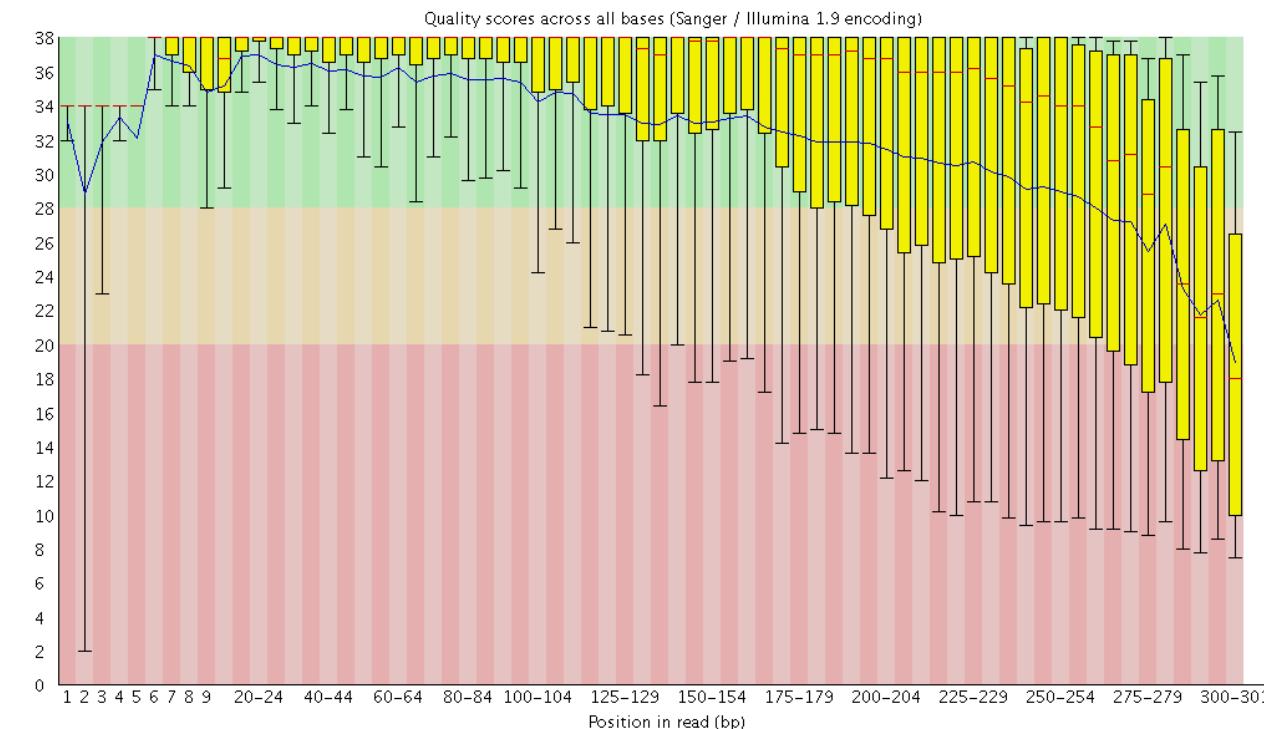
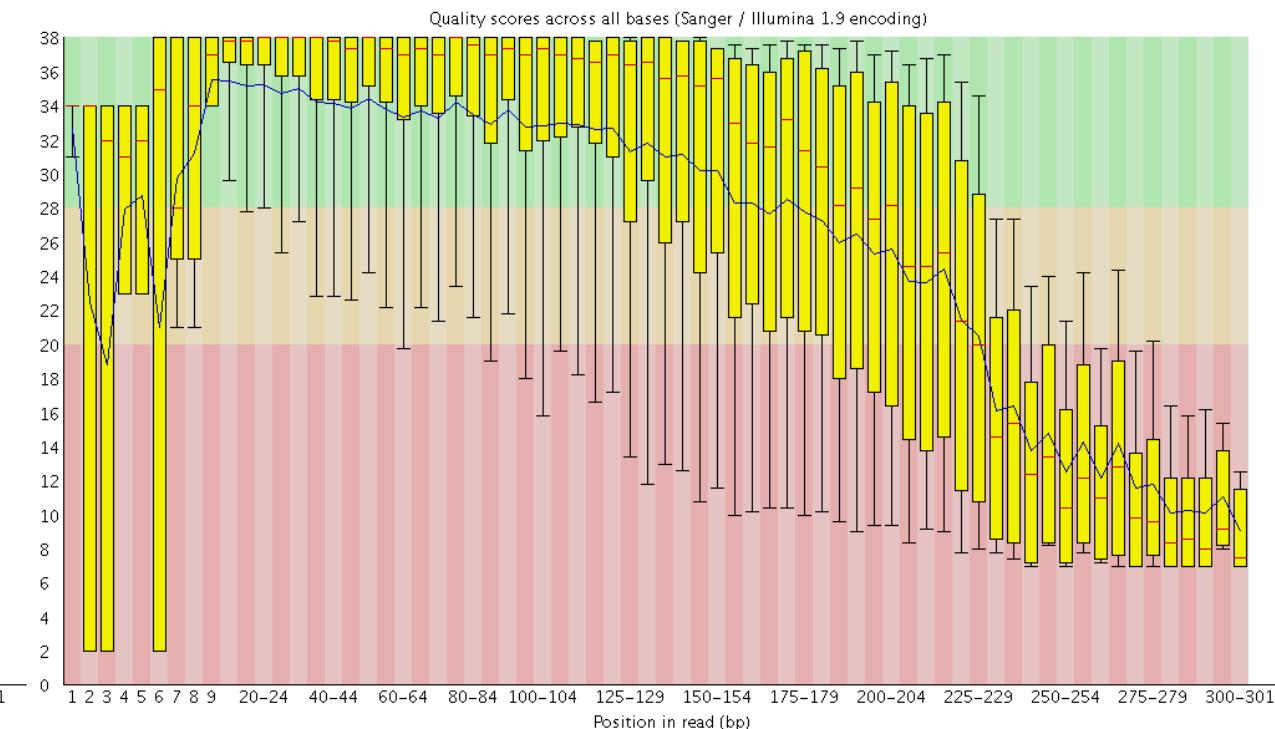


RAWDATA

gut_sample1_pond1_R1.fastq.gz

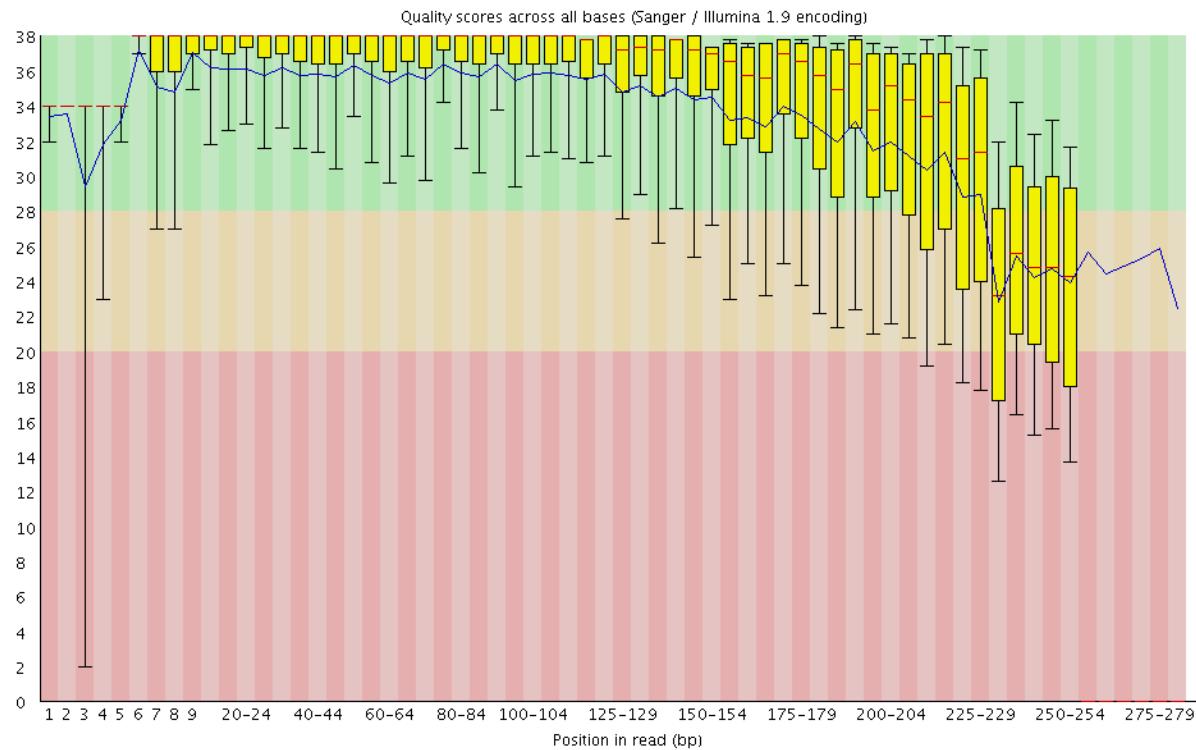


gut_sample1_pond1_R2.fastq.gz

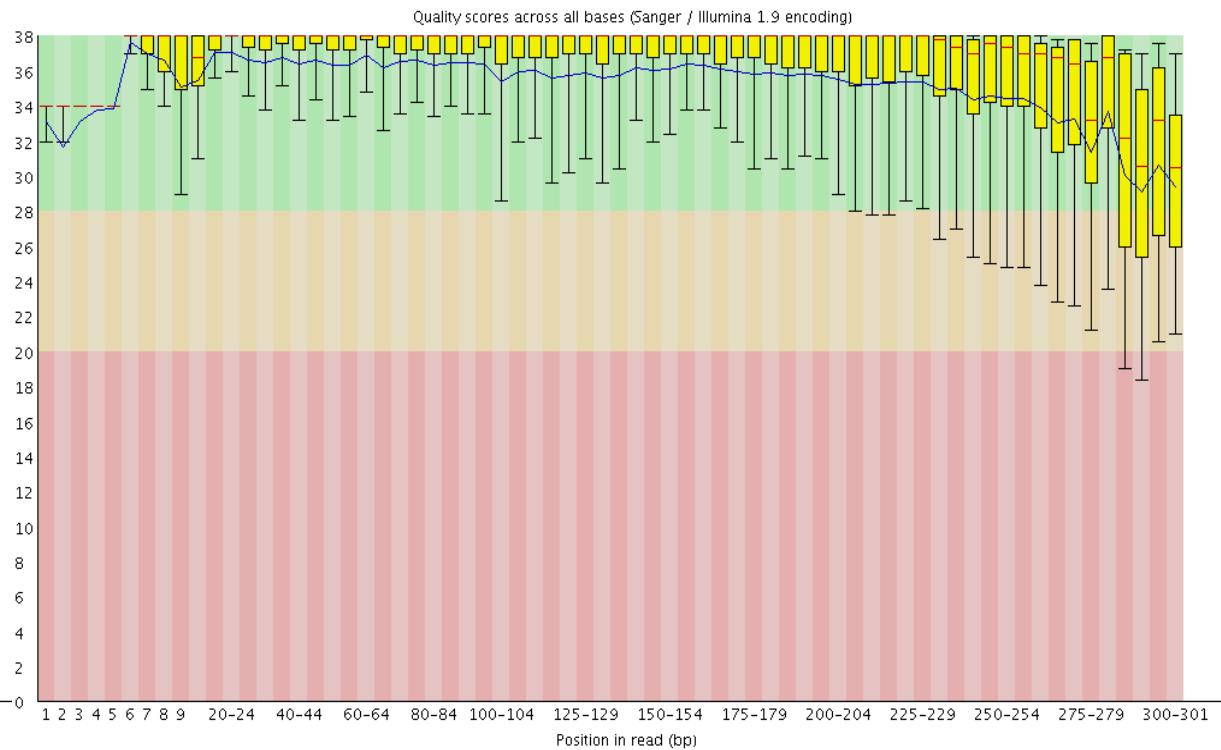


QUALITY

gut_sample1_pond1_R1_paired.fastq.gz

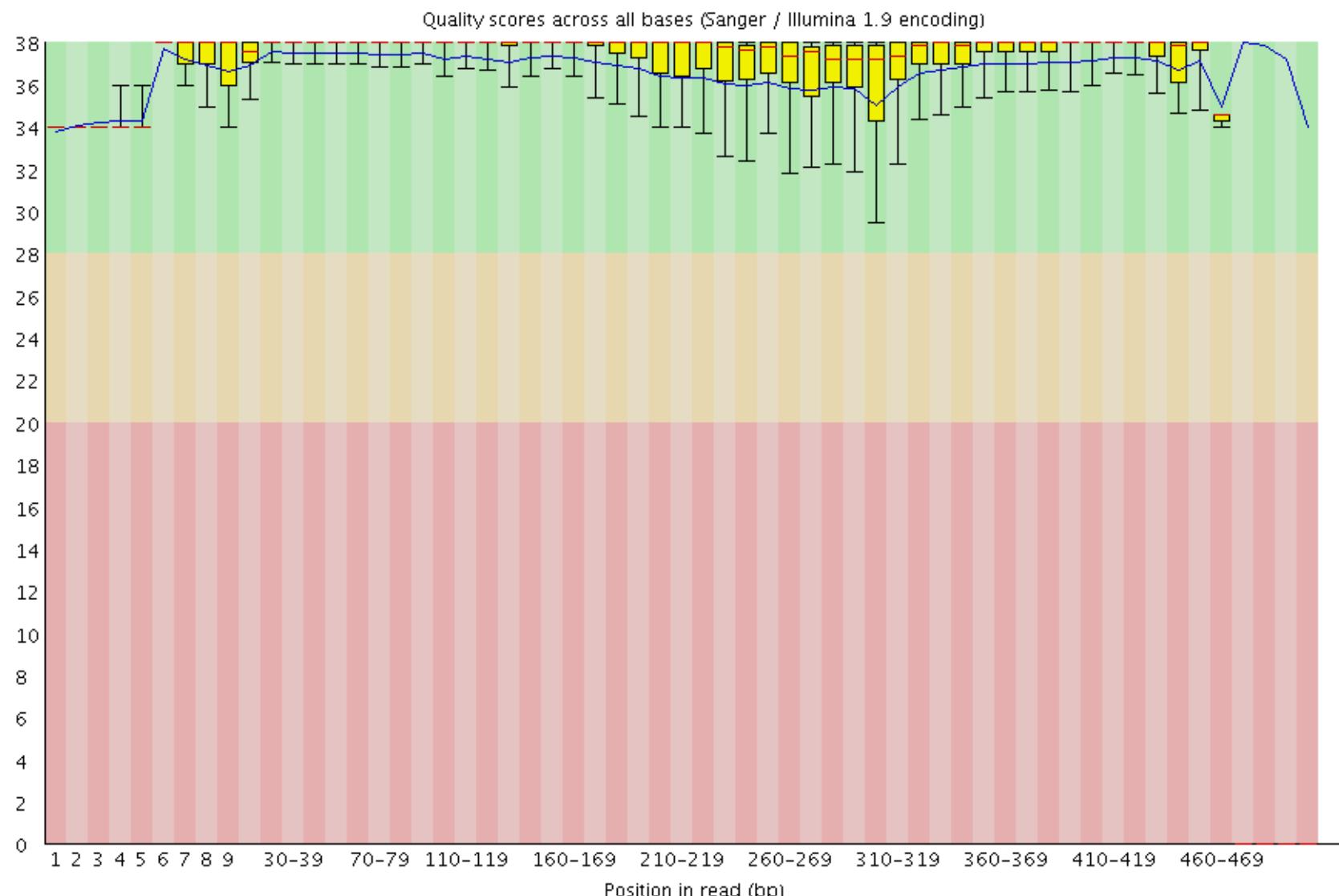


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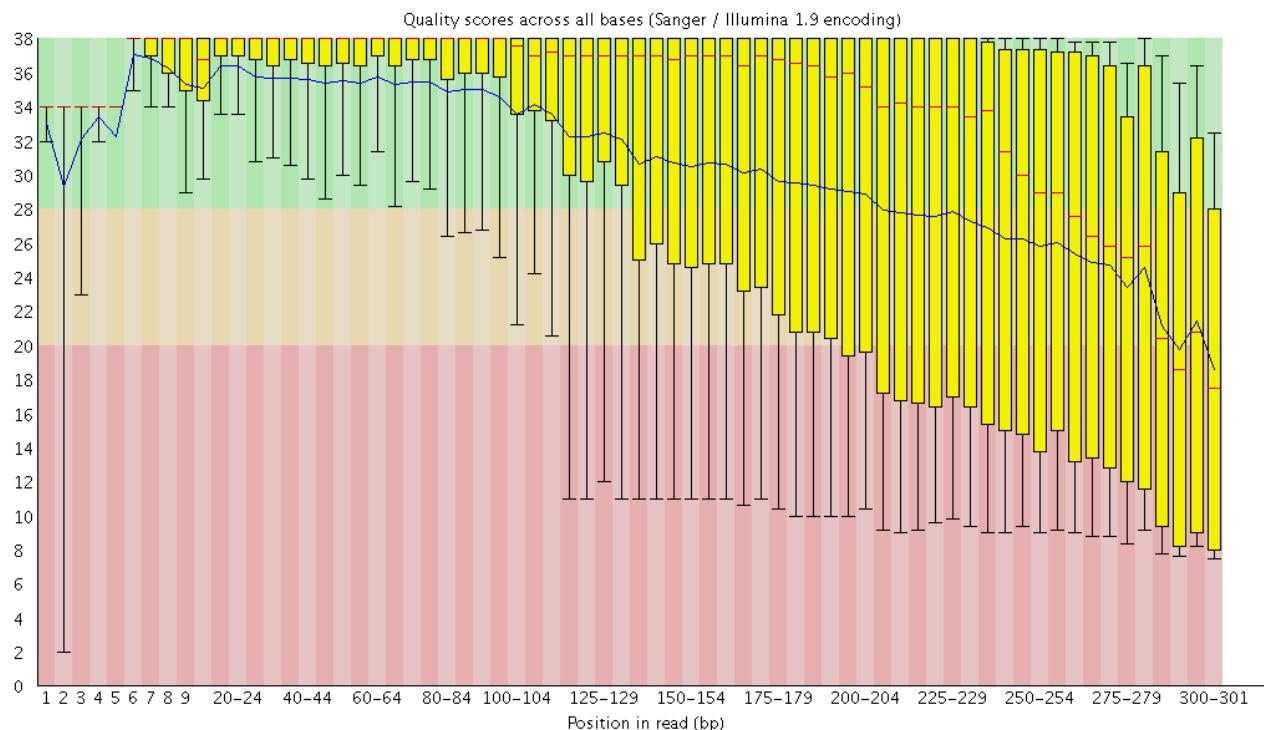
JOIN

gjoinedsample1p1.fastqjoin

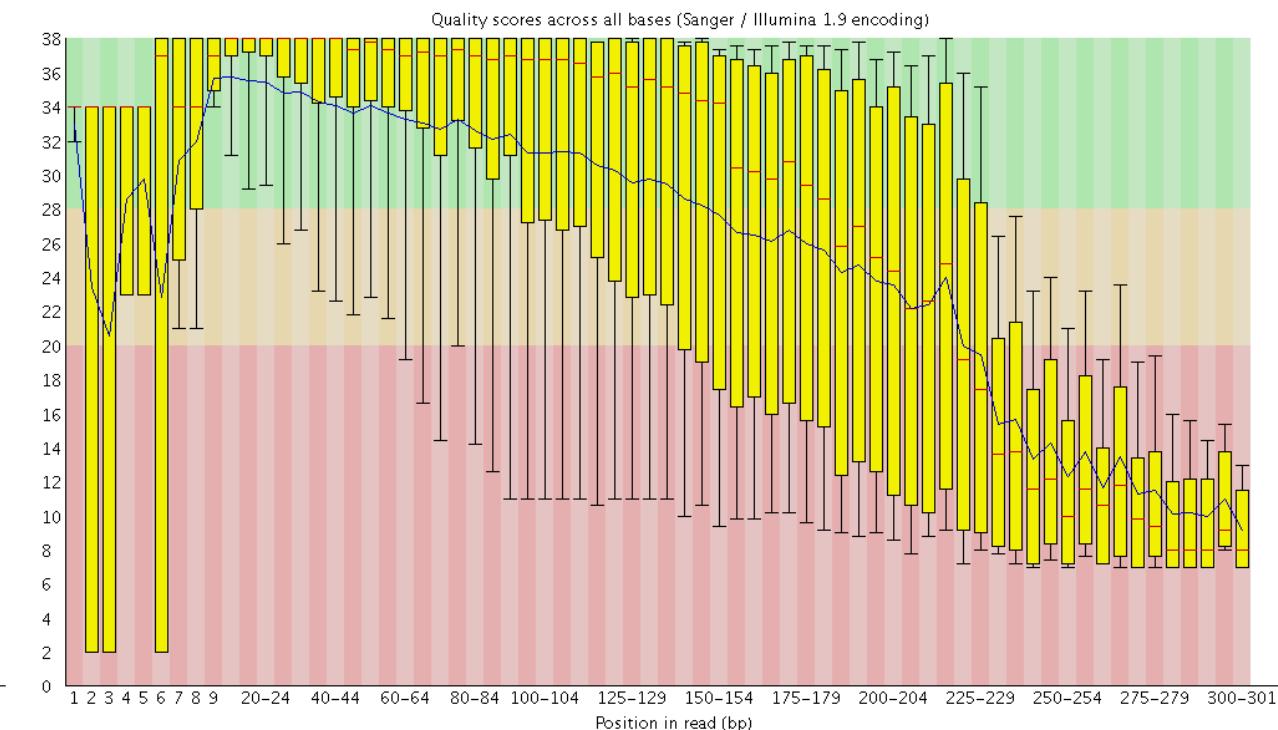


RAWDATA

gut_sample1_pond2_R1.fastq.gz

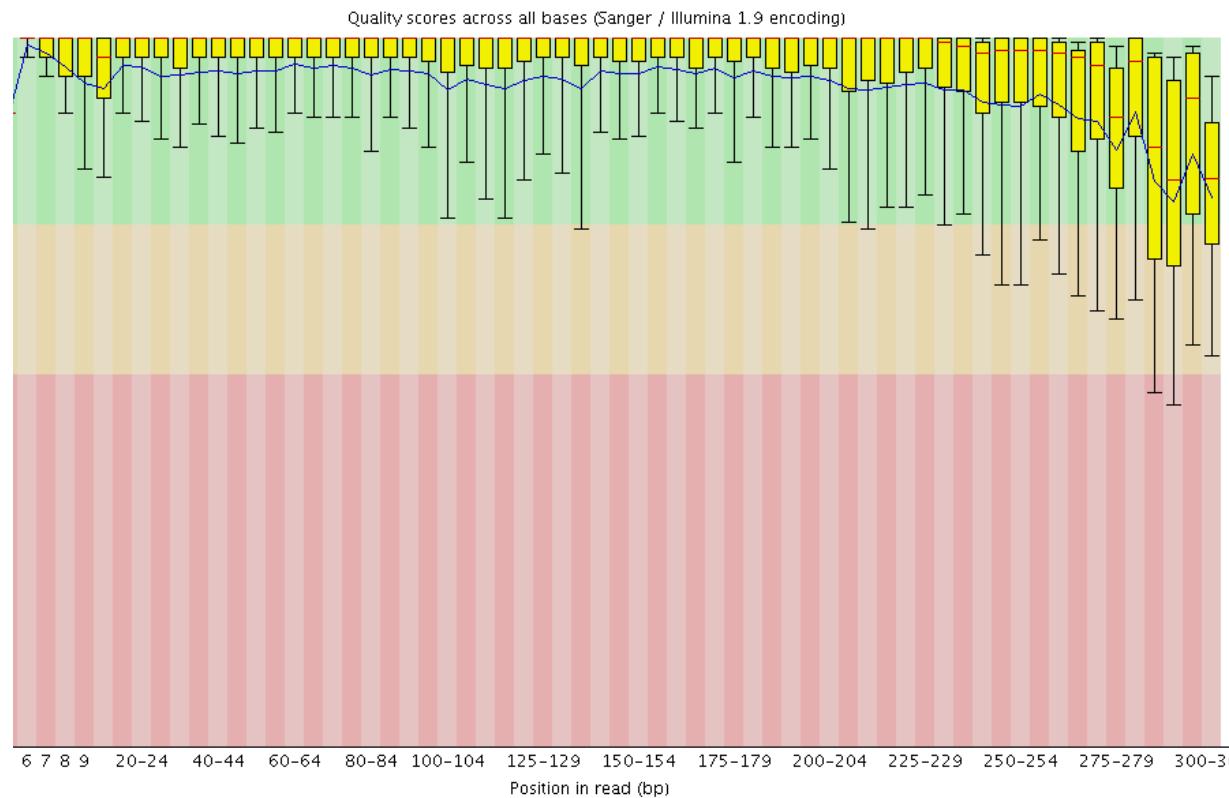


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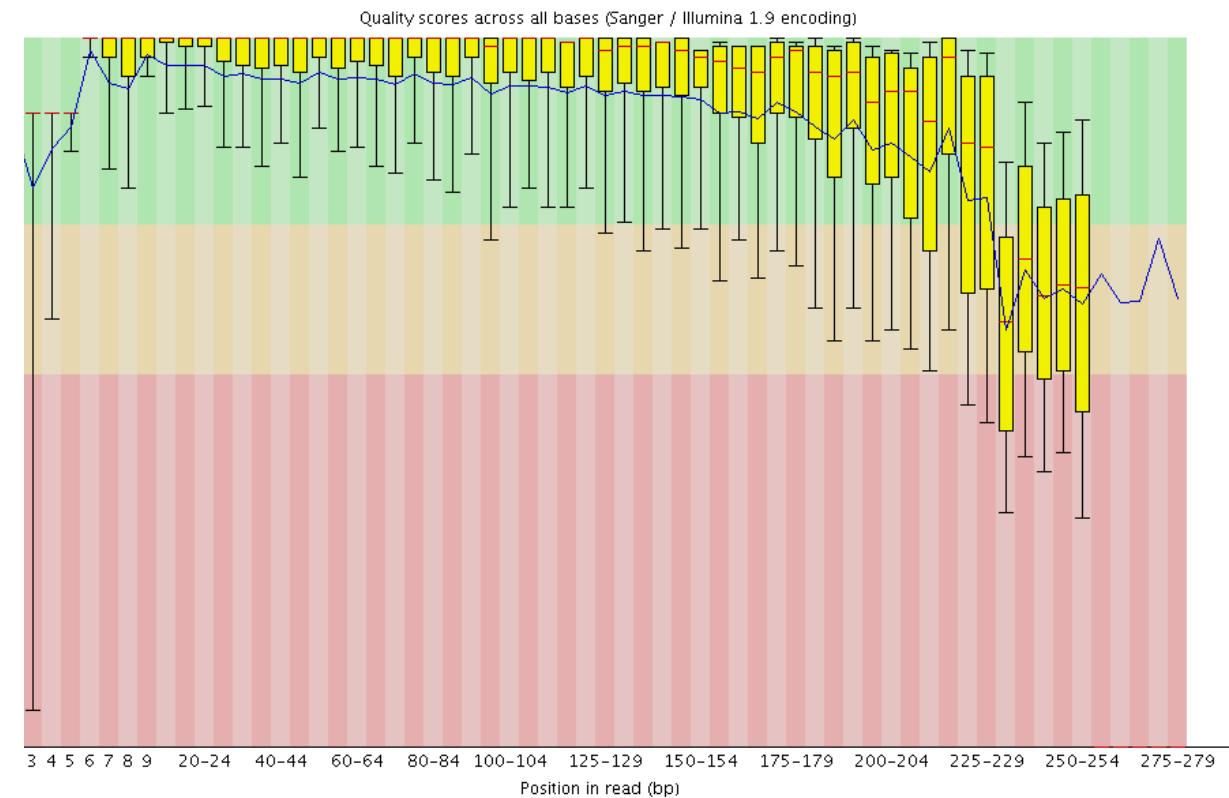


QUALITY

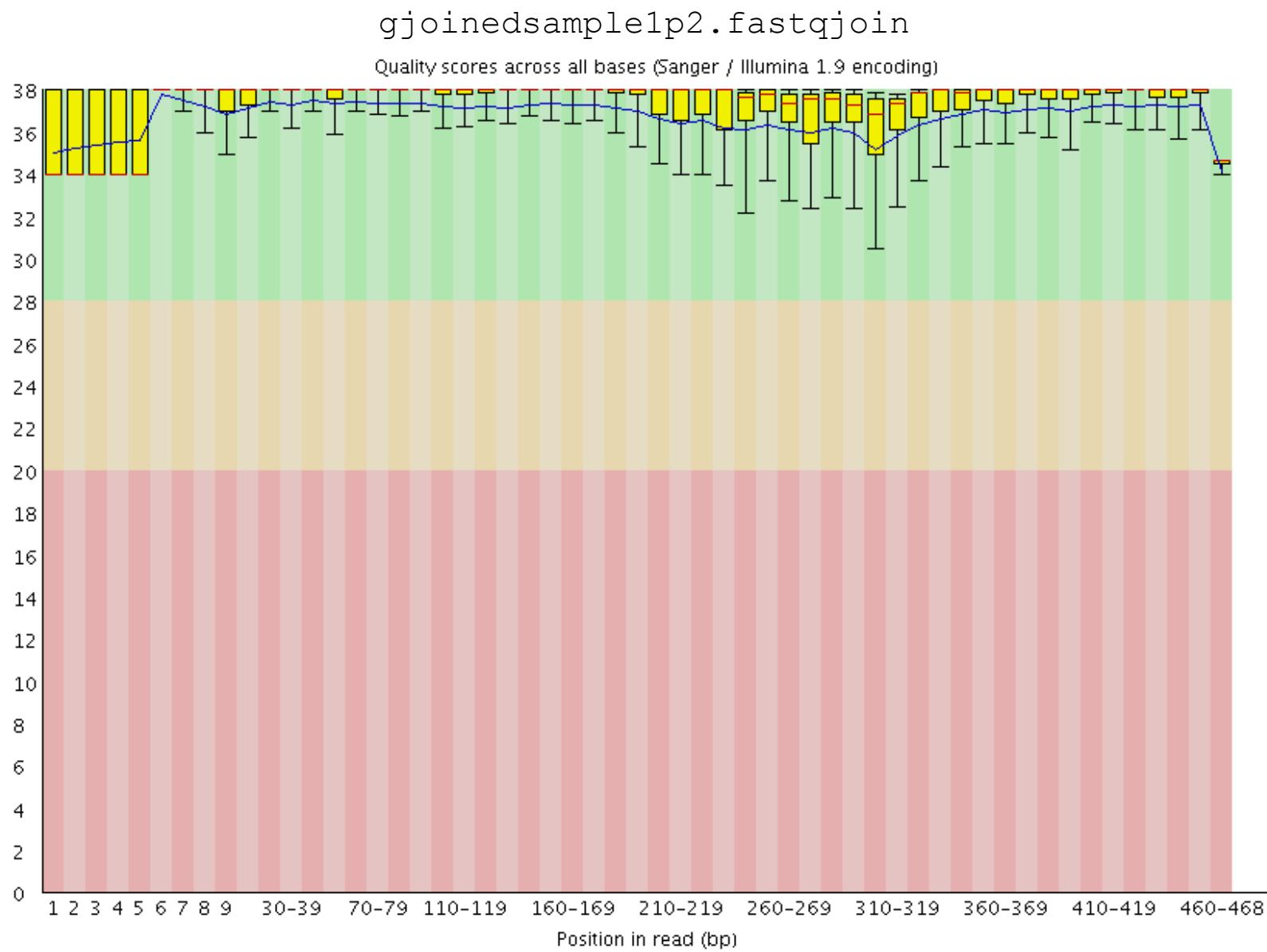
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gut_sample1_pond2_R2_paired.fastq.gz

