

0.2

0.4

0.6

Normalized Score

8.0

0.0

1.0

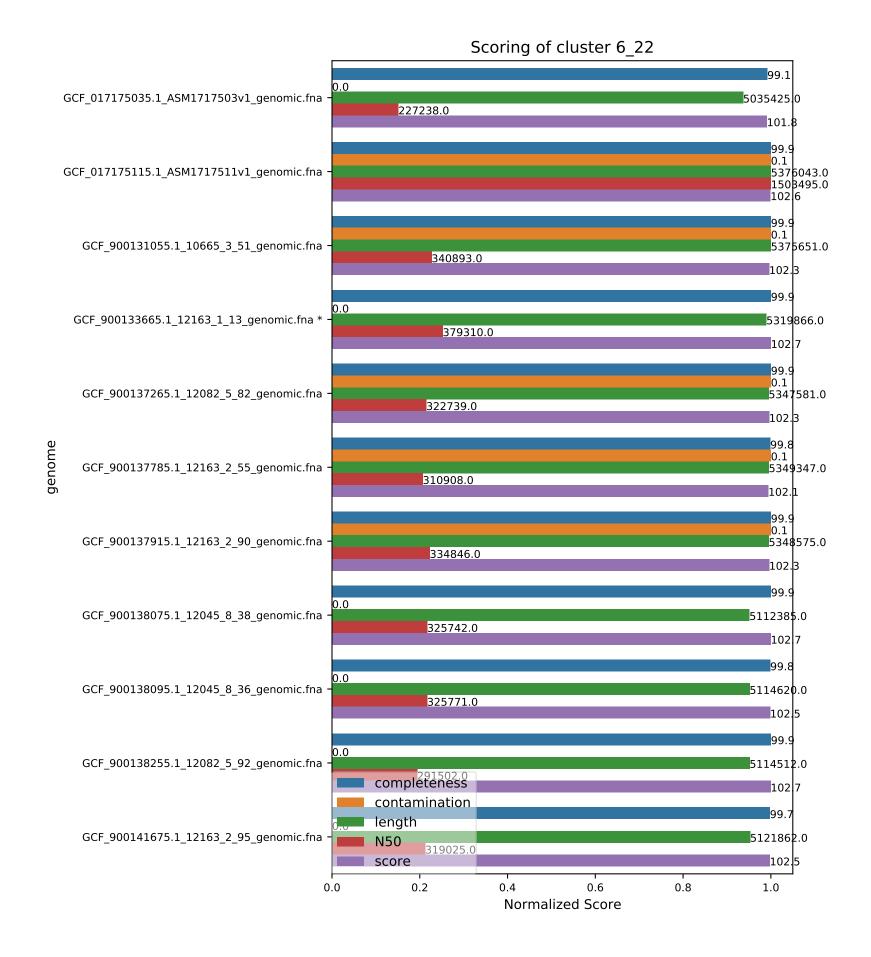
Scoring of cluster 6_21

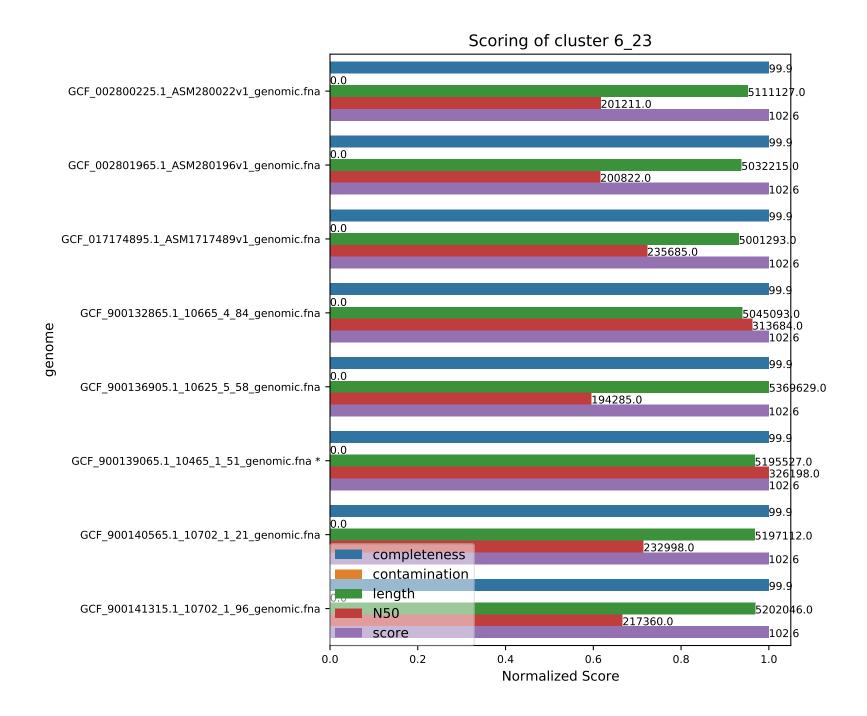
0.3

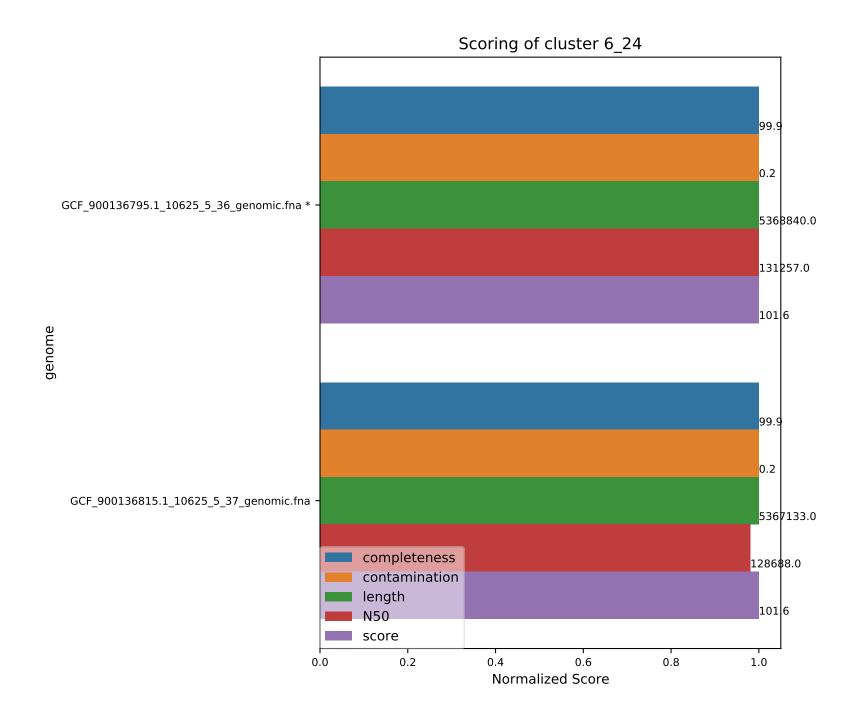
 $\mathsf{GCF}_004025145.1_\mathsf{ASM}402514v1_\mathsf{genomic.fna}$

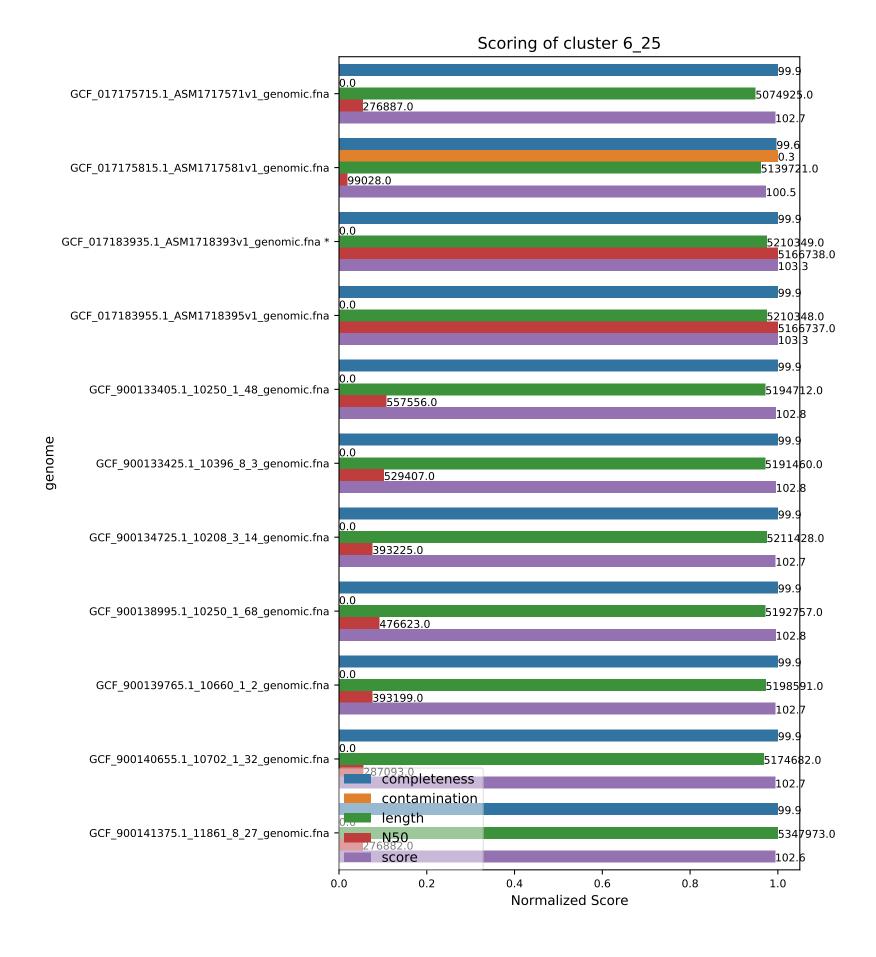
99.9

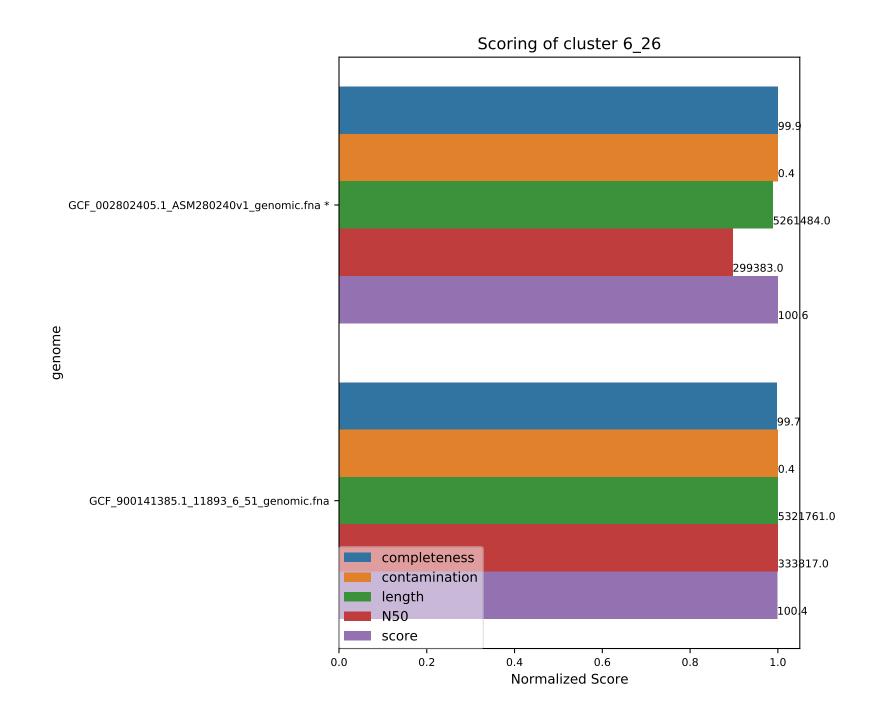
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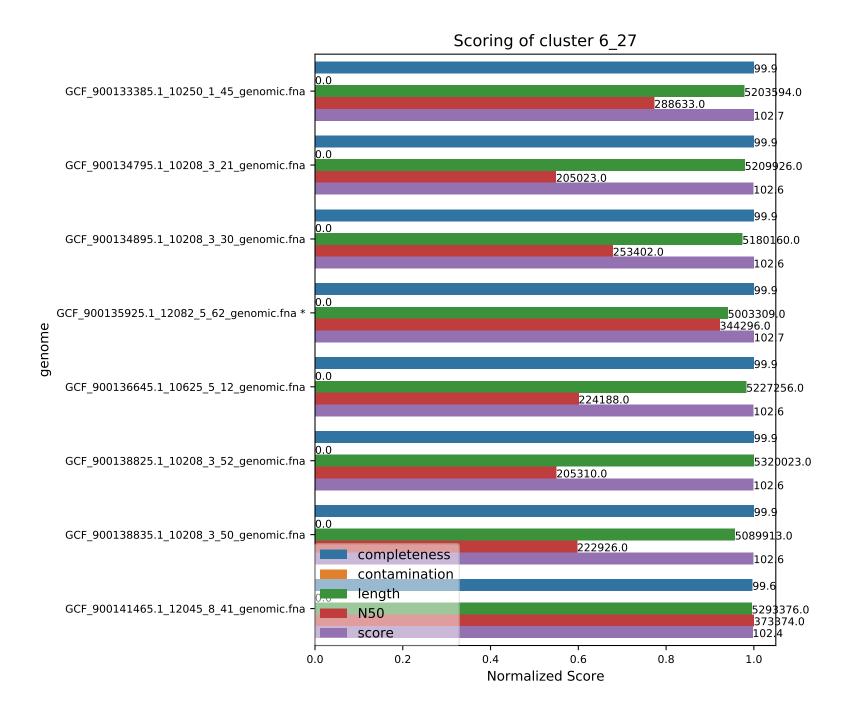


