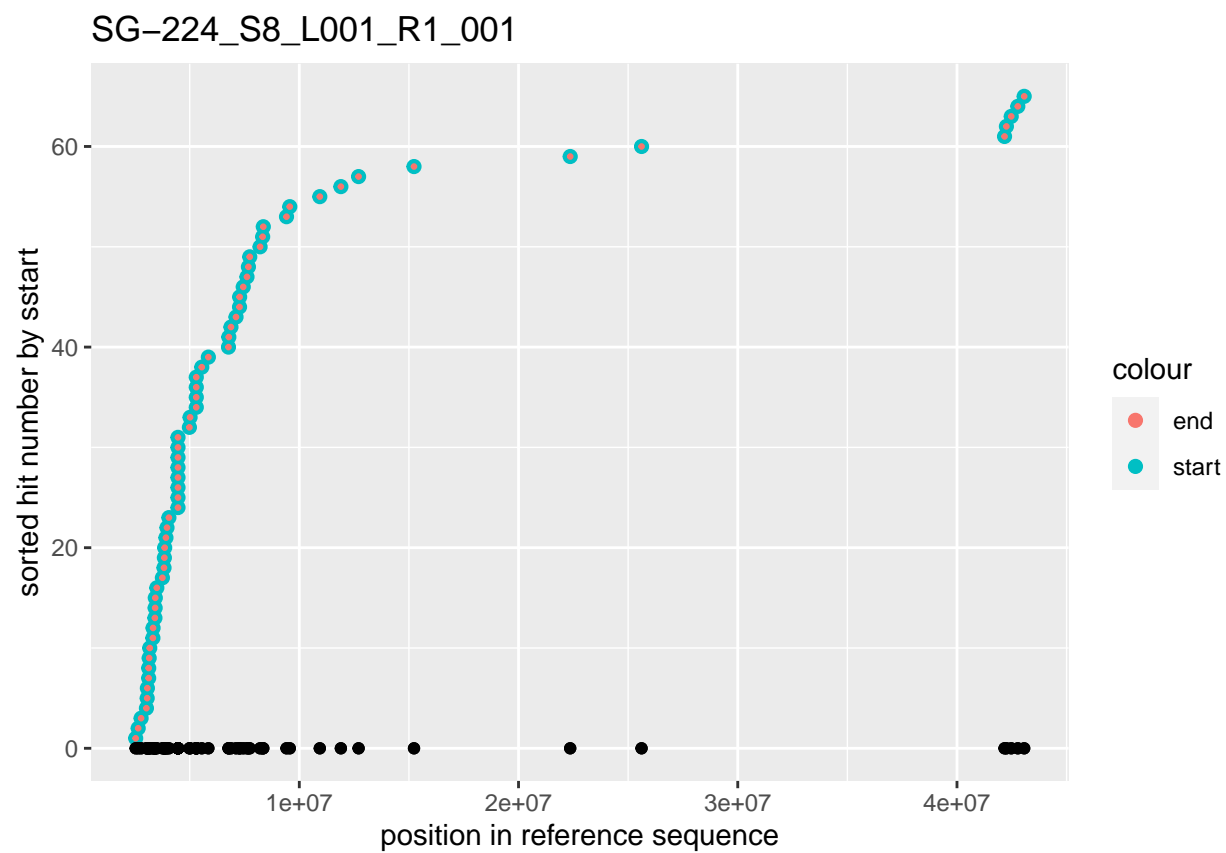


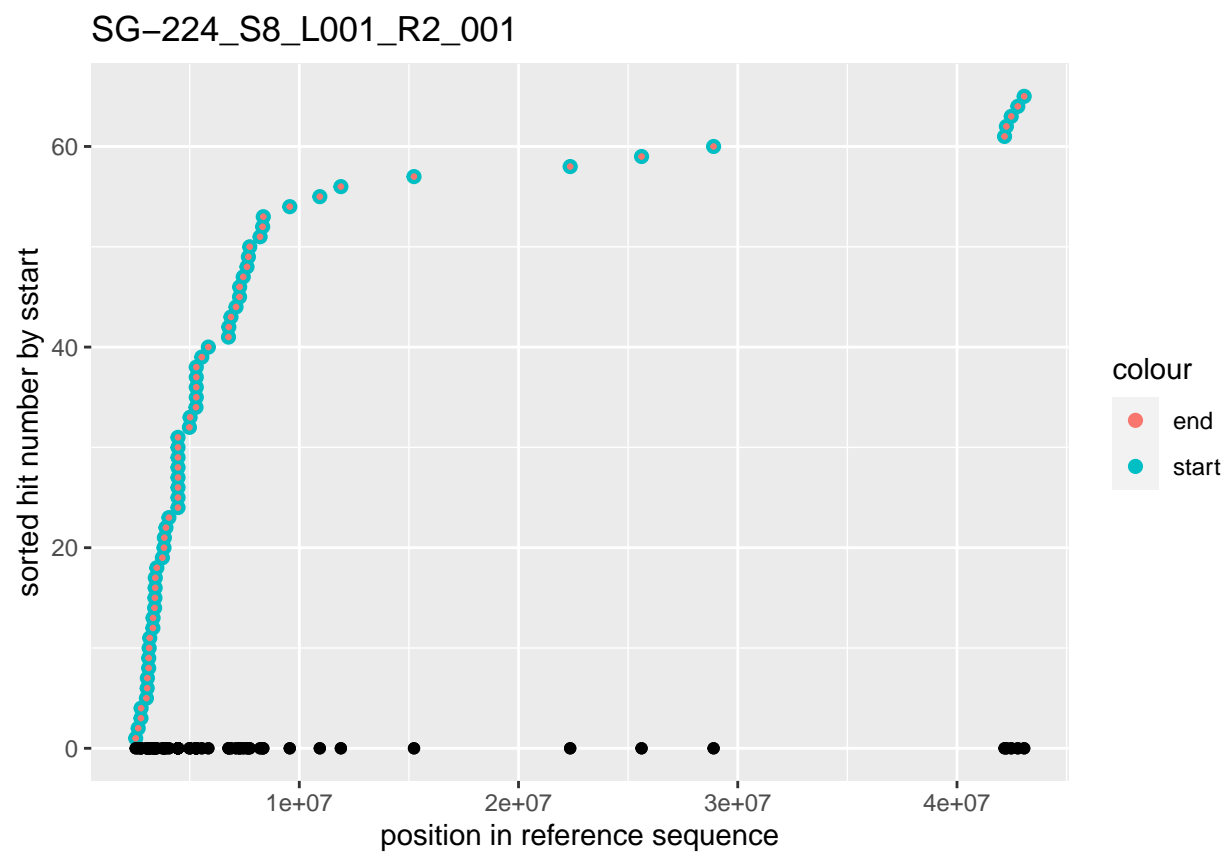