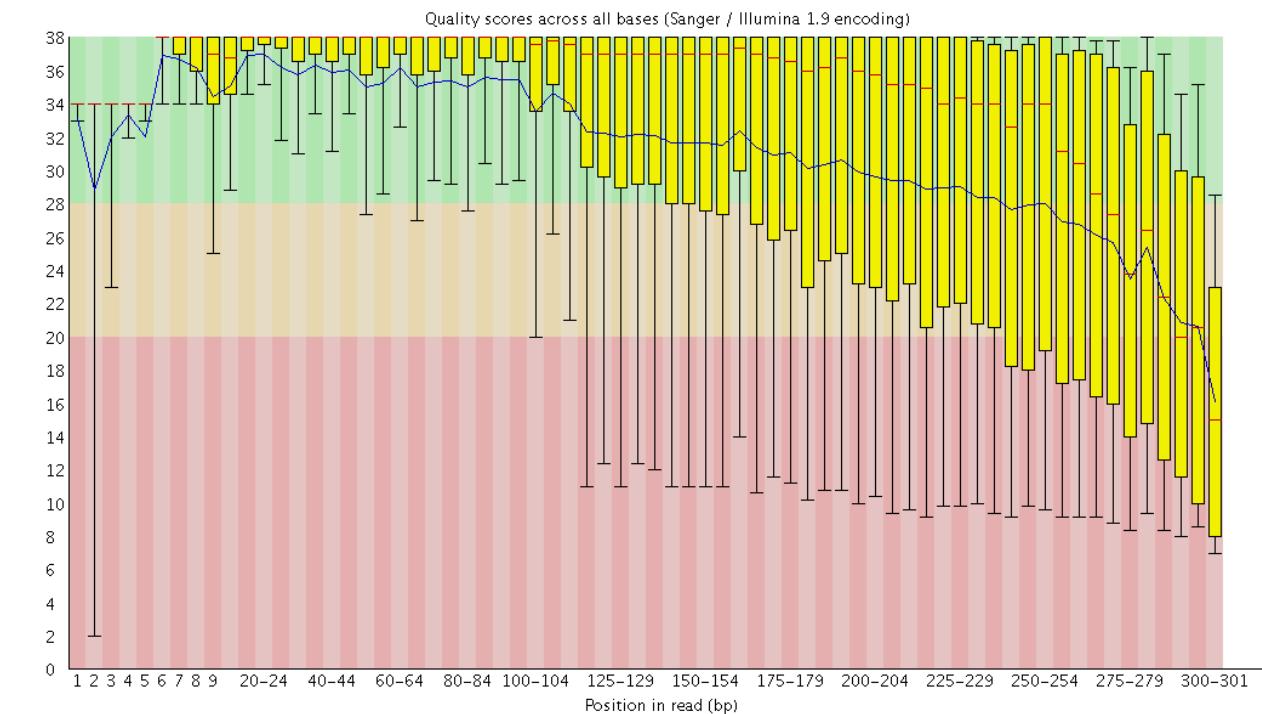


JOIN

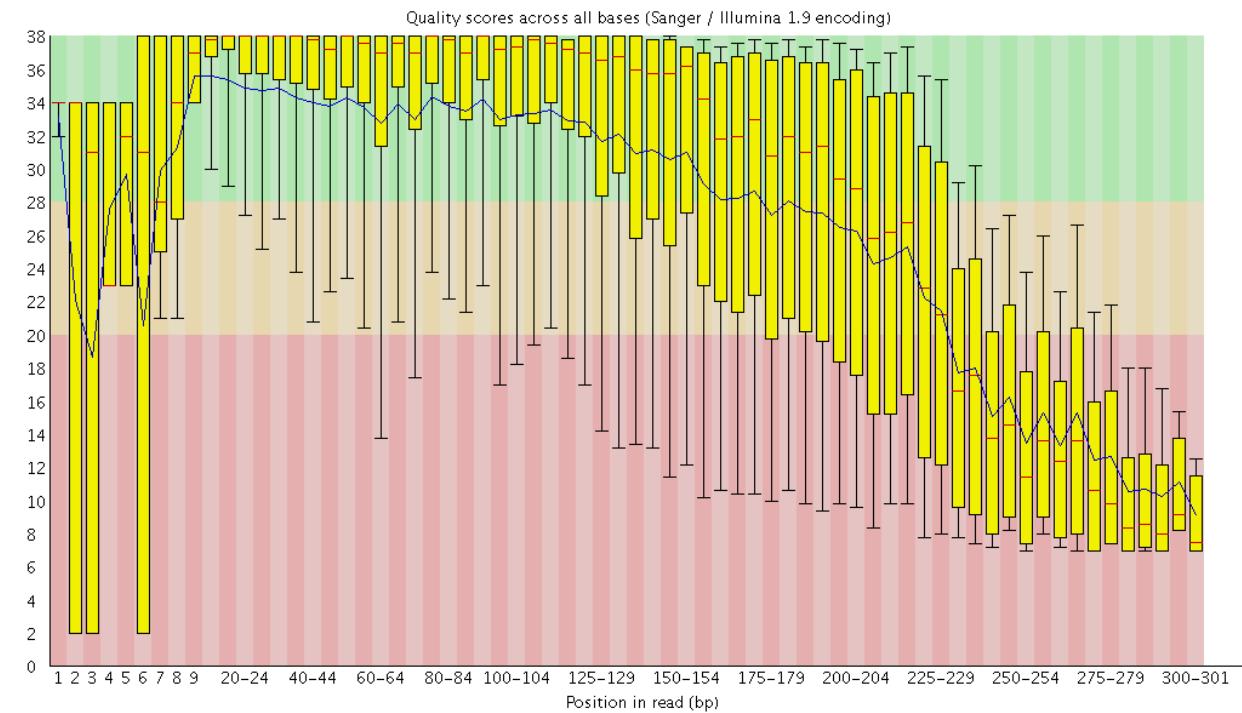


RAWDATA

gut_sample1_pond3_R1.fastq.gz

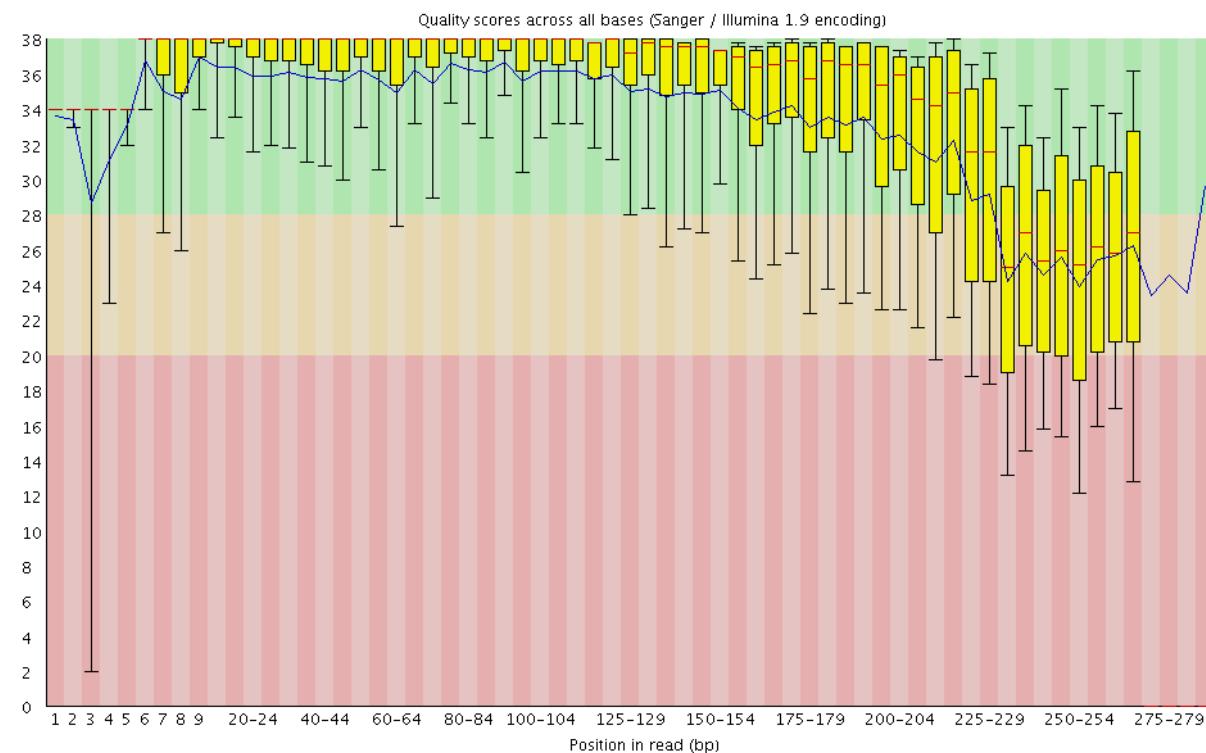


gut_sample1_pond3_R2.fastq.gz

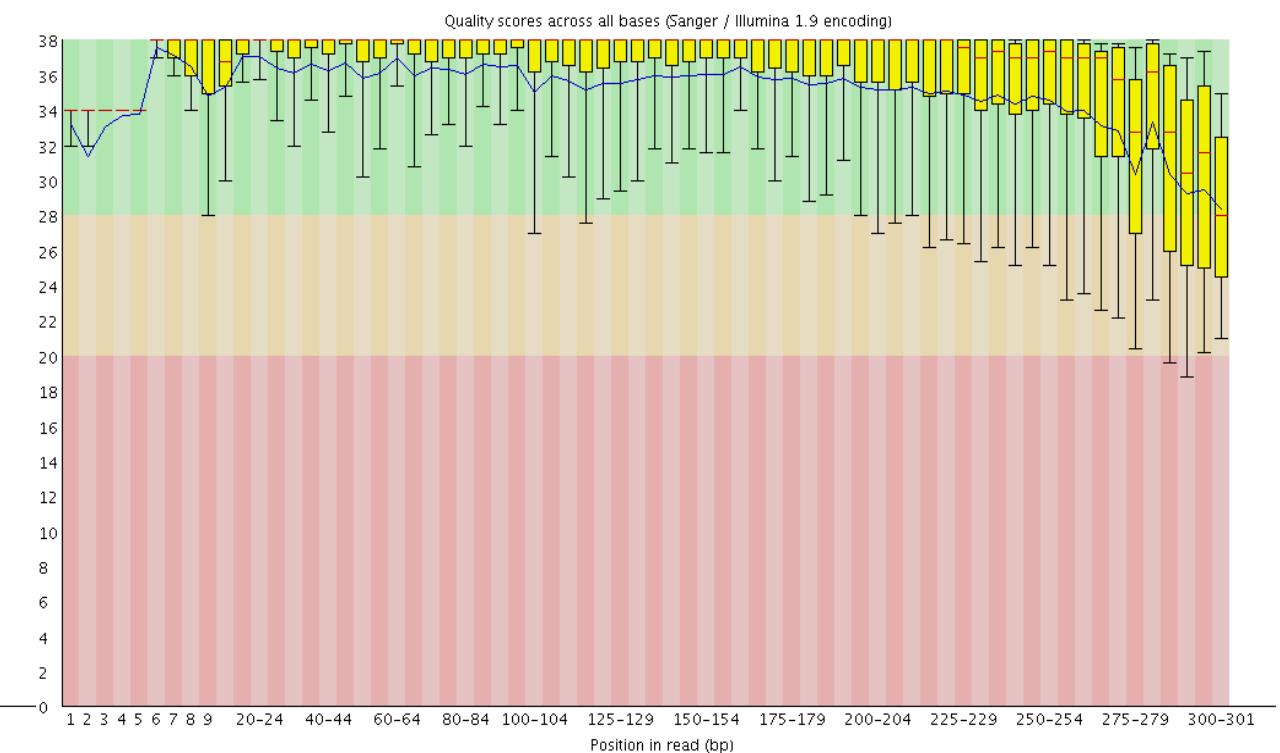


QUALITY

gut_sample1_pond3_R1_paired.fastq.gz

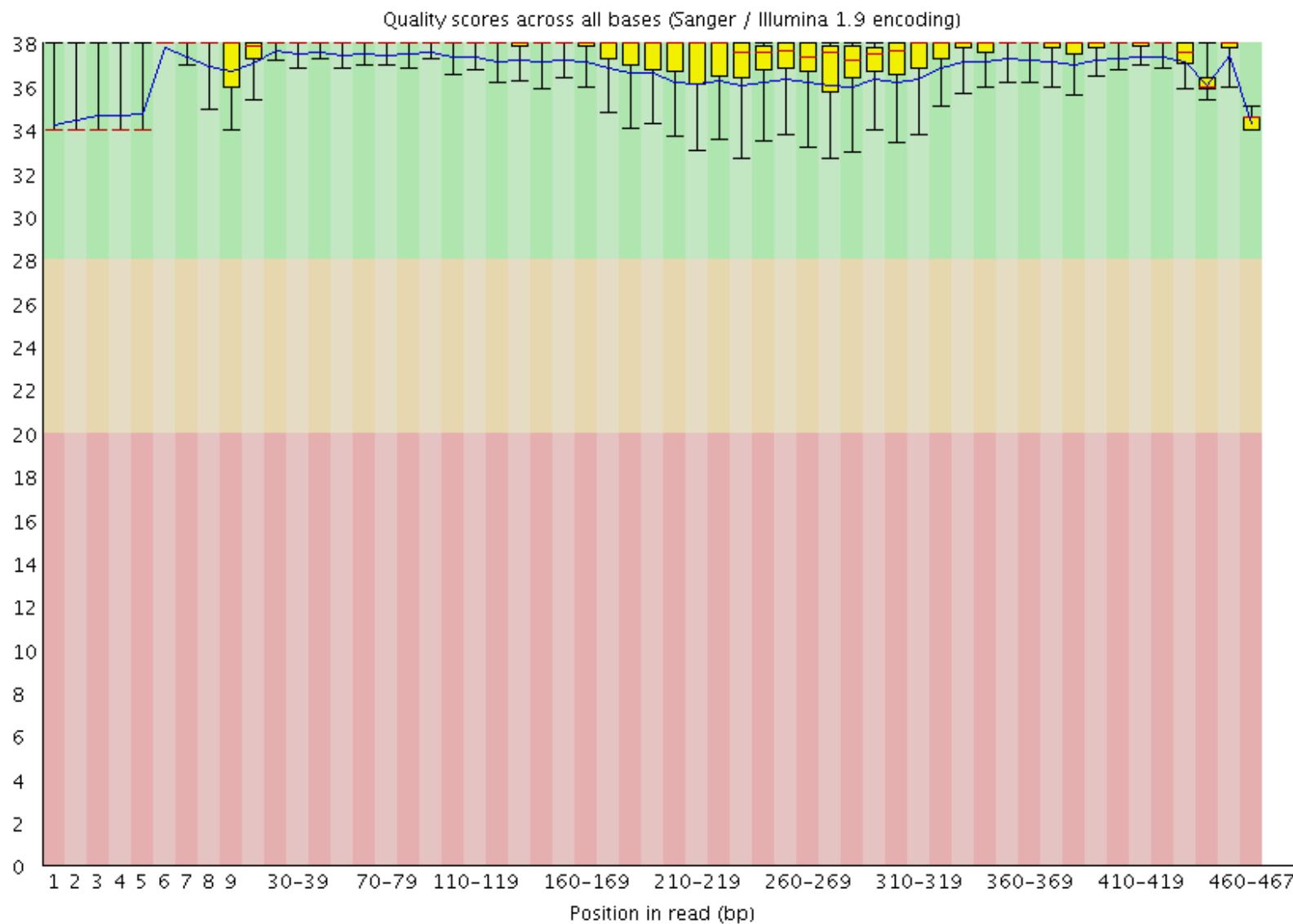


gut_sample1_pond3_R2_paired.fastq.gz



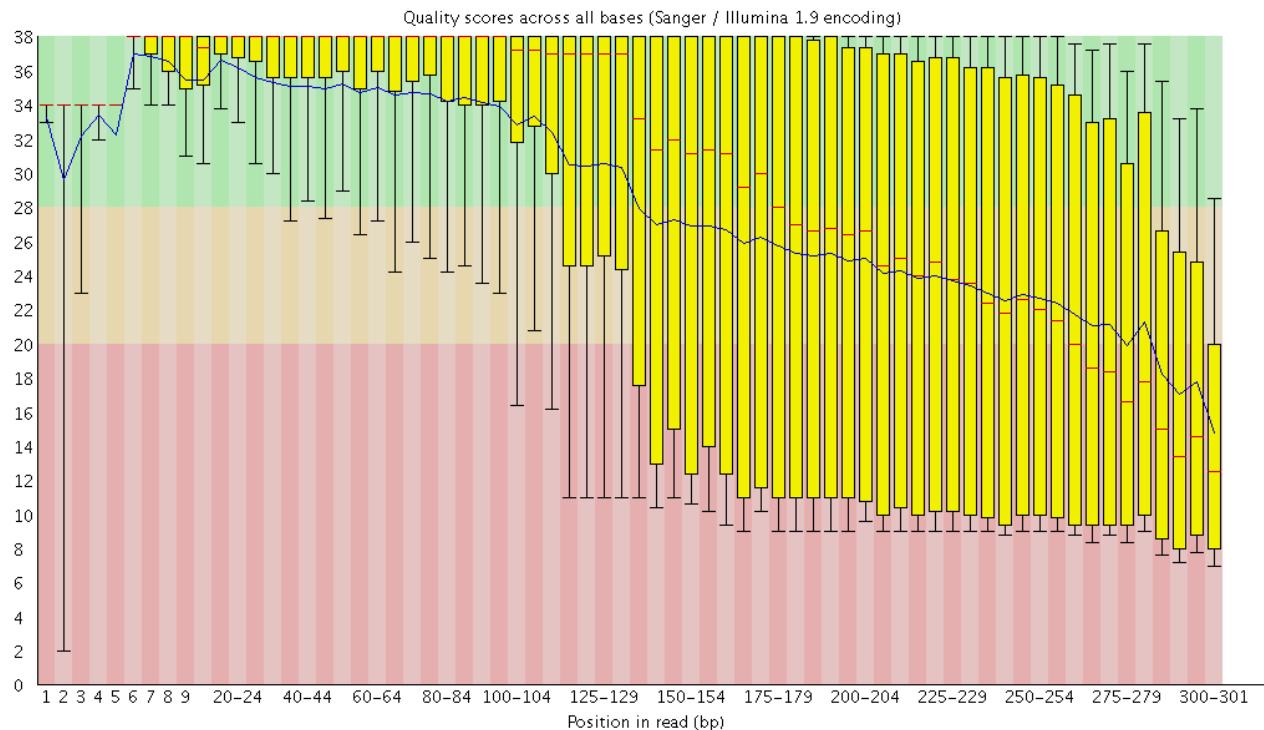
JOIN

gjoinedsample1p3.fastqjoin

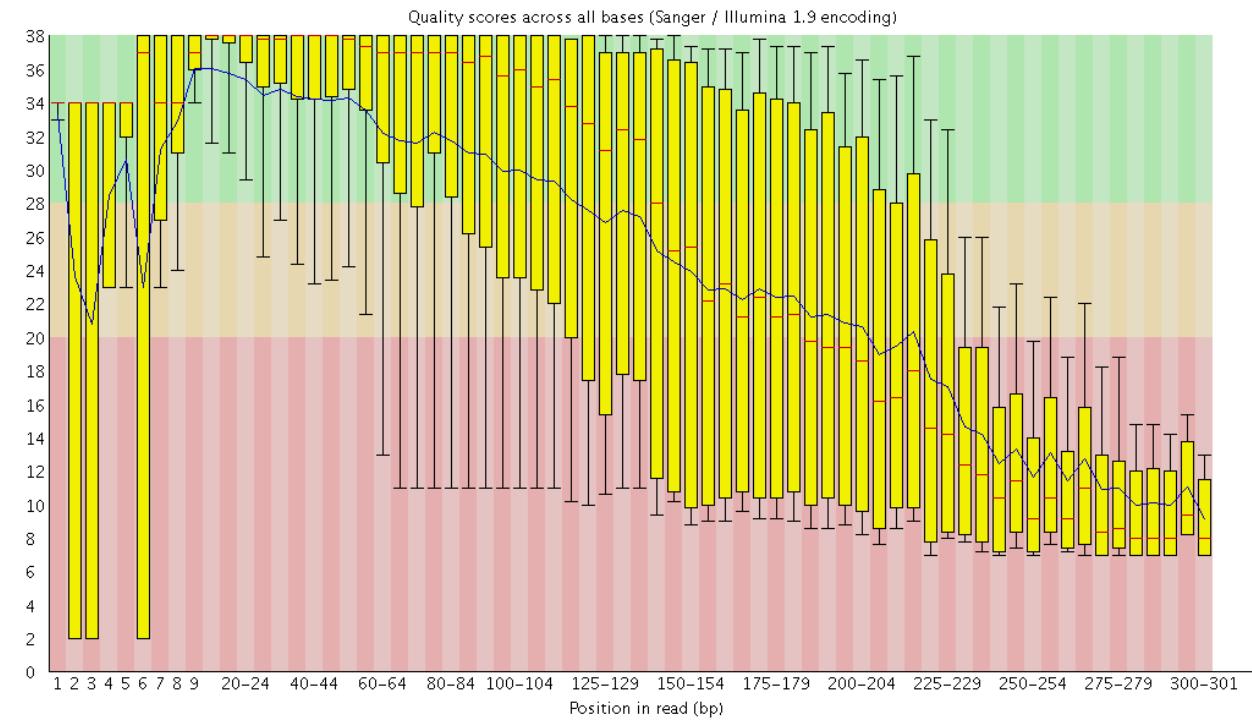


RAWDATA

gut_sample2_pond1_R1.fastq.gz

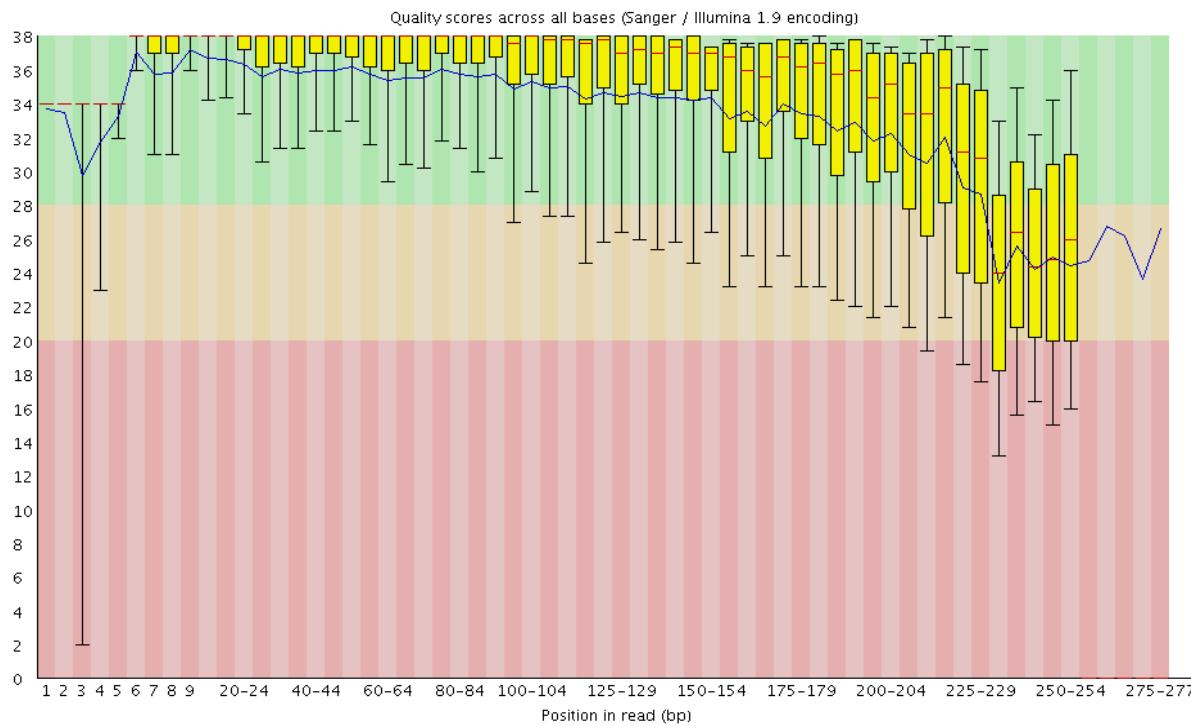


gut_sample2_pond1_R2.fastq.gz

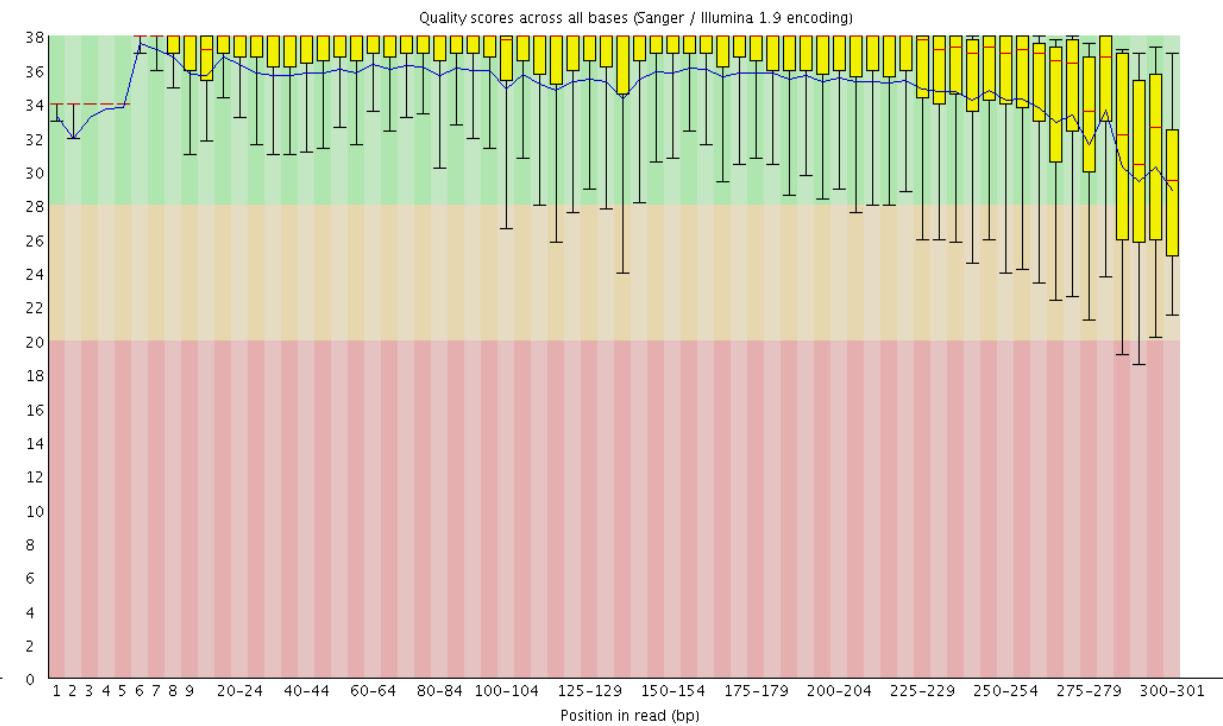


QUALITY

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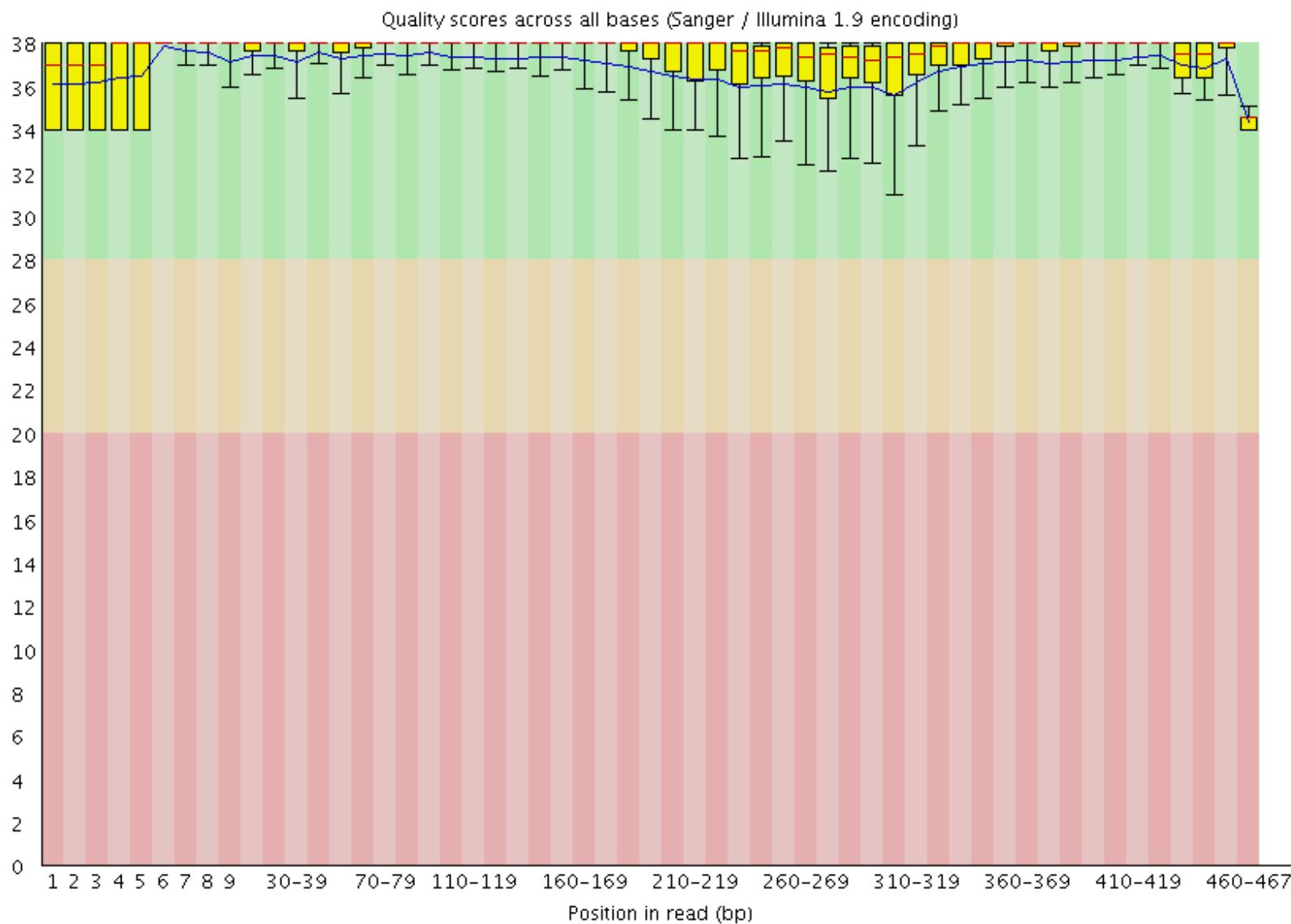


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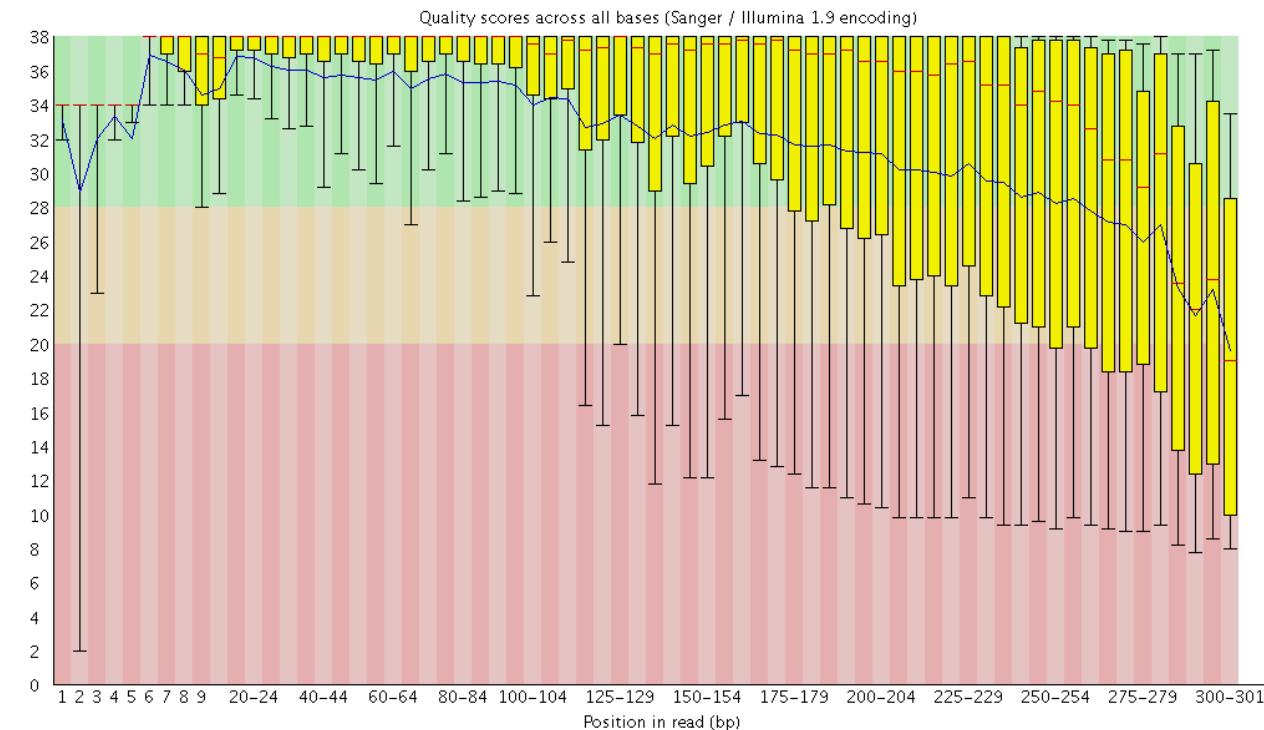
JOIN

