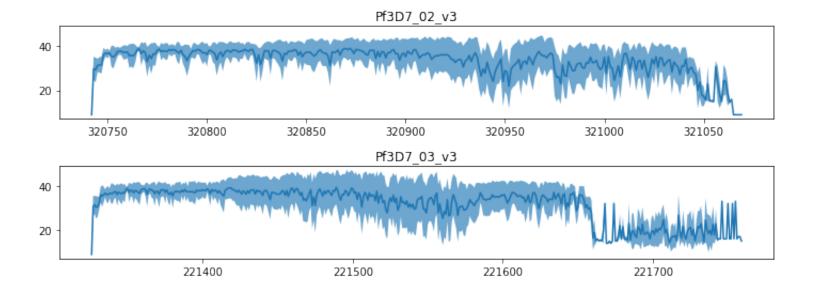
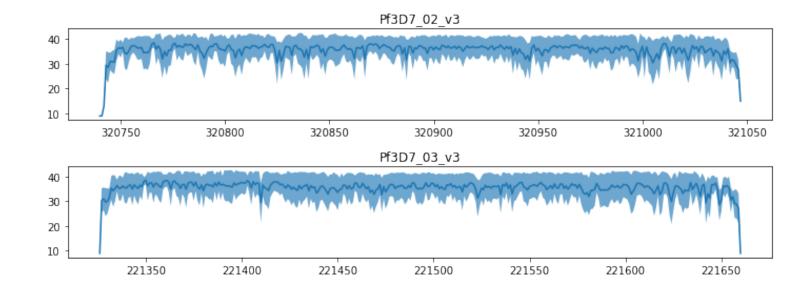
rtss amplicon

raw reads analysis – samtools pileup 20180720

read base quality

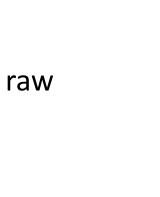






aligned

read mismatch



1.0

0.5

0.0

1.0

0.5

0.0

1.0

0.5

320750

320800

221400

320850

221500





Pf3D7_02_v3

320900

Pf3D7 03 v3

Pf3D7 02 v3

320950

221600

321000

221700

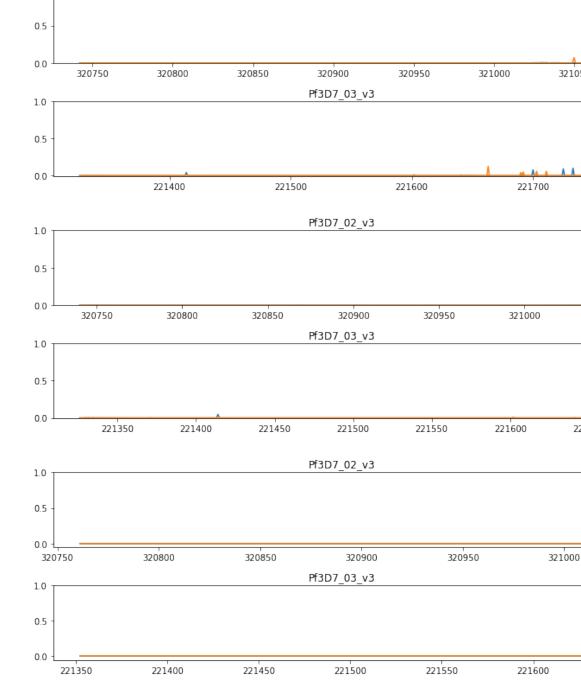
321050



read indel (unscaled)

raw

1.0



Pf3D7 02 v3

321050

321050

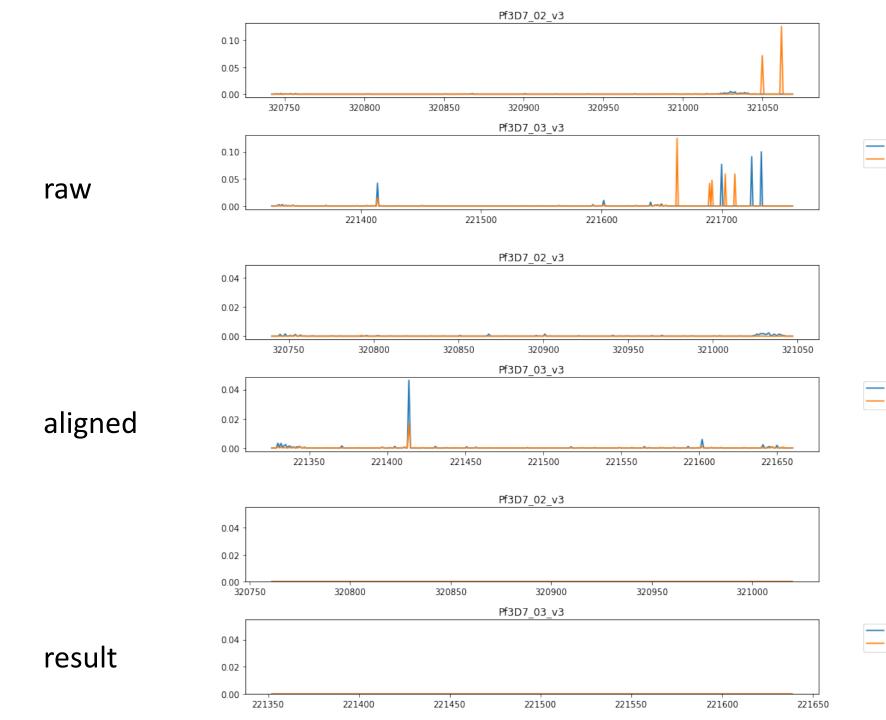
221650

221650

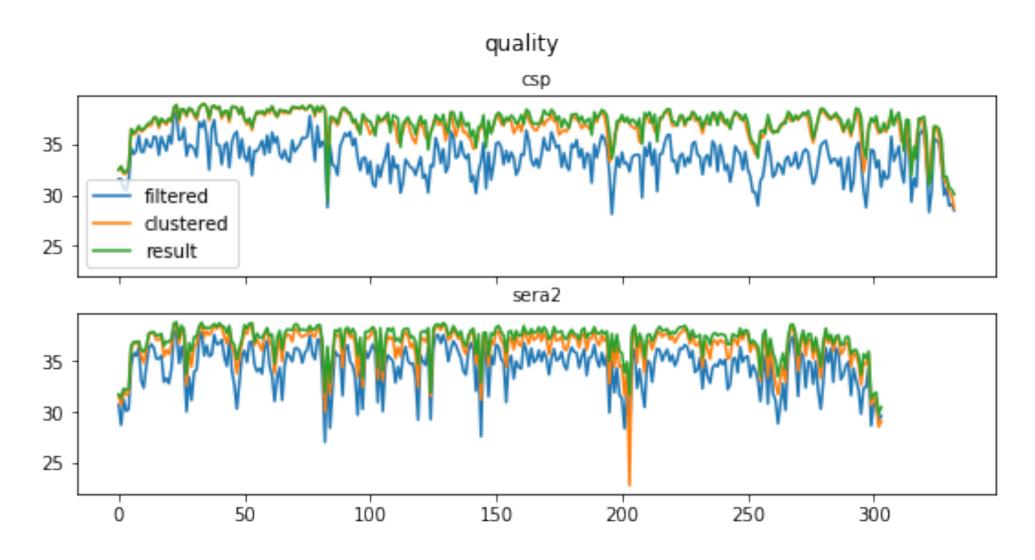
aligned

result

read indel (scaled)



read base quality



read mismatches

