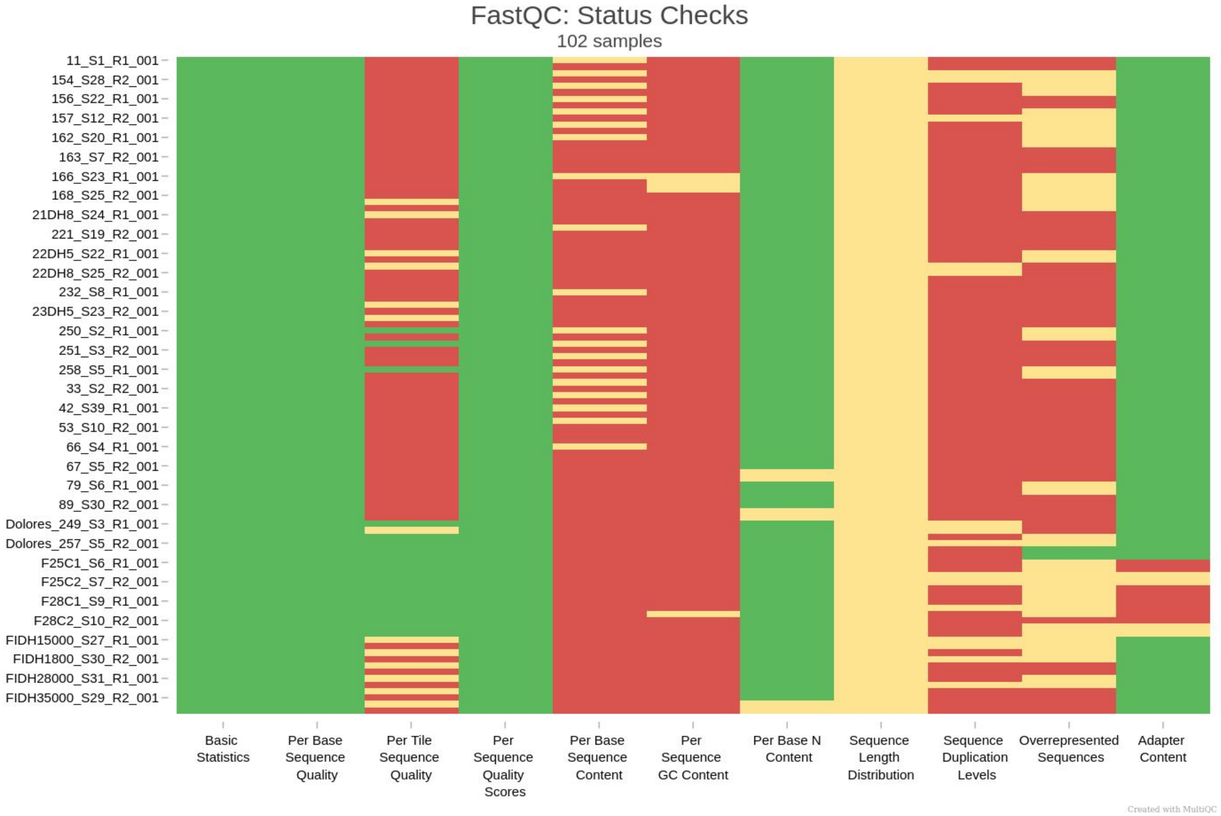
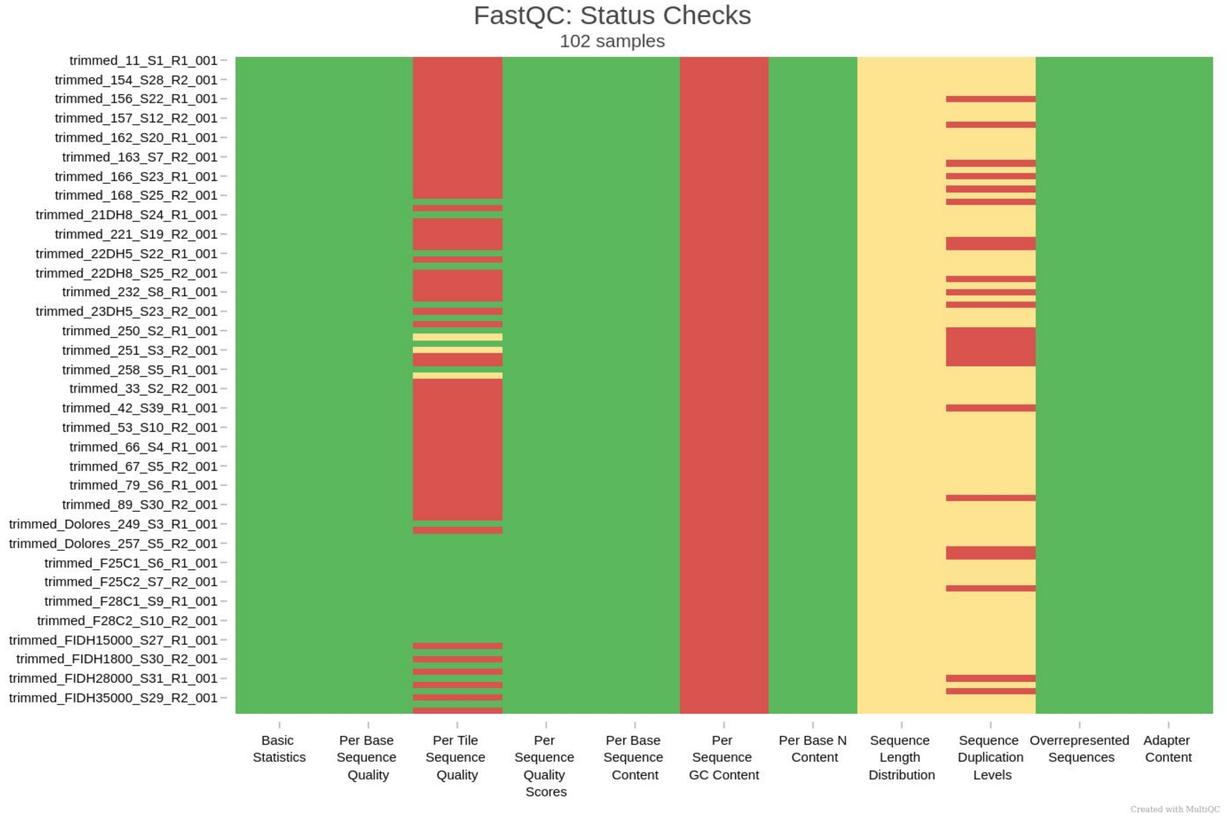
A.