gjoinedsample2p1.fastqjoin

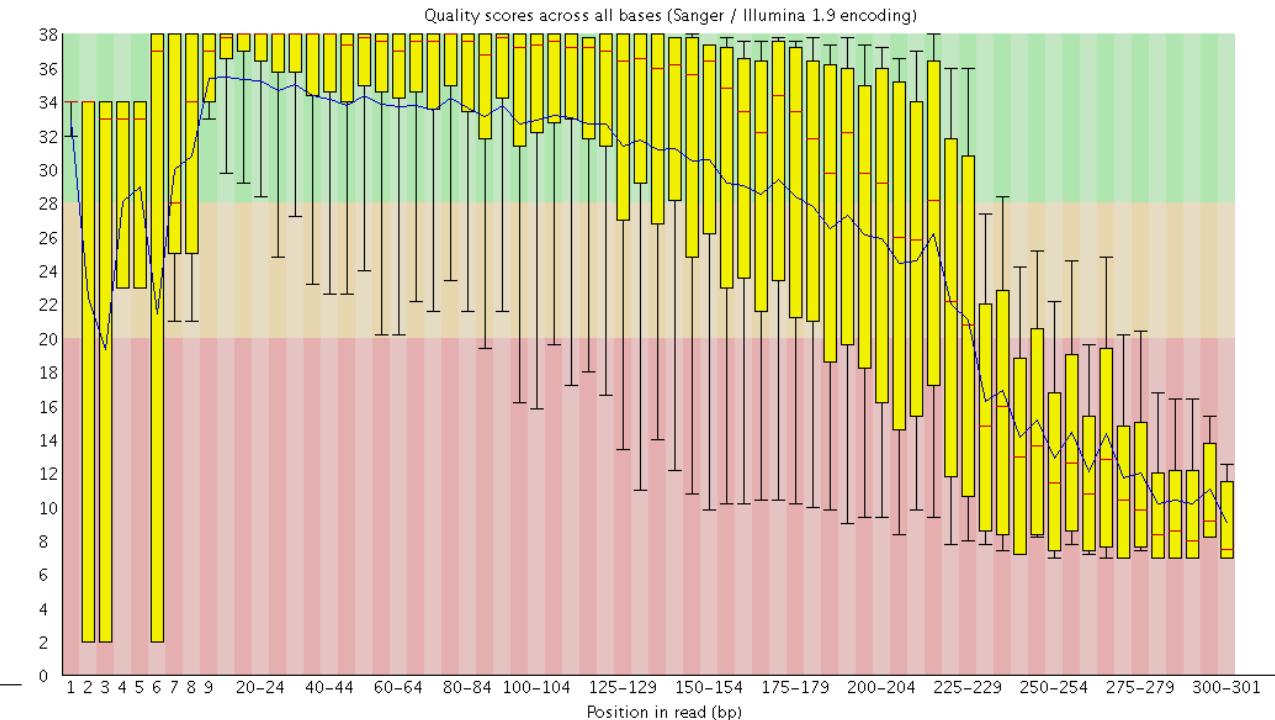


RAWDATA

gut_sample2_pond2_R1.fastq.gz

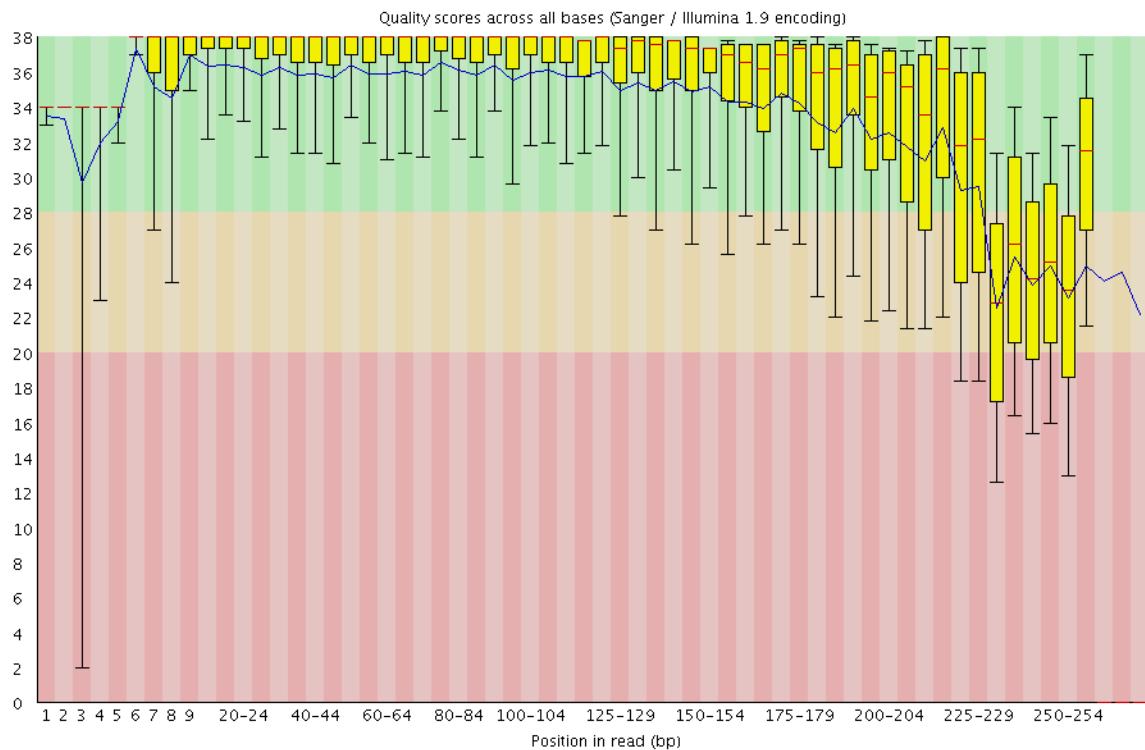


gut_sample2_pond2_R2.fastq.gz

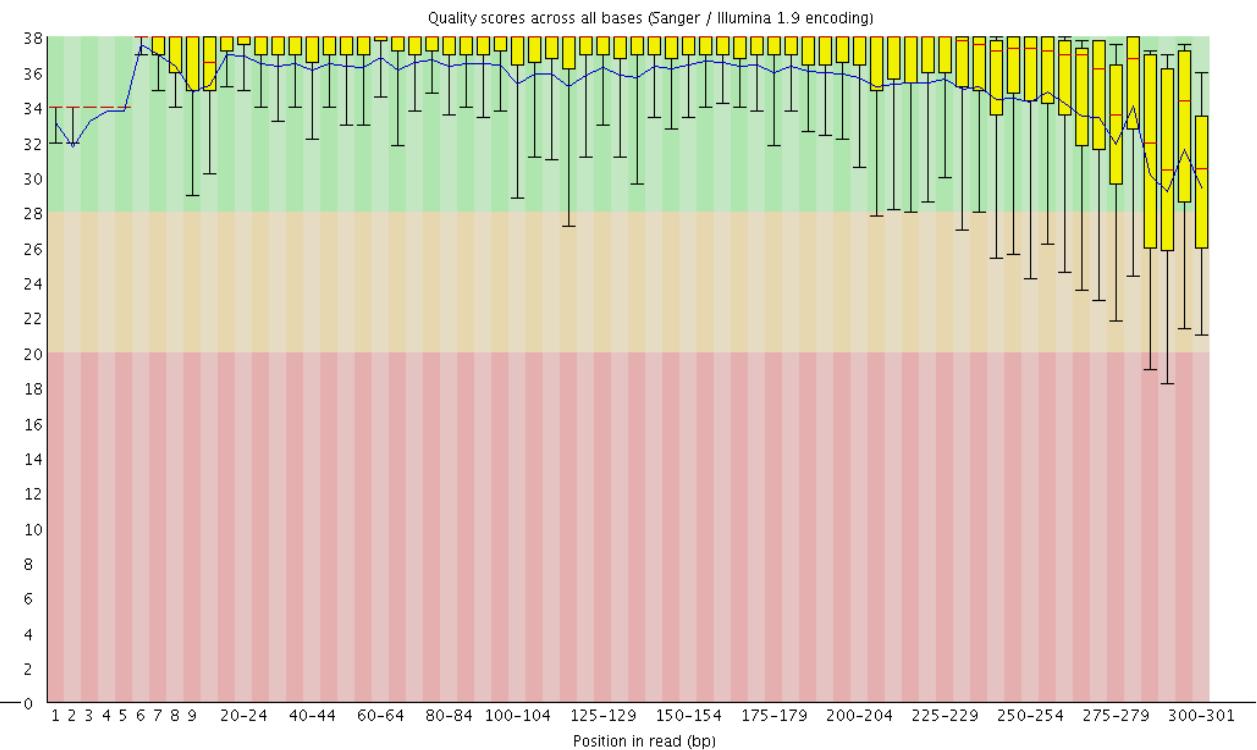


QUALITY

gut_sample2_pond2_R1_paired.fastq.gz

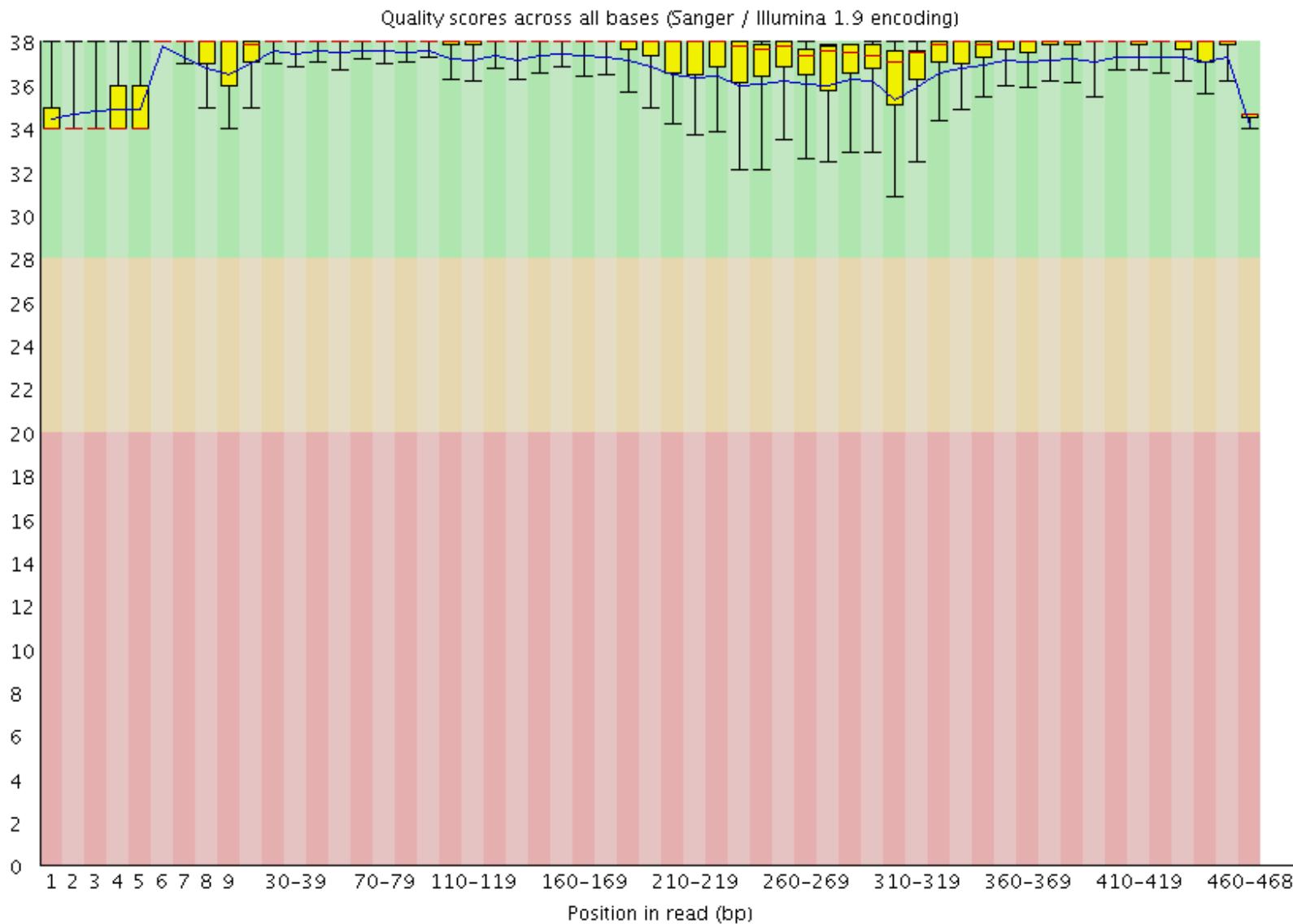


gut_sample2_pond2_R2_paired.fastq.gz



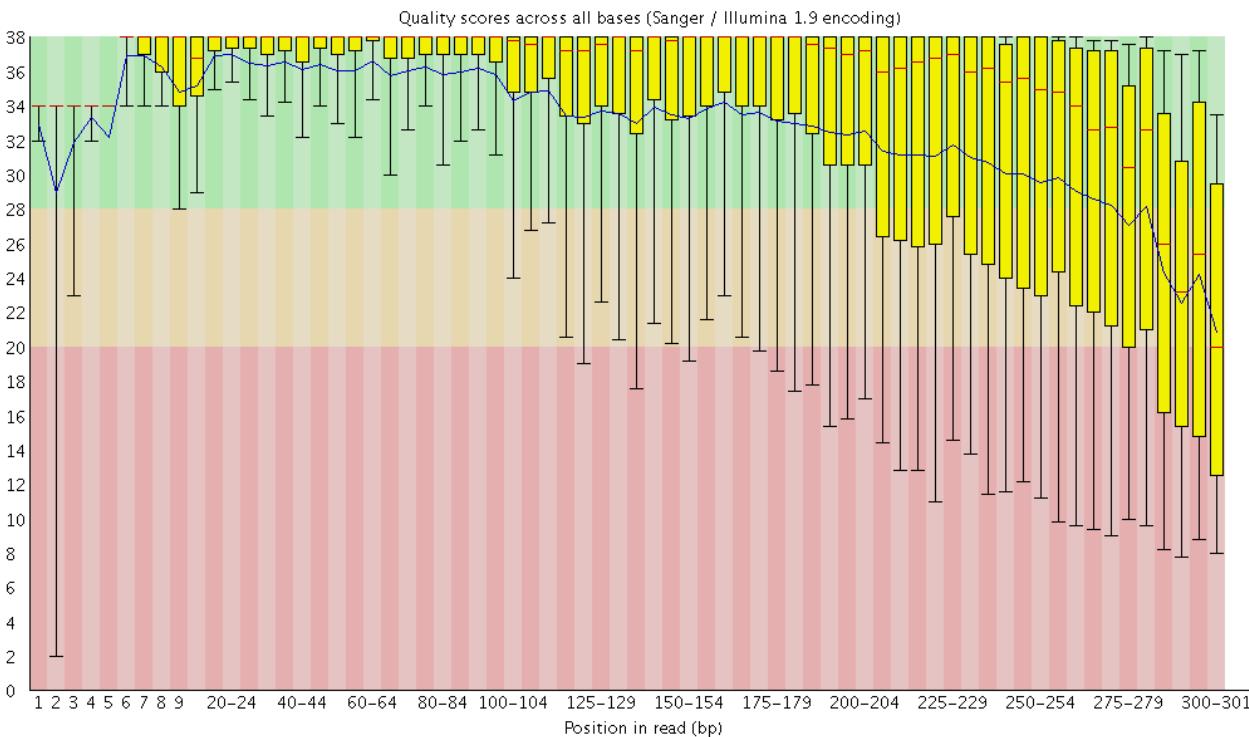
JOIN

gjoinedsample2p2.fastqjoin

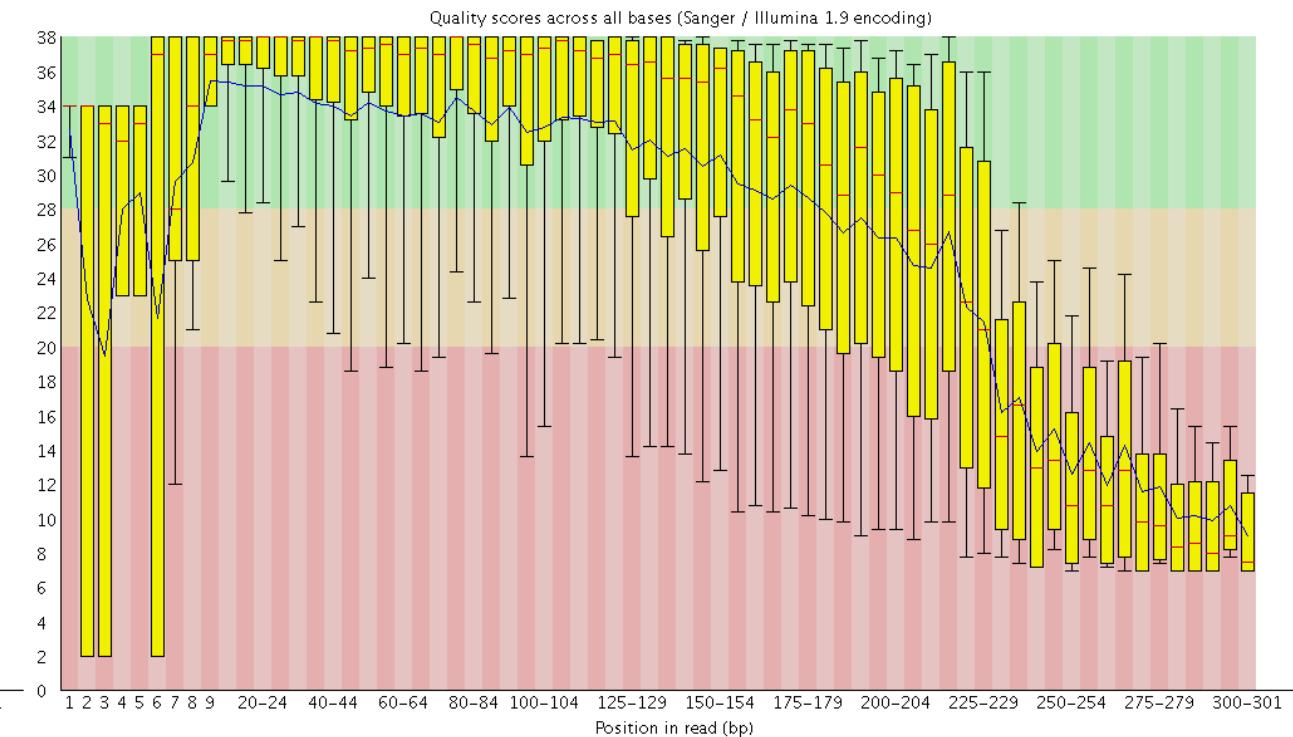


RAWDATA

gut_sample2_pond3_R1.fastq.gz

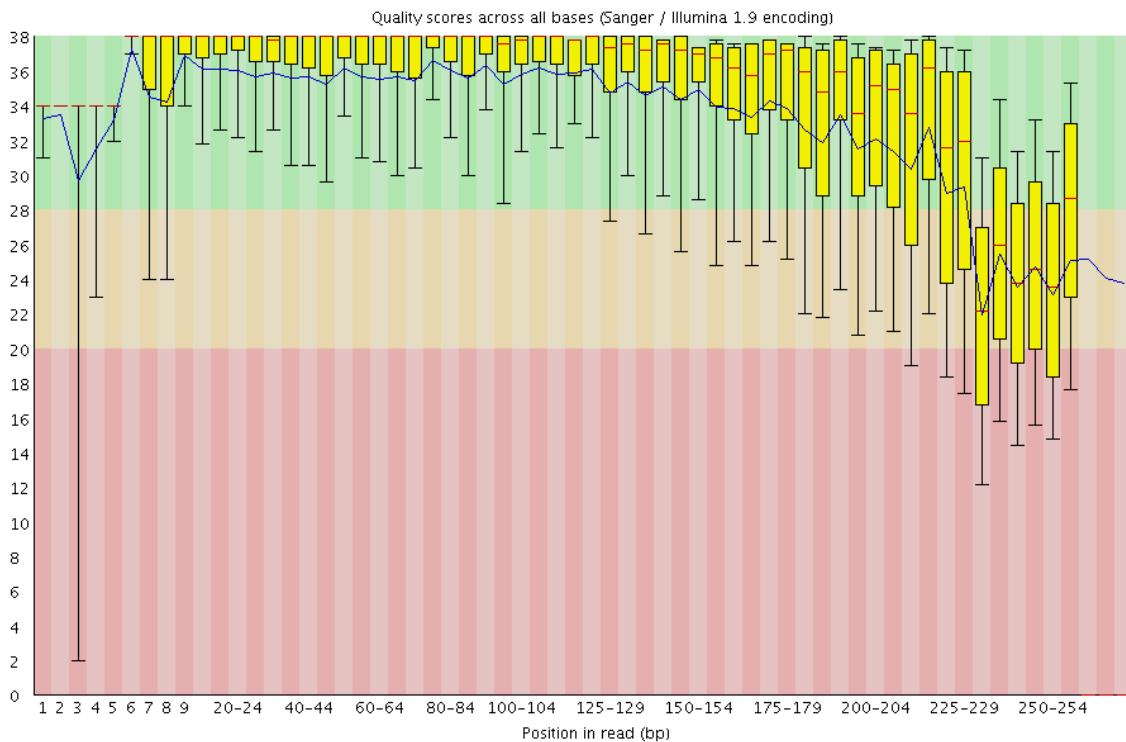


gut_sample2_pond3_R2.fastq.gz

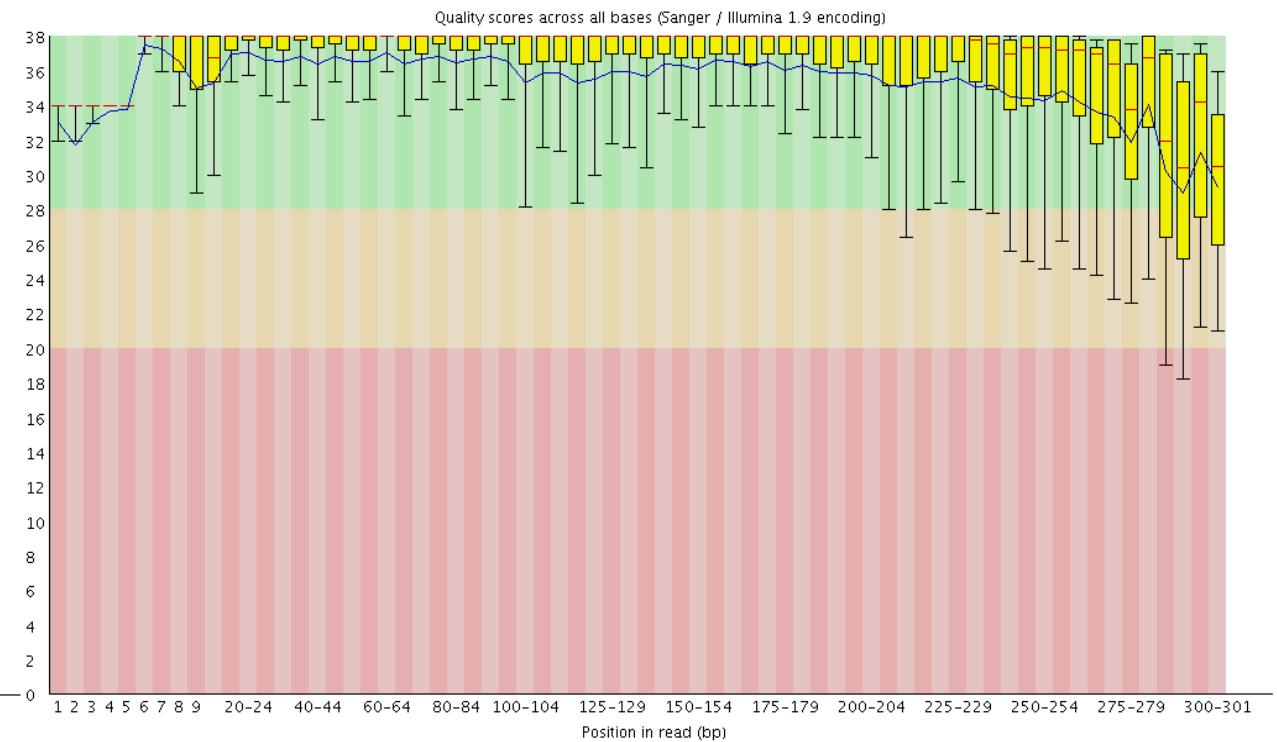


QUALITY

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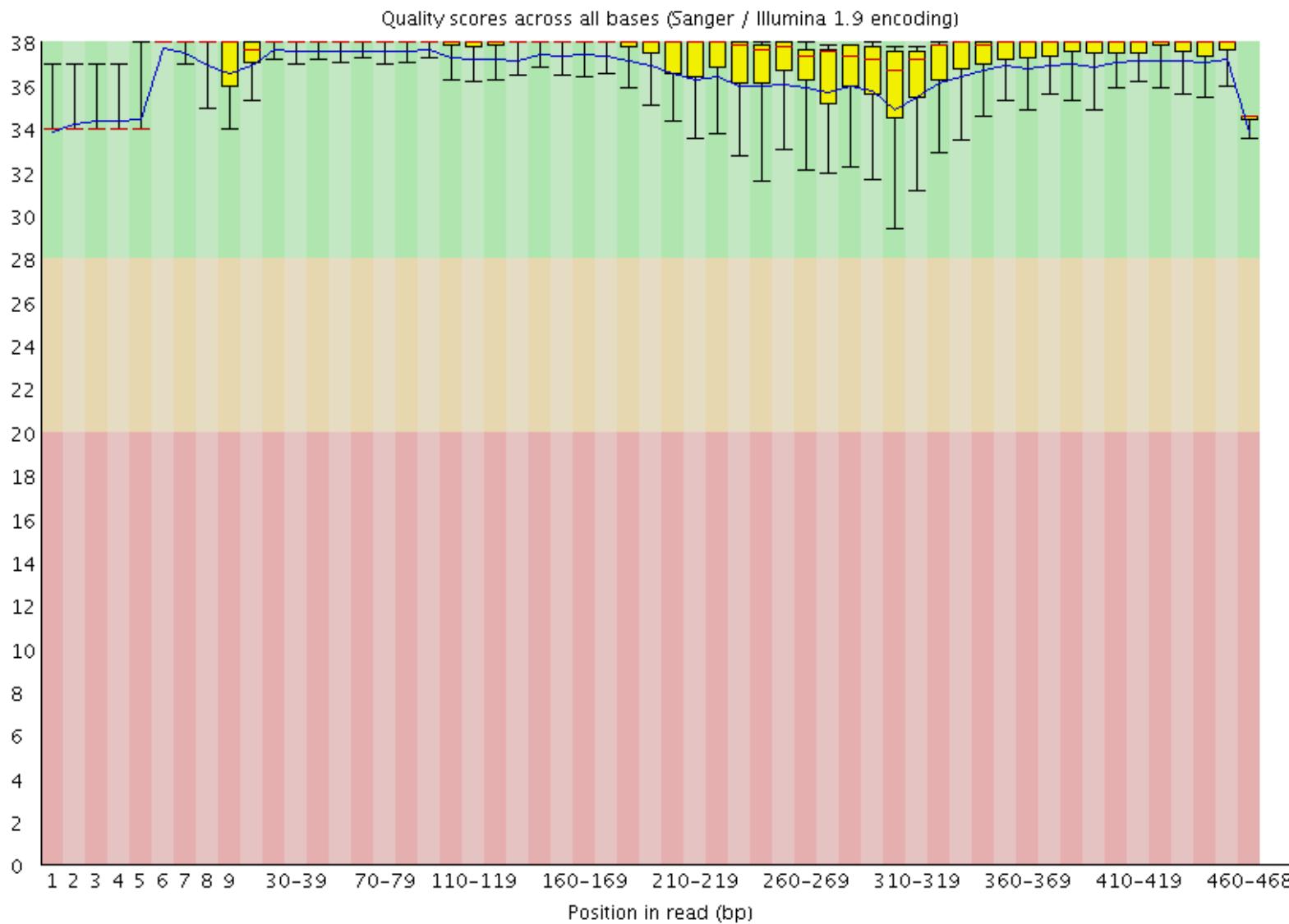