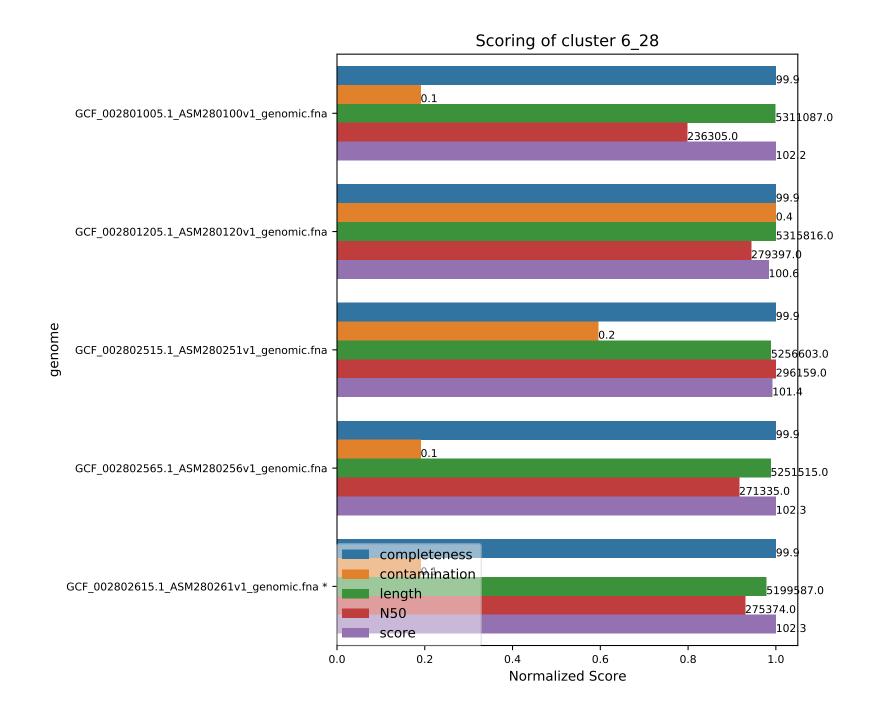


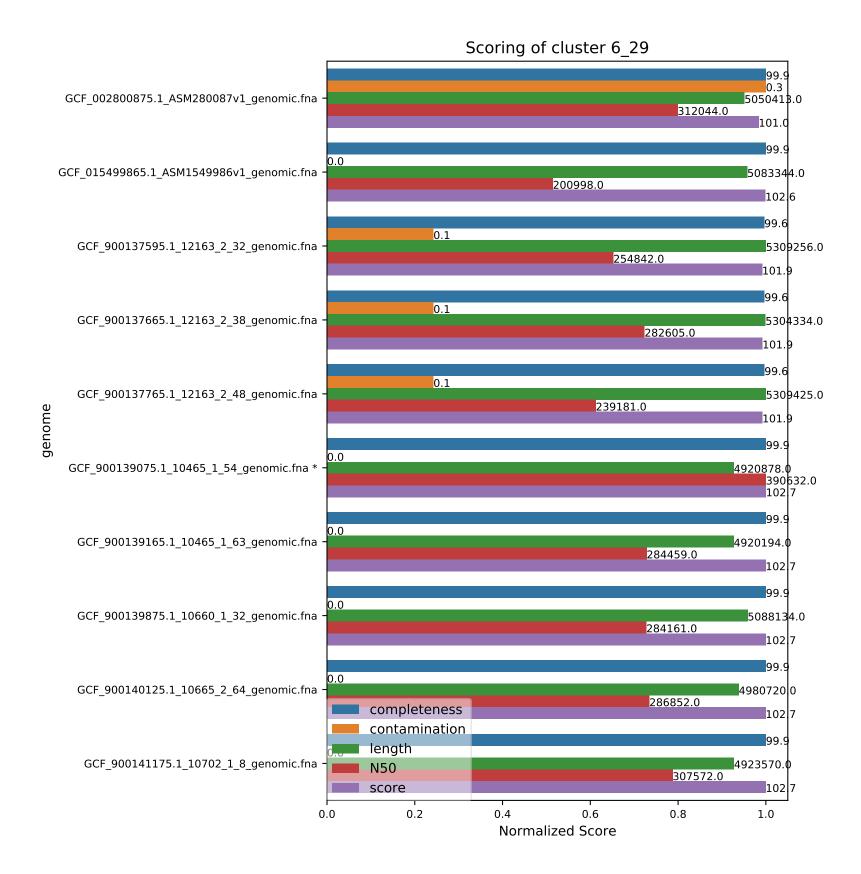


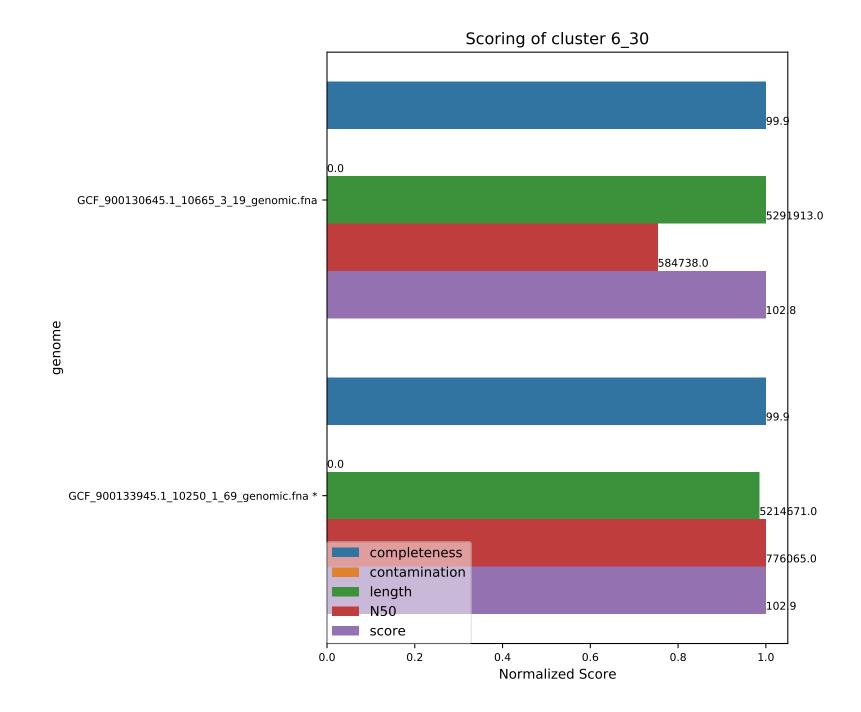


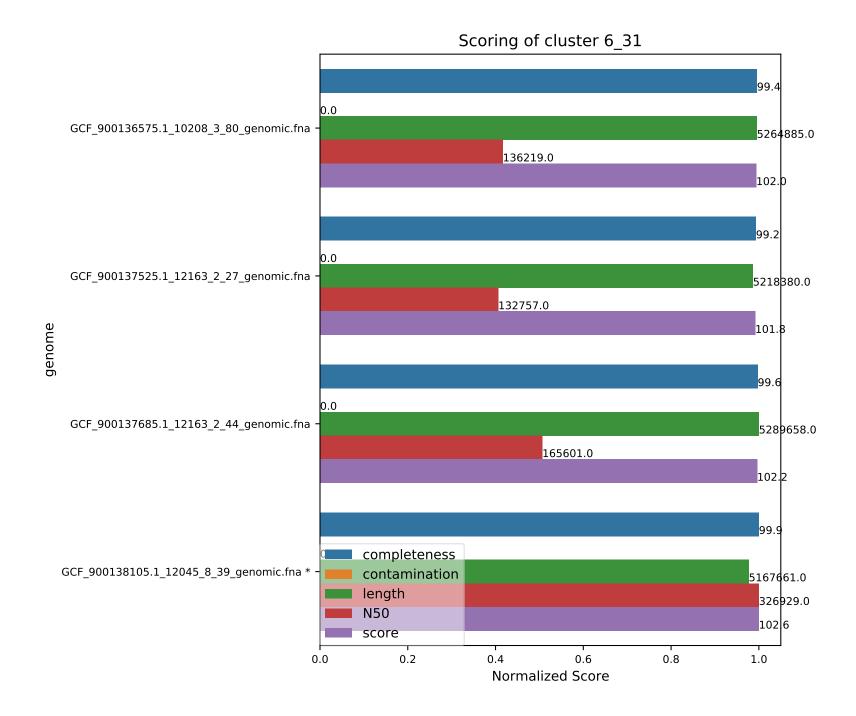


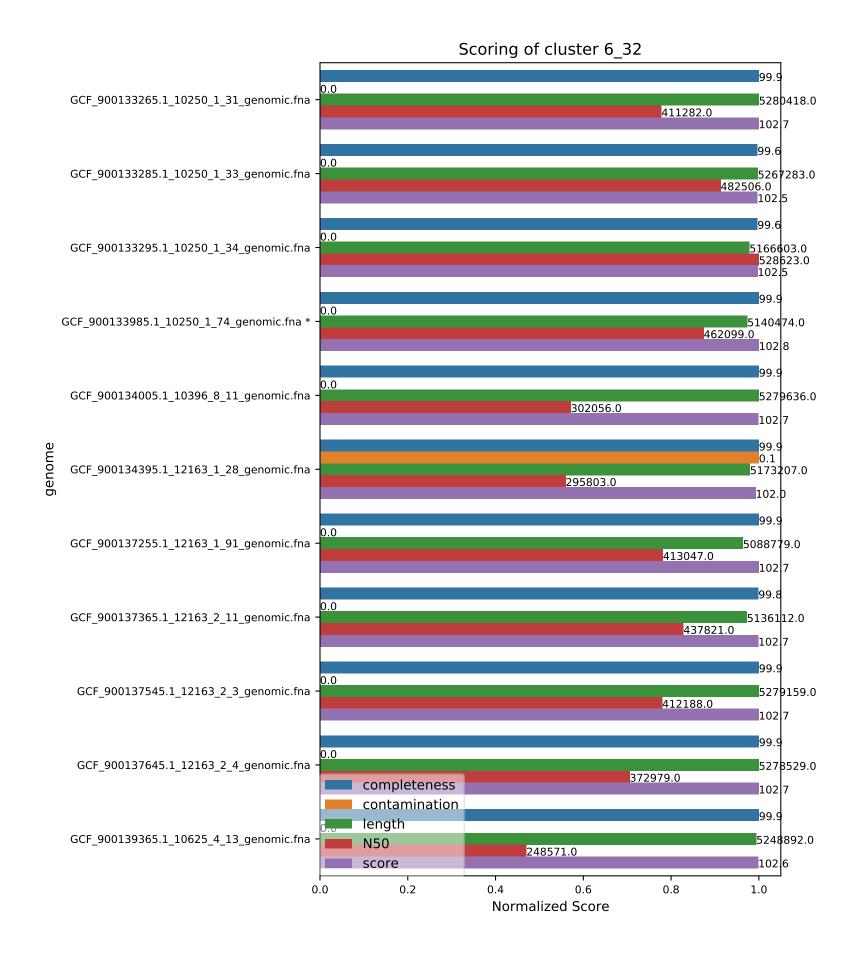


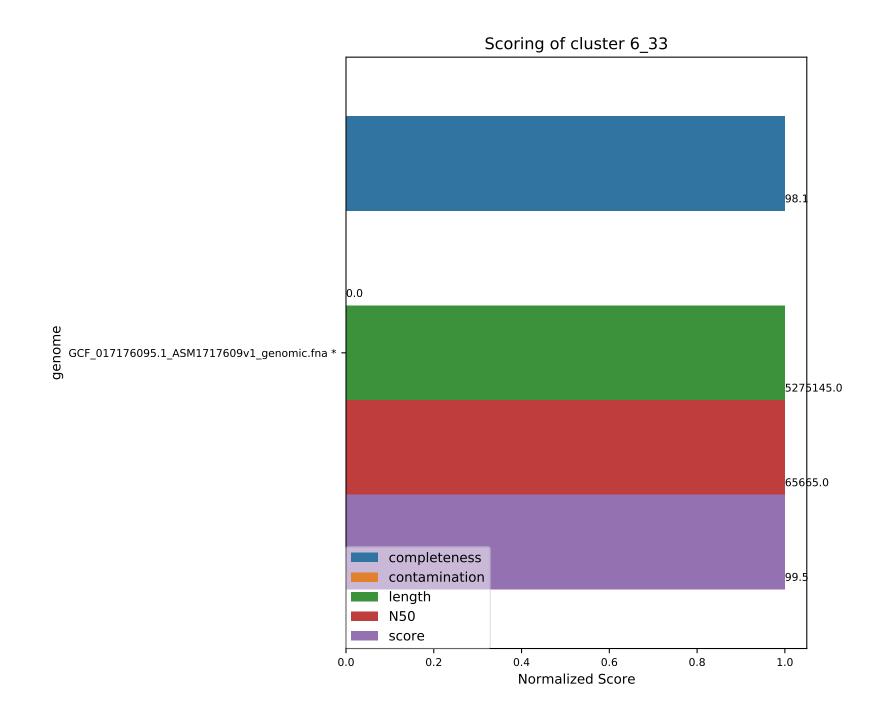


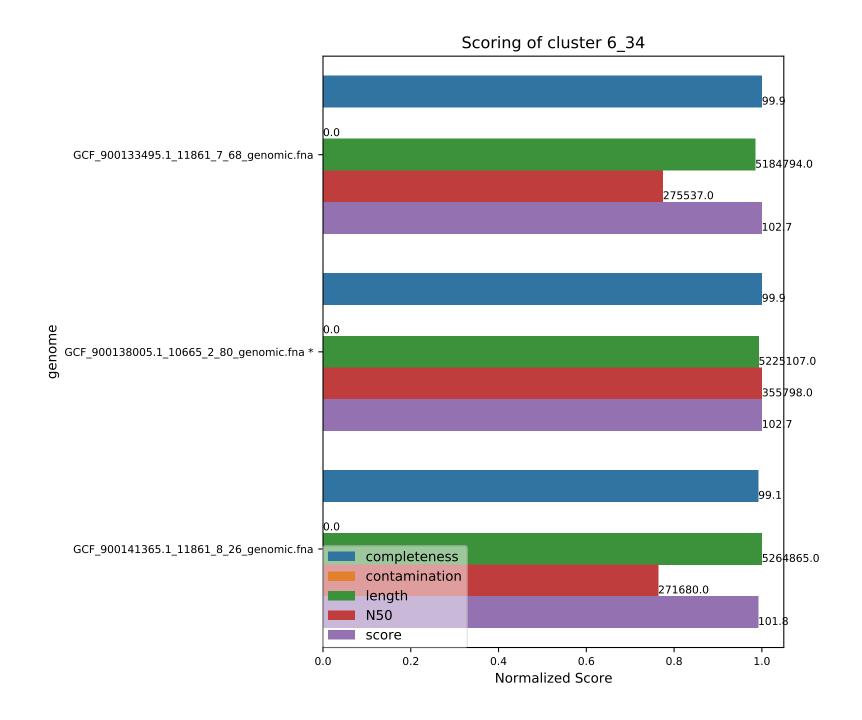


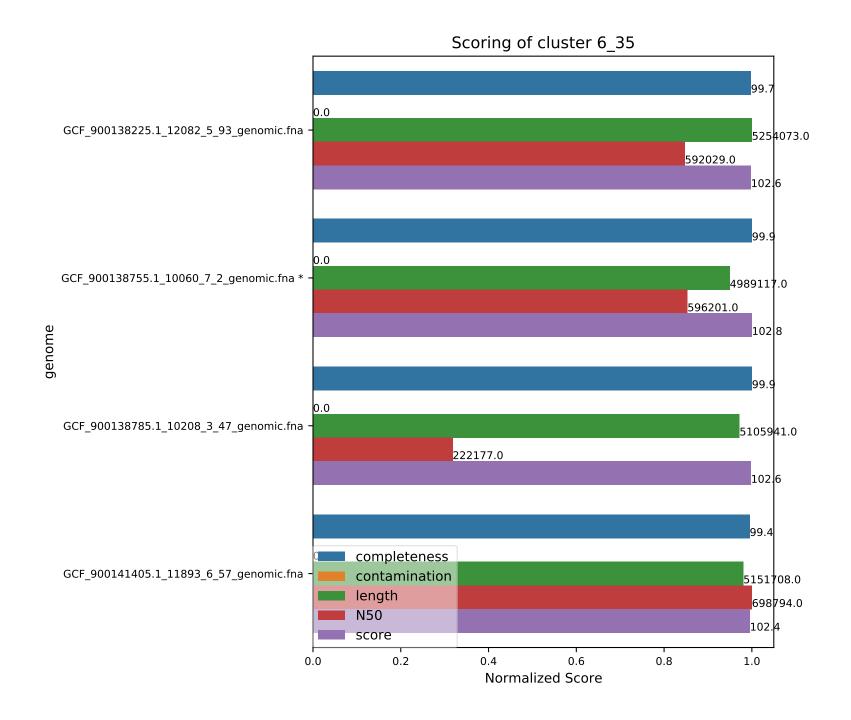


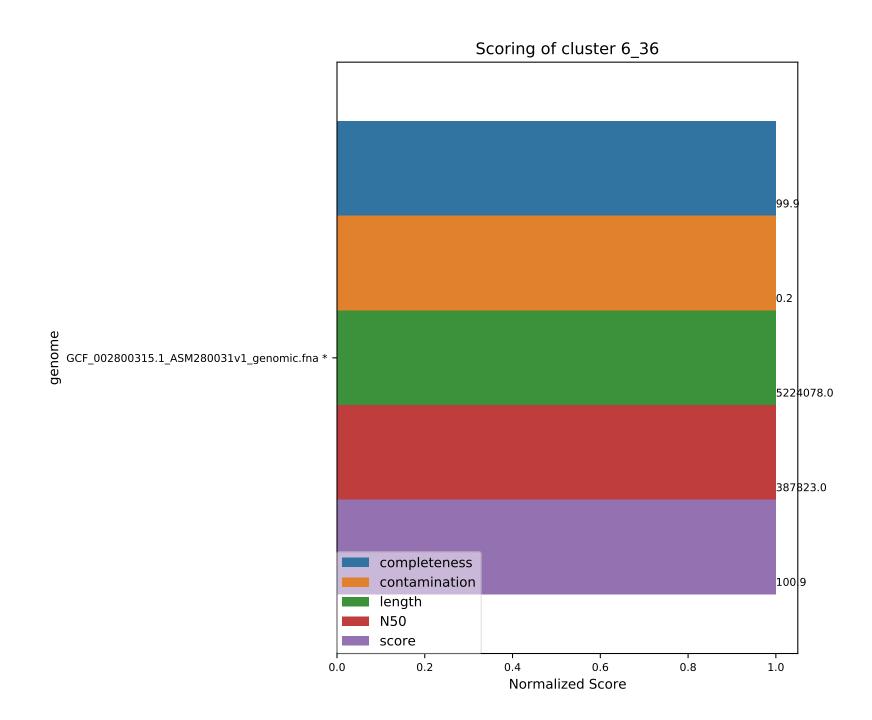


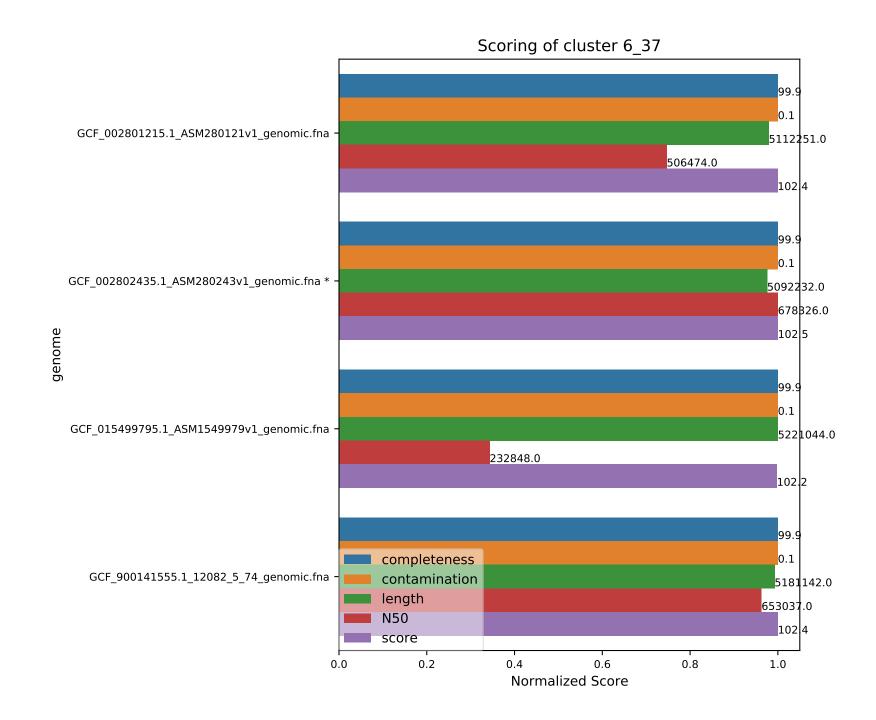


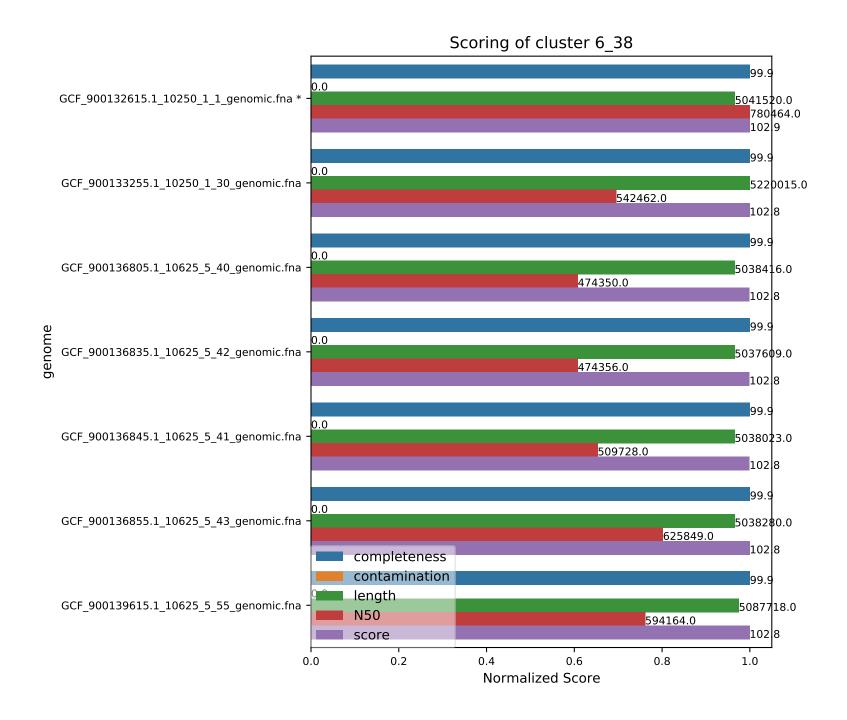


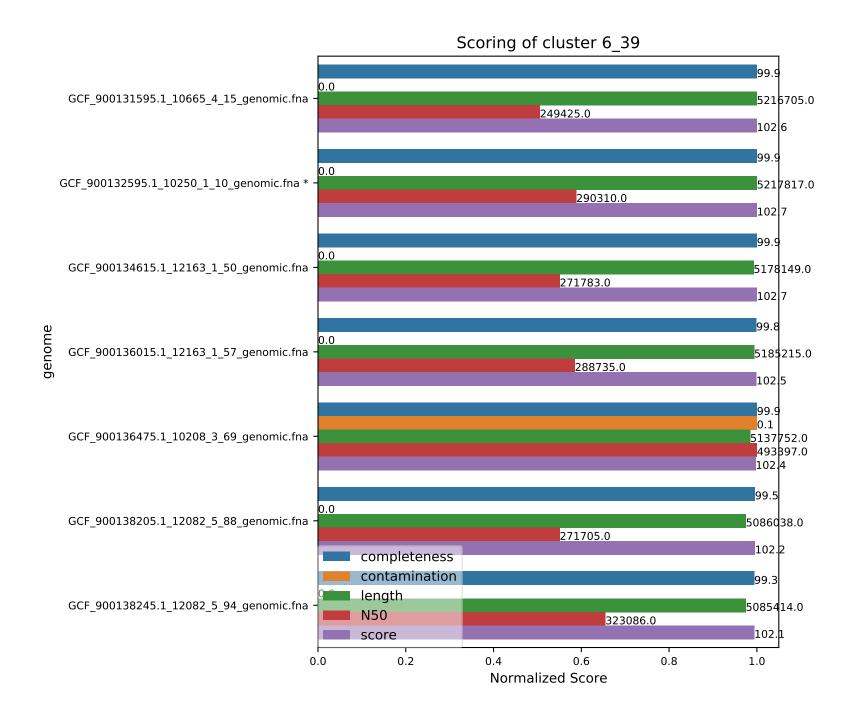


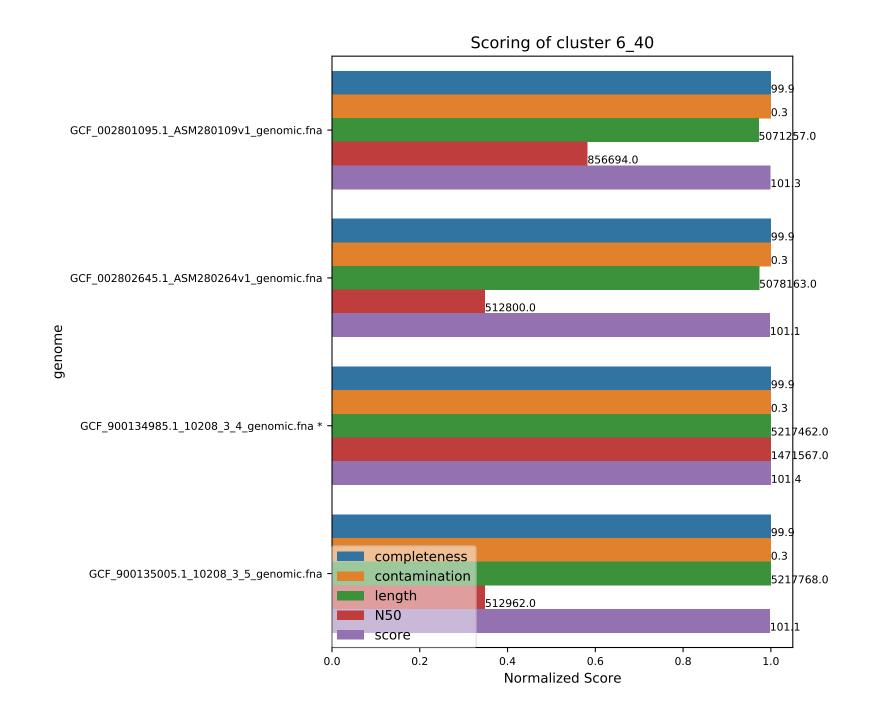