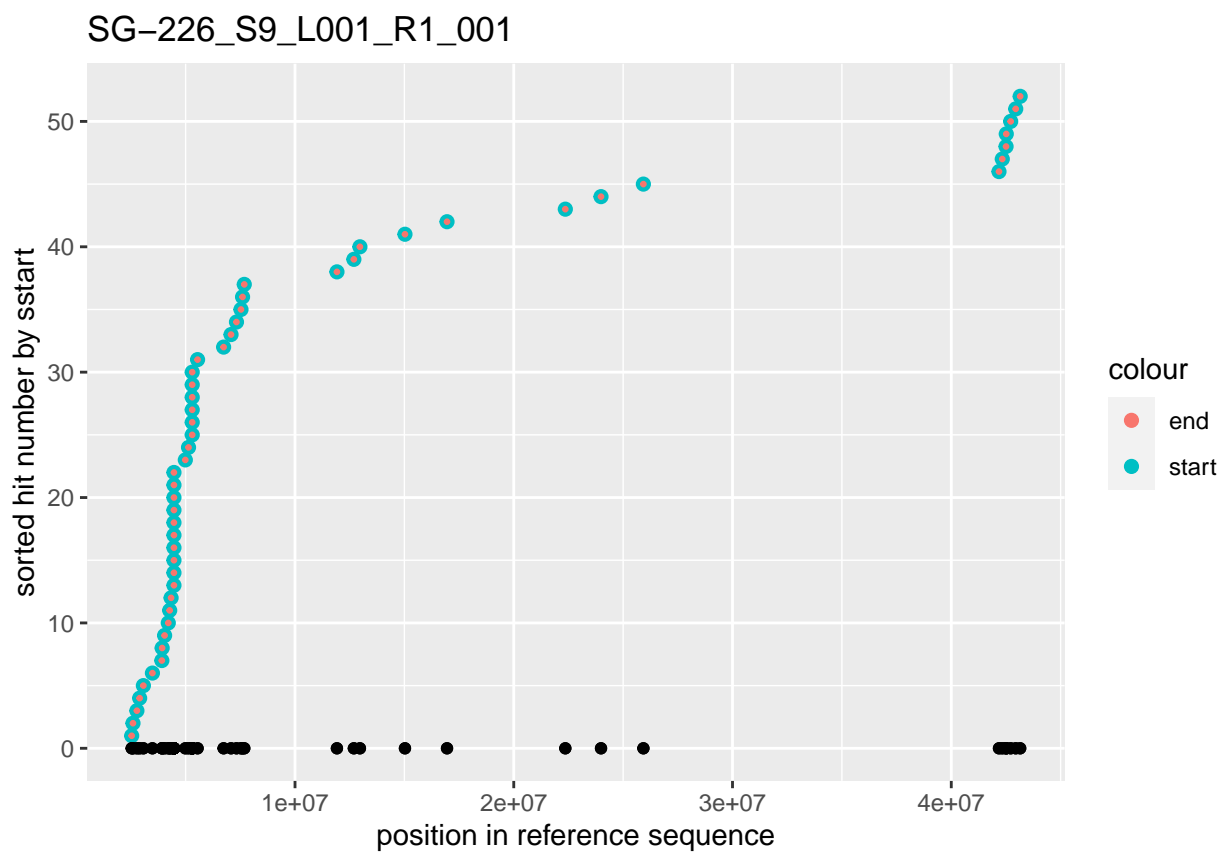


SG-200_S7_L001_R1_001









SG-226_S9_L001_R2_001