gut_sample2_pond3_R2_paired.fastq.gz



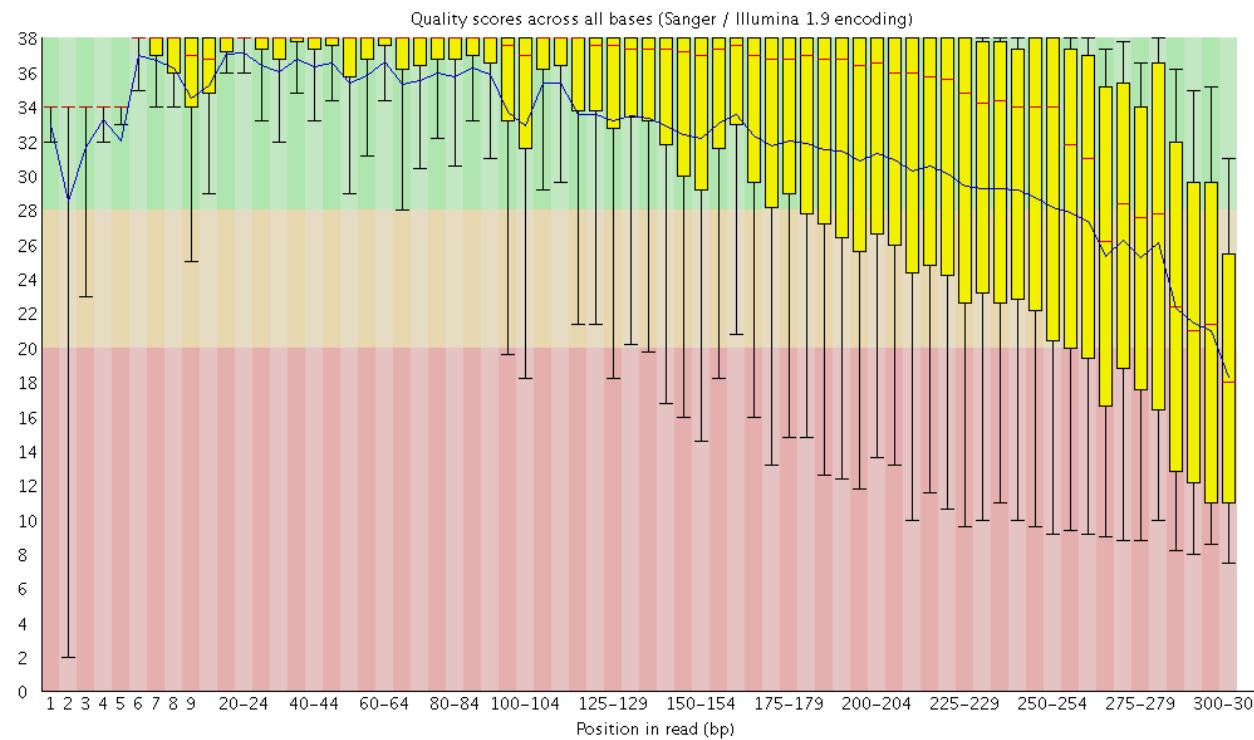
JOIN

gjoinedsample2p3.fastqjoin

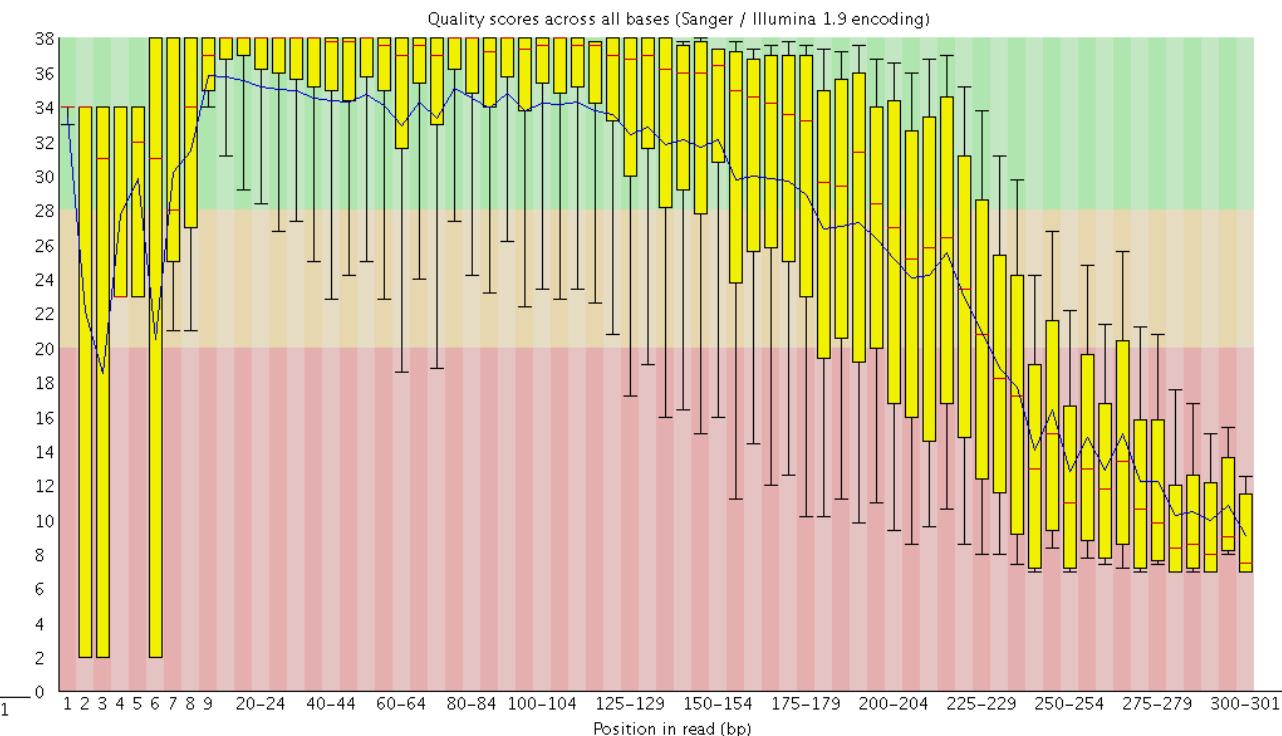


RAWDATA

gut_sample3_pond1_R1.fastq.gz

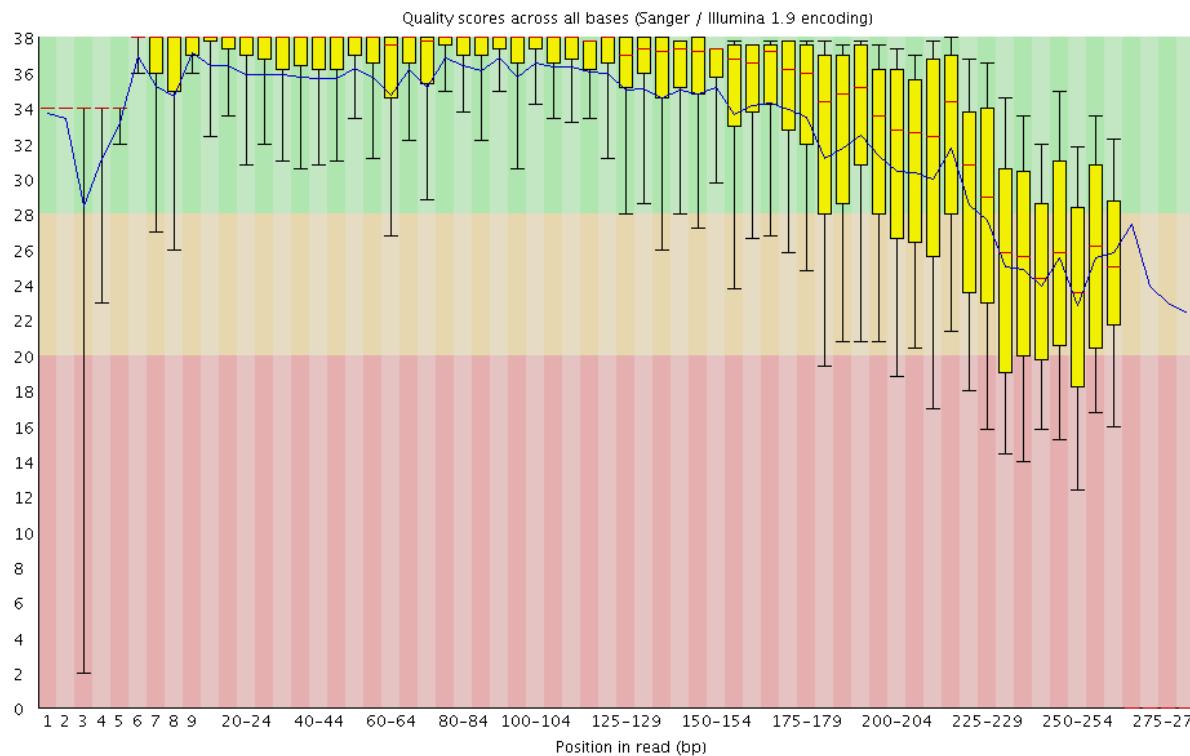


gut_sample3_pond1_R2.fastq.gz

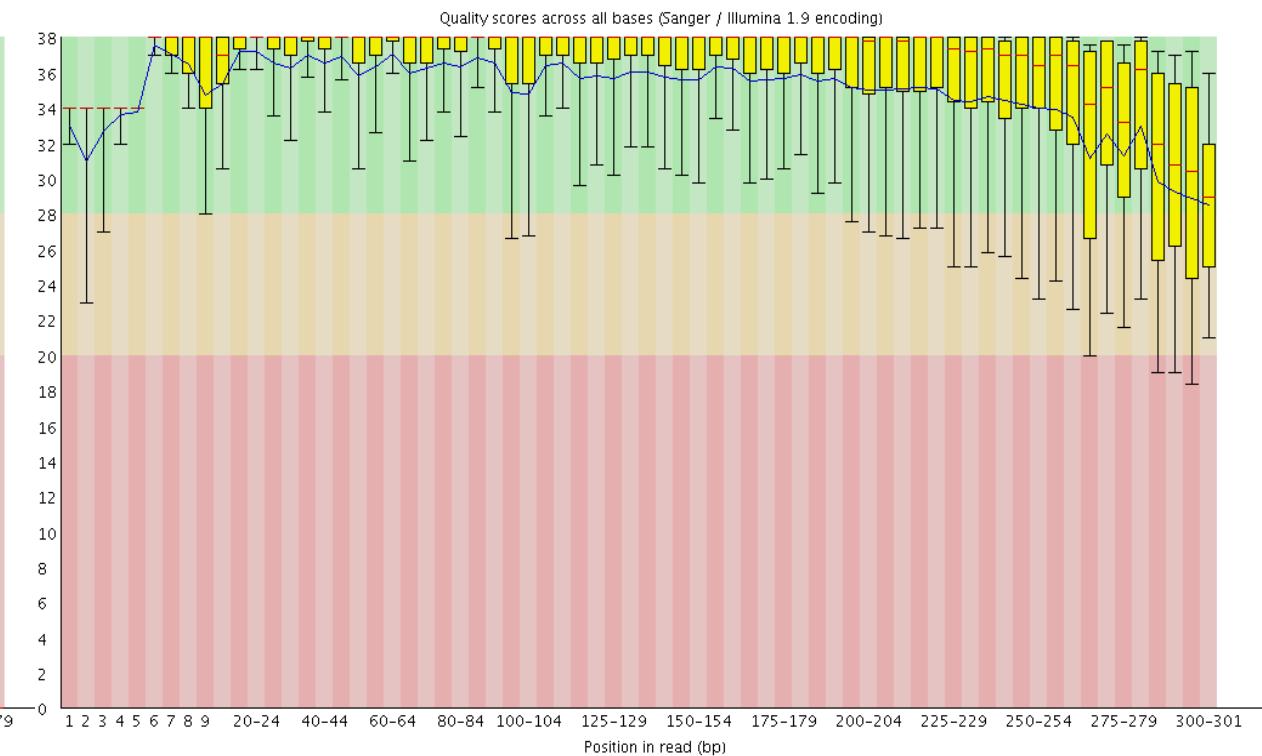


QUALITY

gut_sample3_pond1_R1_paired.fastq.gz

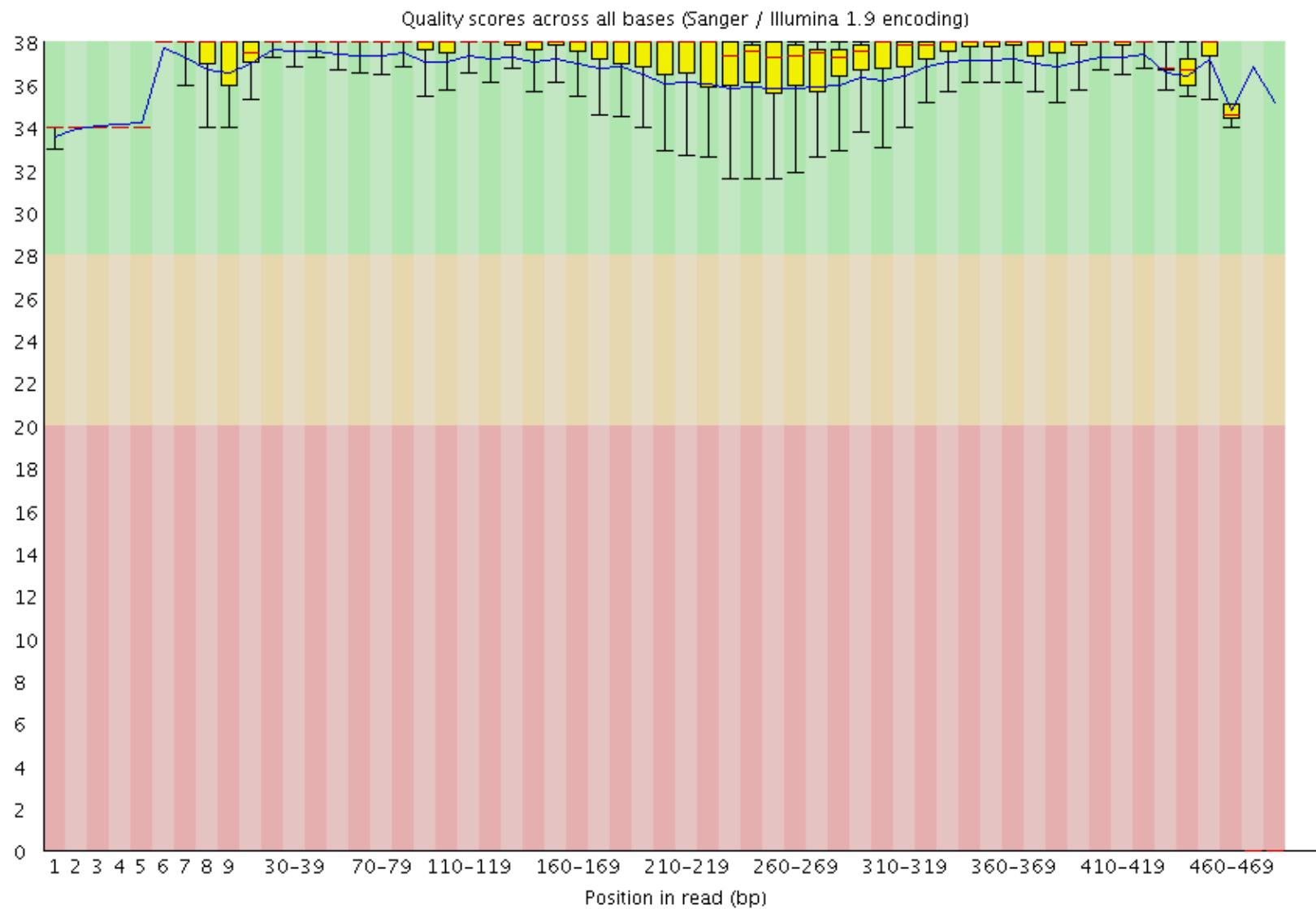


gut_sample3_pond1_R2_paired.fastq.gz



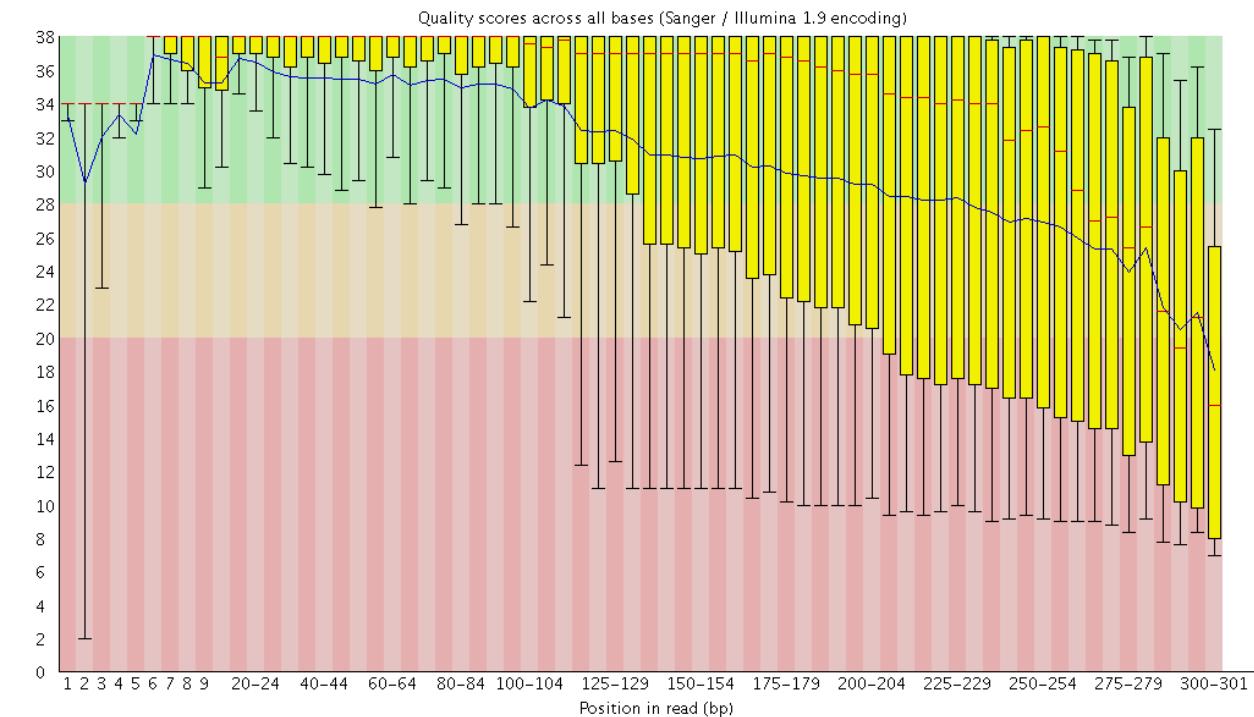
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gjoinedsample3p1.fastqjoin

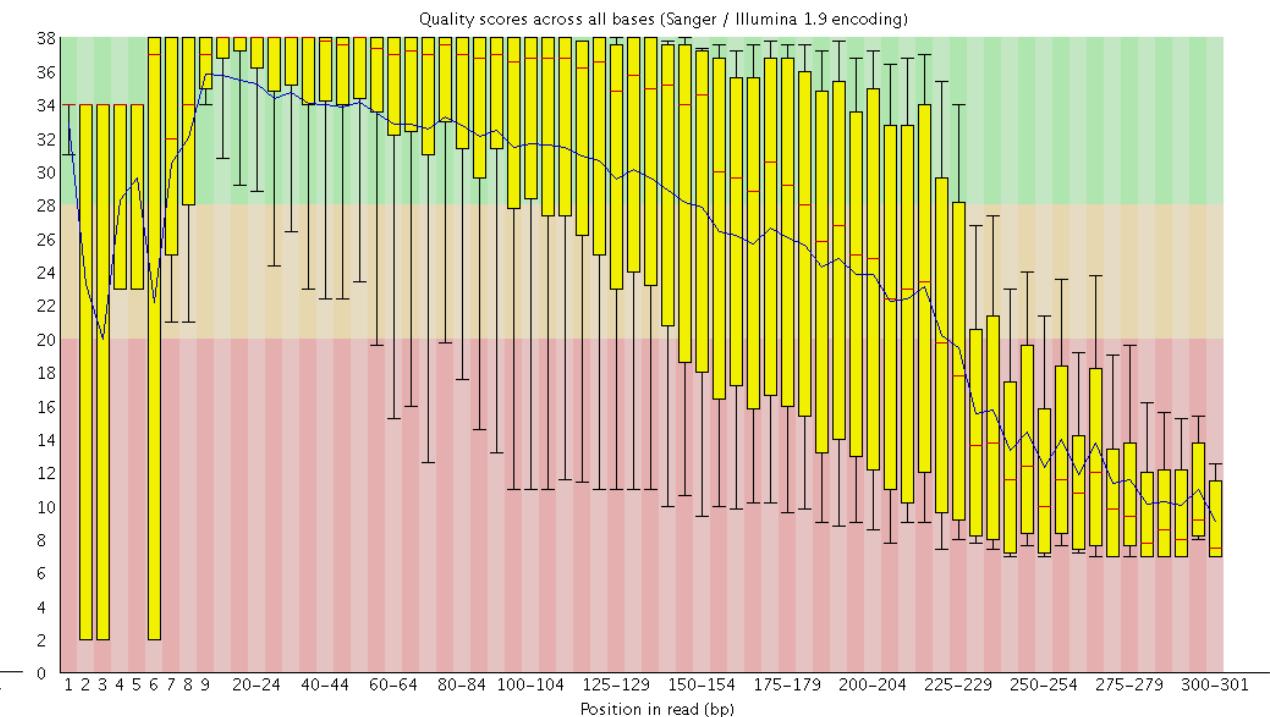


RAWDATA

gut_sample3_pond2_R1.fastq.gz

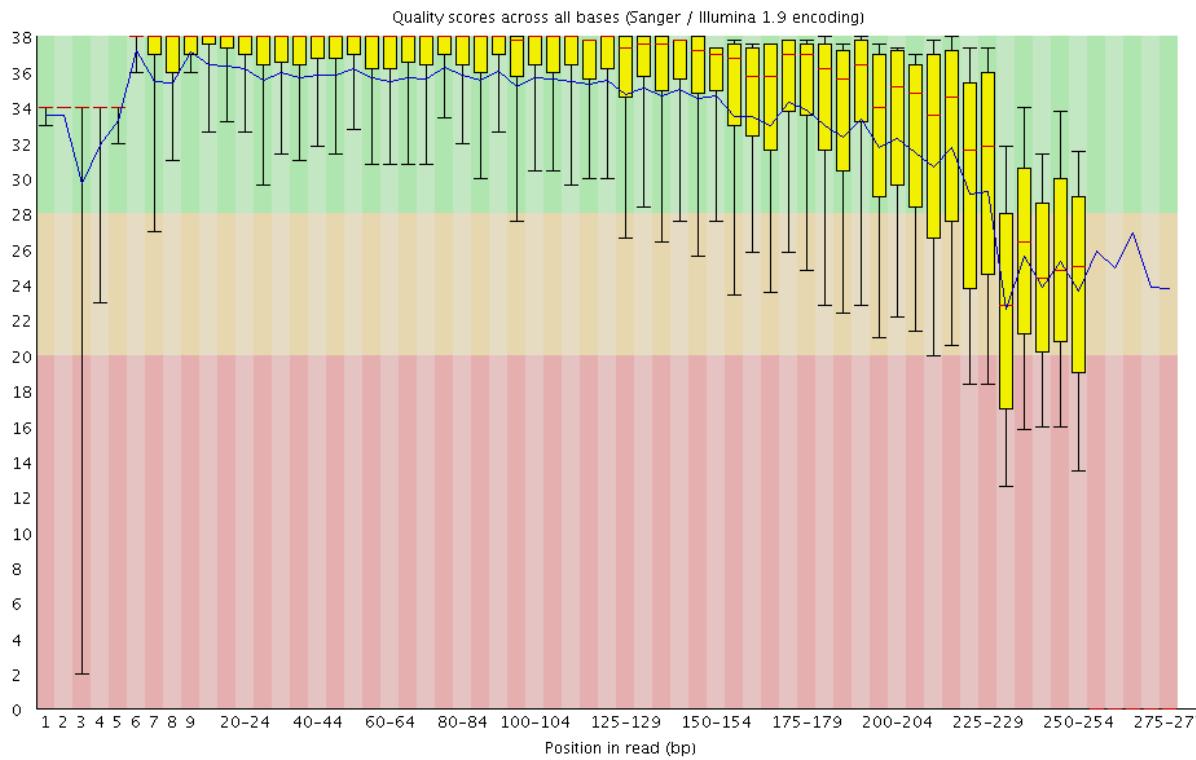


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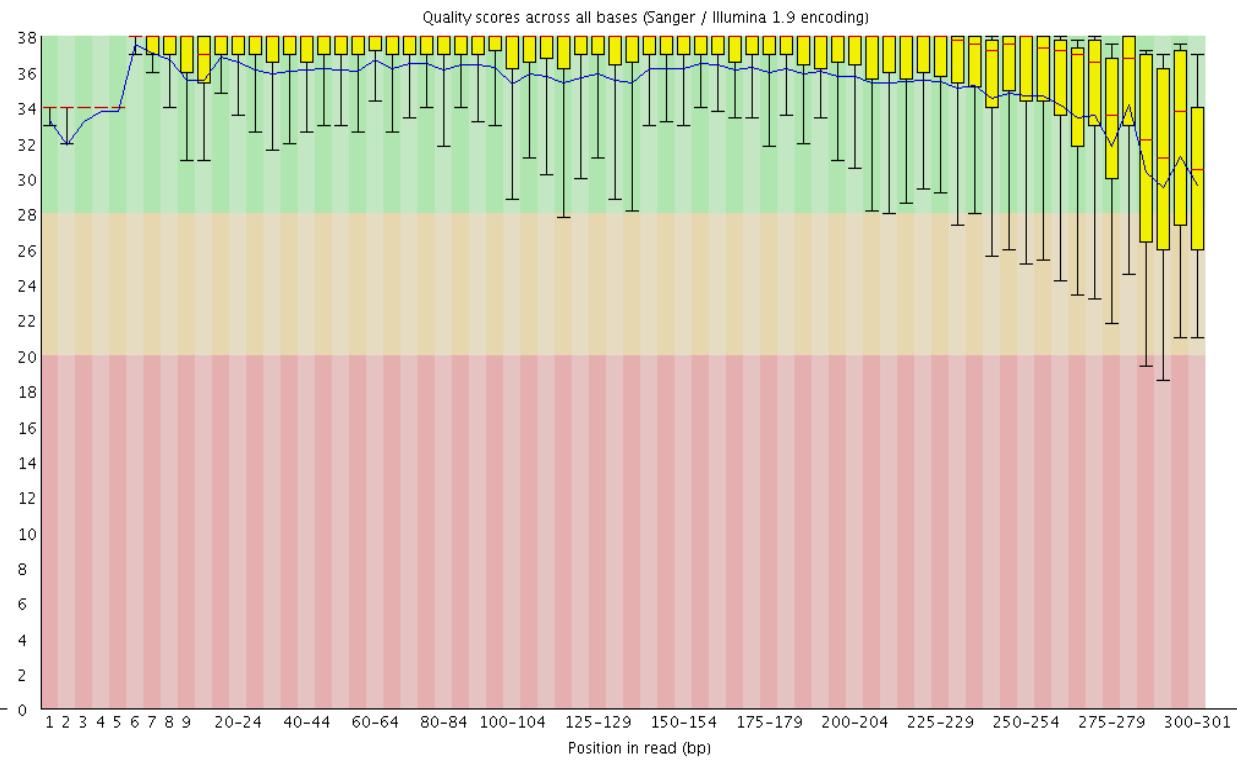


QUALITY

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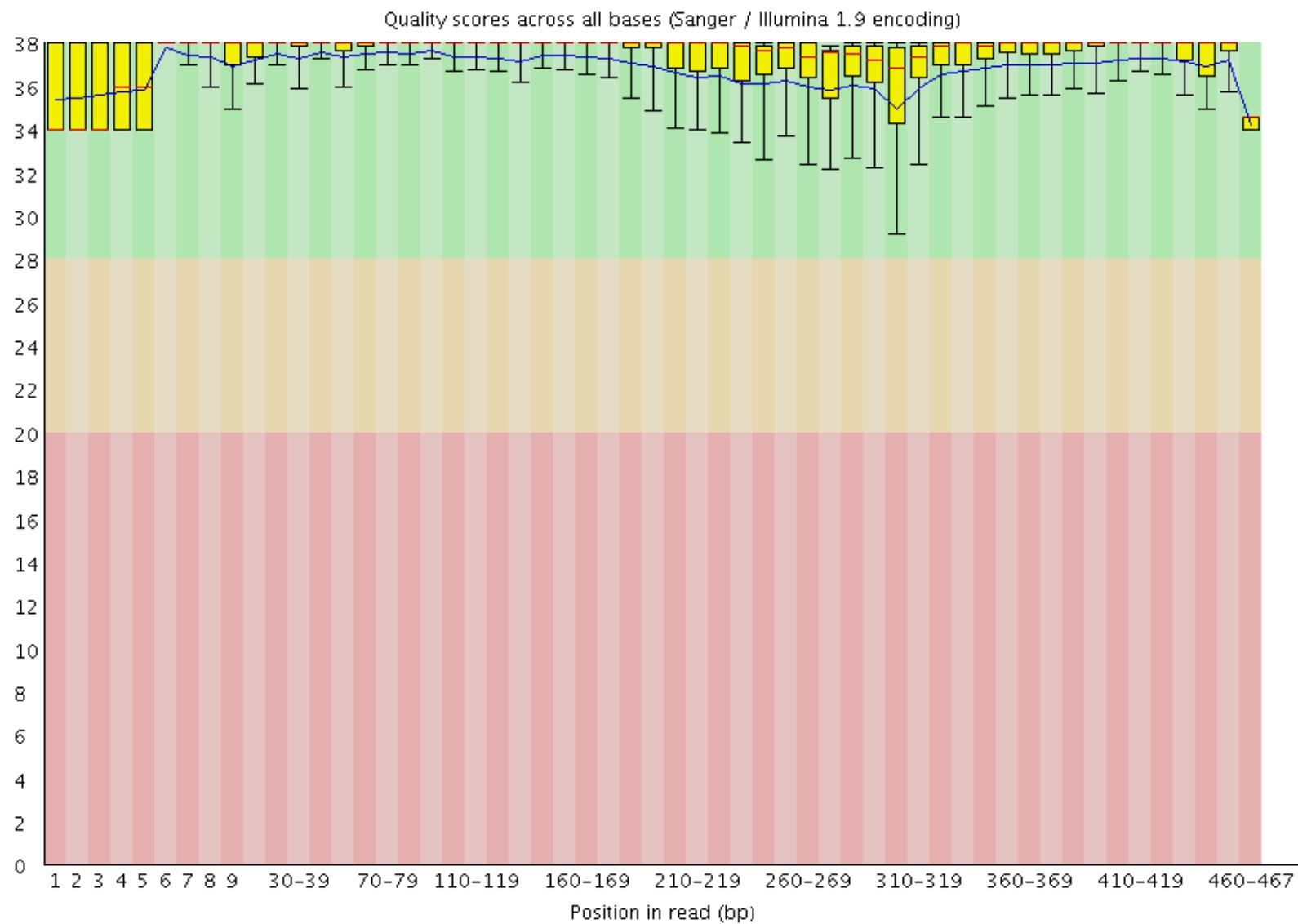


gut_sample3_pond2_R2_paired.fastq.gz



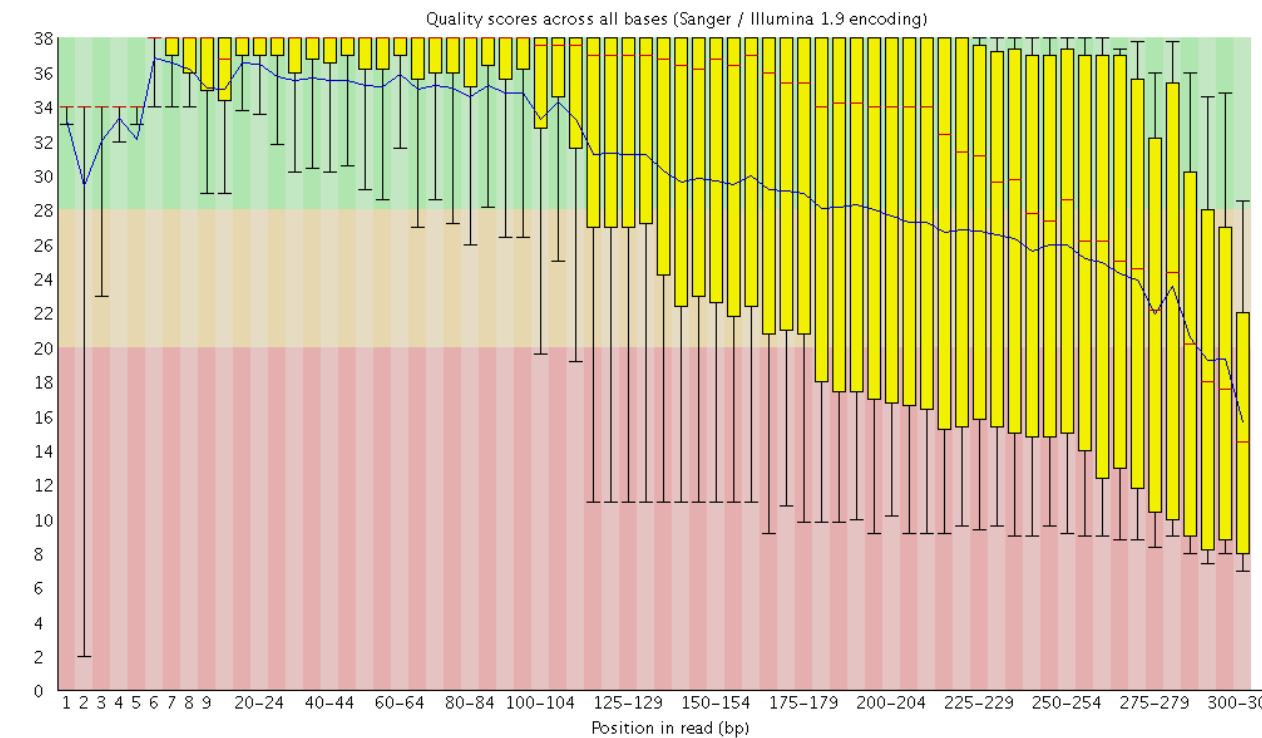
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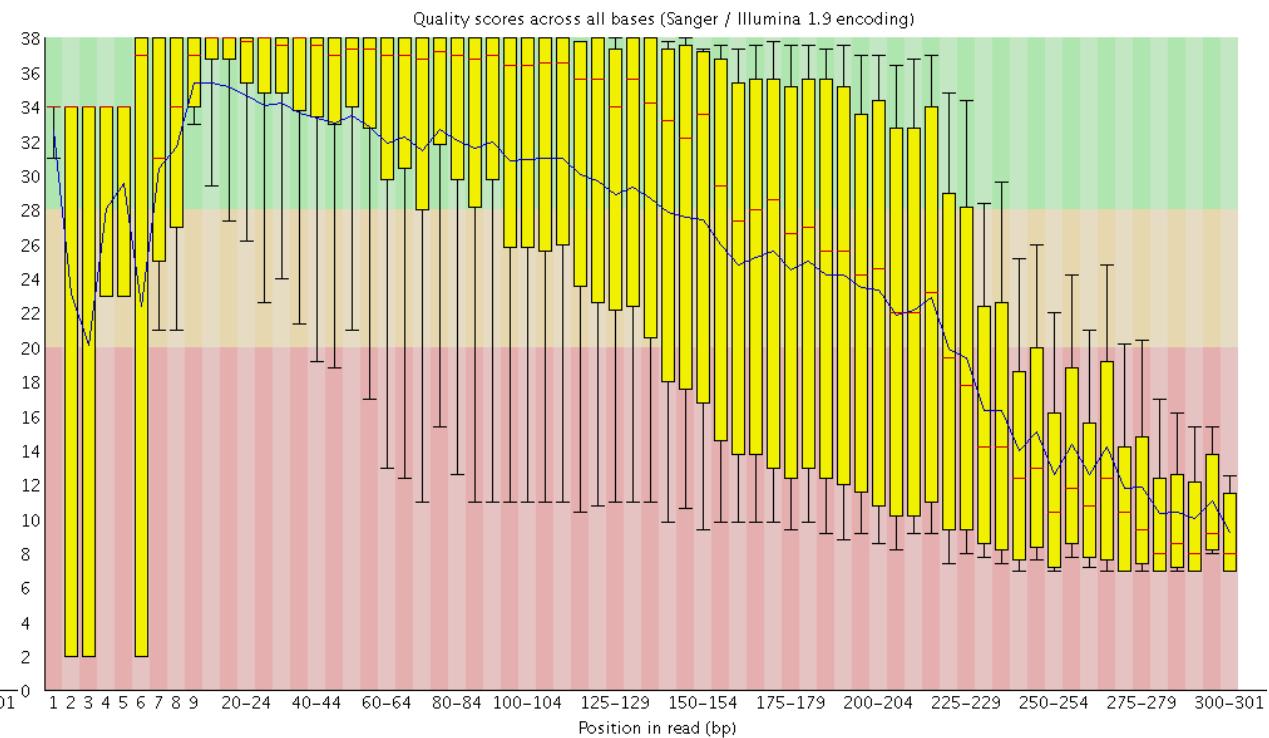