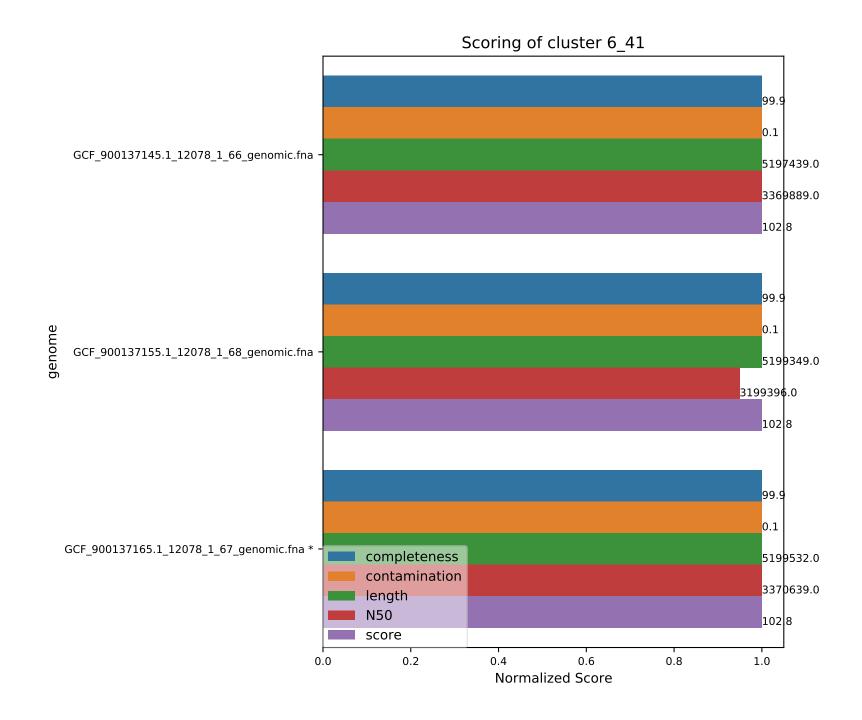


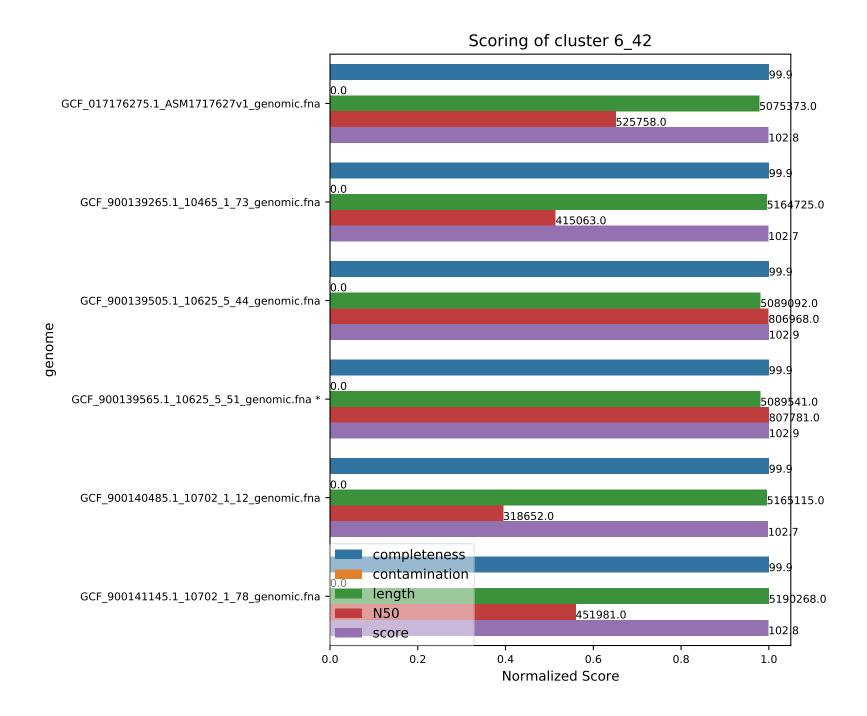


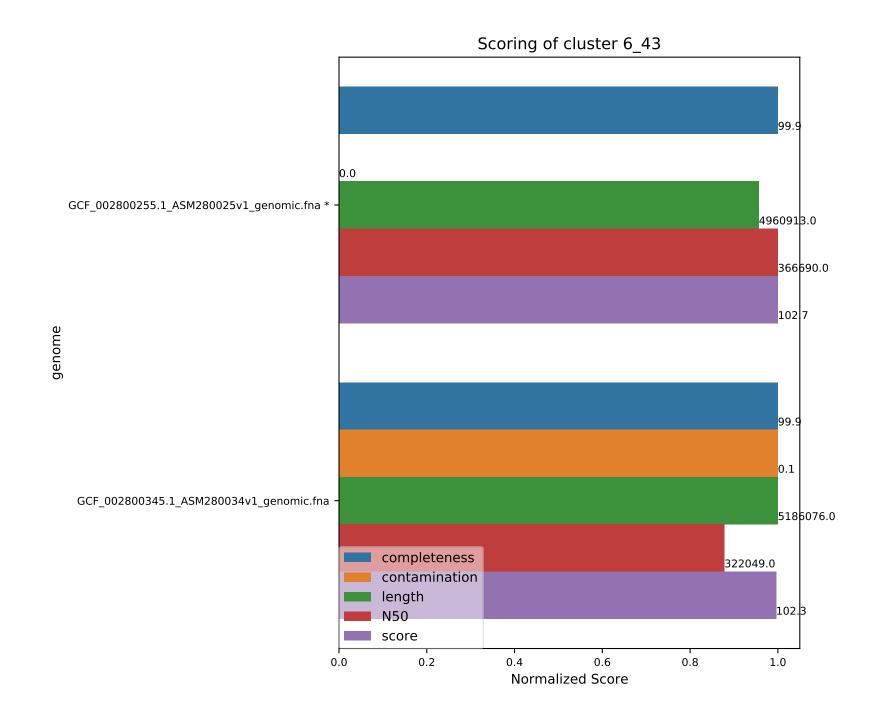


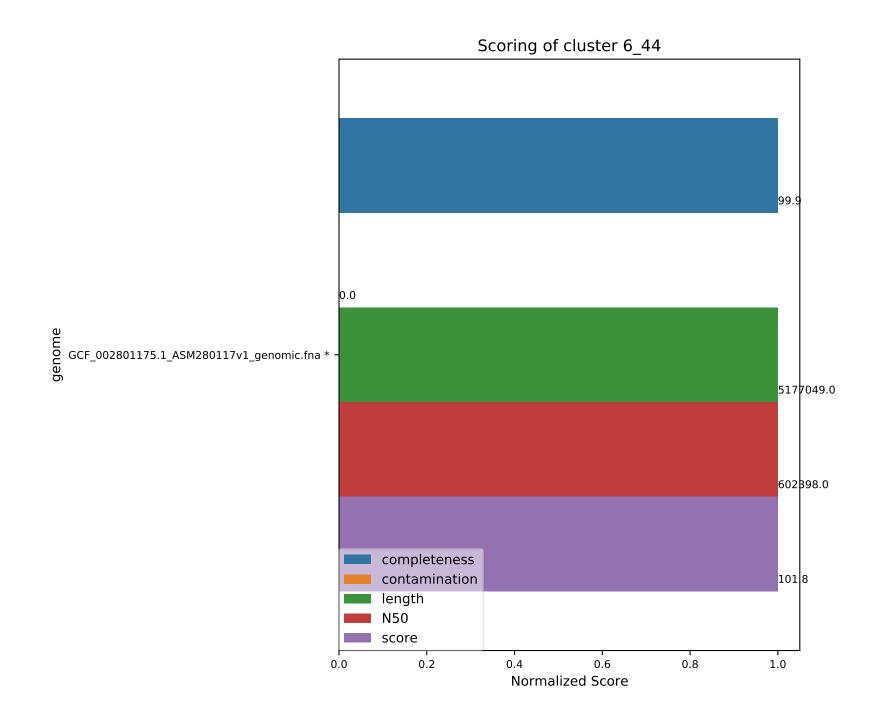


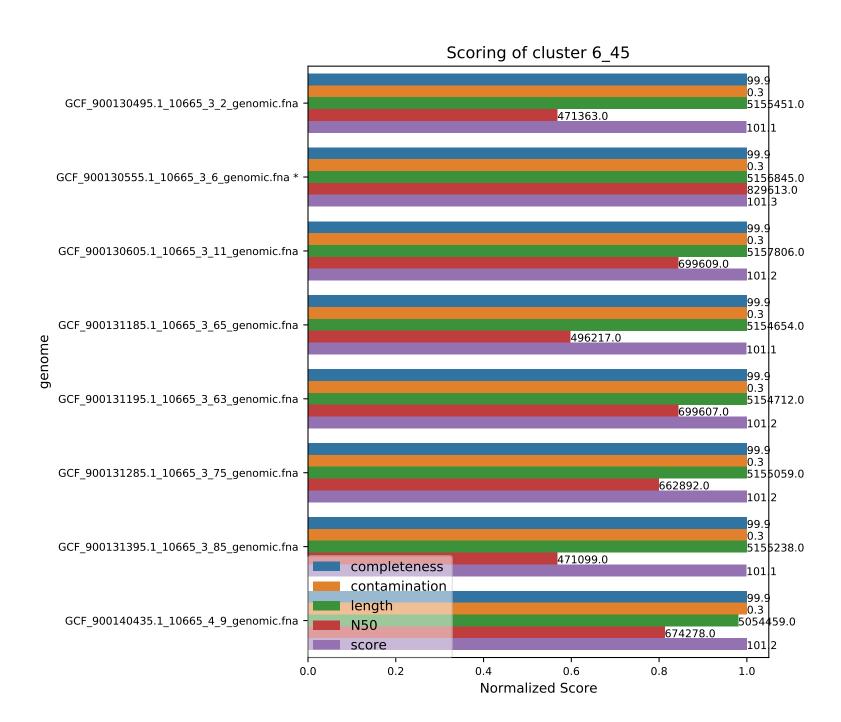


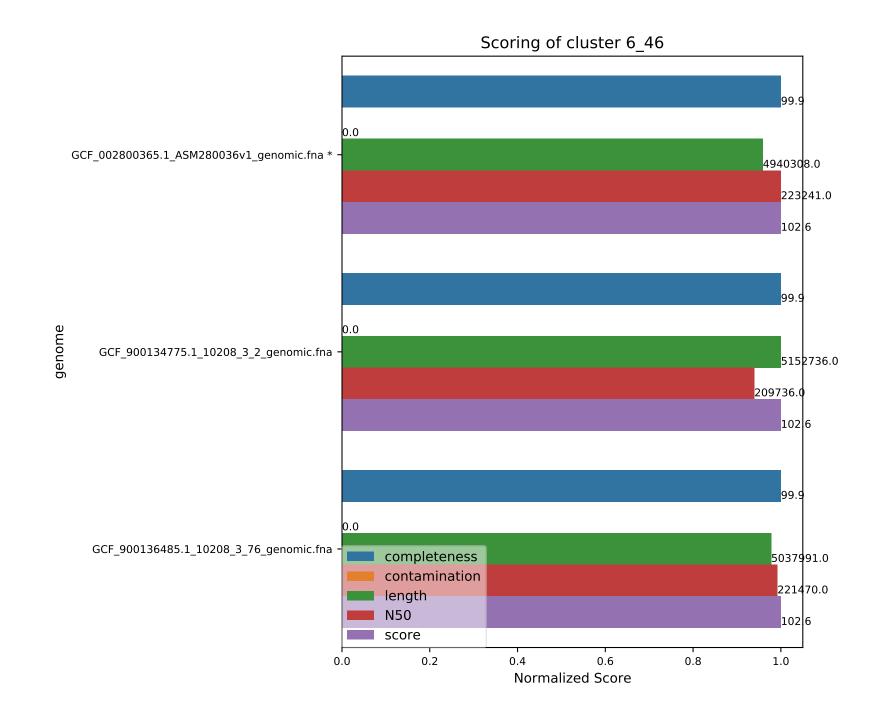


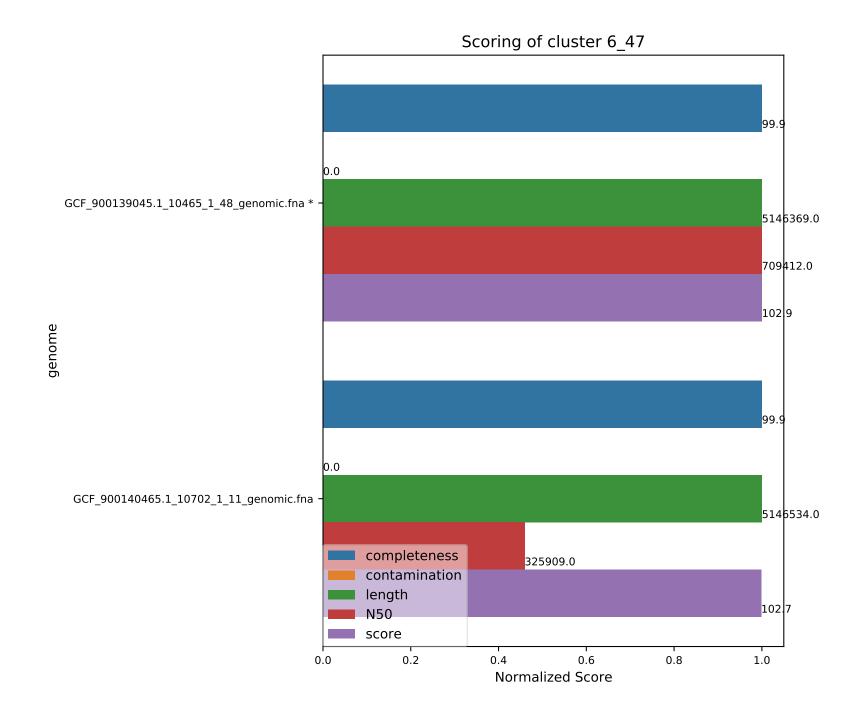


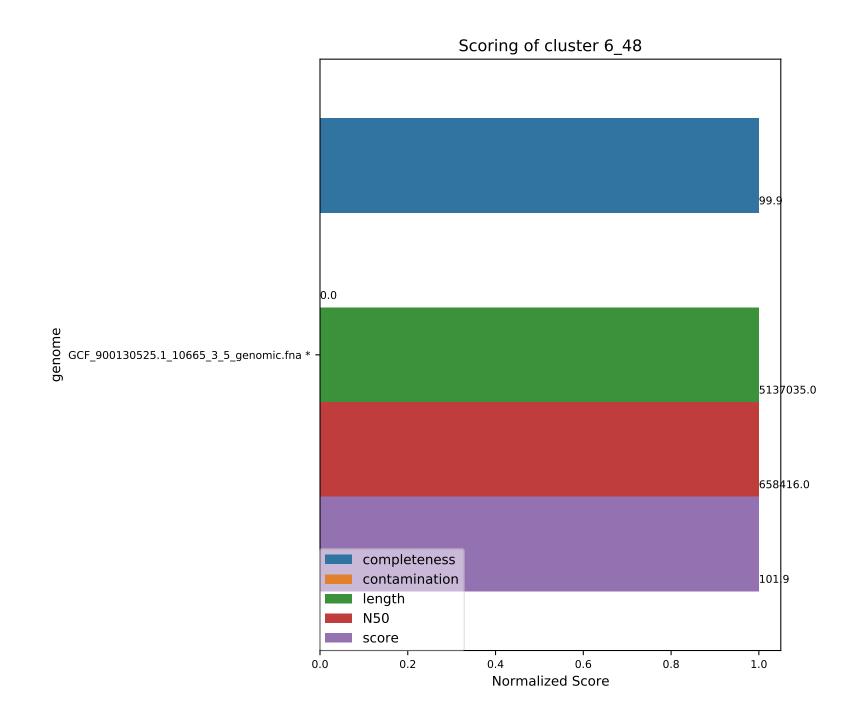


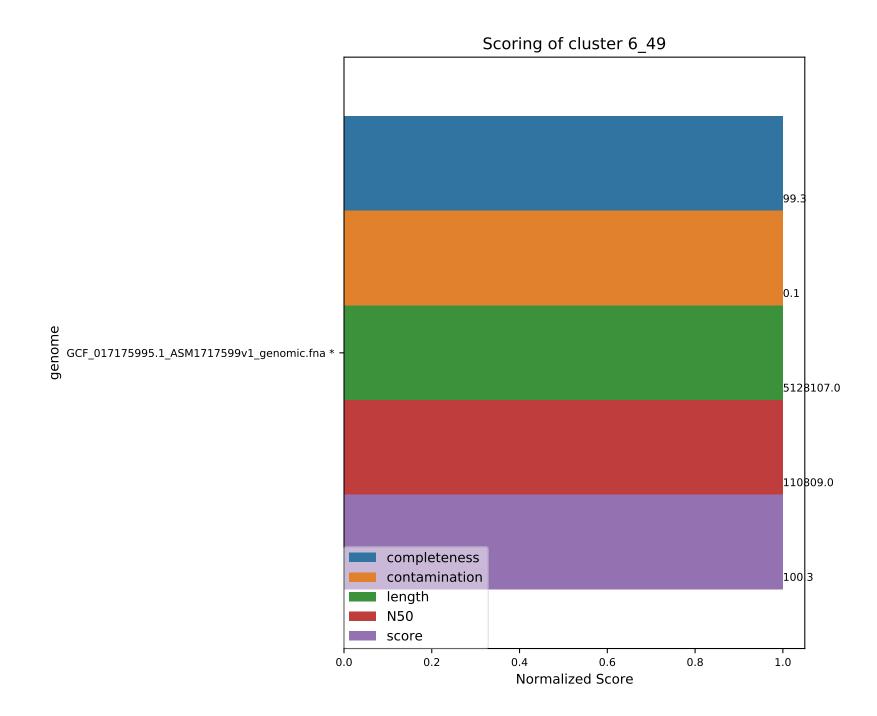


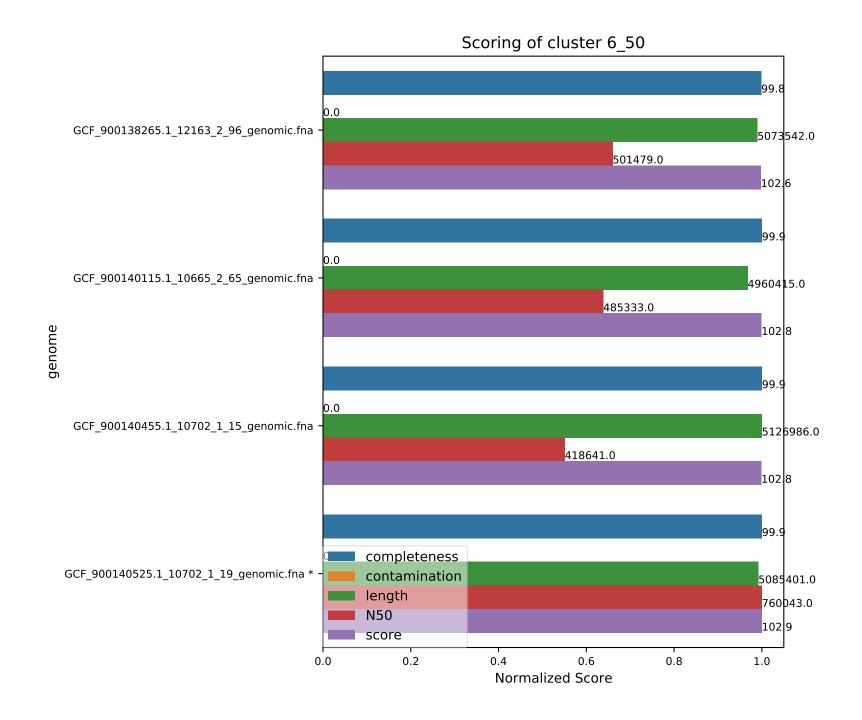


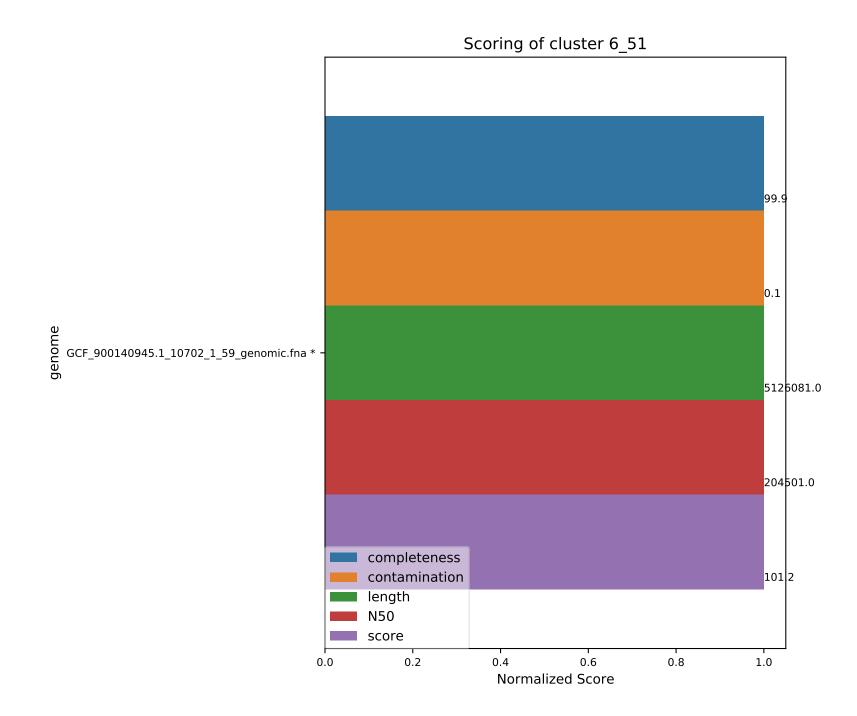


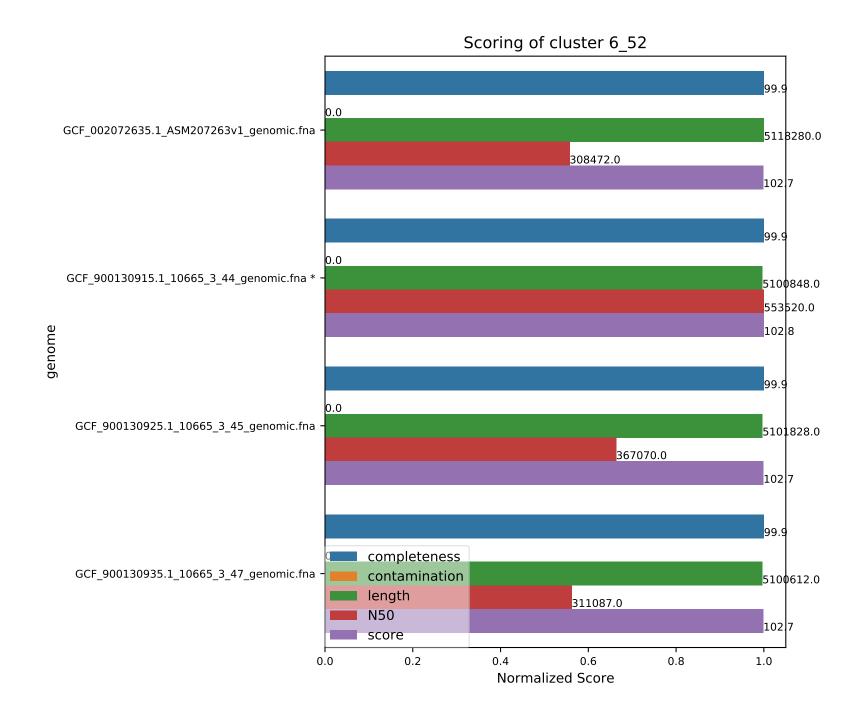


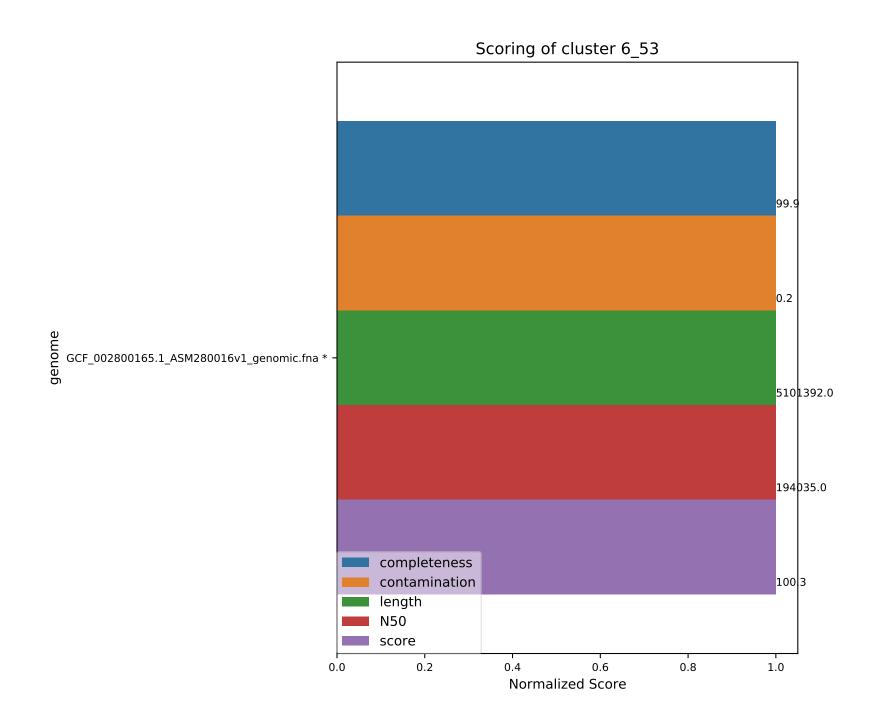


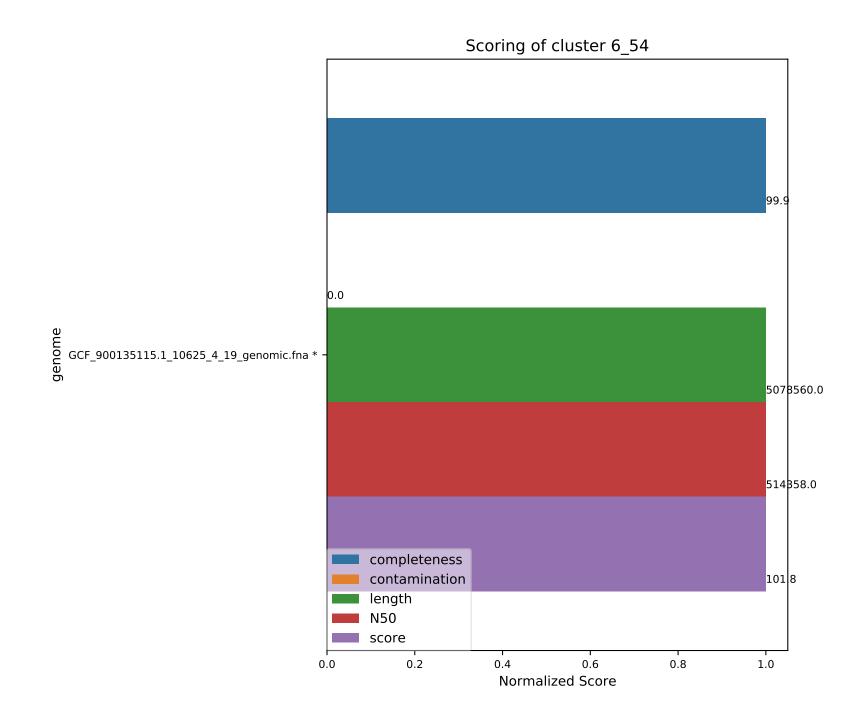