FastQC: Status Checks
102 samples
11_S1_R1_001
154_S28_R2_001
156_S22_R1_001
157_S12_R2_001
162_S20_R1_001
163_S7_R2_001
166_S23_R1_001
168_S25_R2_001
21DH8_S24_R1_001
221_S19_R2_001
22DH5_S22_R1_001
22DH8_S25_R2_001
232_S8_R1_001
23DH5_S23_R2_001
250_S2_R1_001
251_S3_R2_001
258_S5_R1_001
33_S2_R2_001
42_S39_R1_001
53_S10_R2_001
66_S4_R1_001
67_S5_R2_001
79_S6_R1_001
89_S30_R2_001
Dolores_249_S3_R1_001
Dolores_257_S5_R2_001
F25C1_S6_R1_001
F25C2_S7_R2_001
F28C1_S9_R1_001
F28C2_S10_R2_001
FIDH15000_S27_R1_001
FIDH1800_S30_R2_001
FIDH28000_S31_R1_001
FIDH35000_S29_R2_001
Basic Statistics
Per Base Sequence Quality
Per Tile Sequence Quality
Per Sequence Quality Scores
Per Base Sequence Content
Per Sequence GC Content
Per Base N Content
Sequence Length Distribution
Sequence Duplication Levels
Overrepresented Sequences
Adapter Content
Created with MultiQC

B.


FastQC: Status Checks
102 samples
trimmed_11_S1_R1_001
trimmed_154_S28_R2_001
trimmed_156_S22_R1_001
trimmed_157_S12_R2_001
trimmed_162_S20_R1_001
trimmed_163_S7_R2_001
trimmed_166_S23_R1_001
trimmed_168_S25_R2_001
trimmed_21DH8_S24_R1_001
trimmed_221_S19_R2_001
trimmed_22DH5_S22_R1_001
trimmed_22DH8_S25_R2_001
trimmed_232_S8_R1_001
trimmed_23DH5_S23_R2_001
trimmed_250_S2_R1_001
trimmed_251_S3_R2_001
trimmed_258_S5_R1_001
trimmed_33_S2_R2_001
trimmed_42_S39_R1_001
trimmed_53_S10_R2_001
trimmed_66_S4_R1_001
trimmed_67_S5_R2_001
trimmed_79_S6_R1_001
trimmed_89_S30_R2_001
trimmed_Dolores_249_S3_R1_001
trimmed_Dolores_257_S5_R2_001
trimmed_F25C1_S6_R1_001
trimmed_F25C2_S7_R2_001
trimmed_F28C1_S9_R1_001
trimmed_F28C2_S10_R2_001
trimmed_FIDH15000_S27_R1_001
trimmed_FIDH1800_S30_R2_001
trimmed_FIDH28000_S31_R1_001
trimmed_FIDH35000_S29_R2_001
Basic Statistics
Per Base Sequence Quality
Per Tile Sequence Quality
Per Sequence Quality Scores
Per Base Sequence Content
Per Sequence GC Content
Per Base N Content
Sequence Length Distribution
Sequence Duplication Levels
Overrepresented Sequences
Adapter Content
Created with MultiQC