RAWDATA

gut_sample3_pond3_R1.fastq.gz

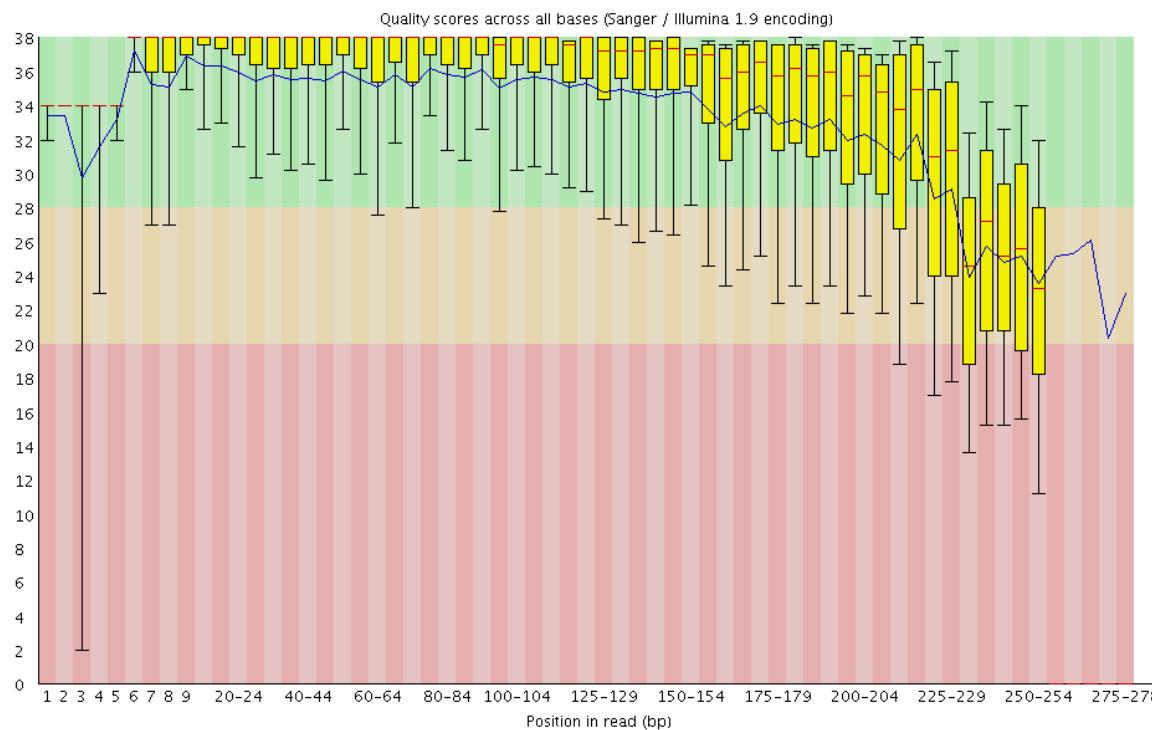


gut_sample3_pond3_R2.fastq.gz

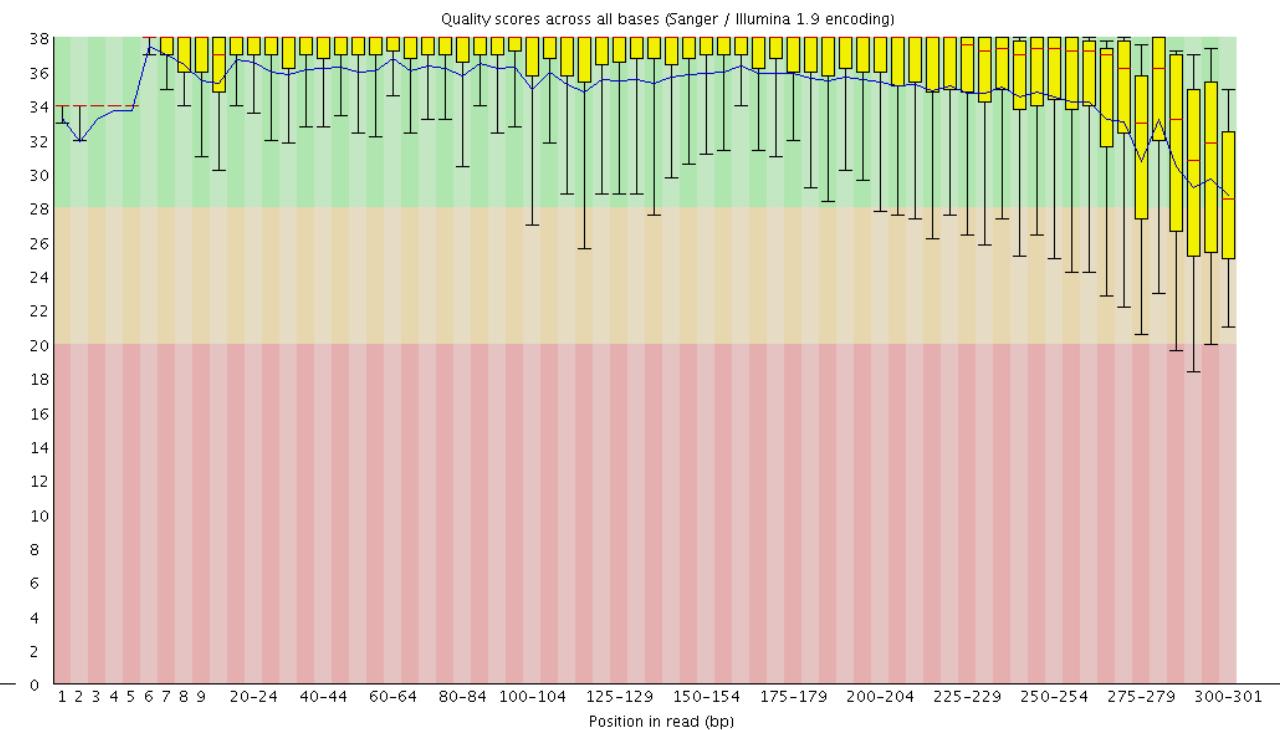


QUALITY

gut_sample3_pond3_R1_paired.fastq.gz

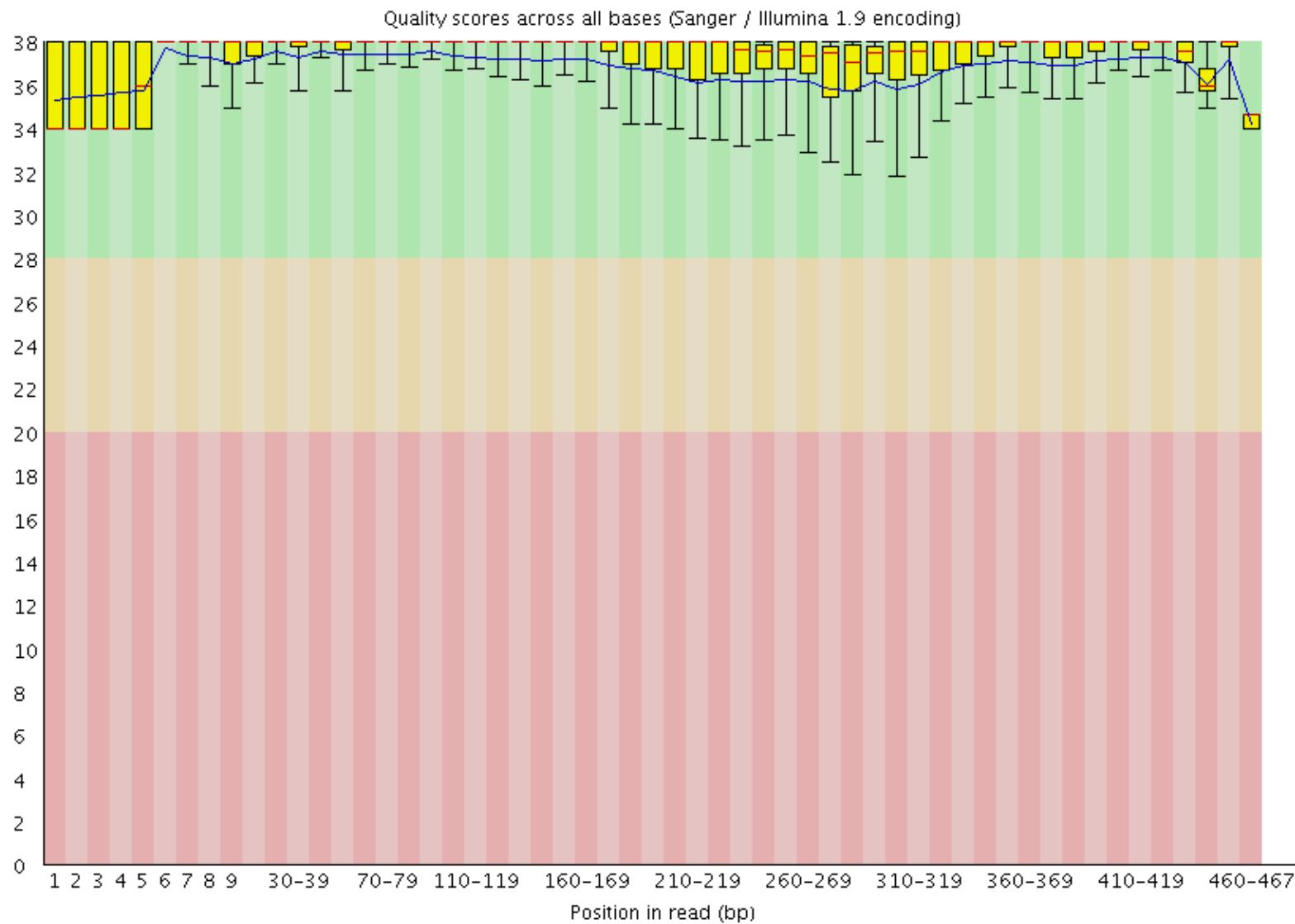


gut_sample3_pond3_R2_paired.fastq.gz



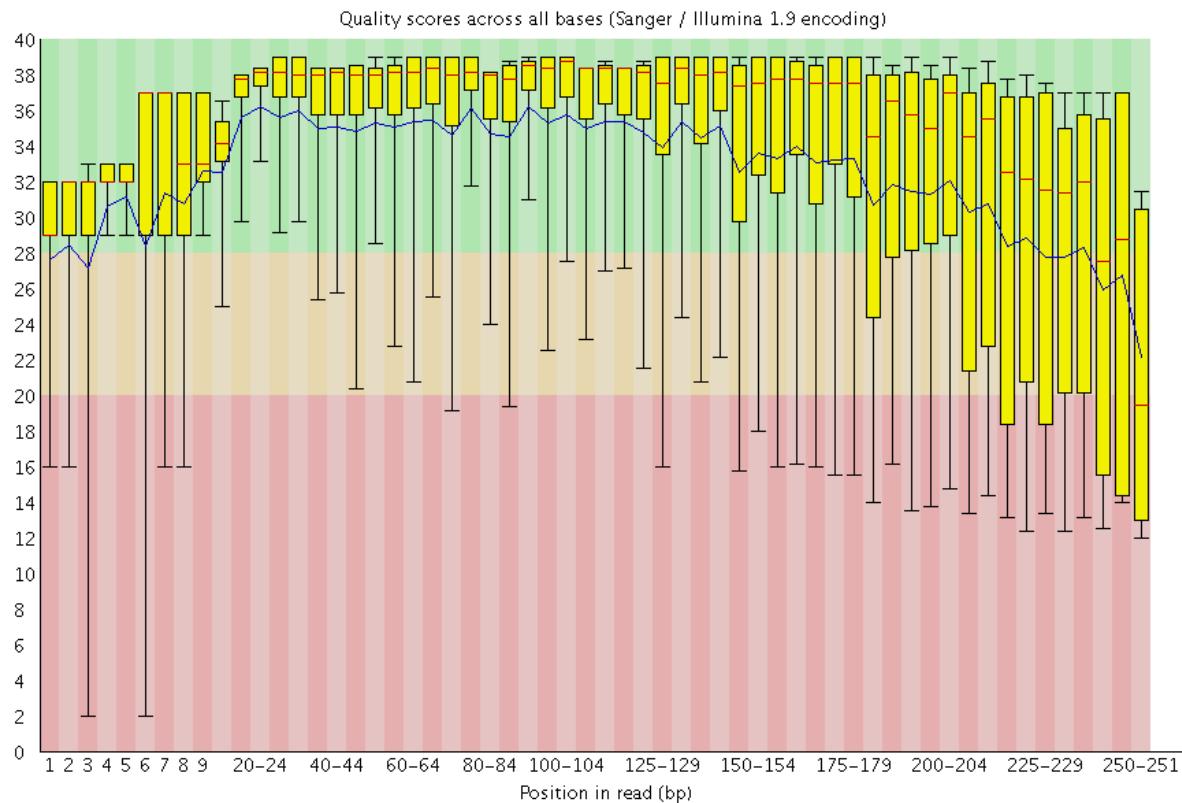
JOIN

gjoinedsample3p3.fastqjoin

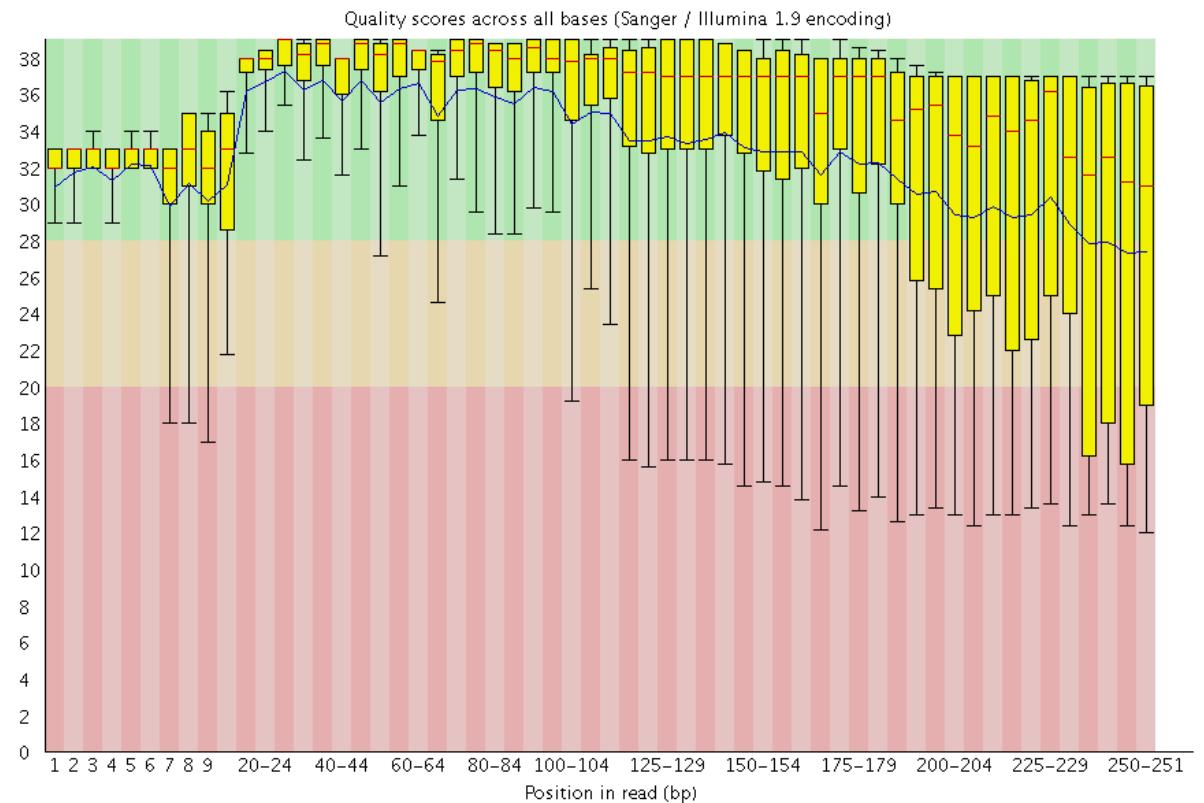


RAWDATA

hep_sample1_pond1_R1.fastq.gz

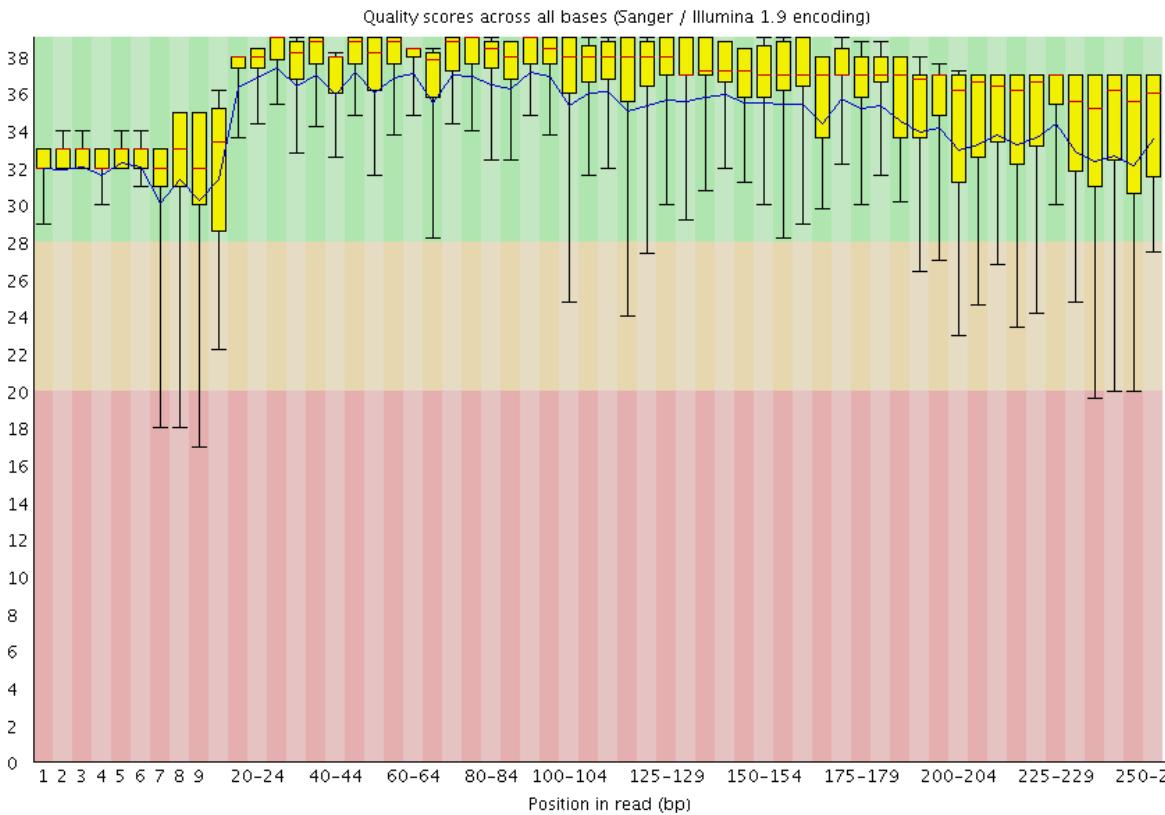


hep_sample1_pond1_R2.fastq.gz

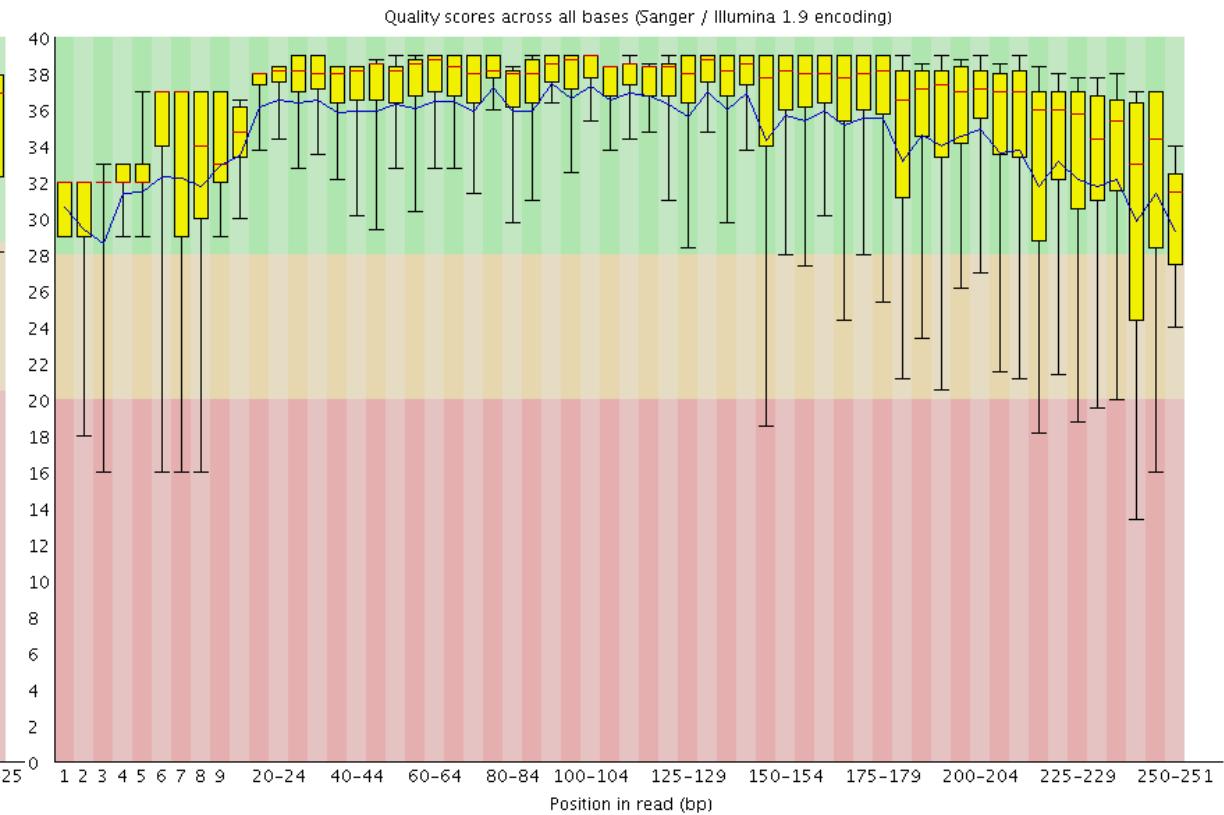


QUALITY

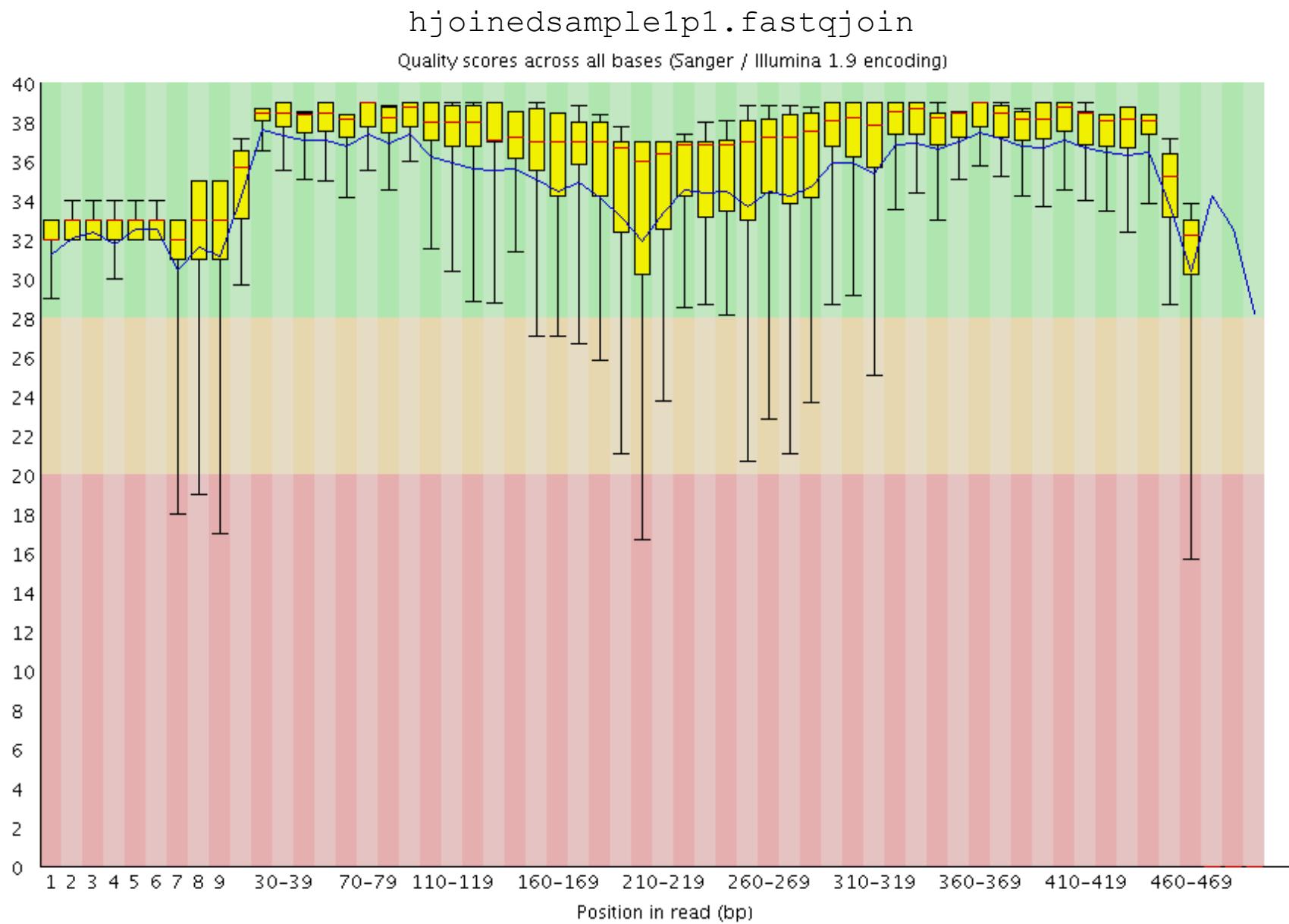
hep_sample1_pond1_R1_paired.fastq.gz



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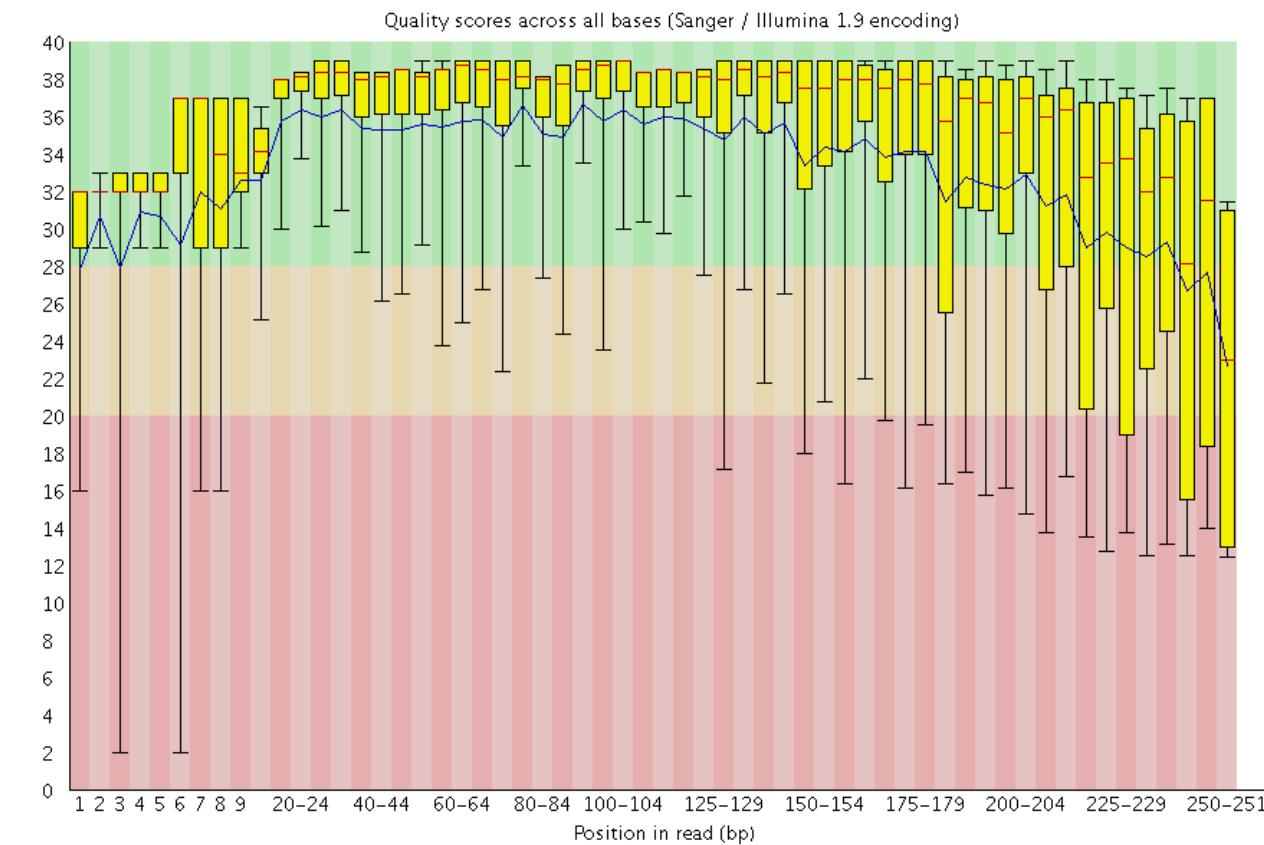


JOIN

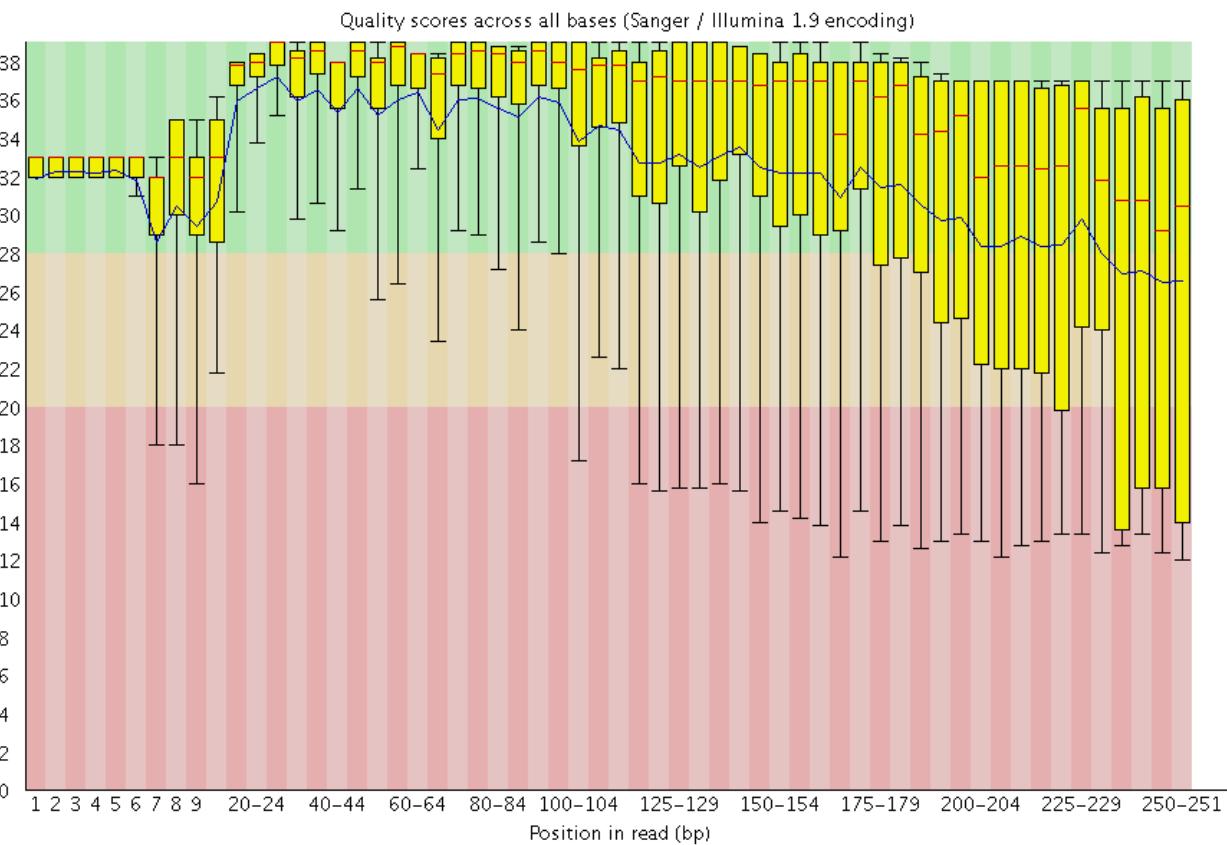


RAWDATA

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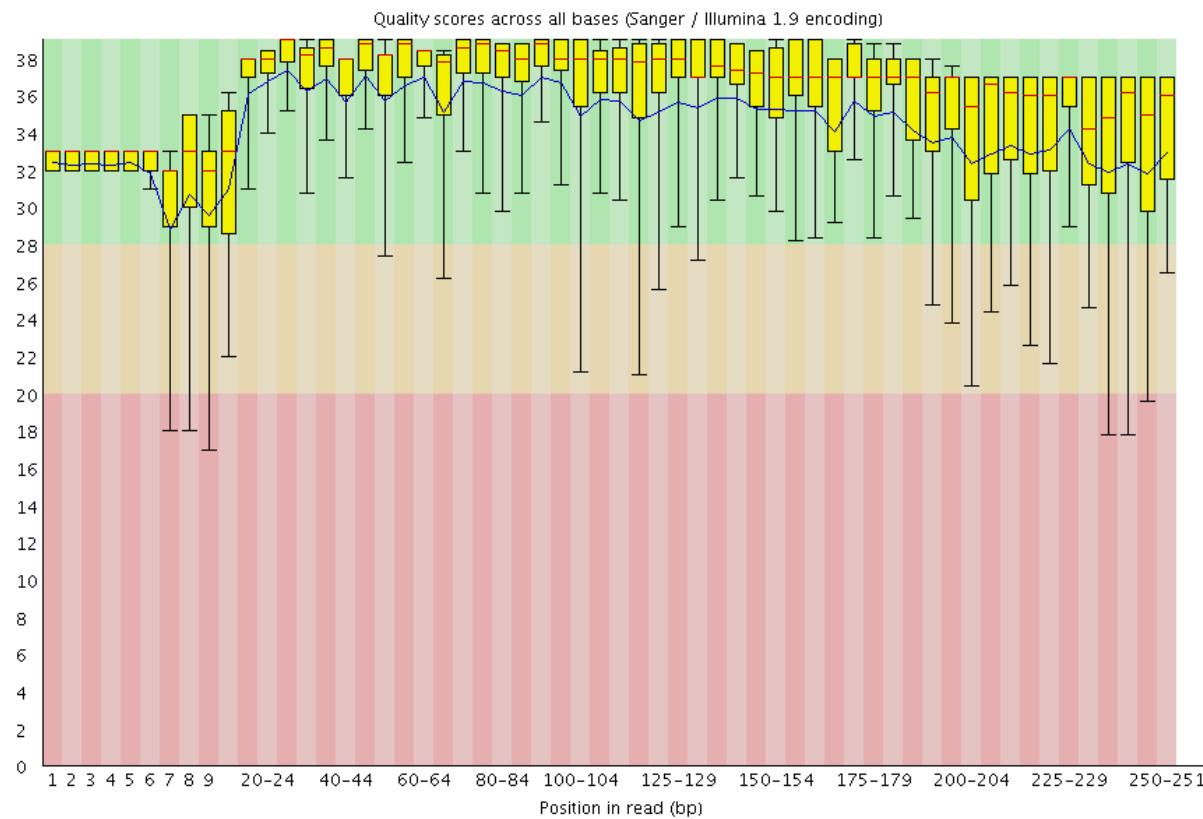


hep_sample1_pond2_R2.fastq.gz

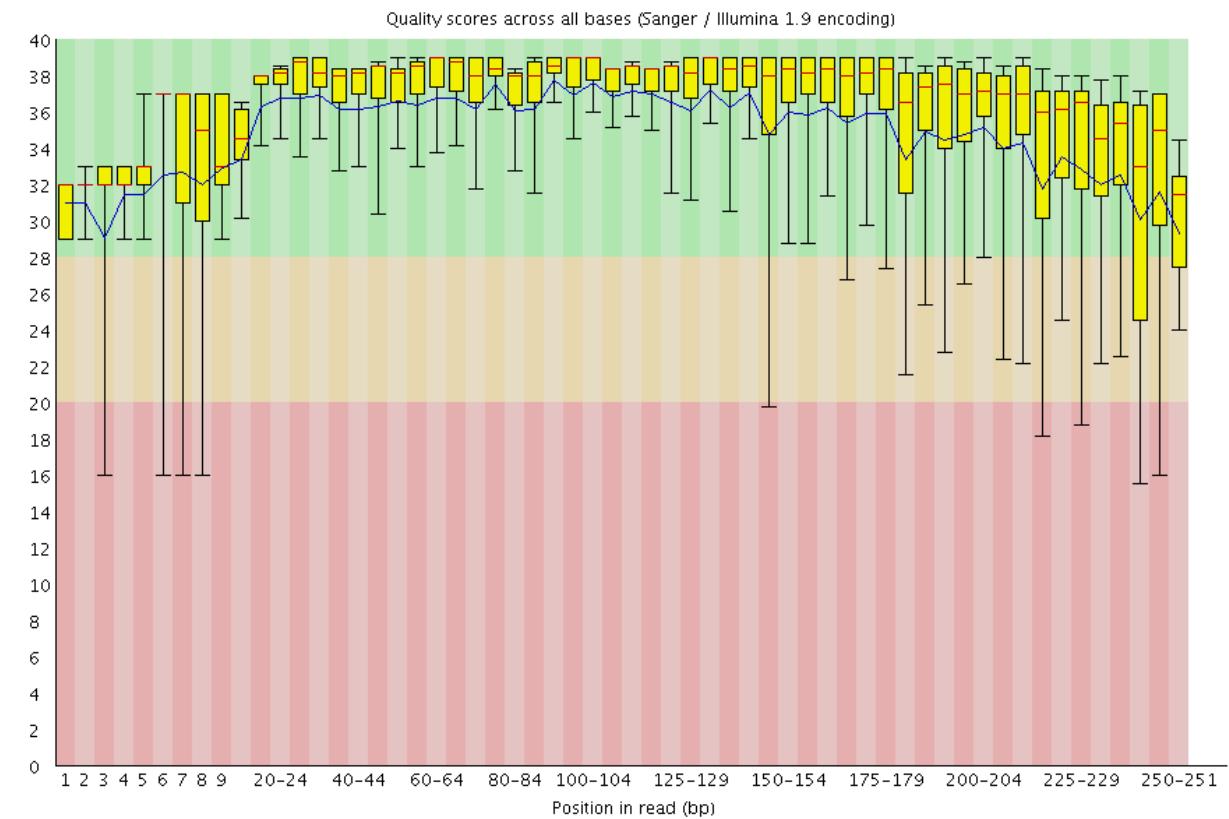


QUALITY

hep_sample1_pond2_R1_paired.fastq.gz



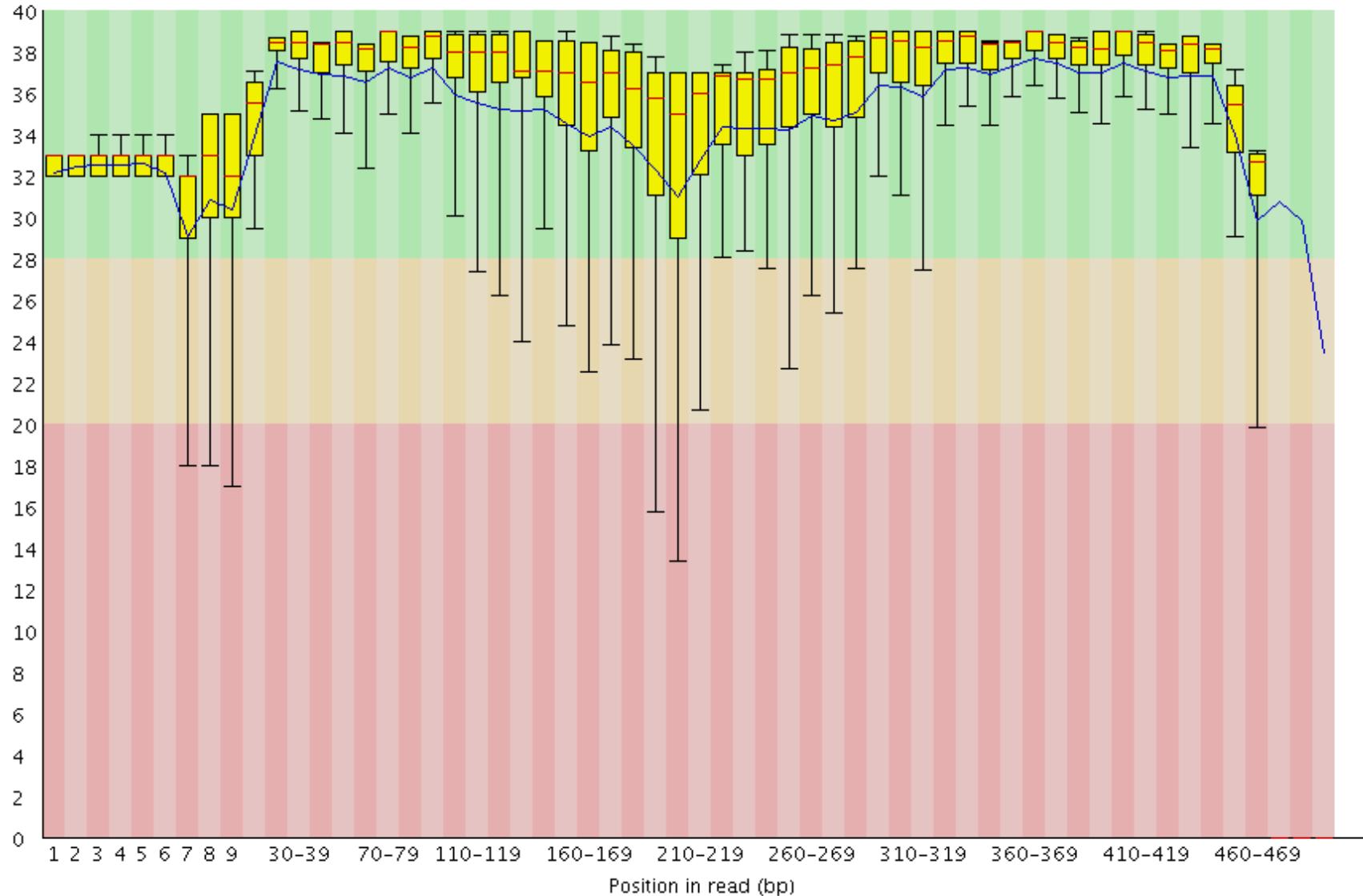
hep_sample1_pond2_R2_paired.fastq.gz



JOIN

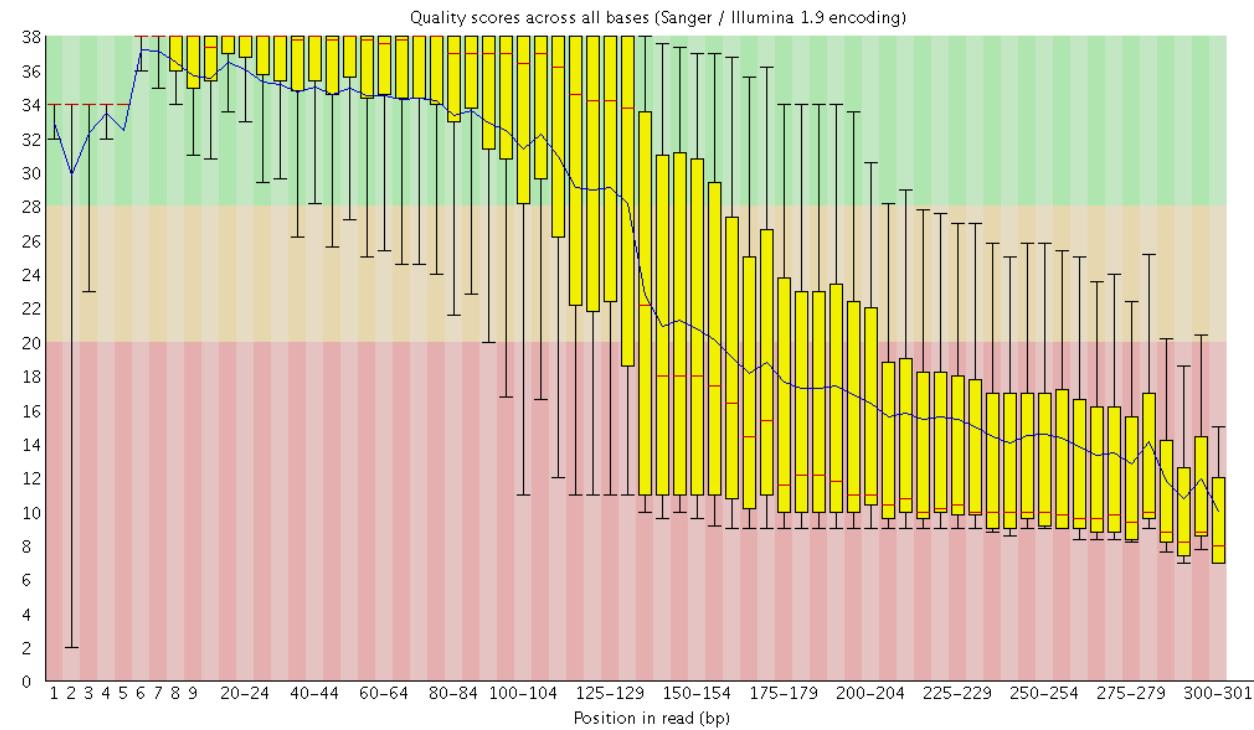
hjoinedsample1p2.fastqjoin

Quality scores across all bases (Sanger / Illumina 1.9 encoding)