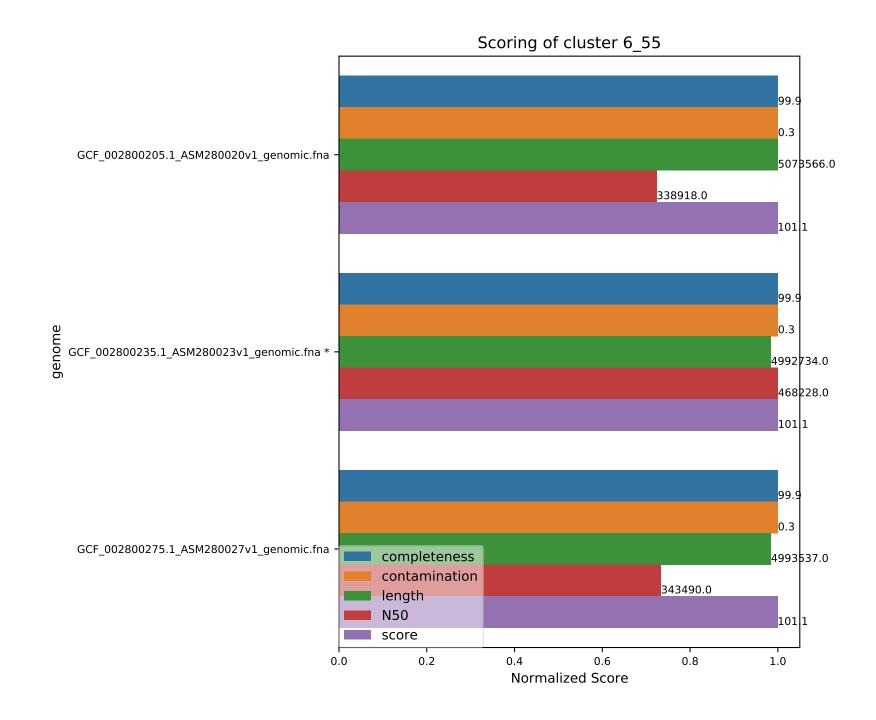


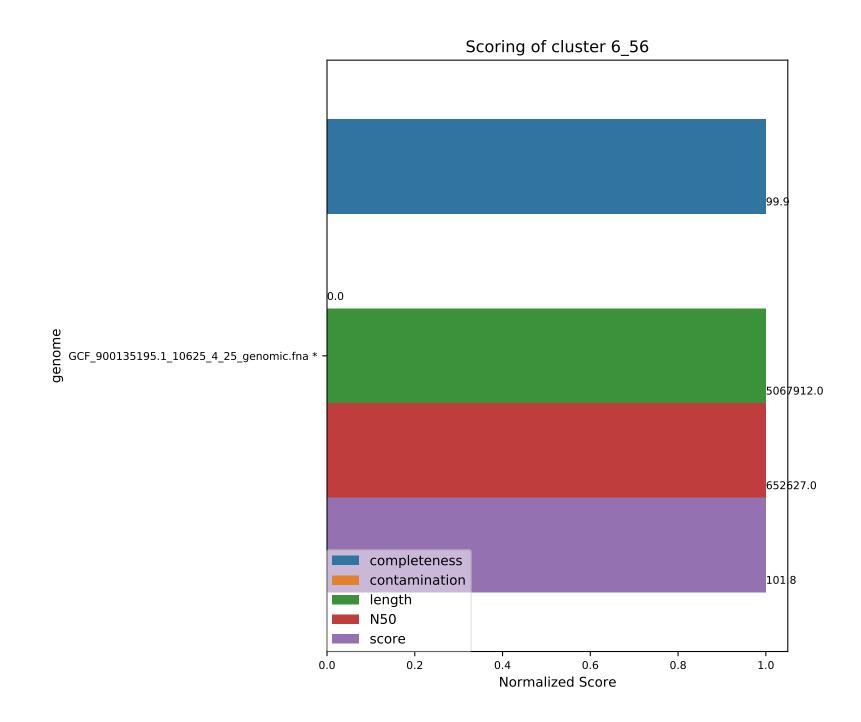


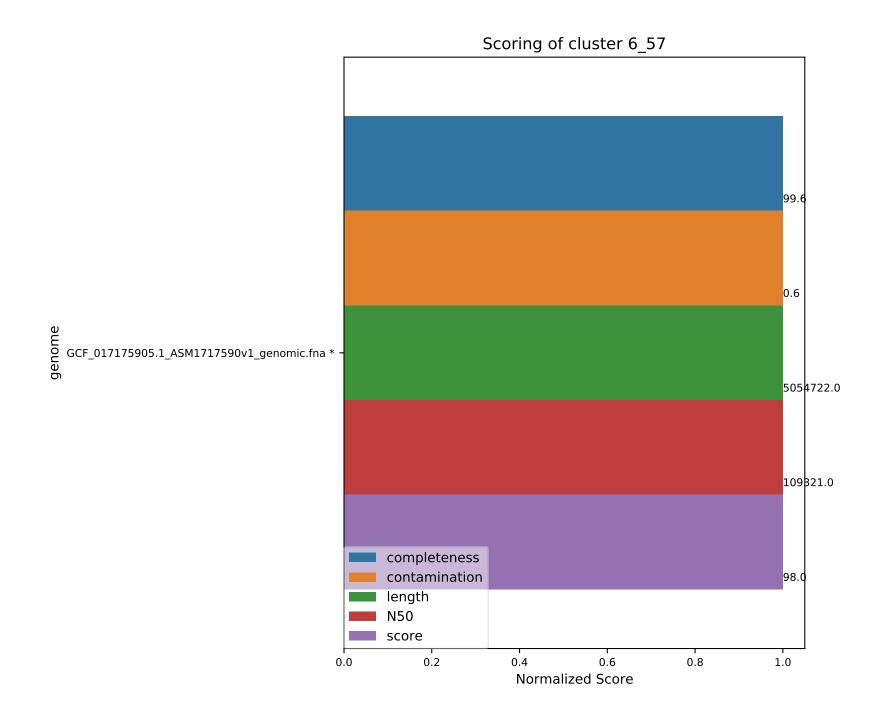


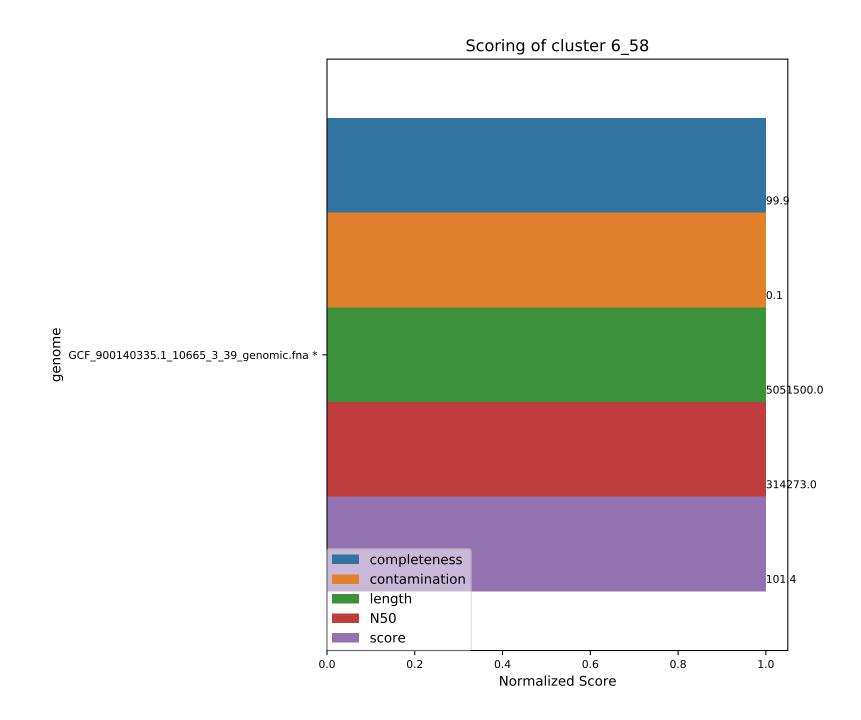


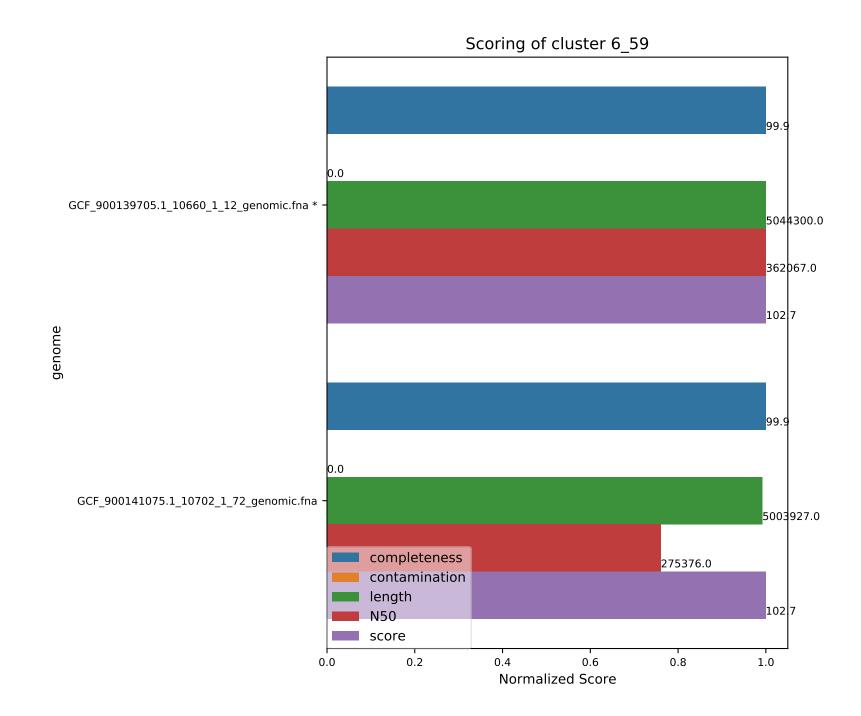


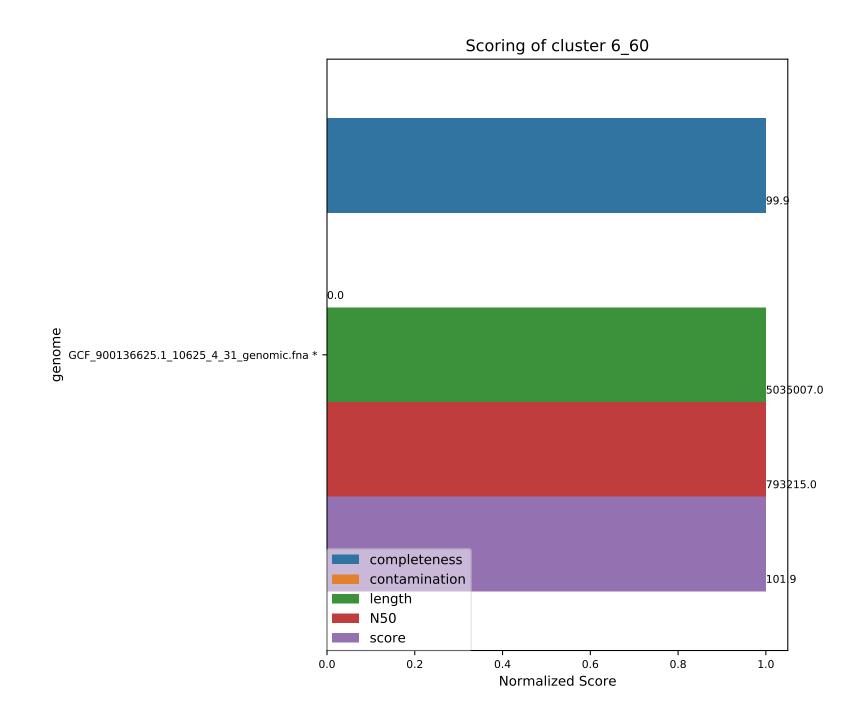


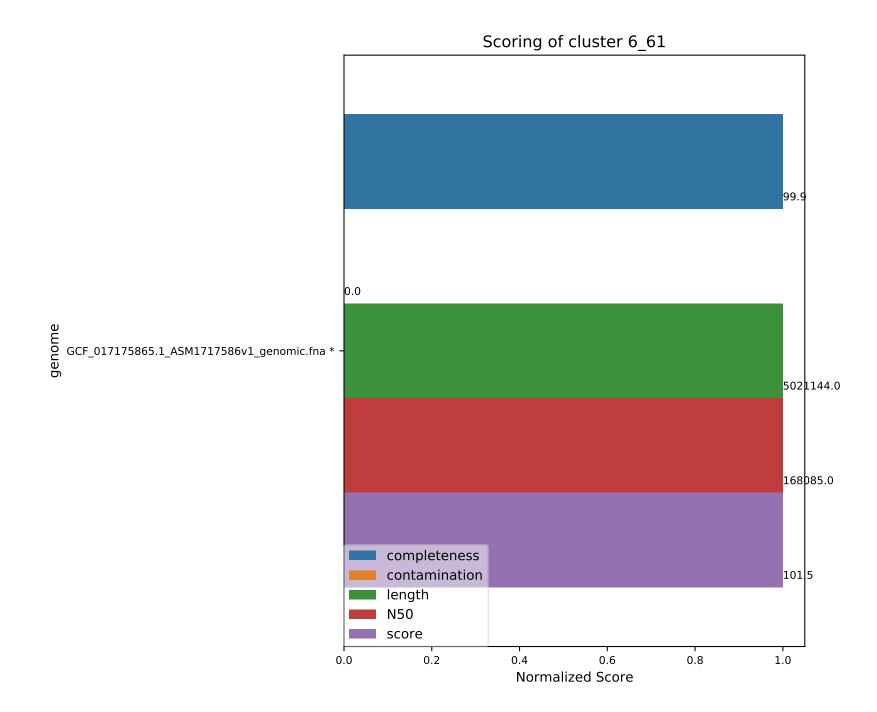


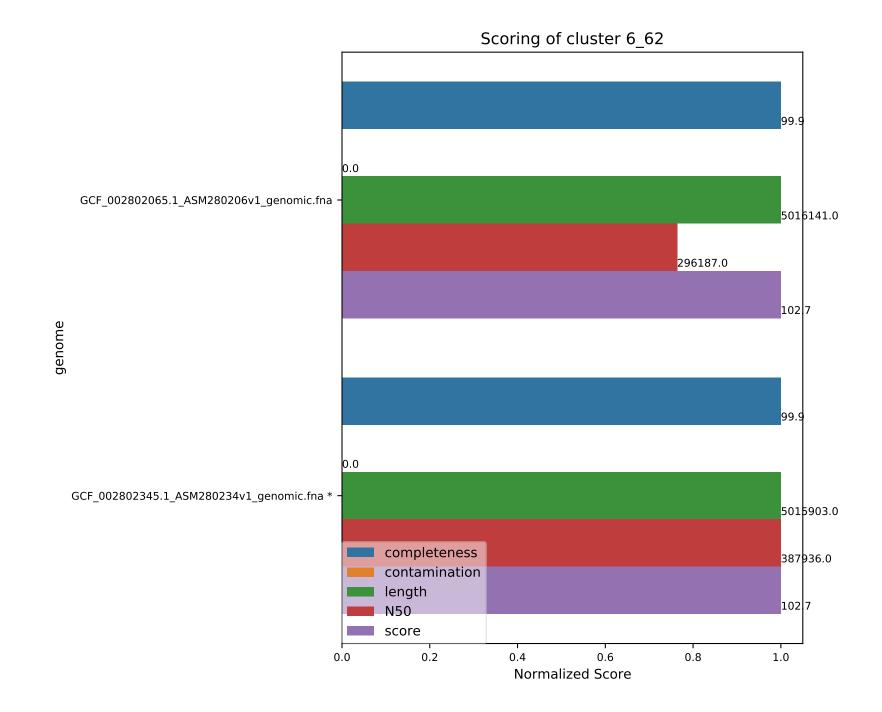


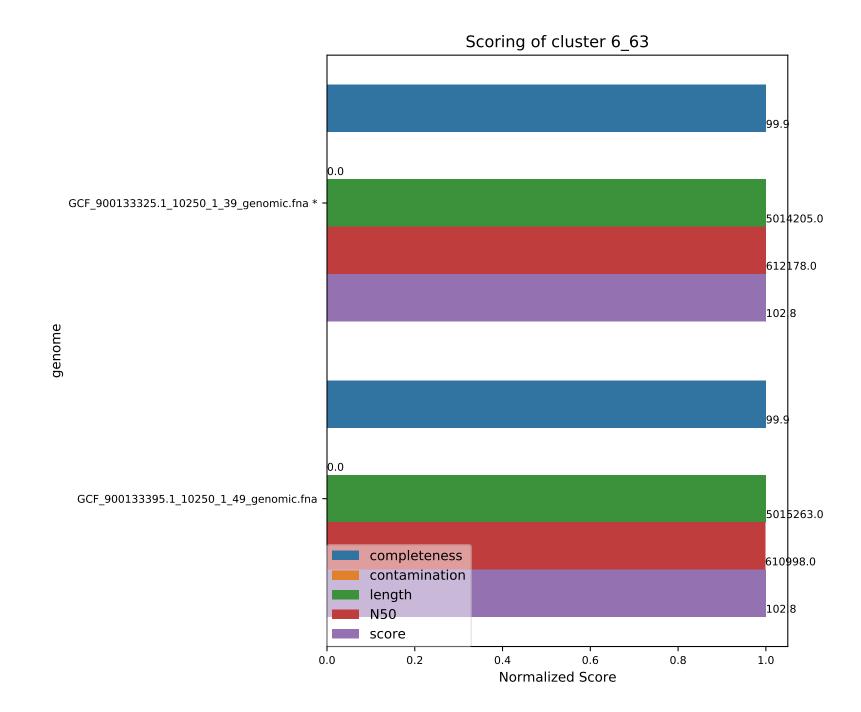


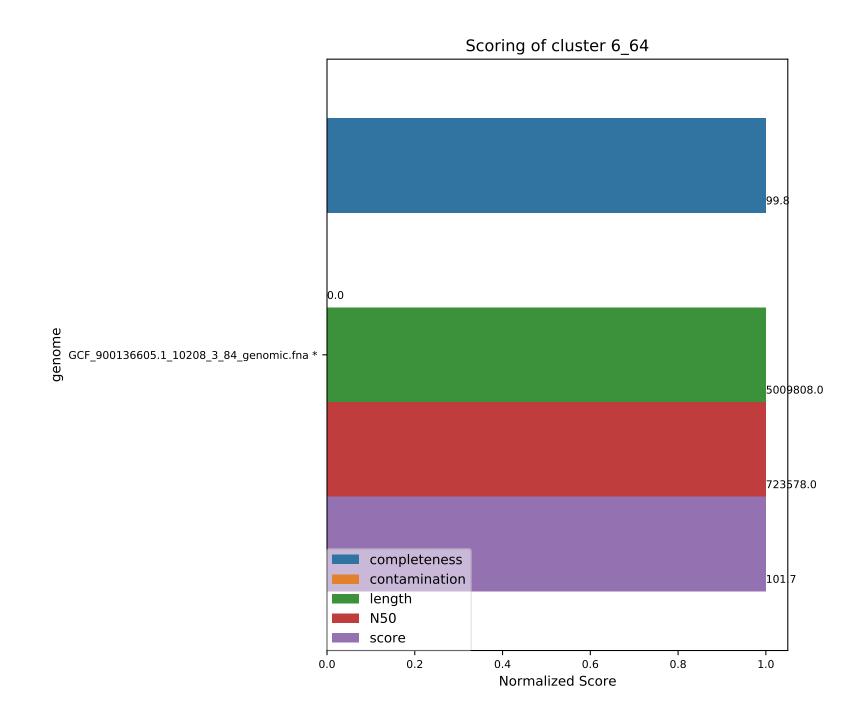


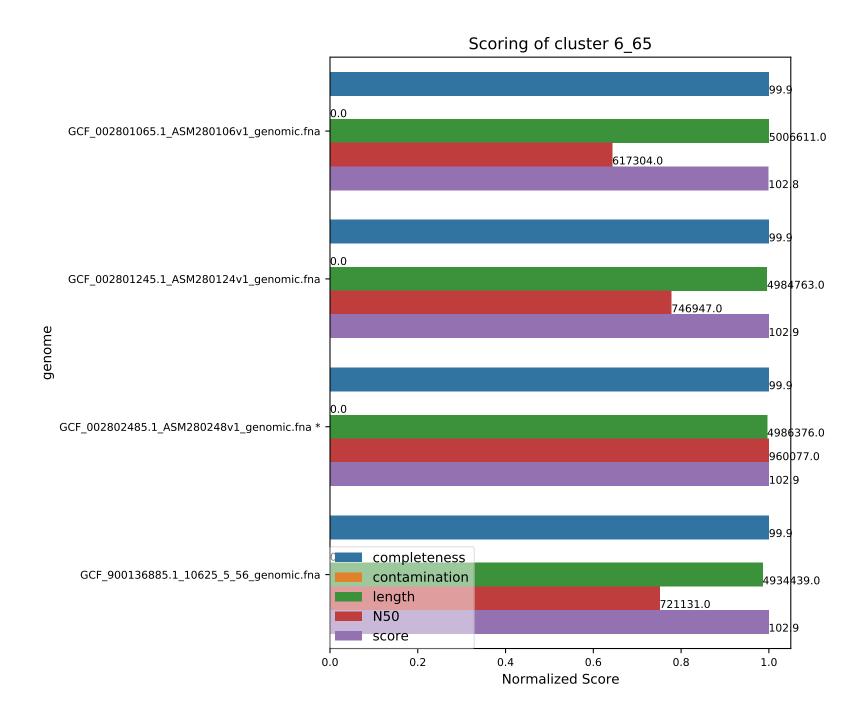


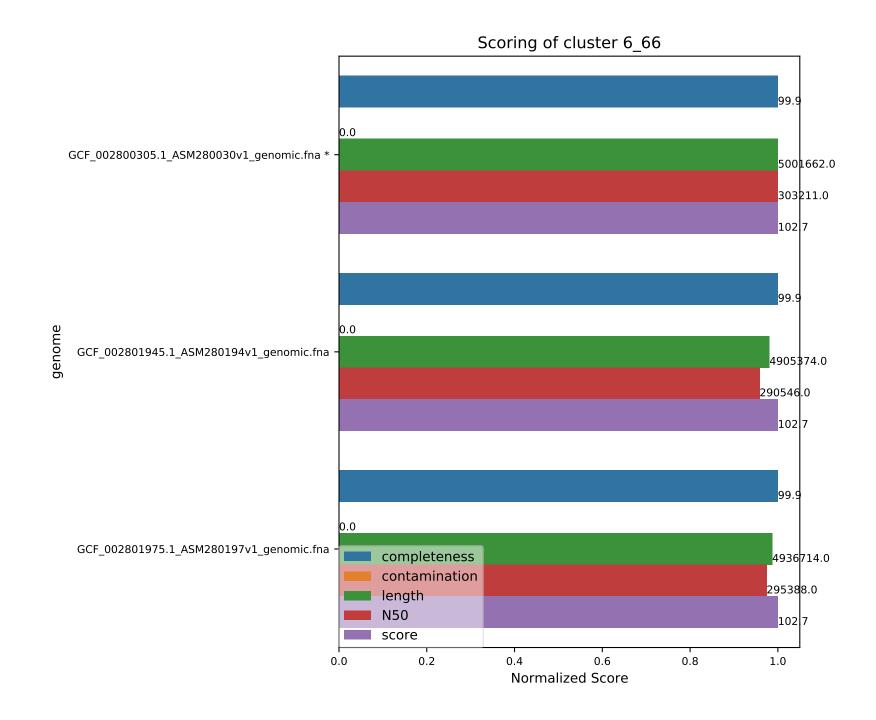


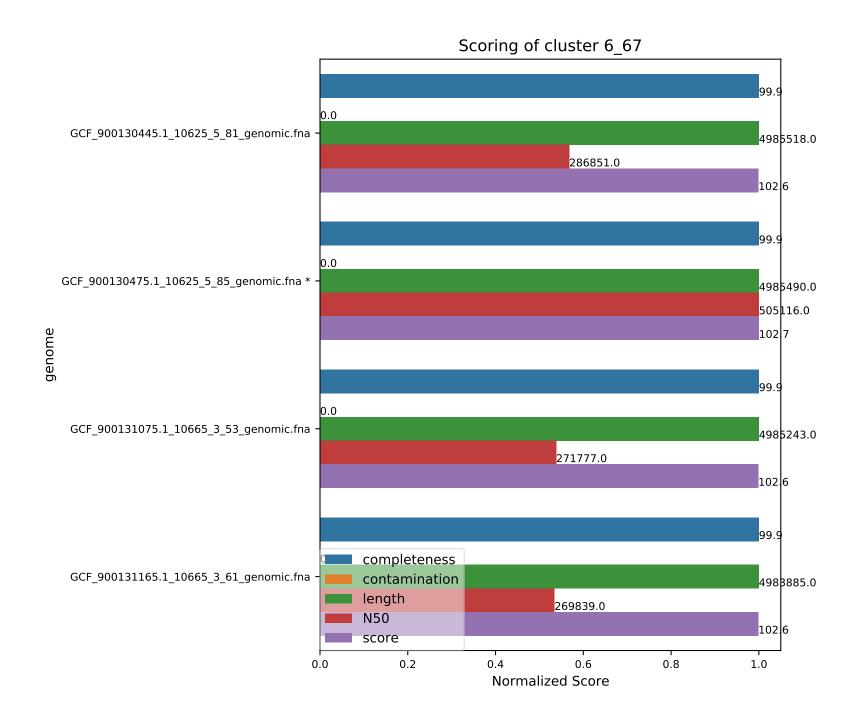