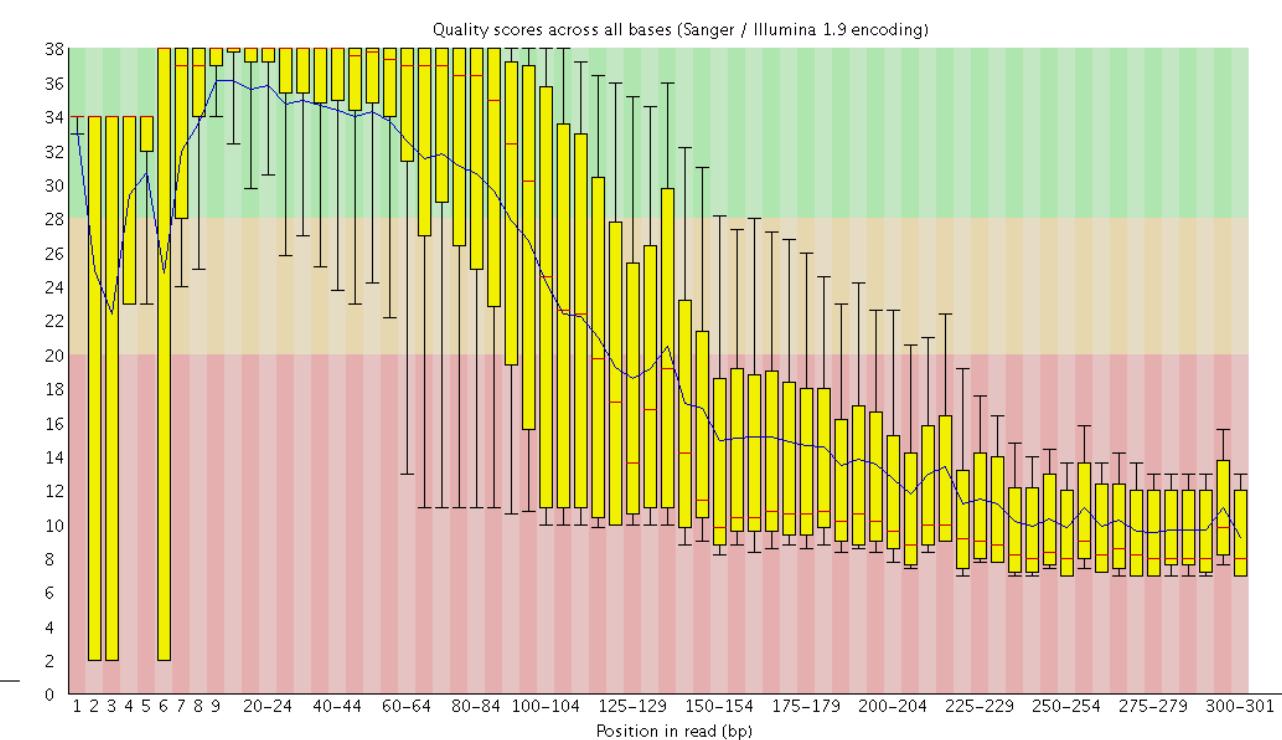


RAWDATA

hep_sample1_pond3_R1.fastq.gz

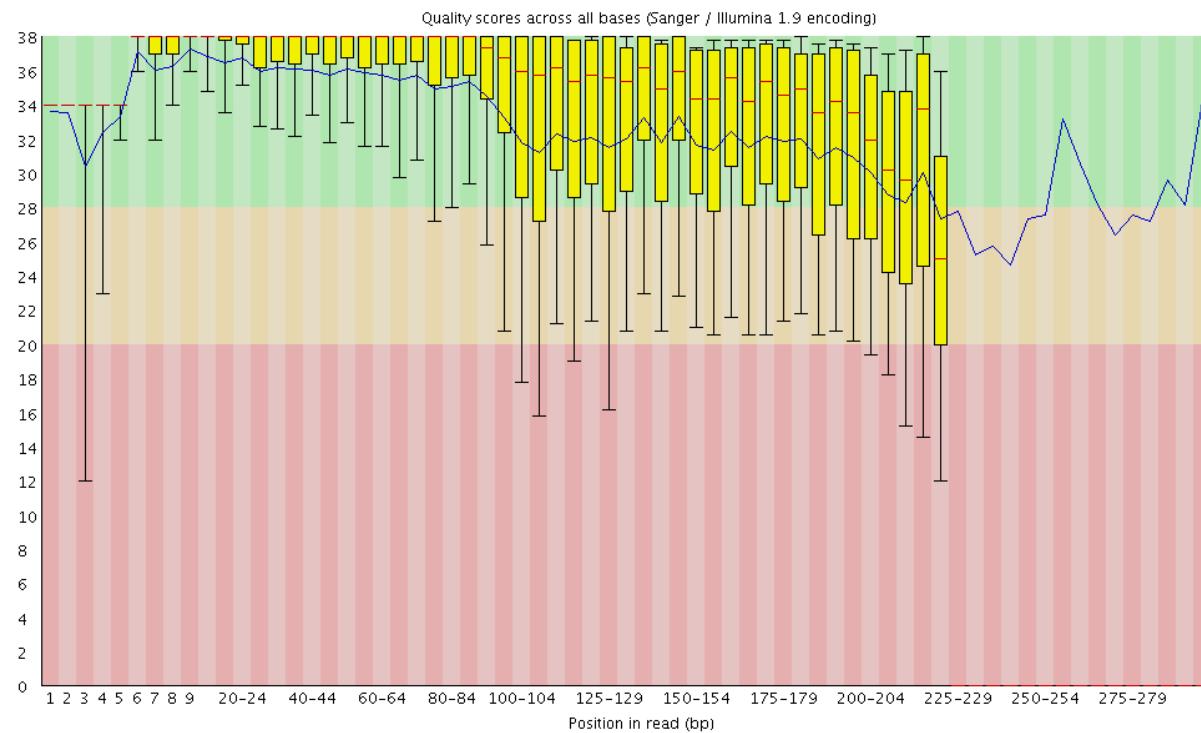


hep_sample1_pond3_R2.fastq.gz

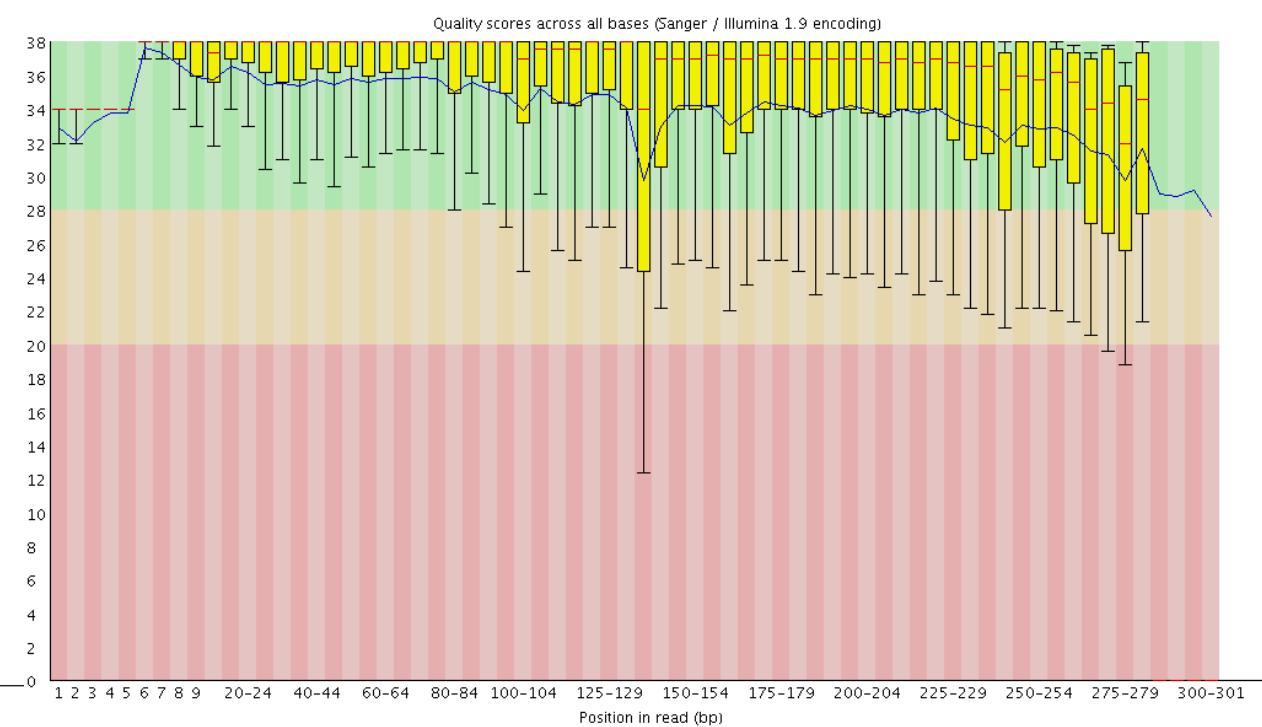


QUALITY

hep_sample1_pond3_R1_paired.fastq.gz

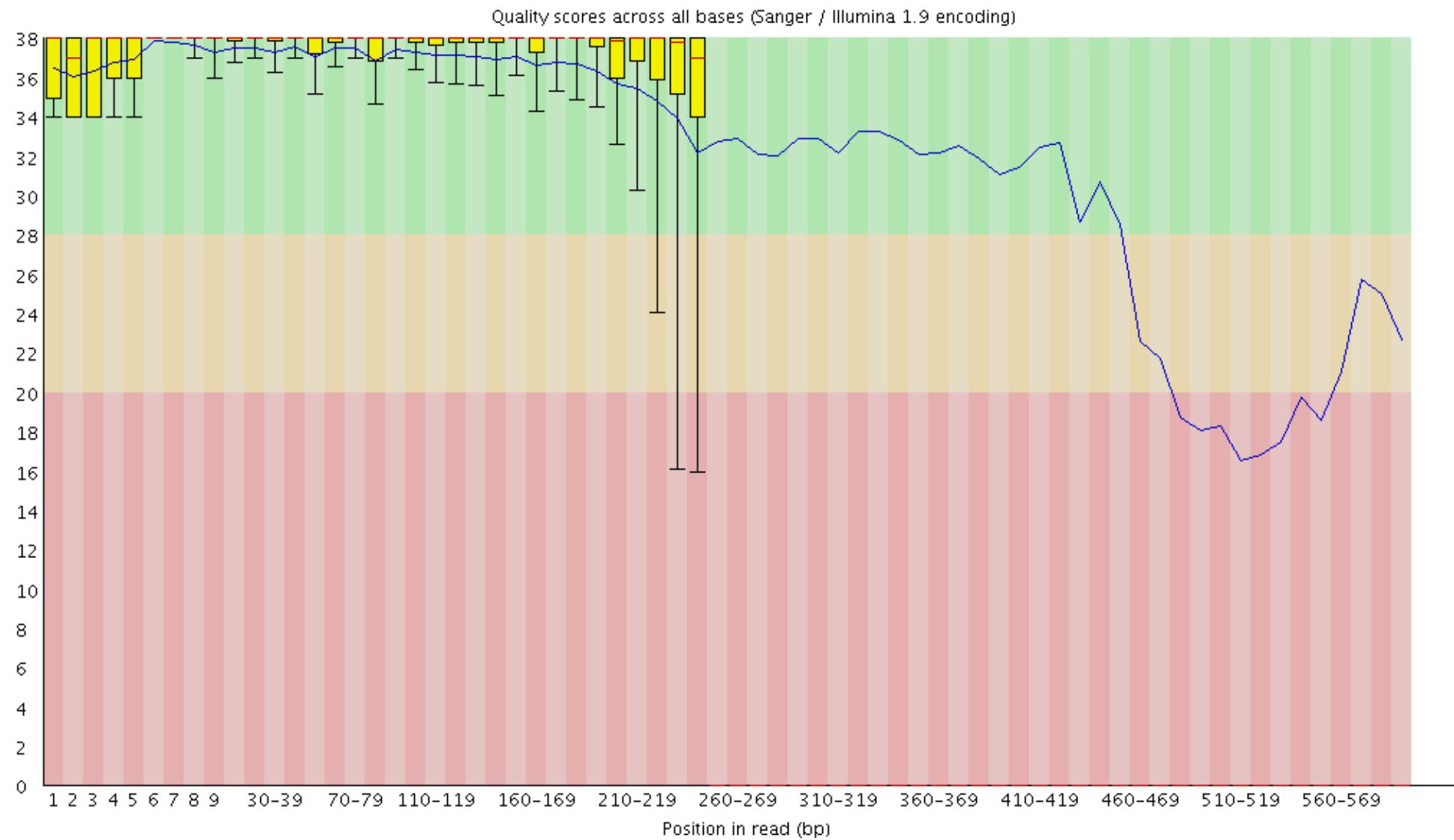


hep_sample1_pond3_R2_paired.fastq.gz



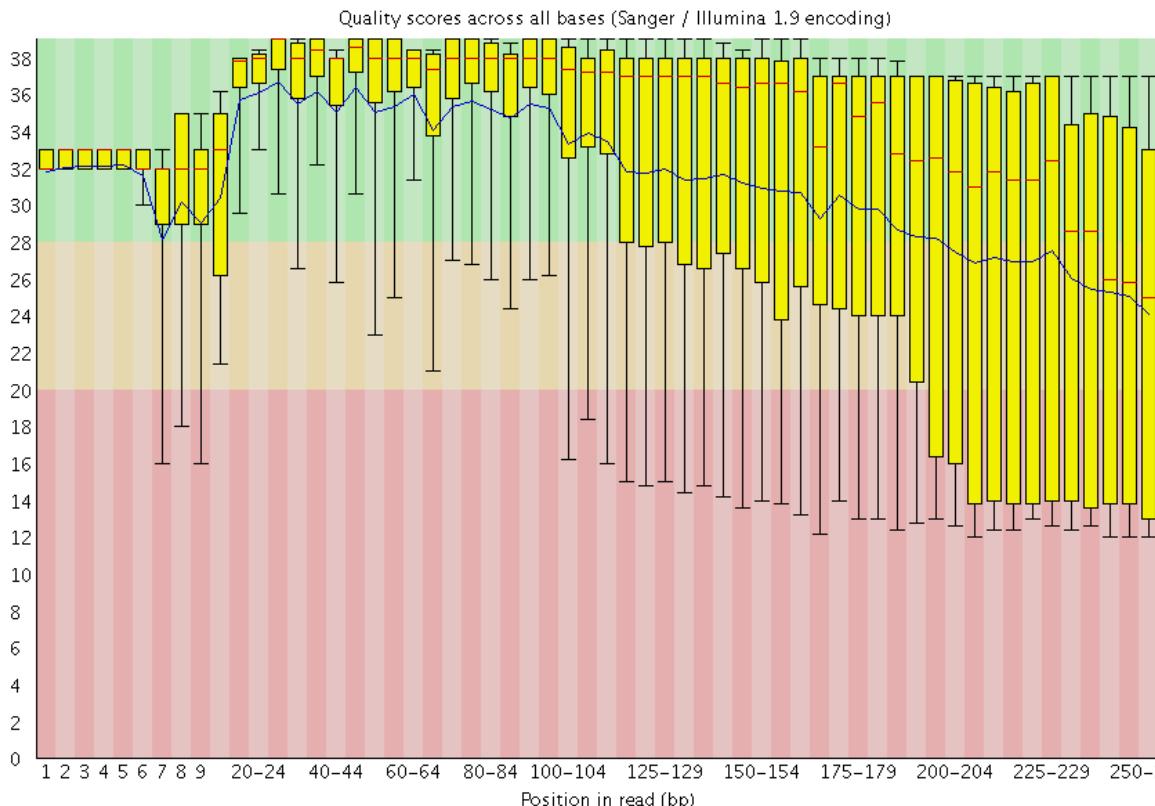
JOIN

hjoinedsample1p3.fastqjoin

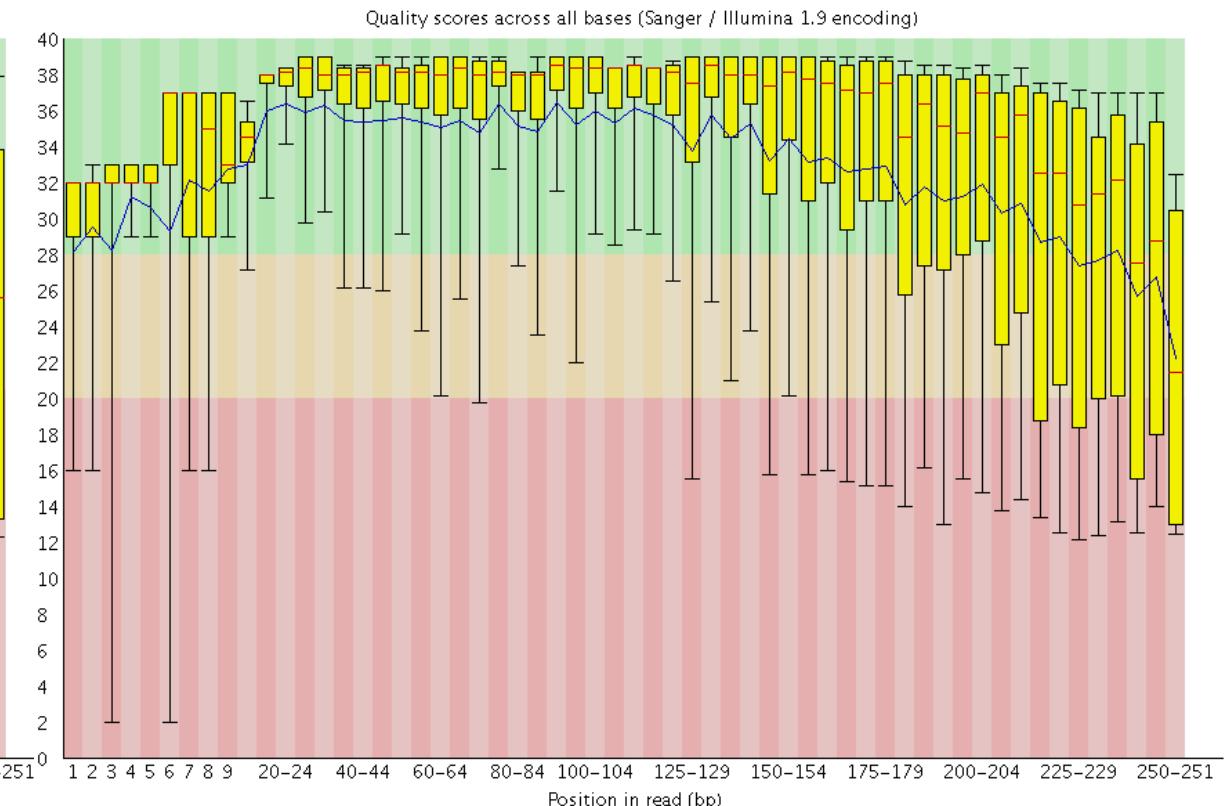


RAWDATA

hep_sample2_pond1_R1.fastq.gz

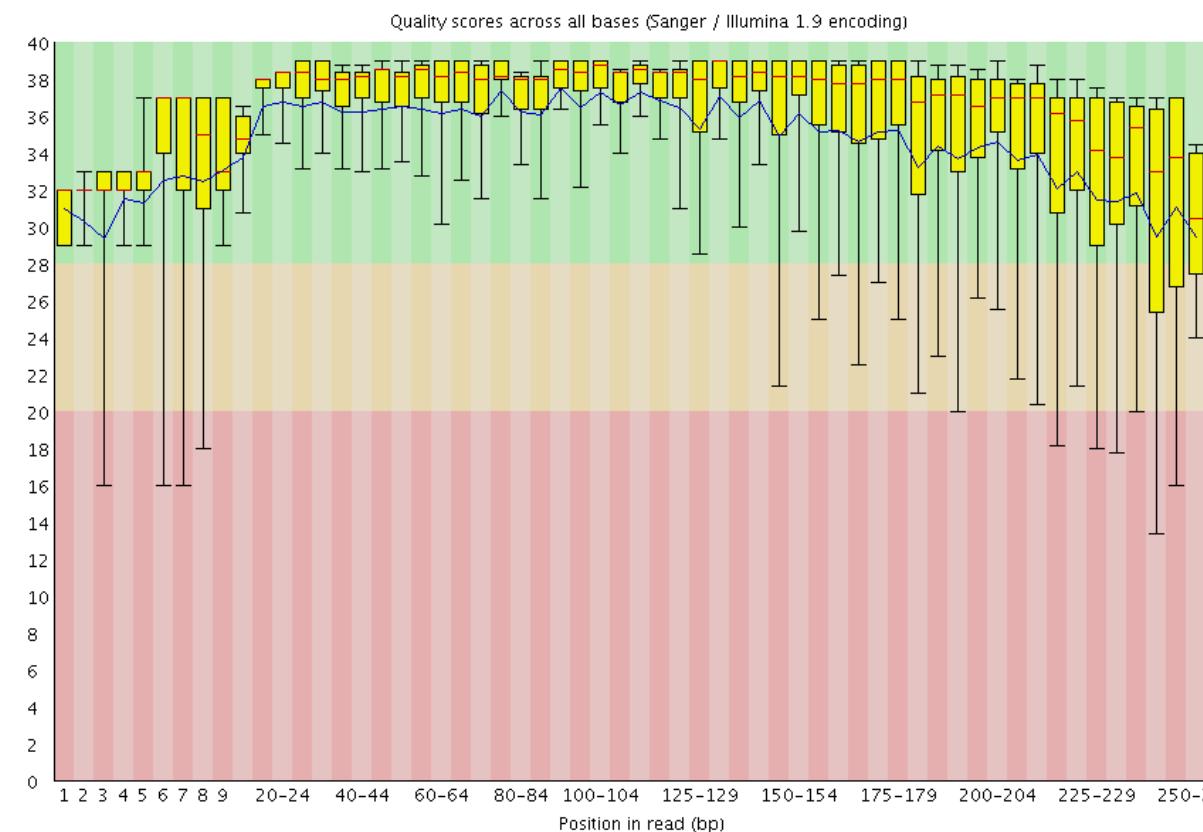


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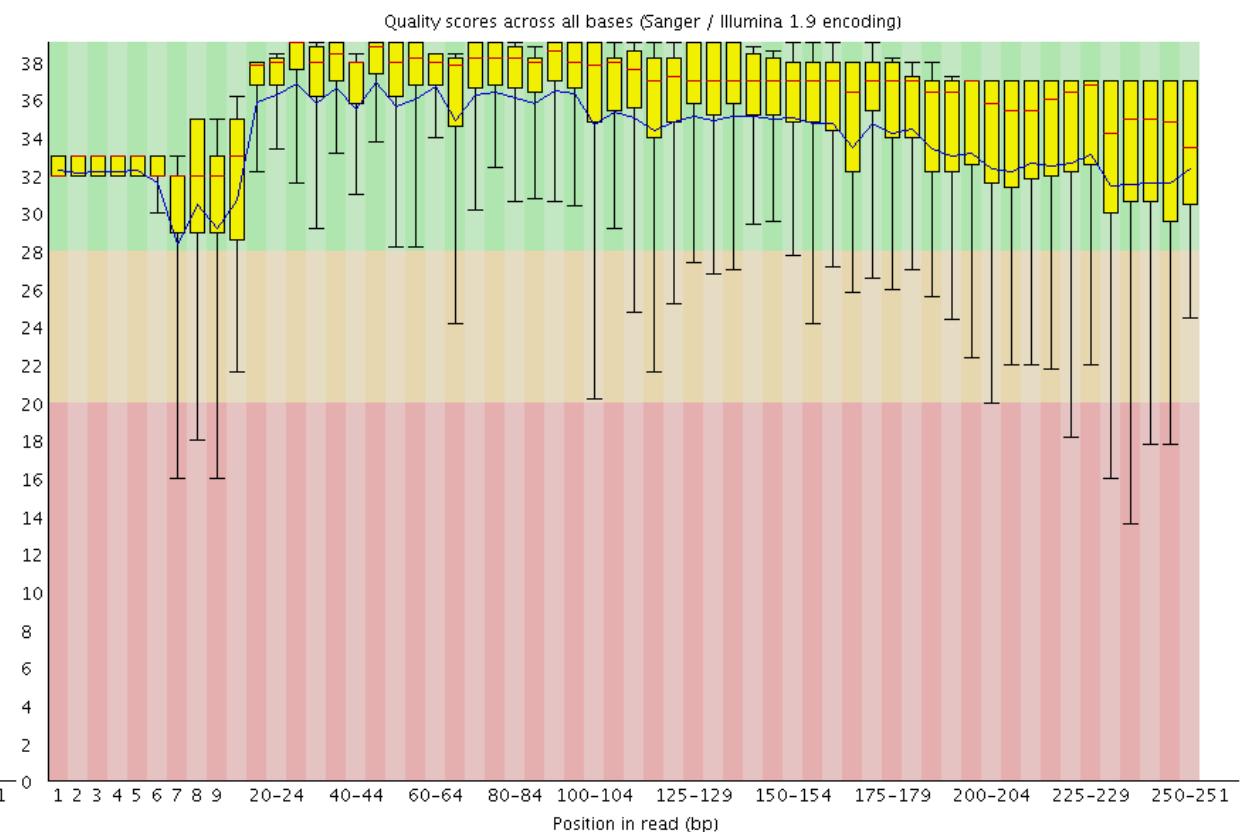


QUALITY

hep_sample2_pond1_R1_paired.fastq.gz

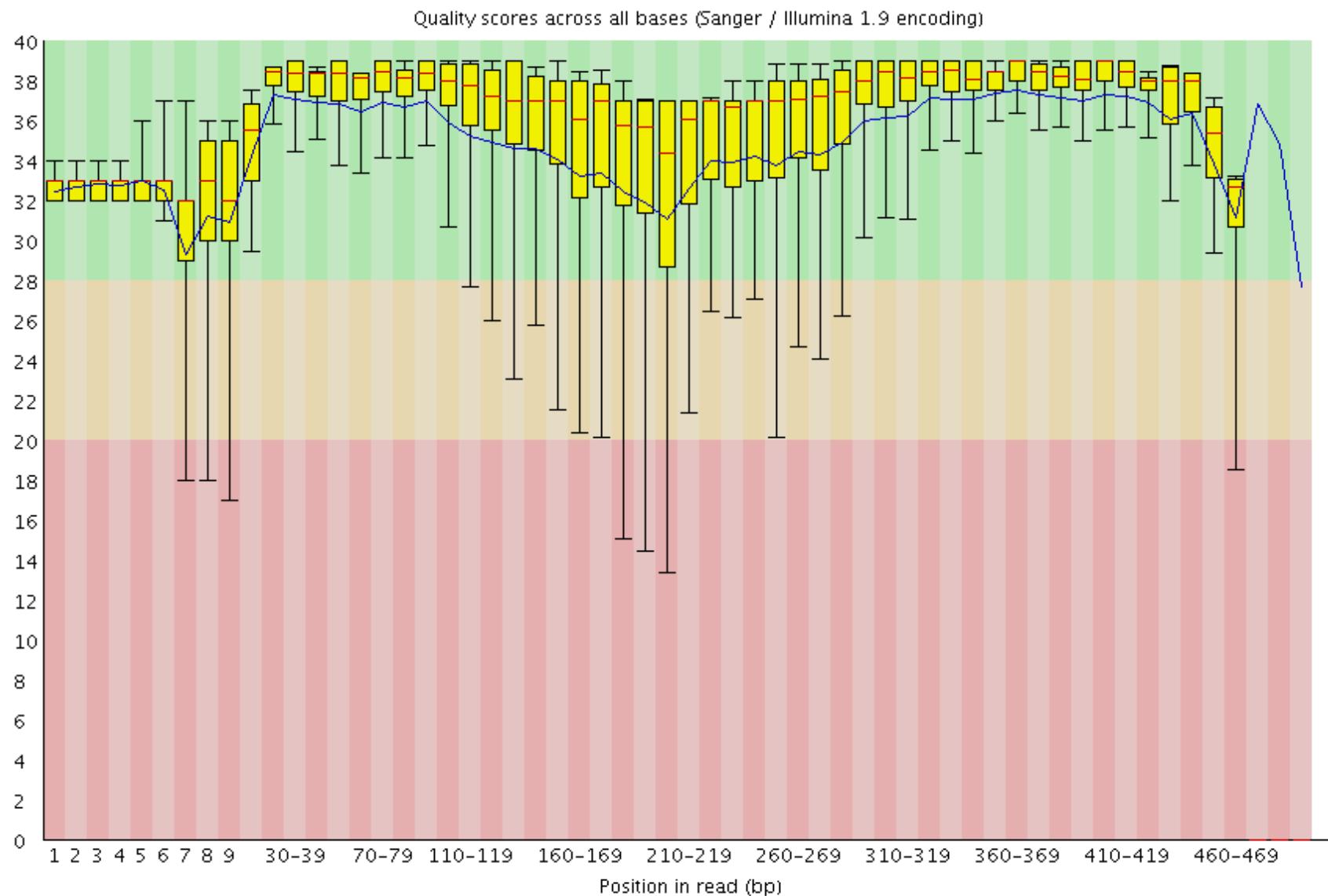


hep_sample2_pond1_R2_paired.fastq.gz



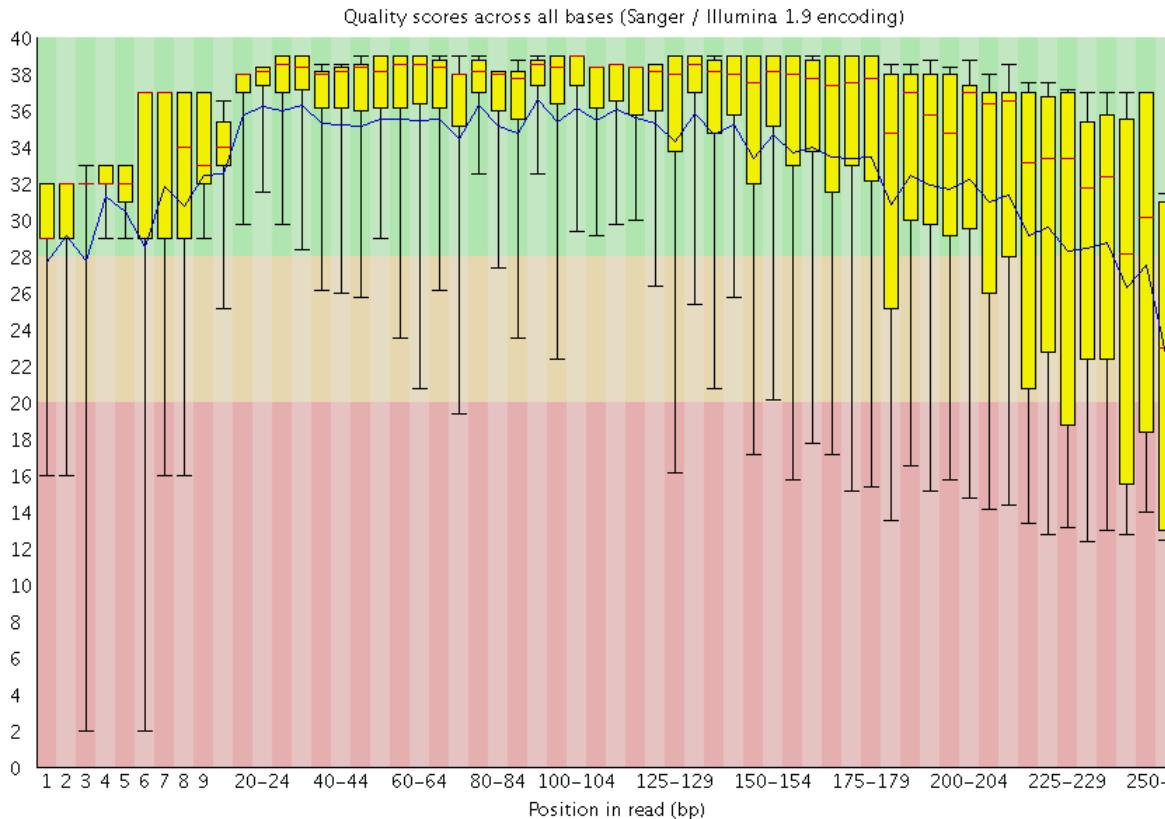
JOIN

hjoinedsample2p1.fastqjoin

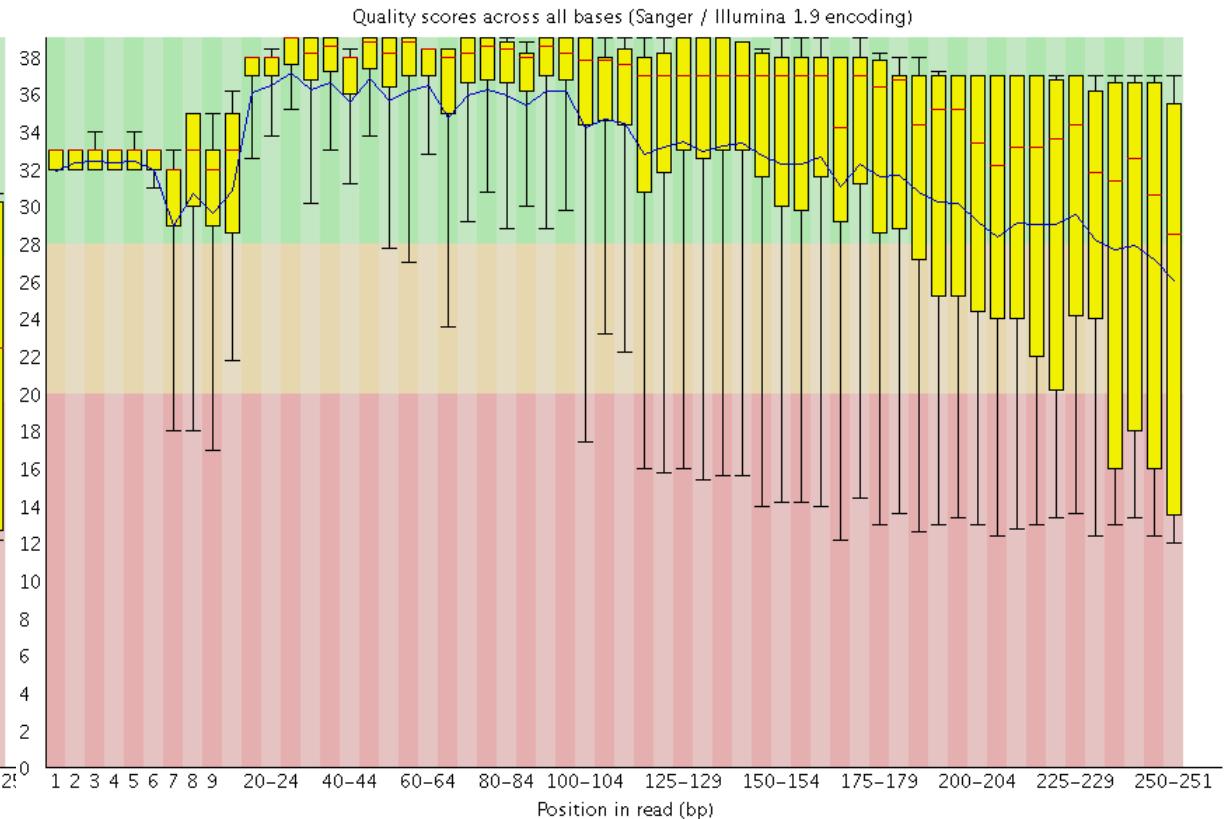


RAWDATA

hep_sample2_pond2_R2.fastq.gz

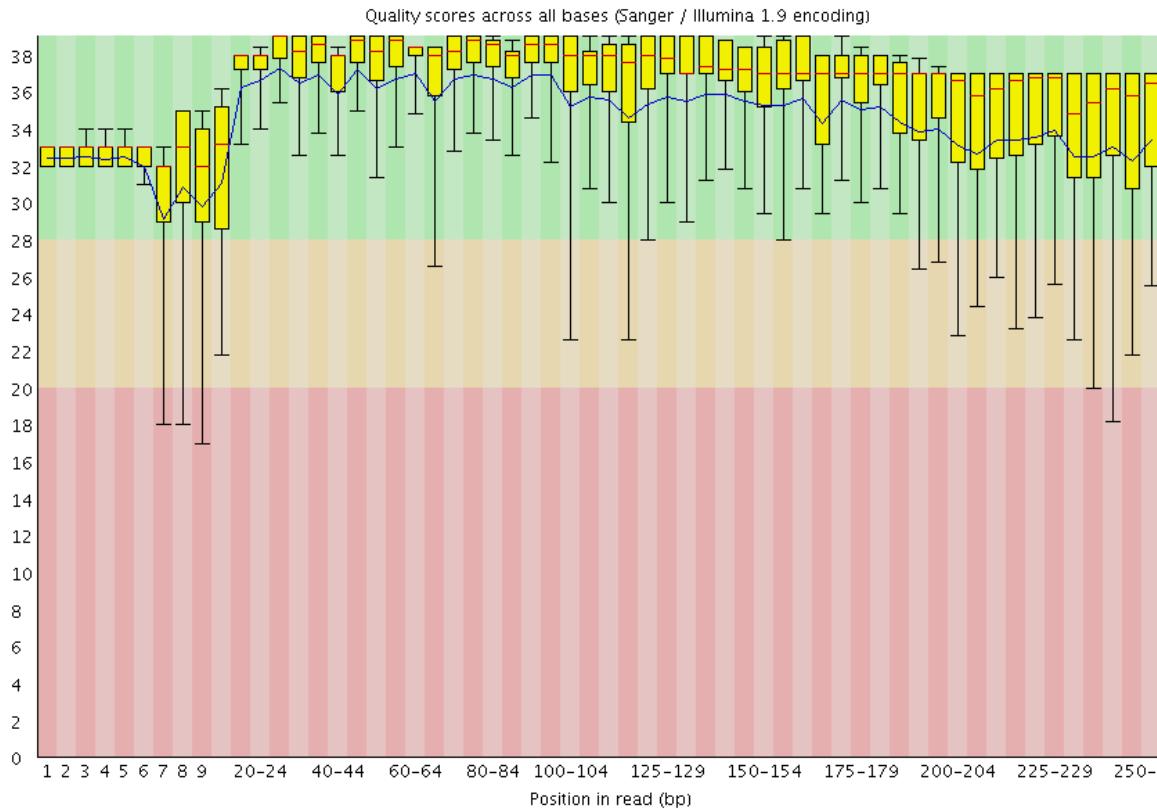


hep_sample2_pond2_R2.fastq.gz

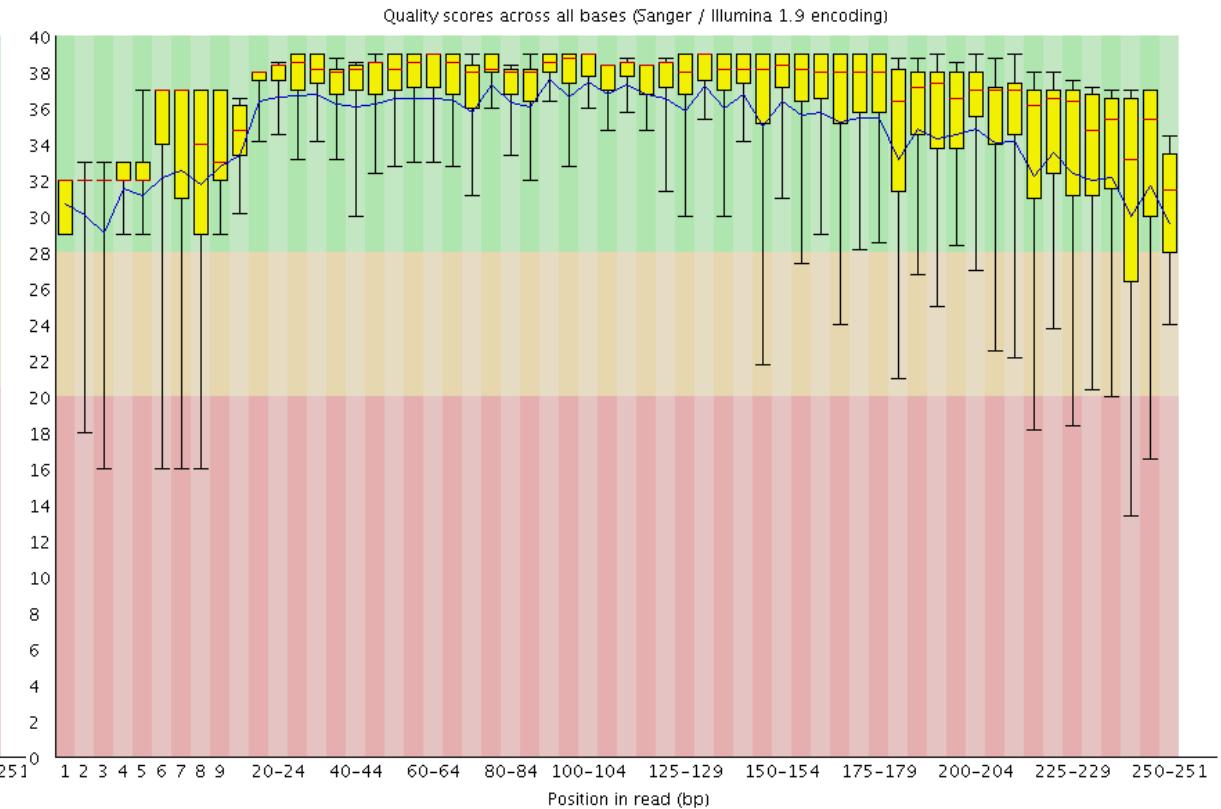


QUALITY

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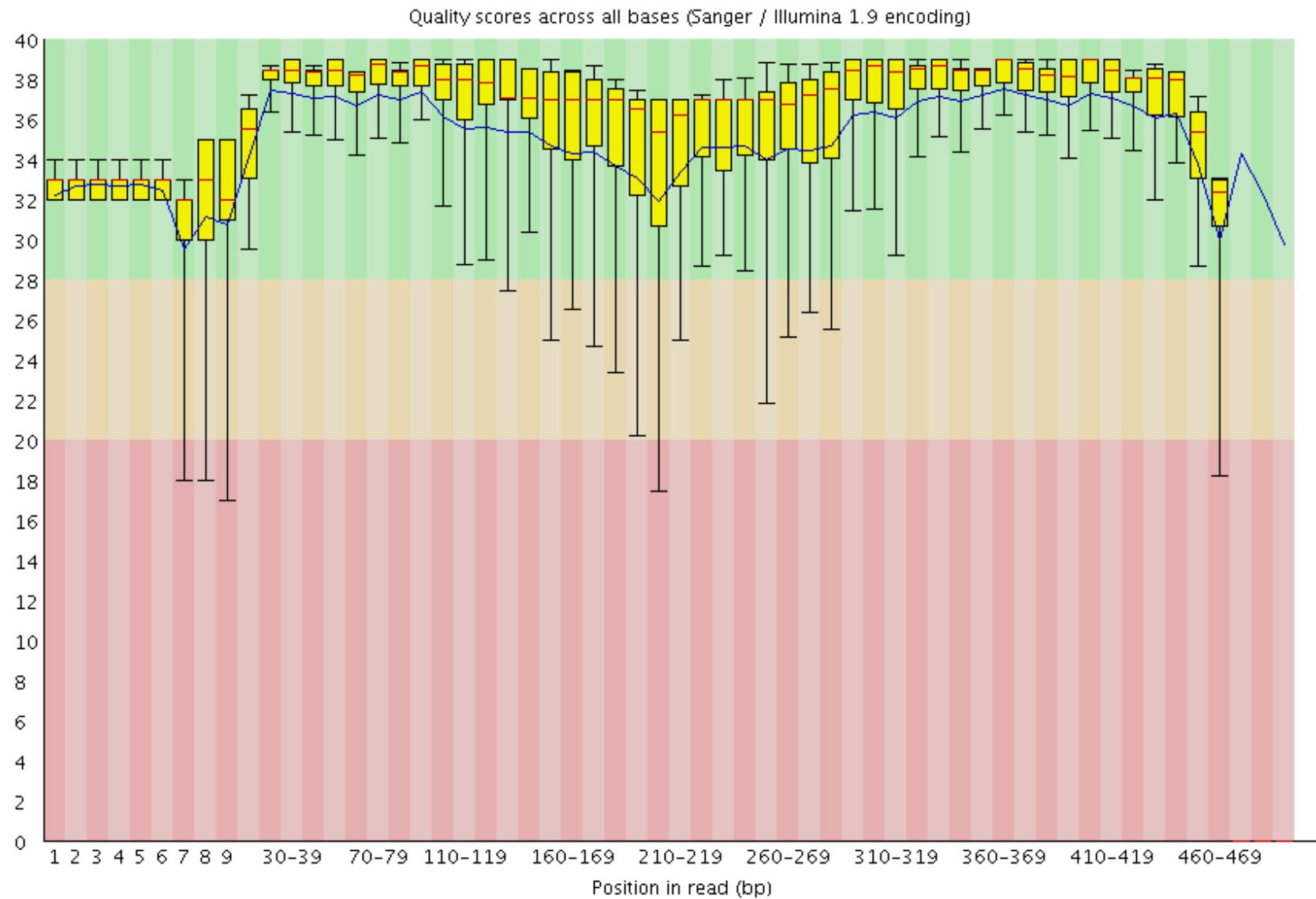


hep_sample2_pond2_R2_paired.fastq.gz



JOIN

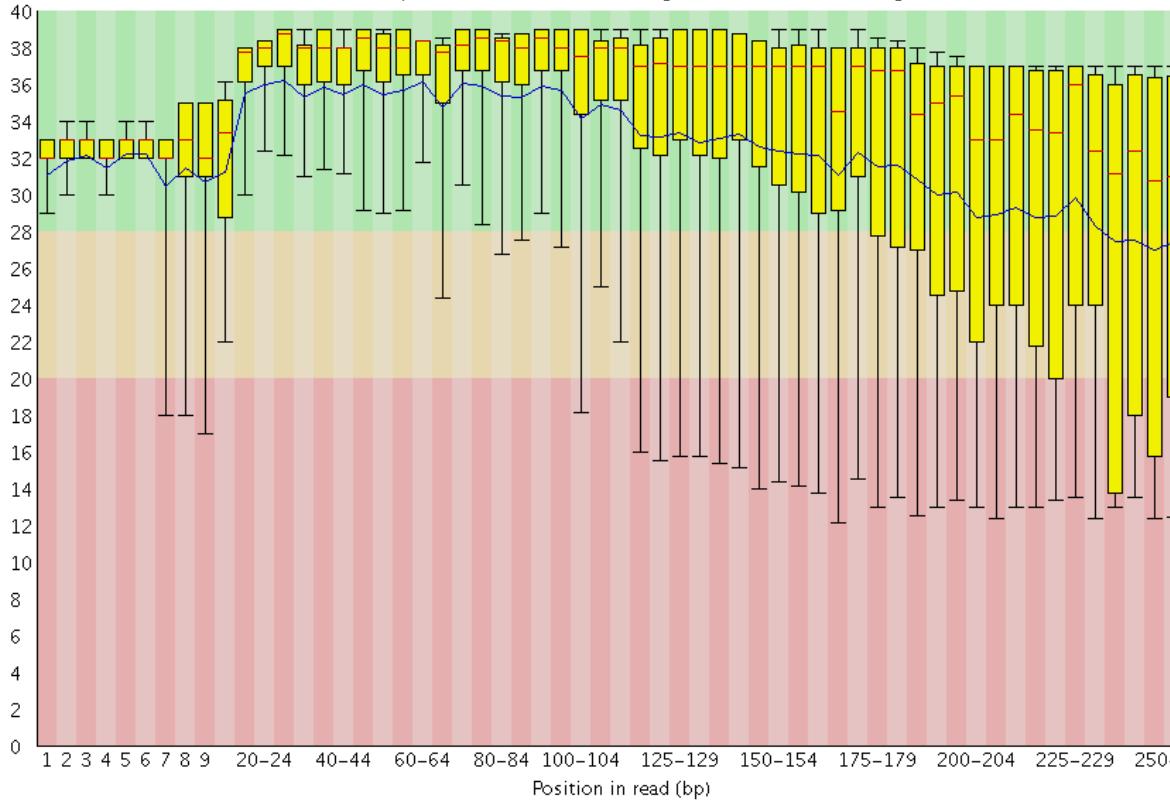
hjoinedsample2p2.fastqjoin



RAWDATA

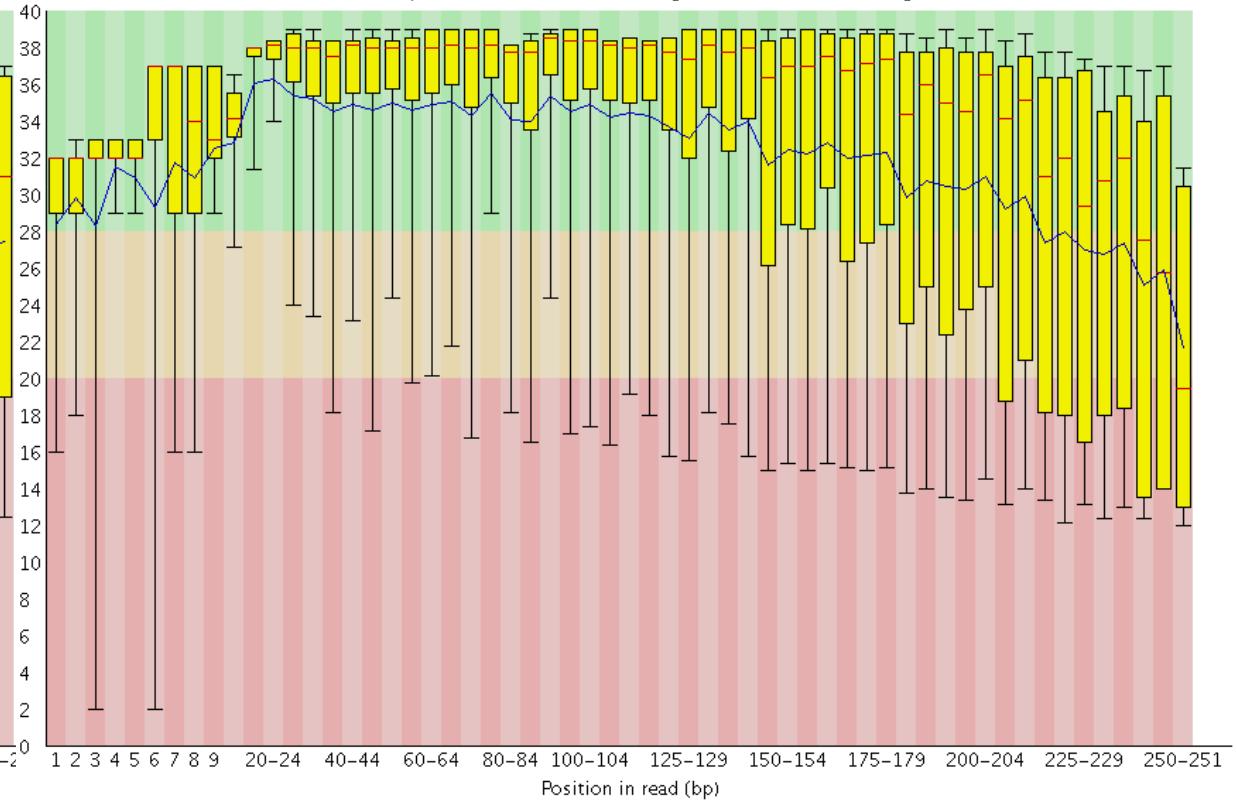
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Quality scores across all bases (Sanger / Illumina 1.9 encoding)



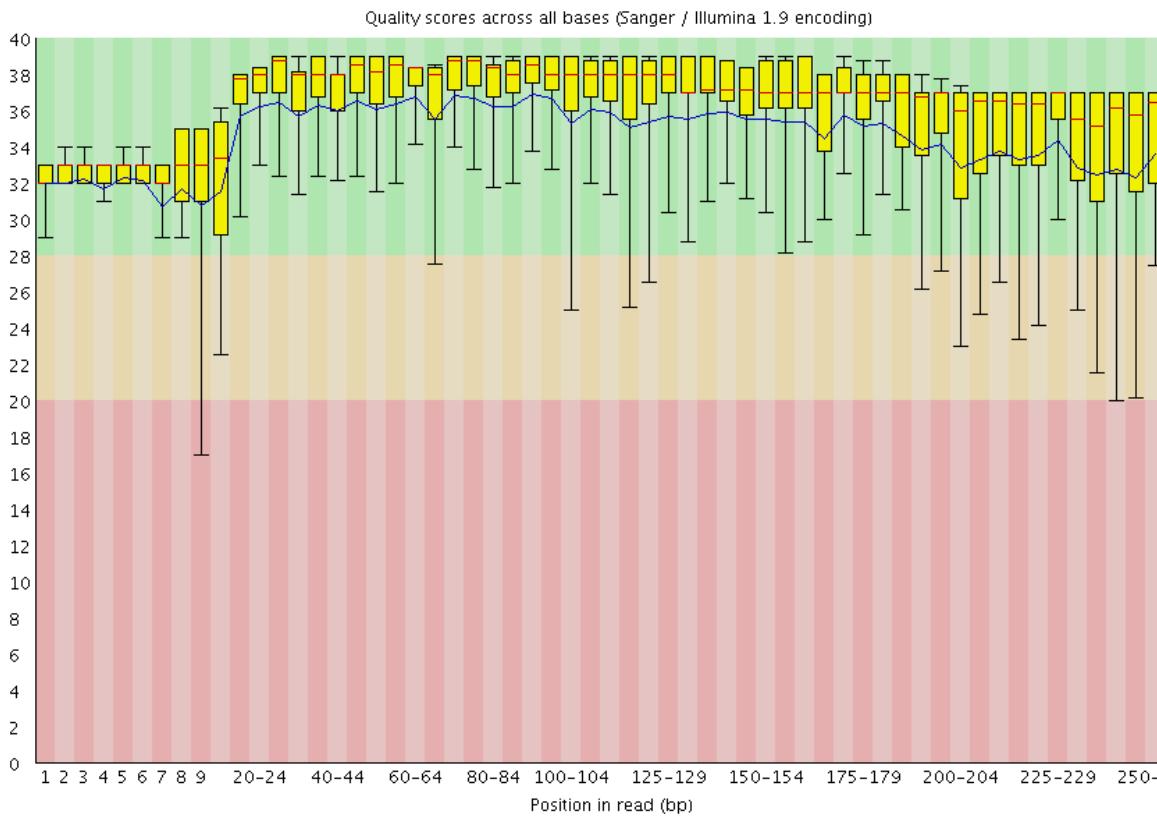
hep_sample2_pond3_R2.fastq.gz

Quality scores across all bases (Sanger / Illumina 1.9 encoding)

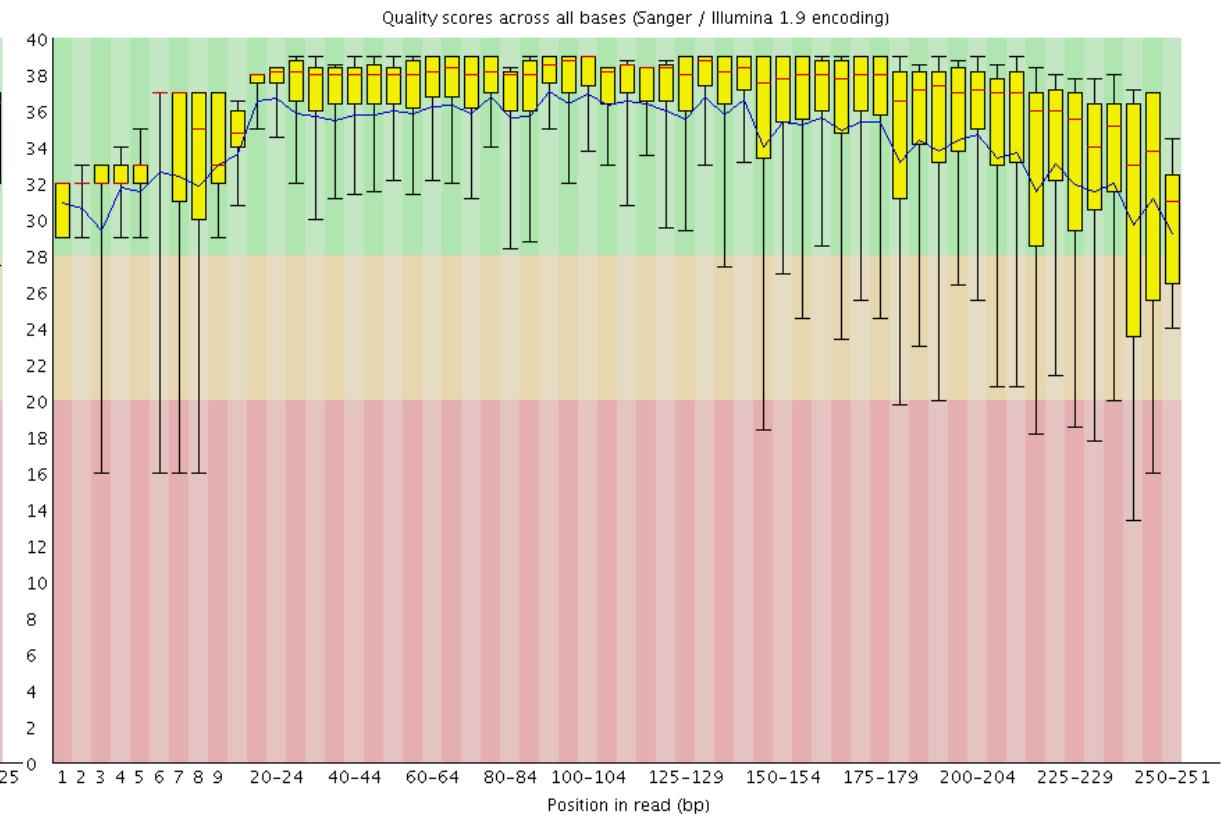


QUALITY

hep_sample2_pond3_R1_paired.fastq.gz

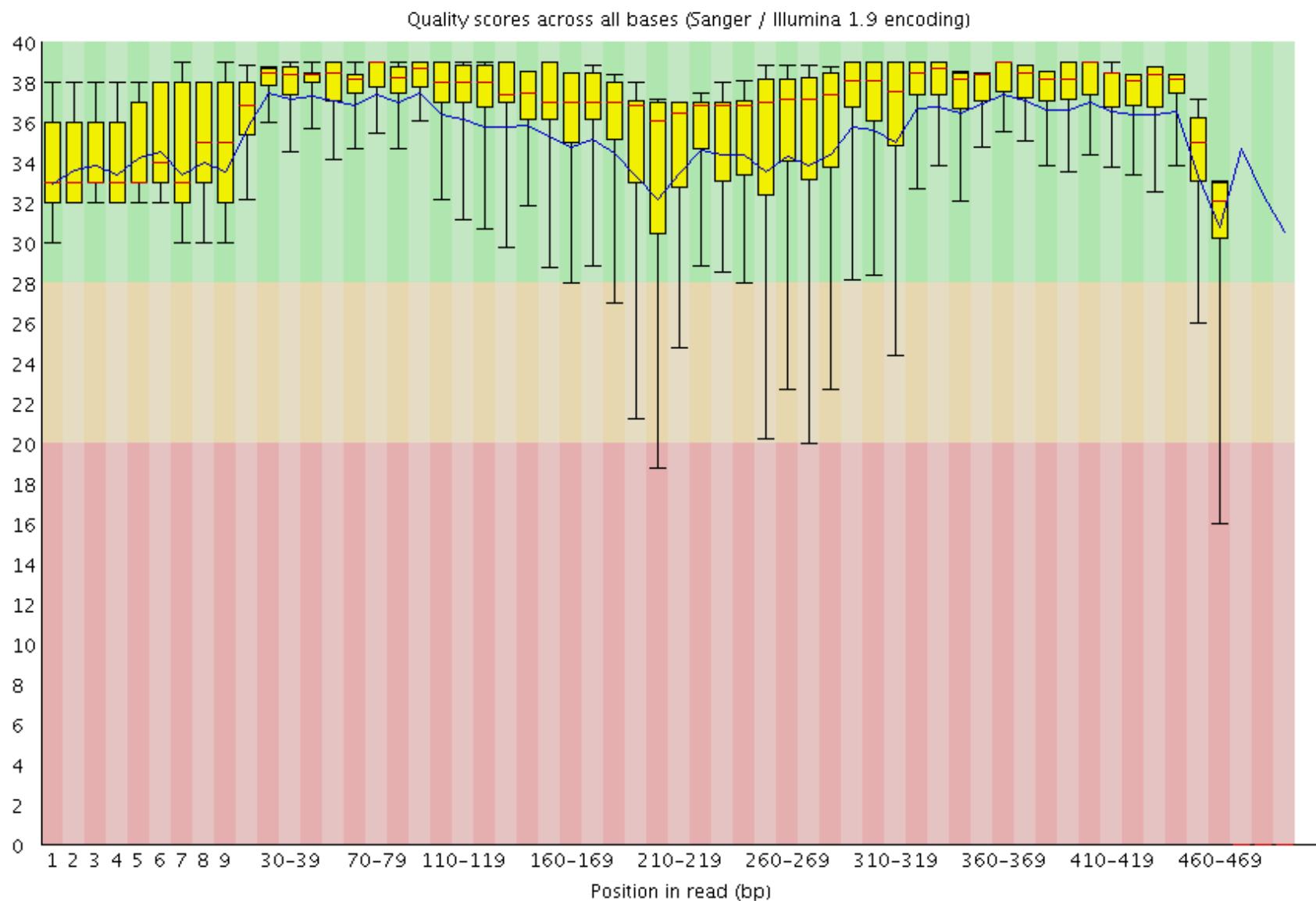


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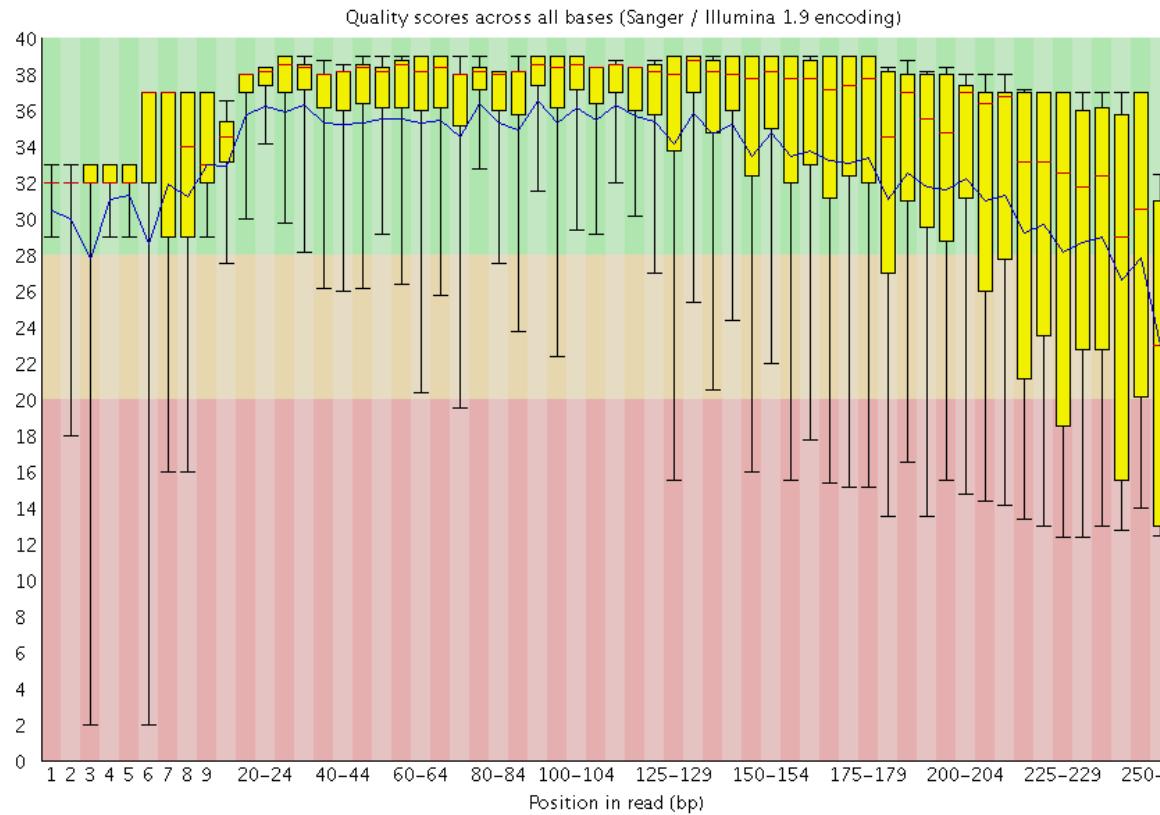
JOIN