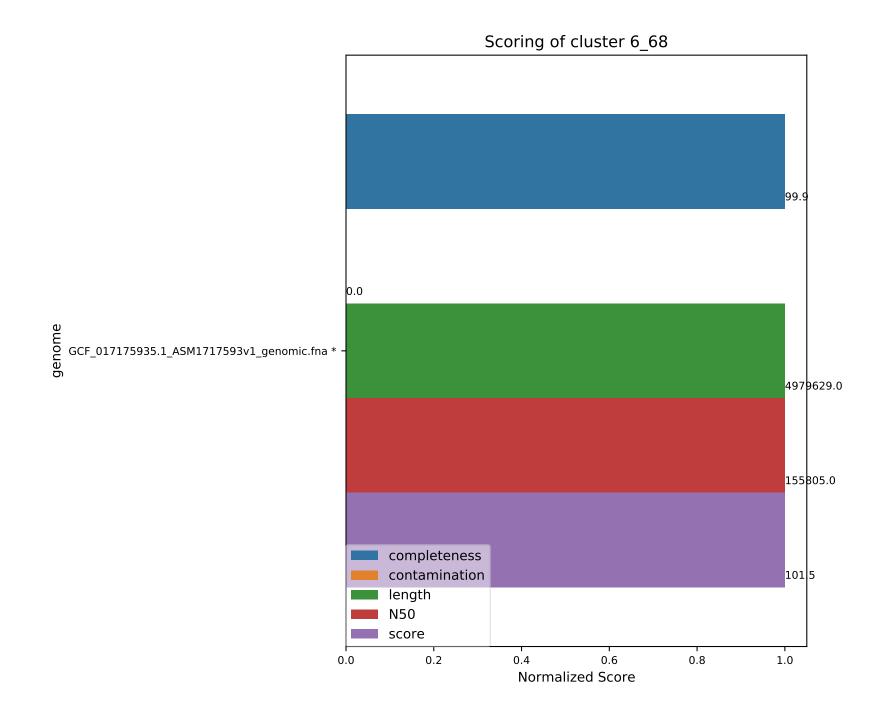


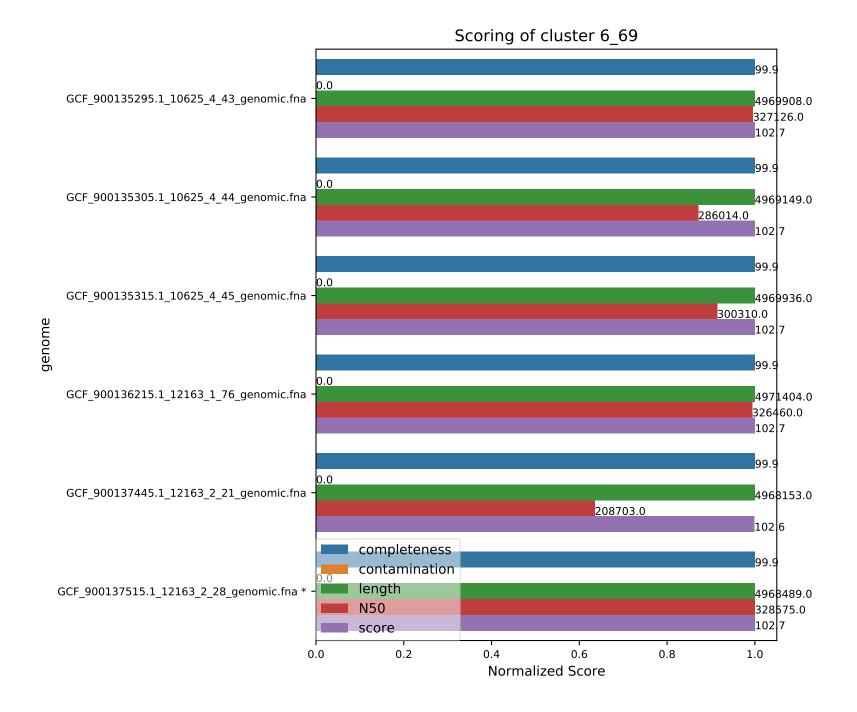


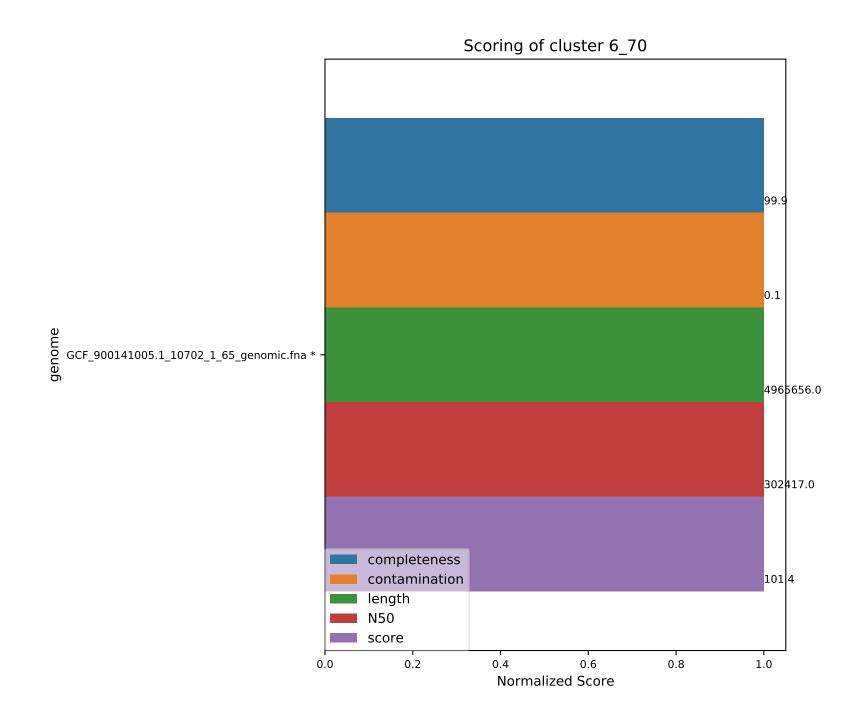


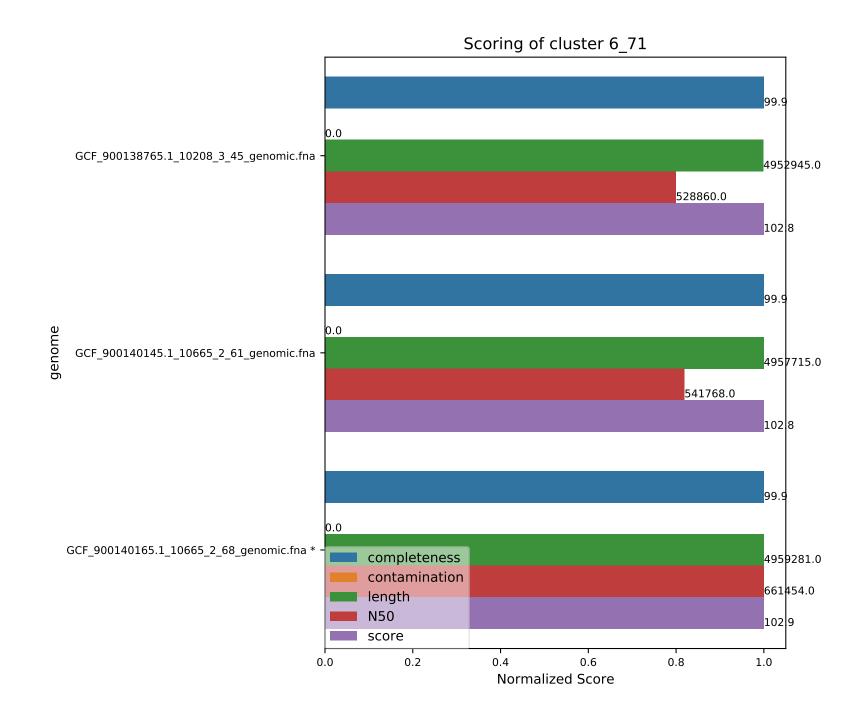


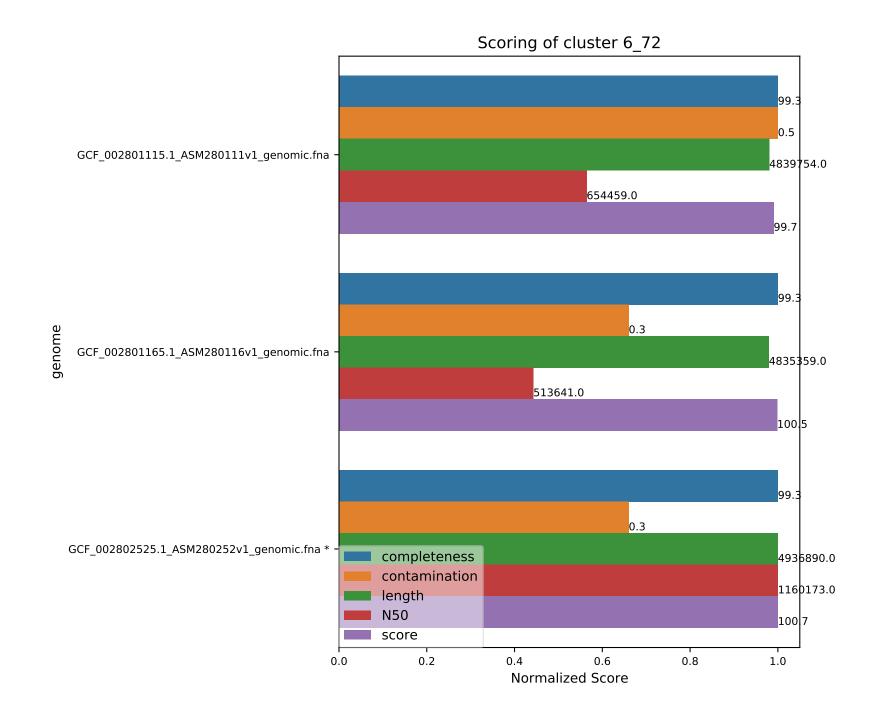


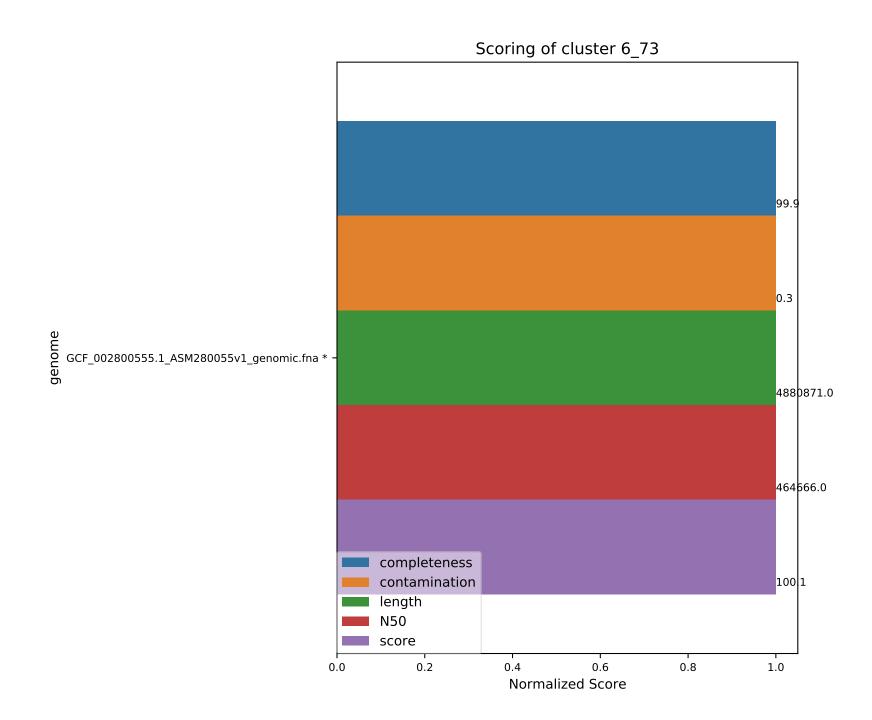


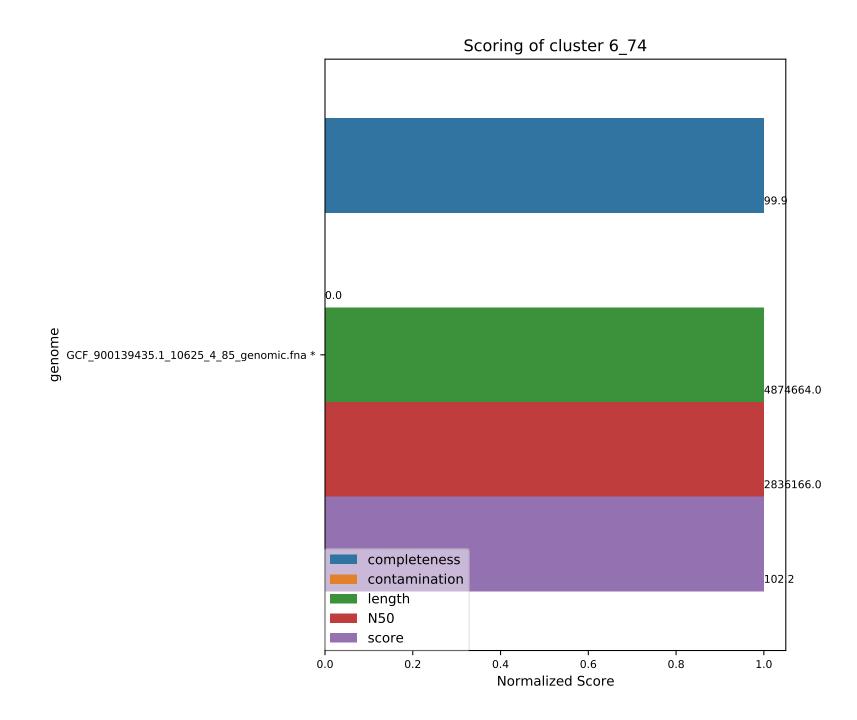


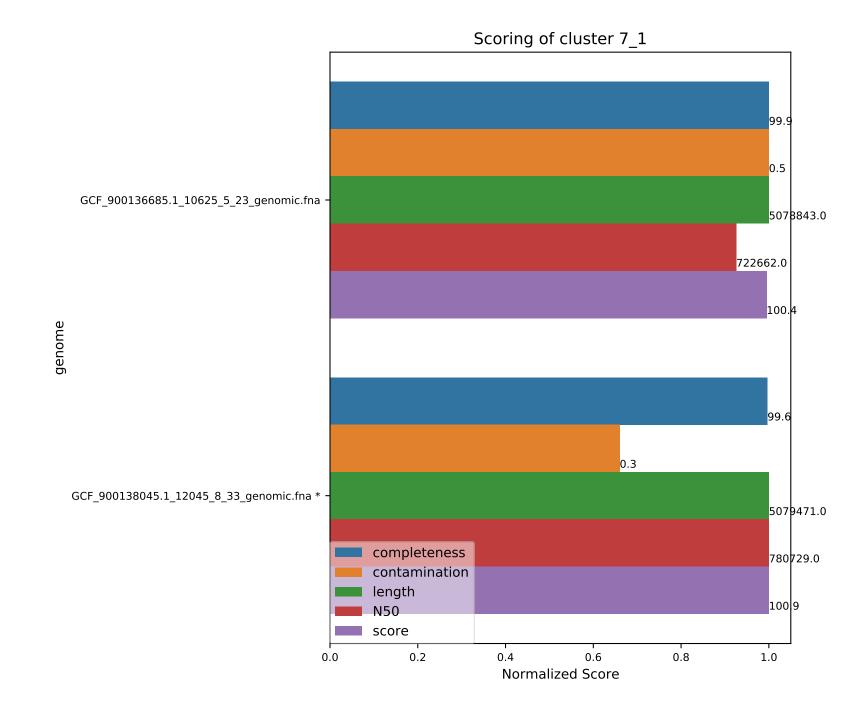












0.0

0.0

436574.0

391147.0

436629.0

0.4

62.6

702438.0

654367.0

0.6

579321.0

Normalized Score

313780.0

completeness contamination

0.2

length

score

■ N50

0.0

5180807.0

5231098.0 1036010.0

5108002.0

5186770.0

5173982.0

5068812.0

5187051.0

8.0

102.8

8.0

99.9

102 9

102.8

102 8

1.0

5874072.0

GCF_900137745.1_12163_2_51_genomic.fna

GCF_900137805.1_12163_2_54_genomic.fna

GCF 900137965.1 10660 1 4 genomic.fna

GCF_900138975.1_10208_3_65_genomic.fna

GCF_900139175.1_10465_1_66_genomic.fna

GCF_900139185.1_10465_1_61_genomic.fna -

GCF_900139275.1_10465_1_74_genomic.fna

GCF_900141095.1_10702_1_76_genomic.fna

Scoring of cluster 8_1

386915.0

390767.0

466559.0

446721.0

427586.0

520627.0

317036.0

342961.0

388618.0

666085.0

GCF_002800405.1_ASM280040v1_genomic.fna -

 ${\sf GCF_002800445.1_ASM280044v1_genomic.fna}$

 ${\sf GCF_002800495.1_ASM280049v1_genomic.fna}$

 ${\sf GCF_002800545.1_ASM280054v1_genomic.fna}$

GCF_002800625.1_ASM280062v1_genomic.fna

 ${\sf GCF_002800665.2_ASM280066v2_genomic.fna}$

GCF_002800715.1_ASM280071v1_genomic.fna

GCF_002802055.1_ASM280205v1_genomic.fna -

GCF_002802255.1_ASM280225v1_genomic.fna -

 ${\sf GCF_017175795.1_ASM1717579v1_genomic.fna} \ \, \textbf{-}$

99.9

102.7

102.4

102 9

102.7

102.7

5280104.0

5232465.0

5231696.0

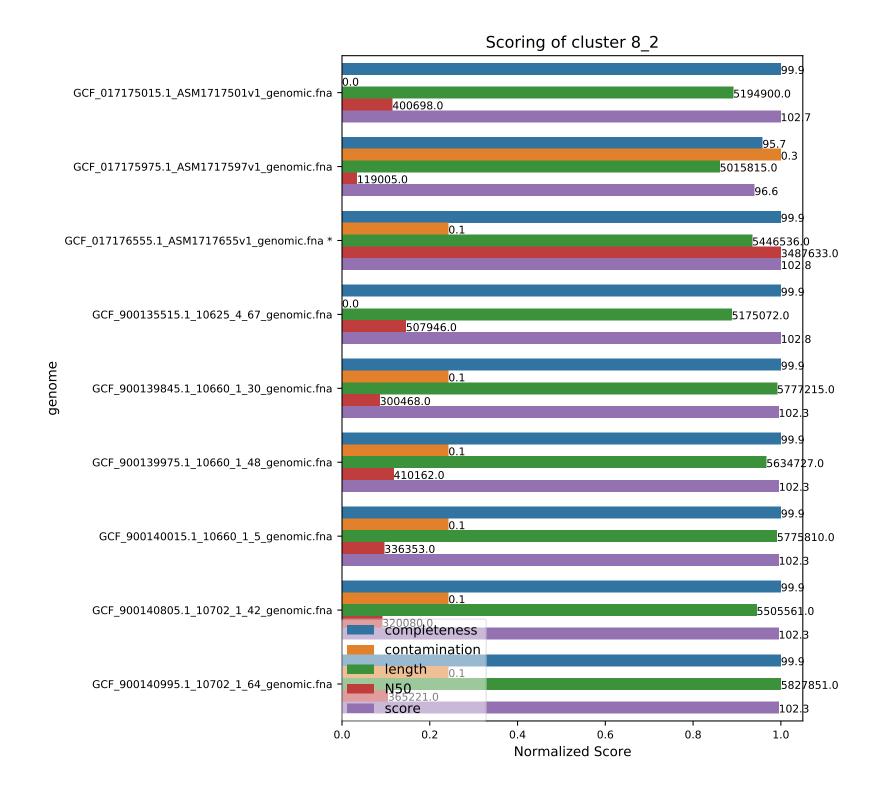
5192777.0