hjoinedsample2p3.fastqjoin

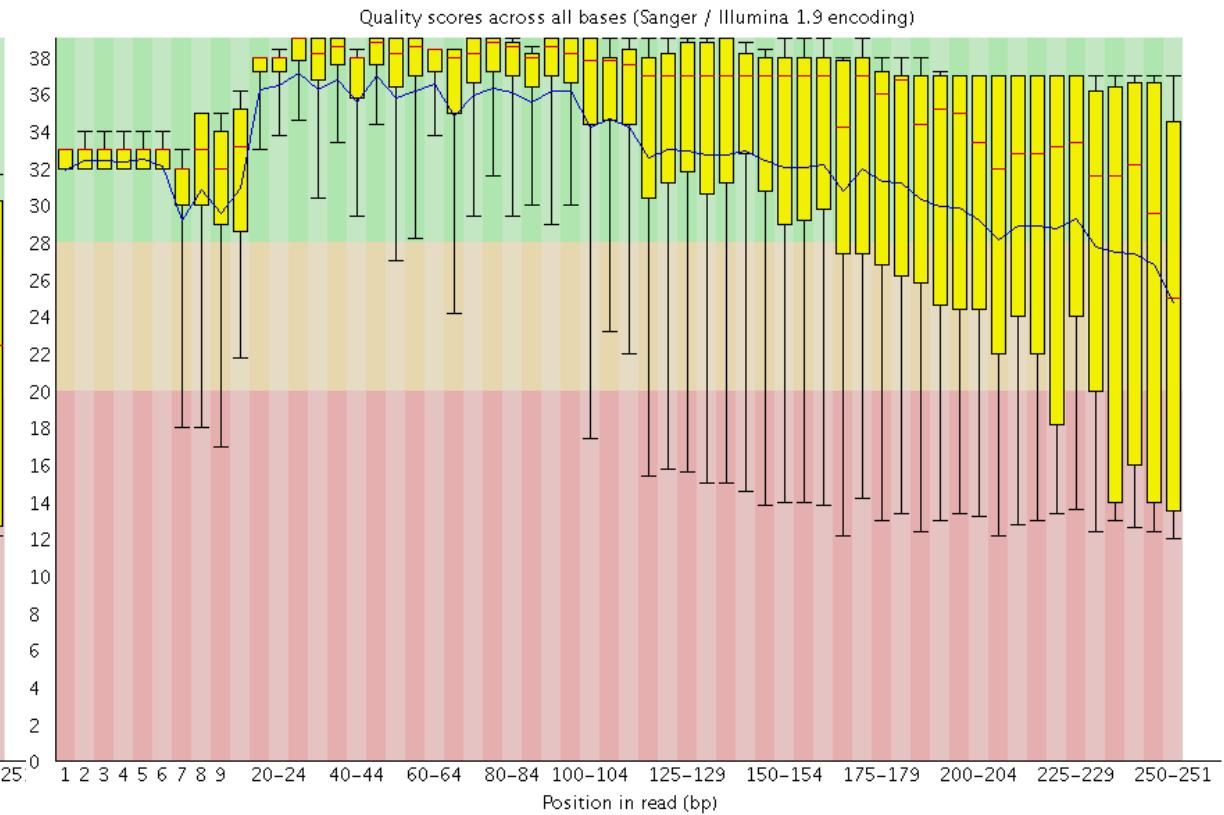


RAWDATA

hep_sample3_pond1_R1.fastq.gz

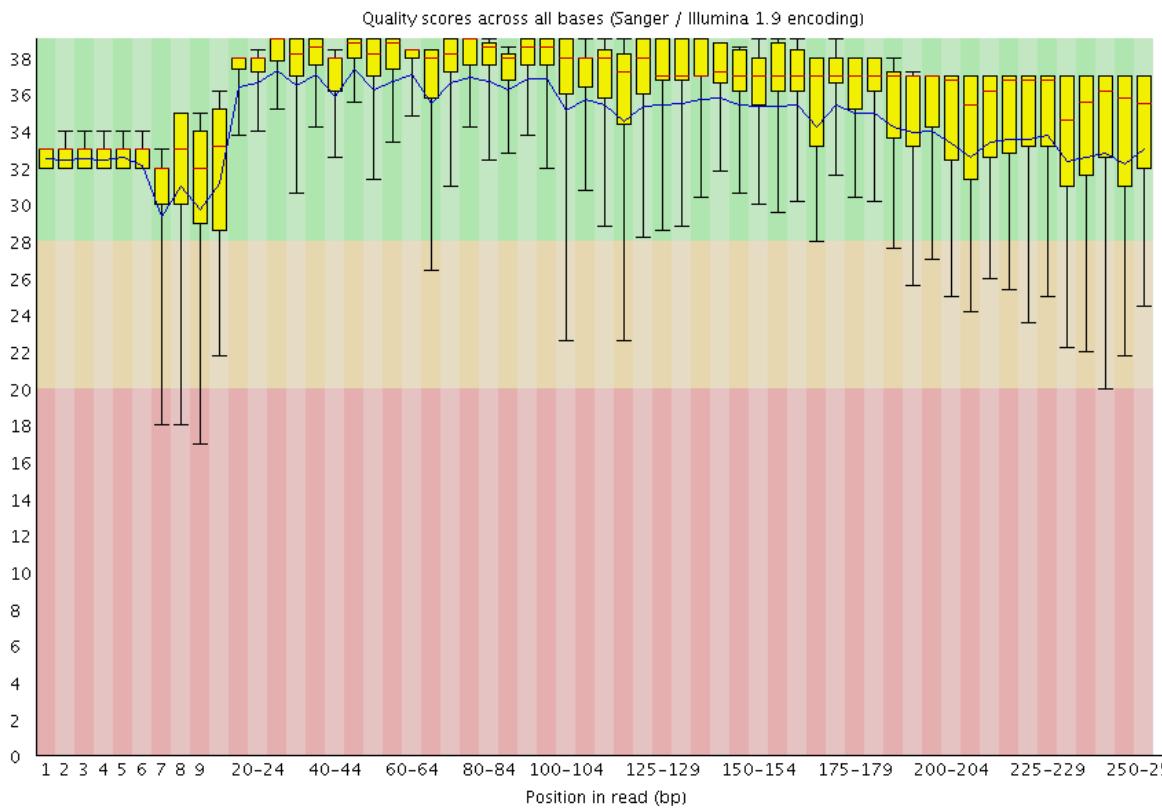


hep_sample3_pond1_R2.fastq.gz

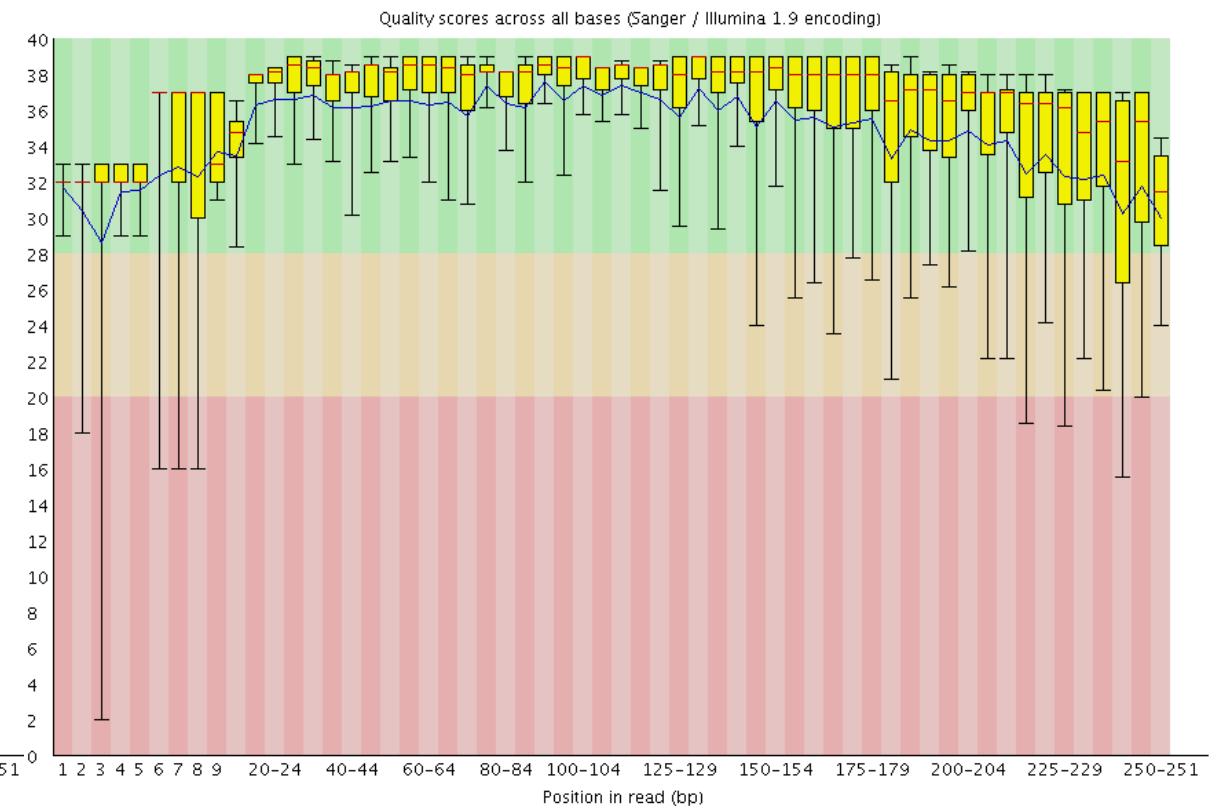


QUALITY

hep_sample3_pond1_R1_paired.fastq.gz

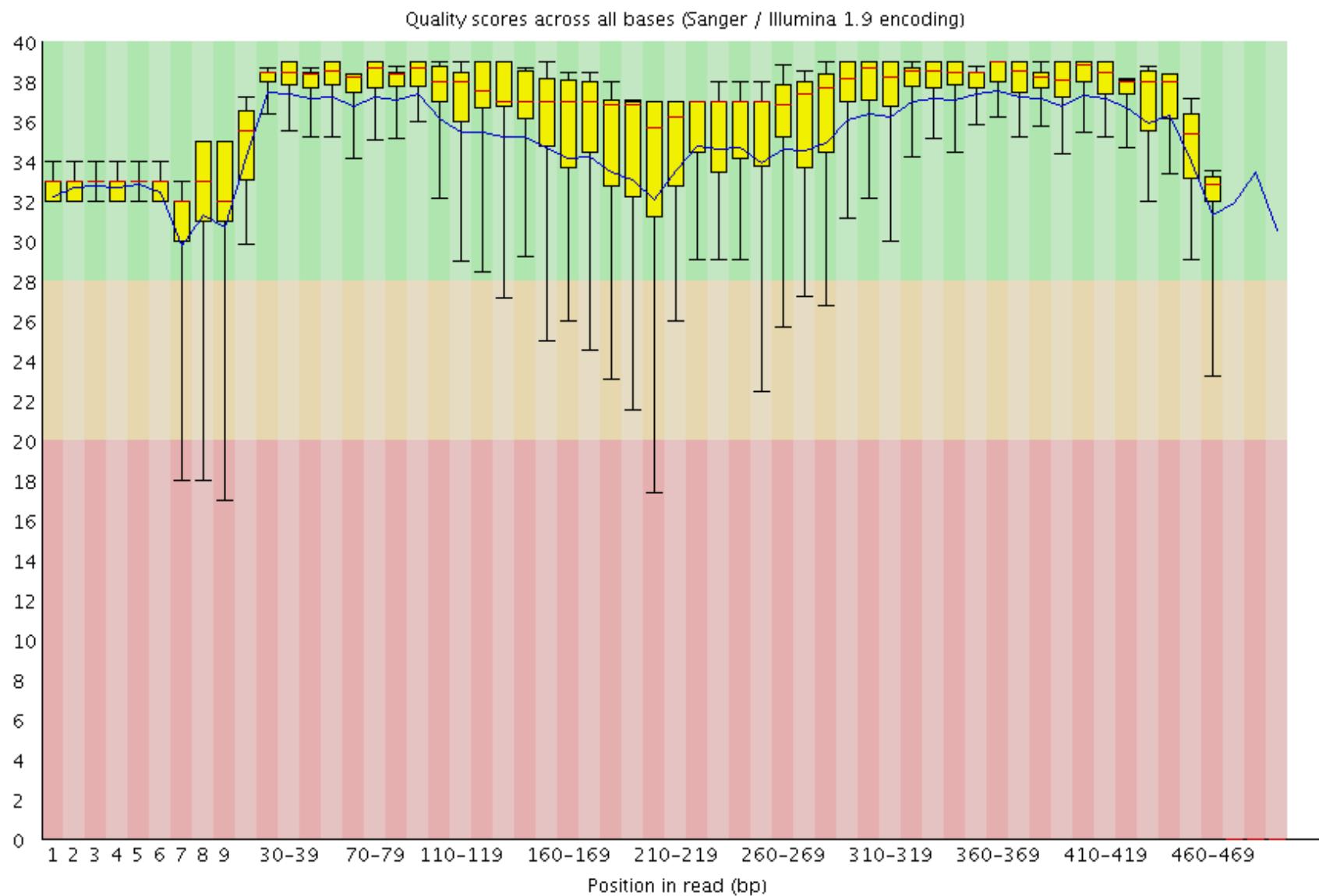


hep_sample3_pond1_R2_paired.fastq.gz



JOIN

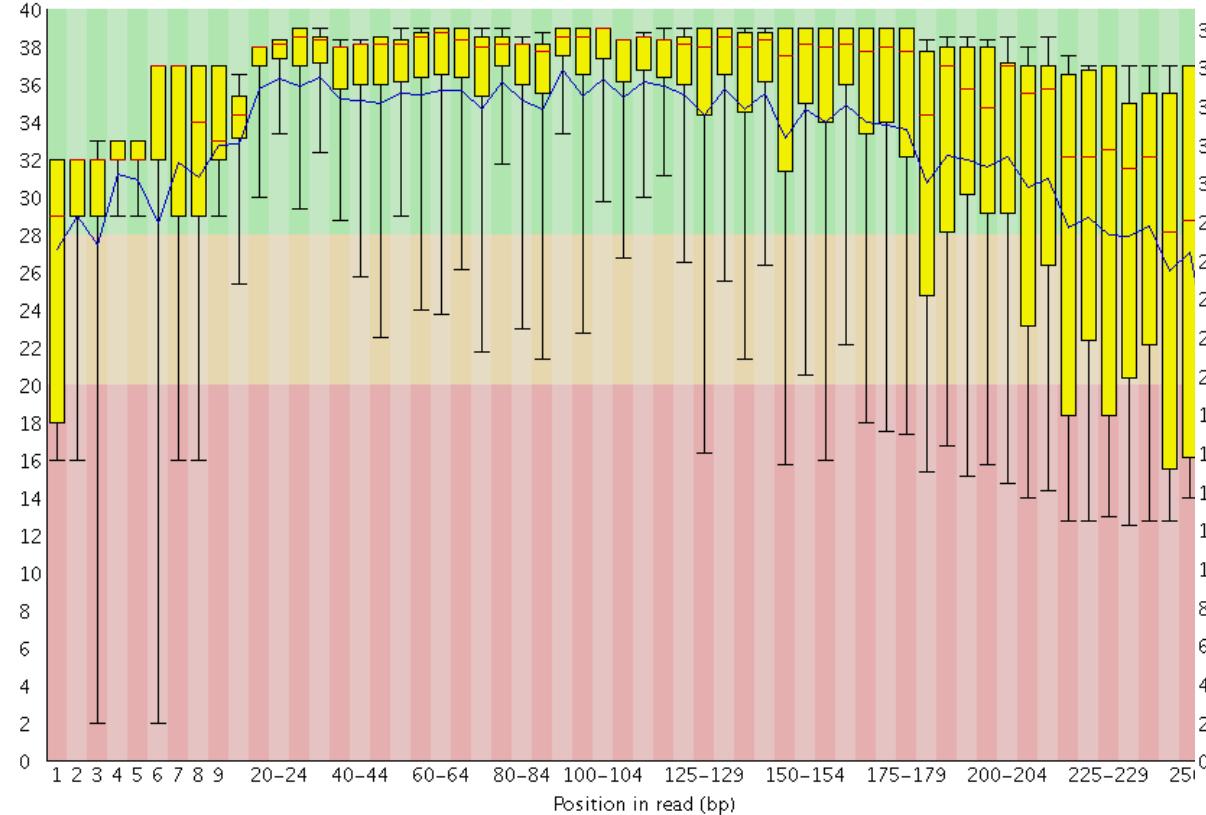
hjoinedsample3p1.fastqjoin



RAWDATA

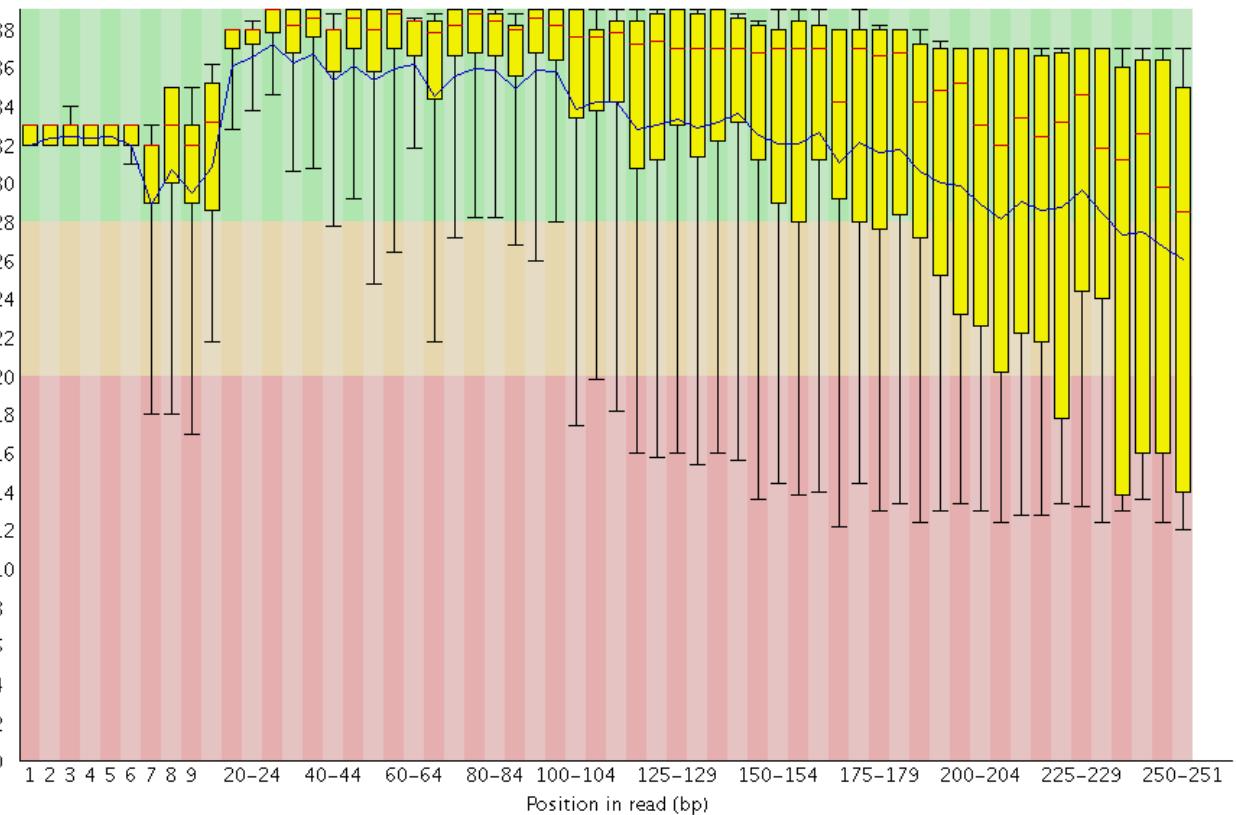
hep_sample3_pond32_R1.fastq.gz

Quality scores across all bases (Sanger / Illumina 1.9 encoding)



hep_sample3_pond2_R2.fastq.gz

Quality scores across all bases (Sanger / Illumina 1.9 encoding)

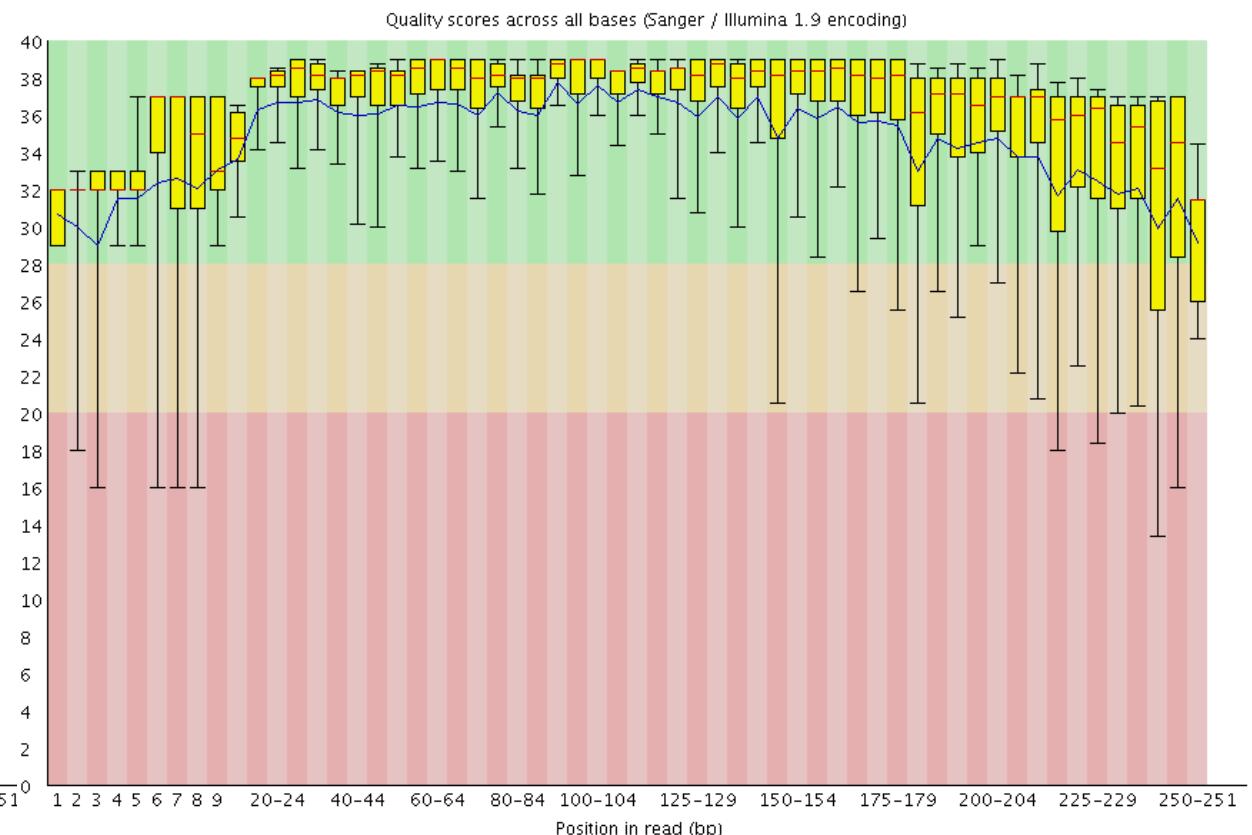


QUALITY

hep_sample3_pond2_R1_paired.fastq.gz

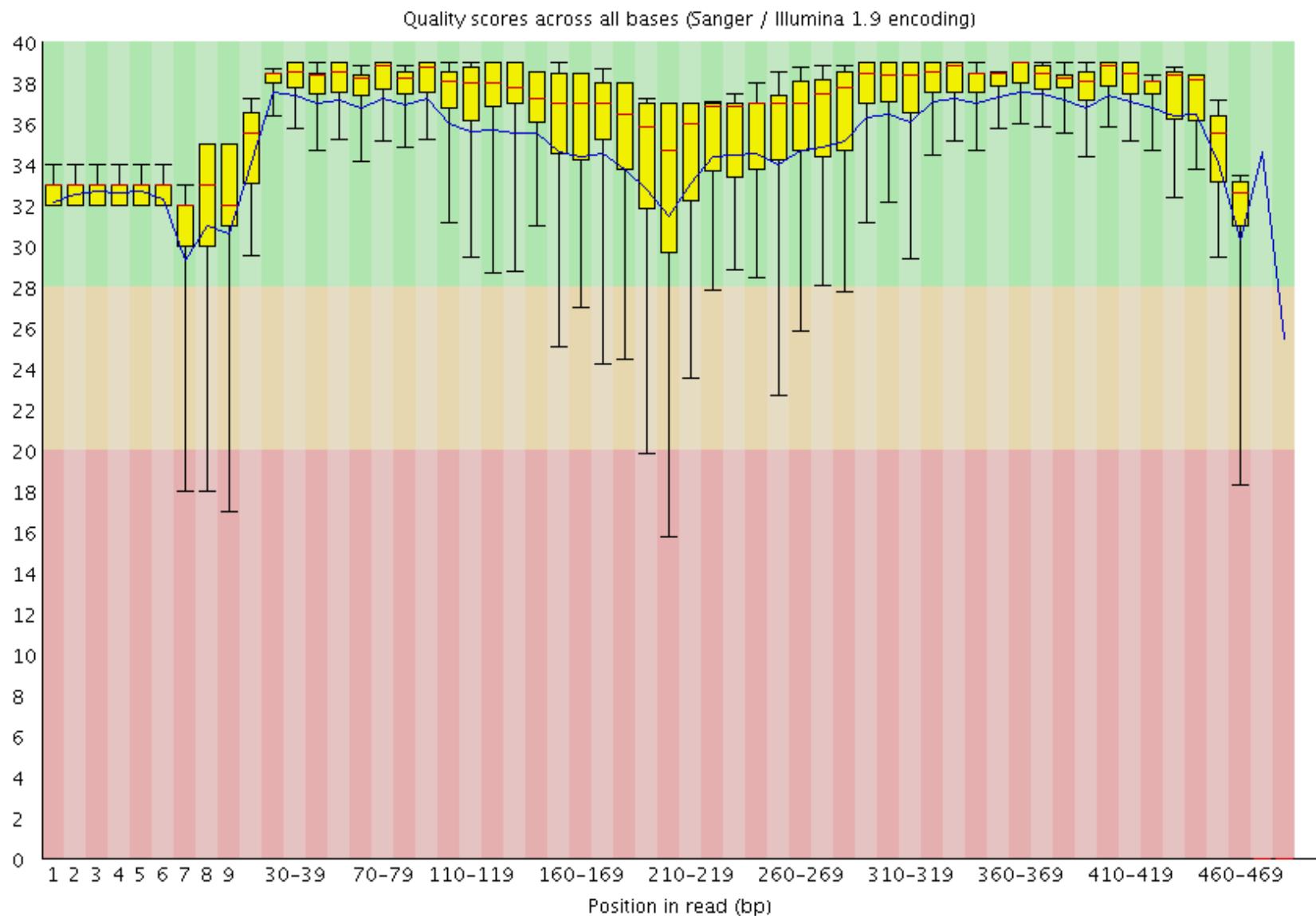


hep_sample3_pond2_R2_paired.fastq.gz



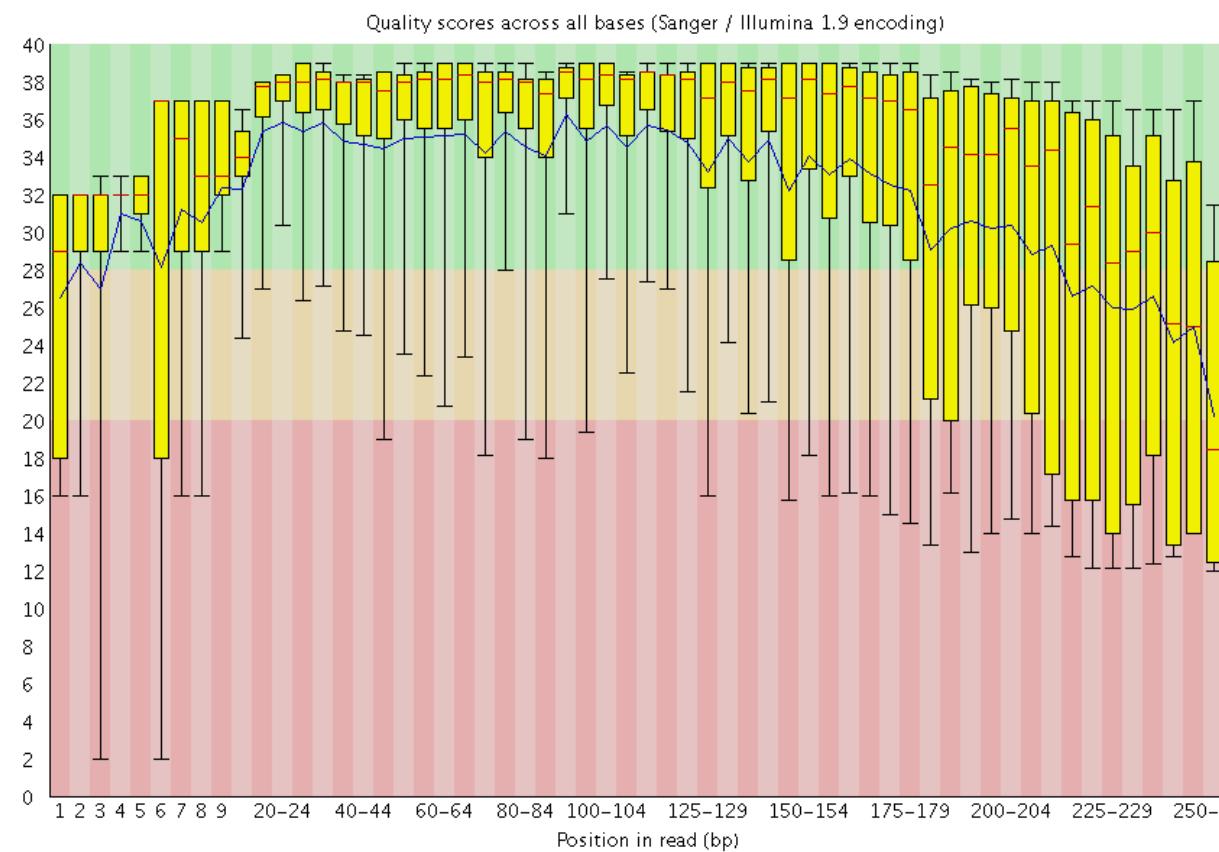
JOIN

hjoinedsample3p2.fastqjoin

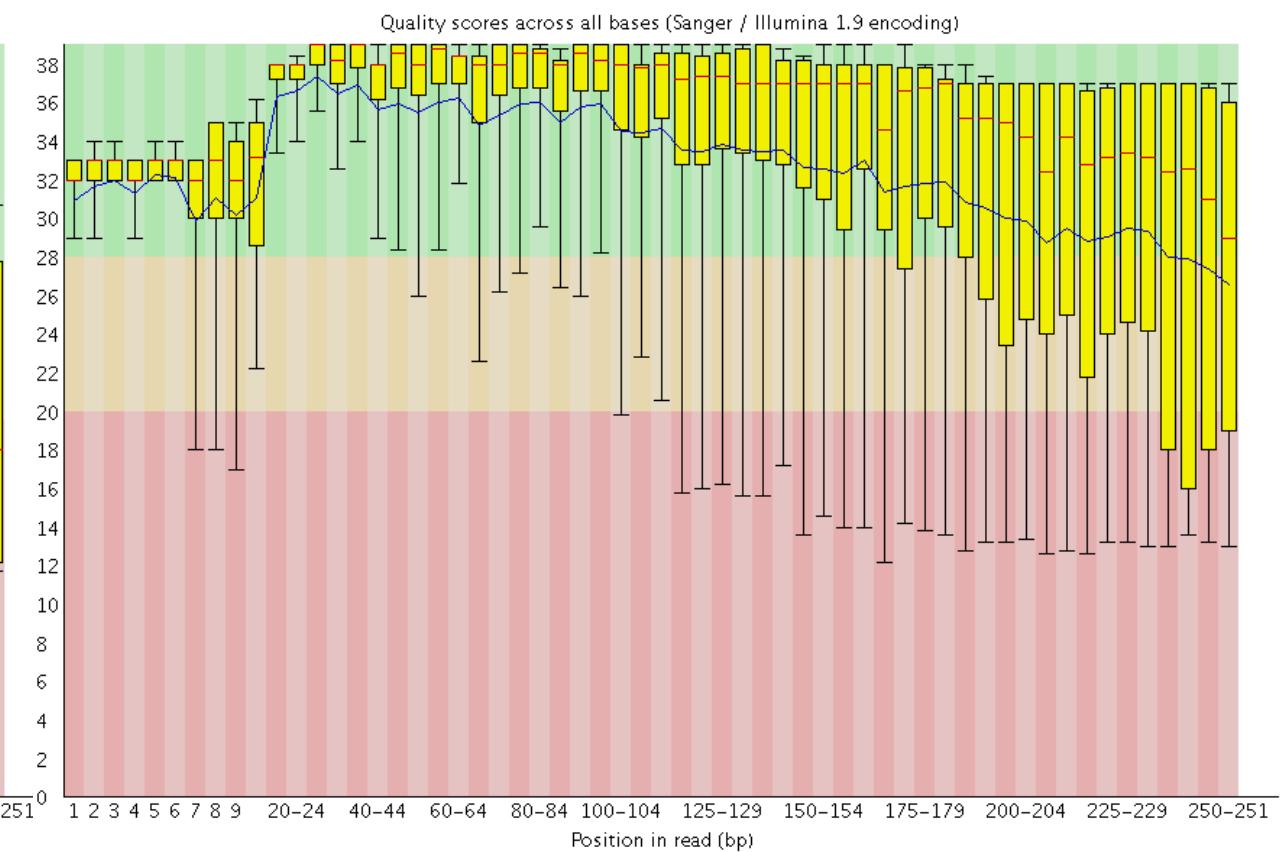


RAWDATA

hep_sample3_pond3_R1.fastq.gz

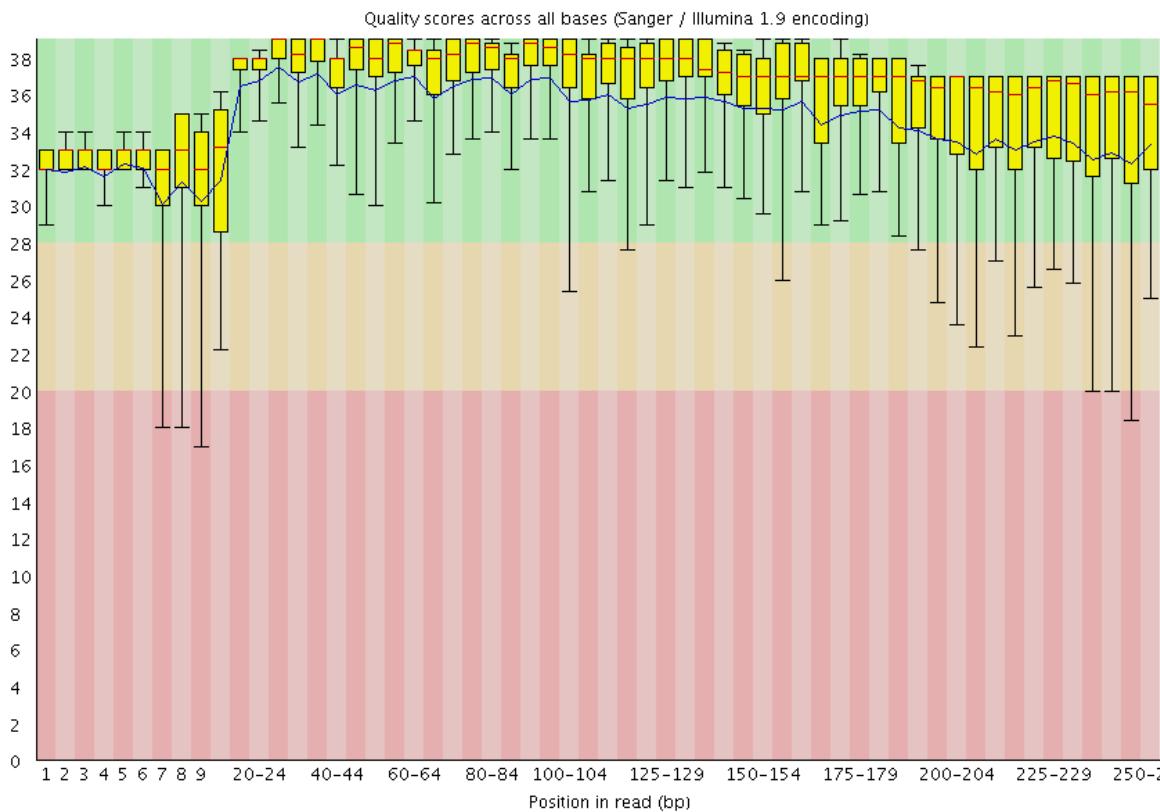


hep_sample3_pond3_R2.fastq.gz

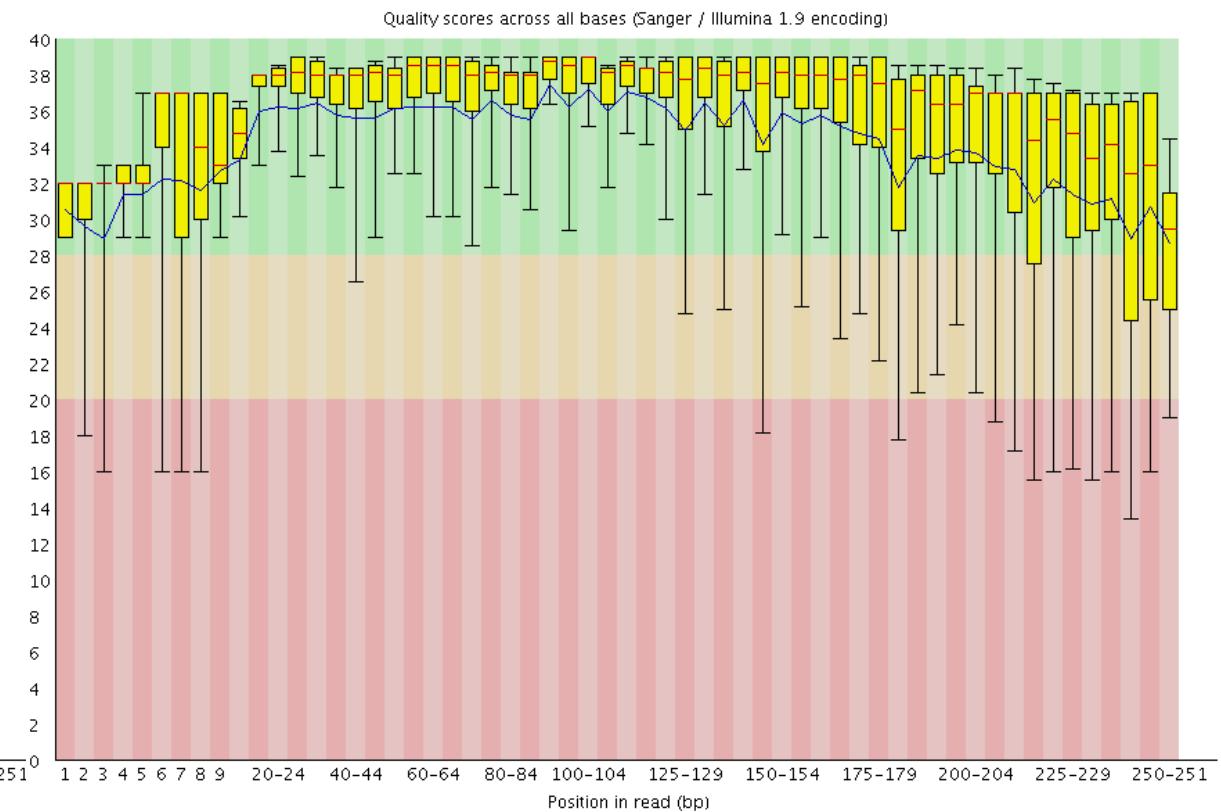


QUALITY

hep_sample3_pond3_R1_paired.fastq.gz



hep_sample3_pond3_R2_paired.fastq.gz



JOIN

hjoinedsample3p3.fastqjoin