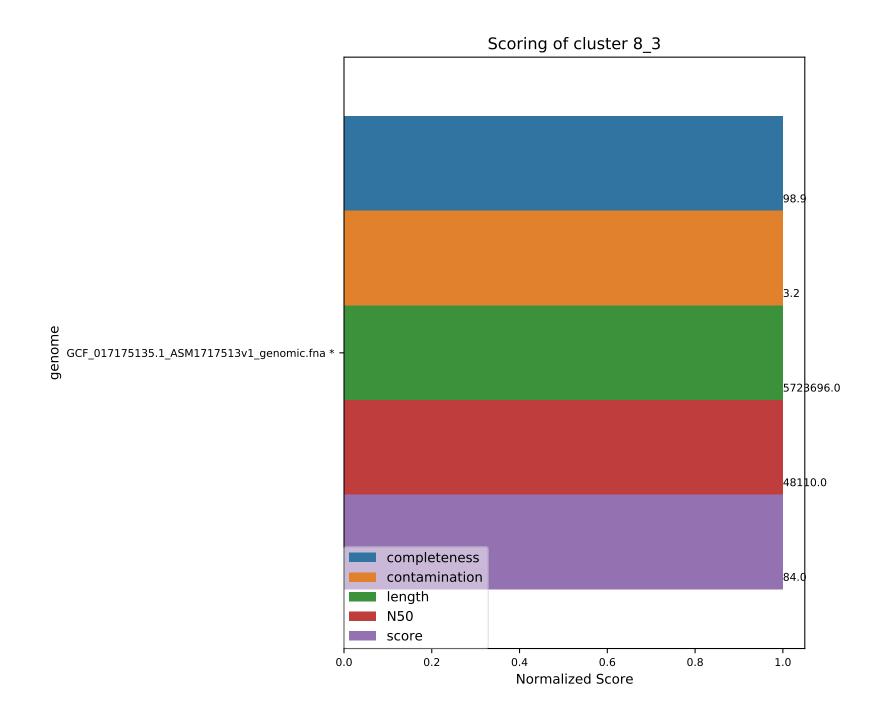
5254828.0

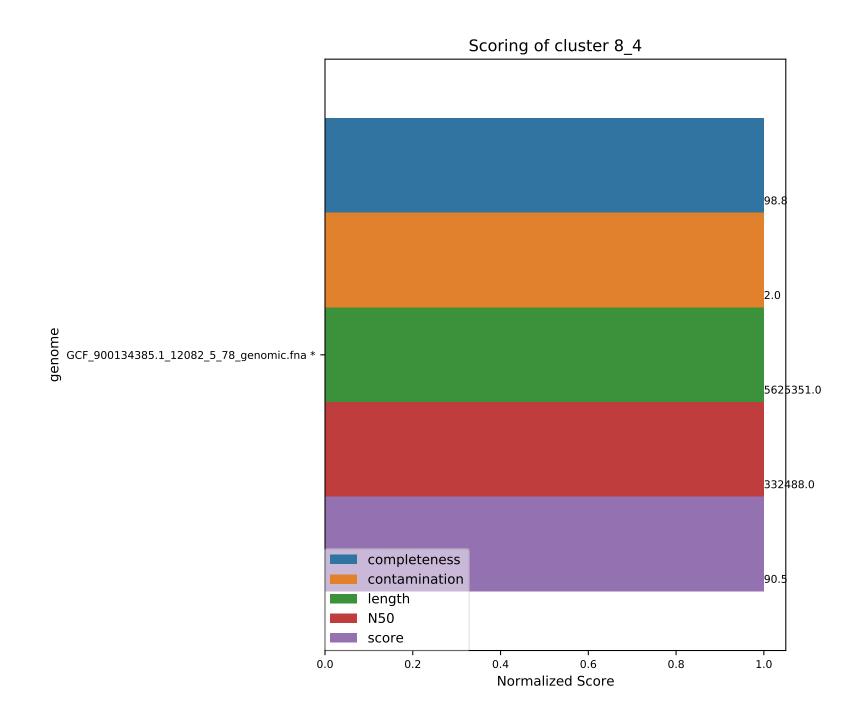
5109395.0

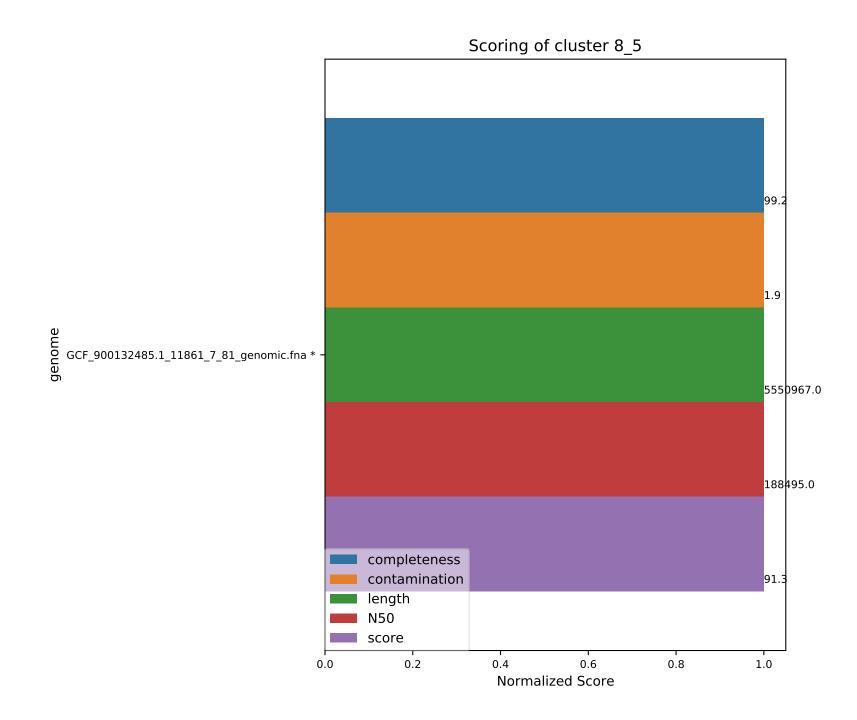
5134600.0

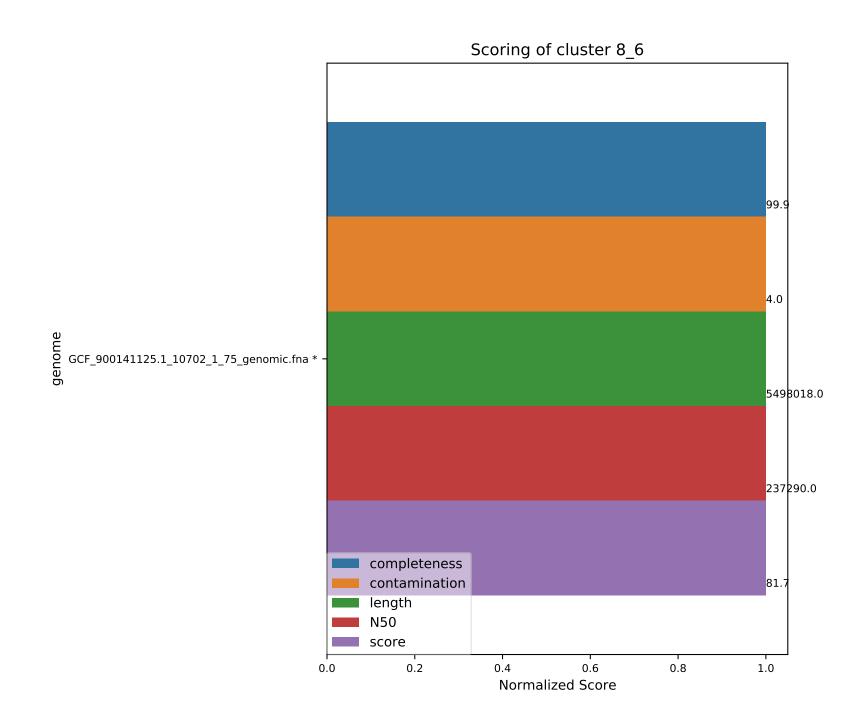
5054963.0

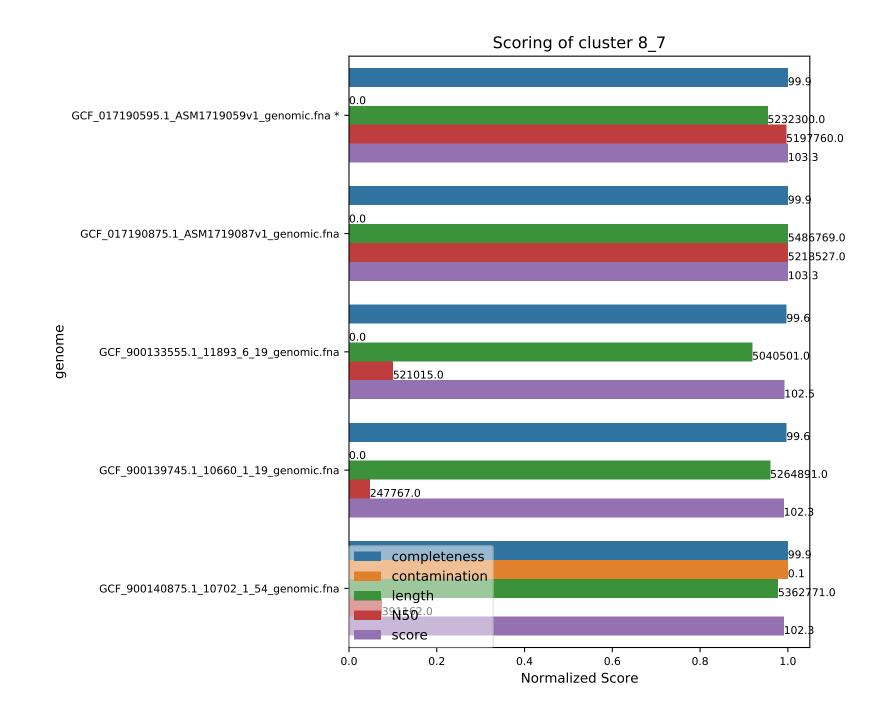


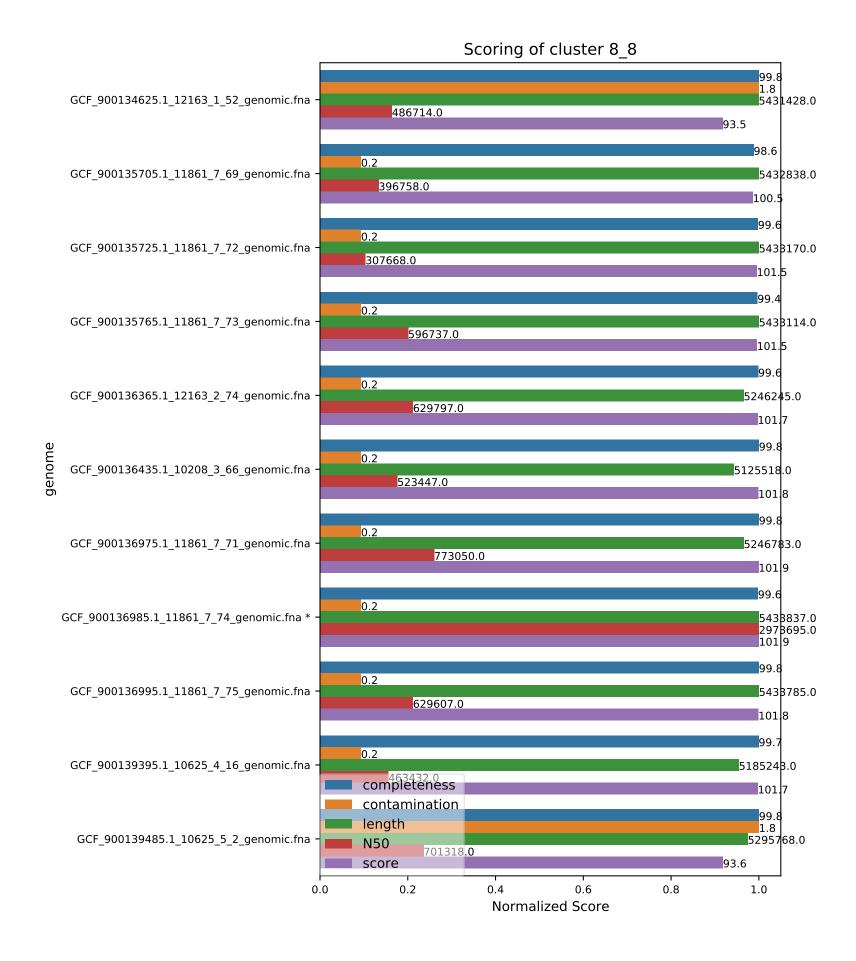


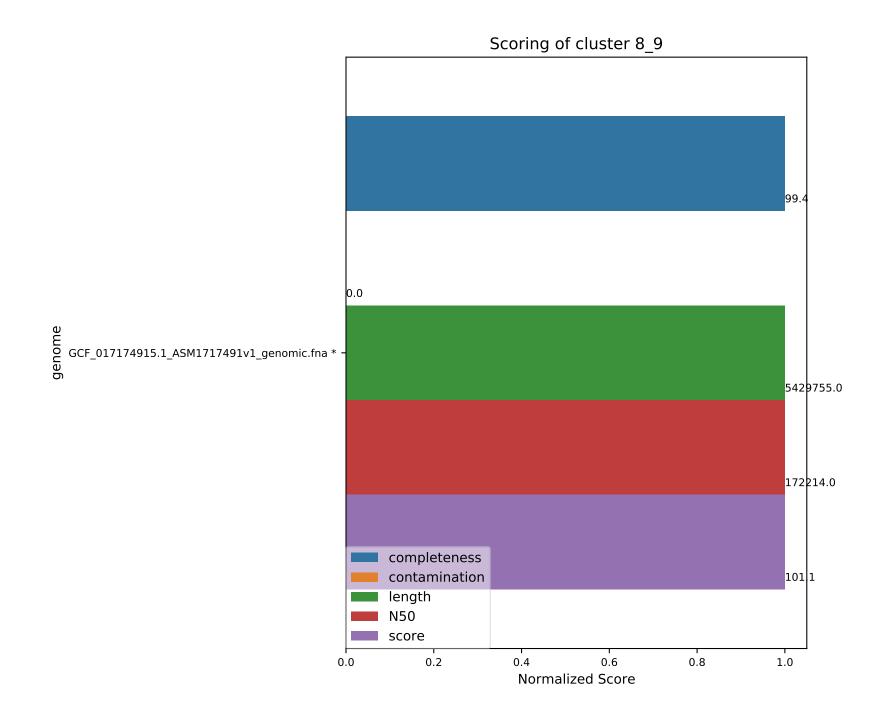


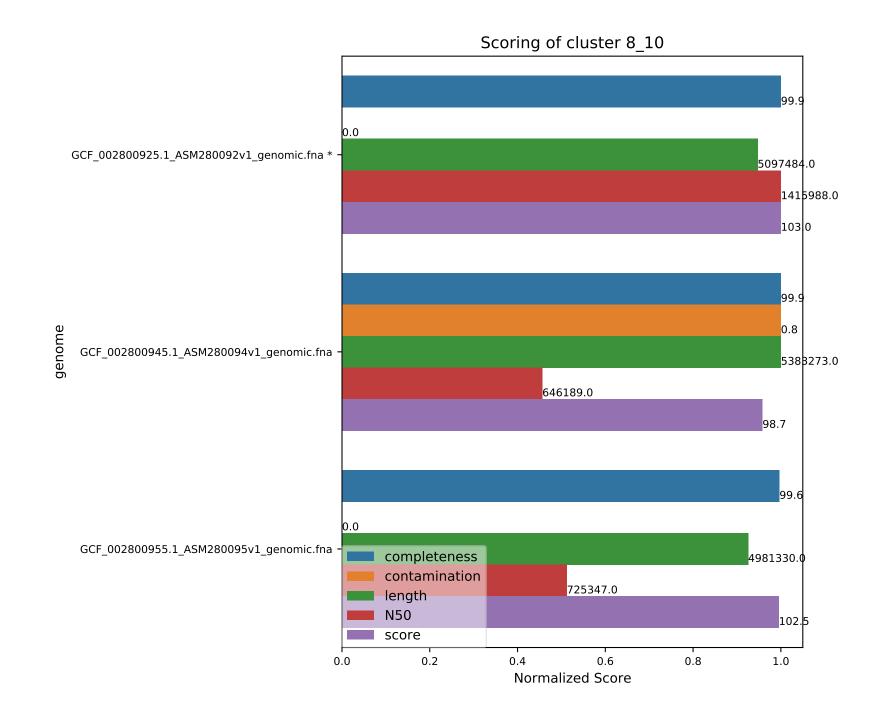


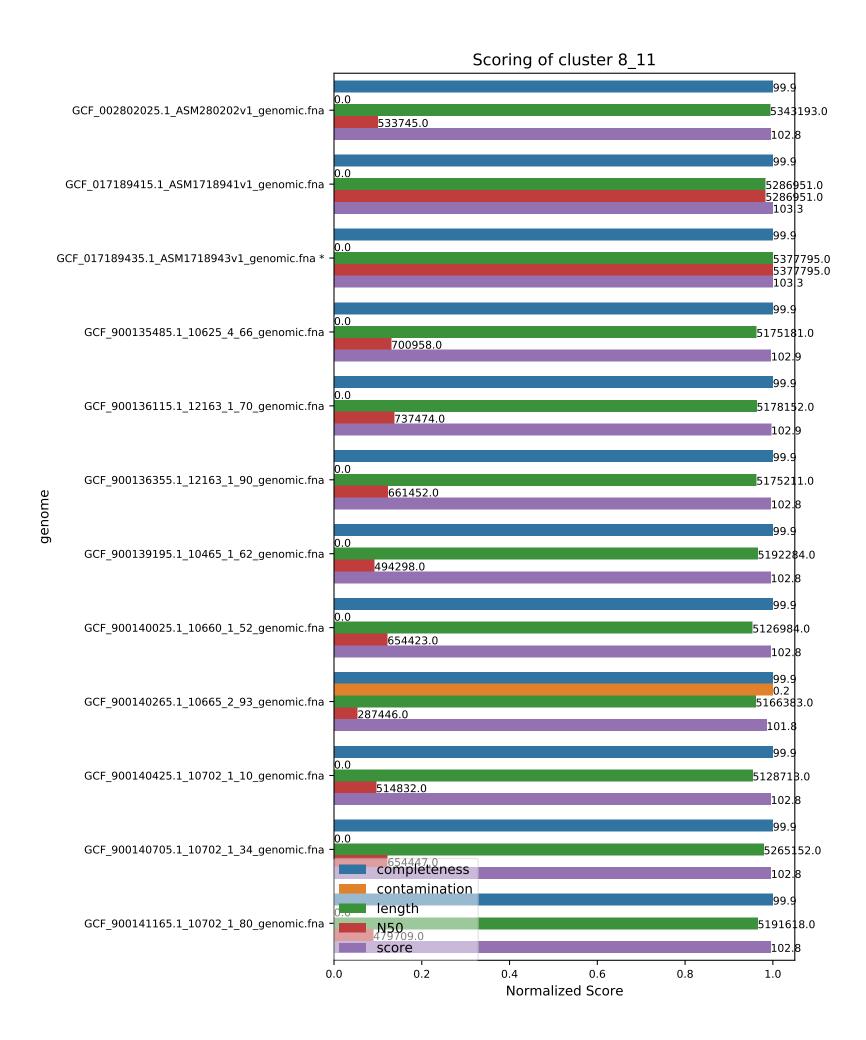


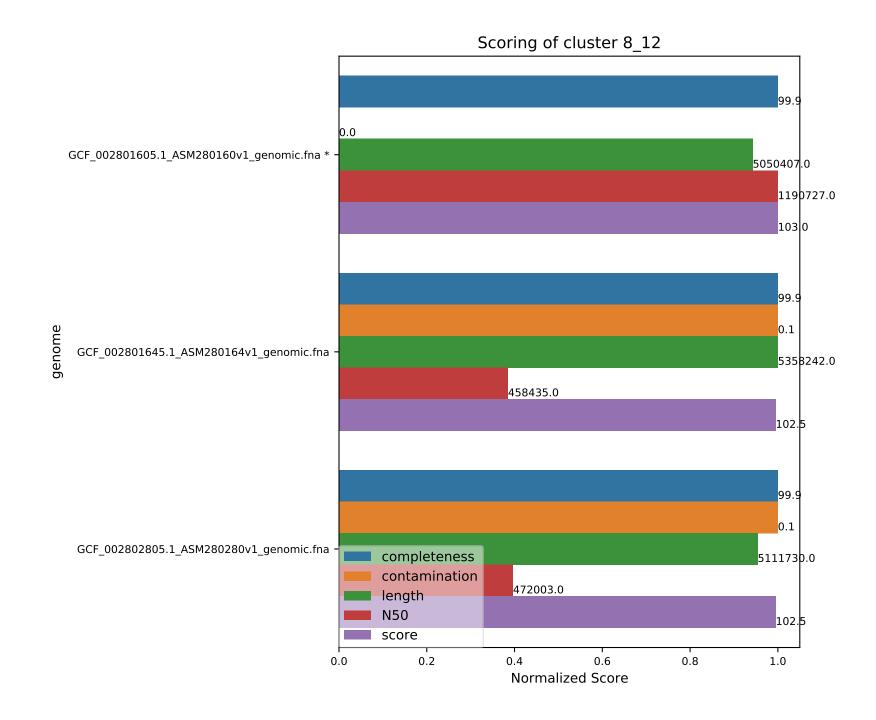


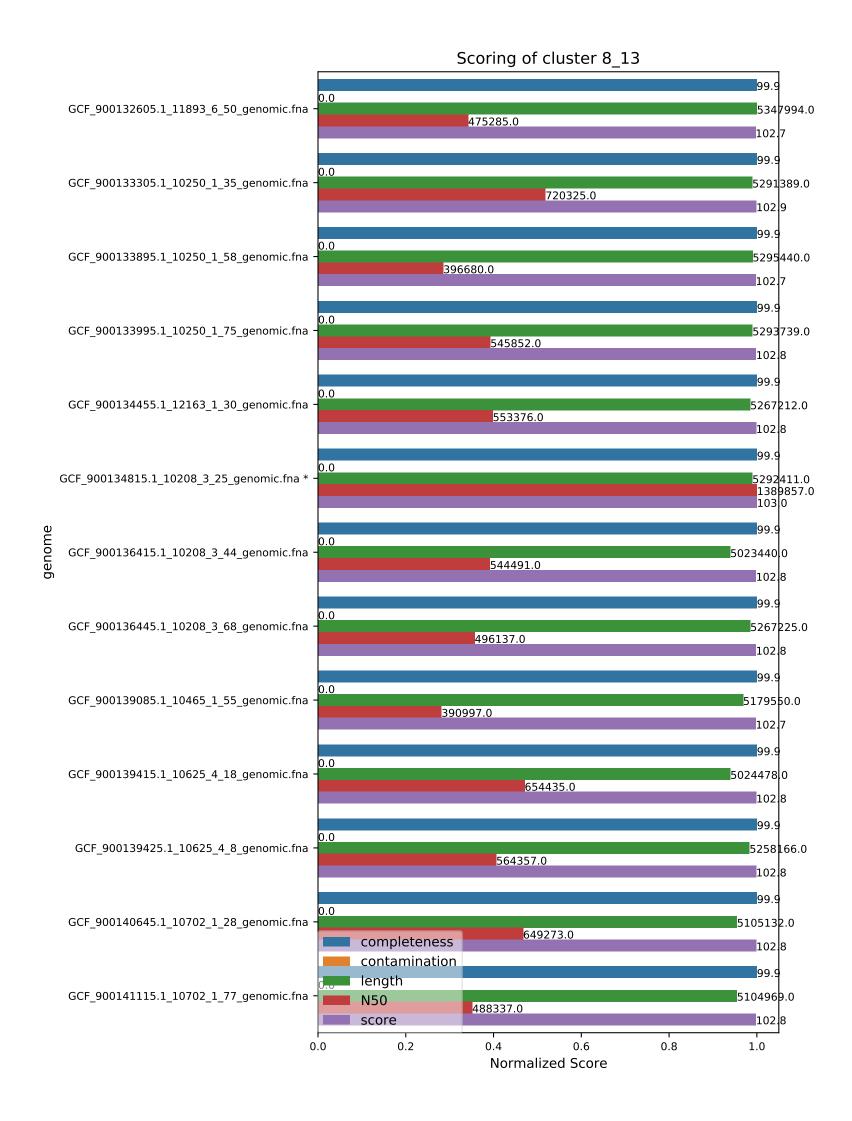


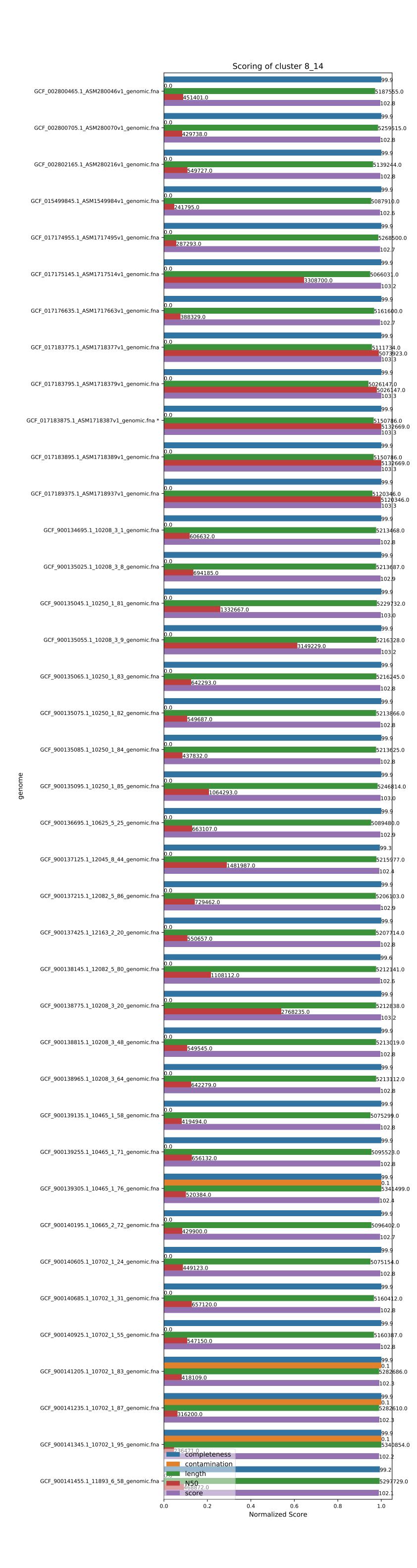


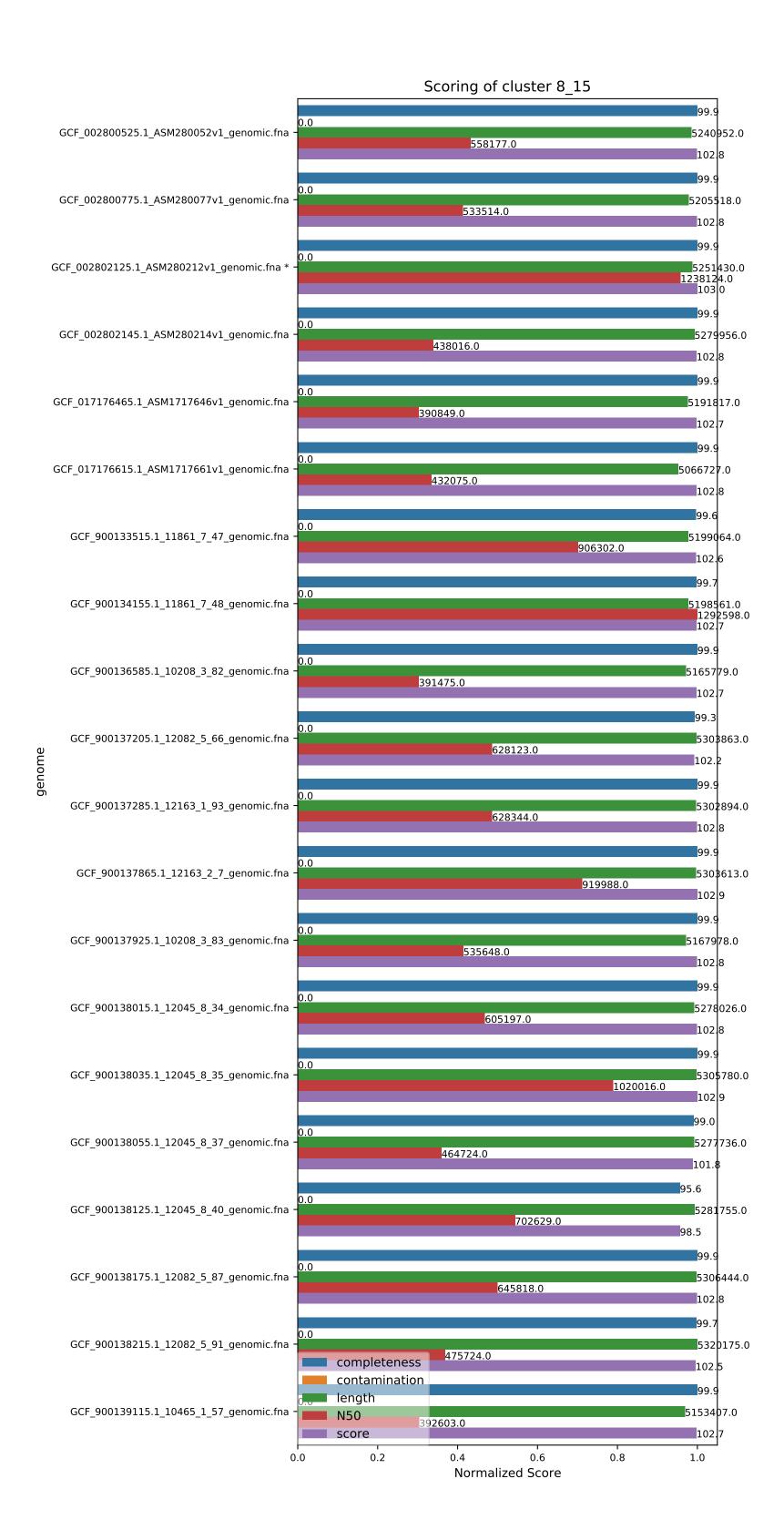


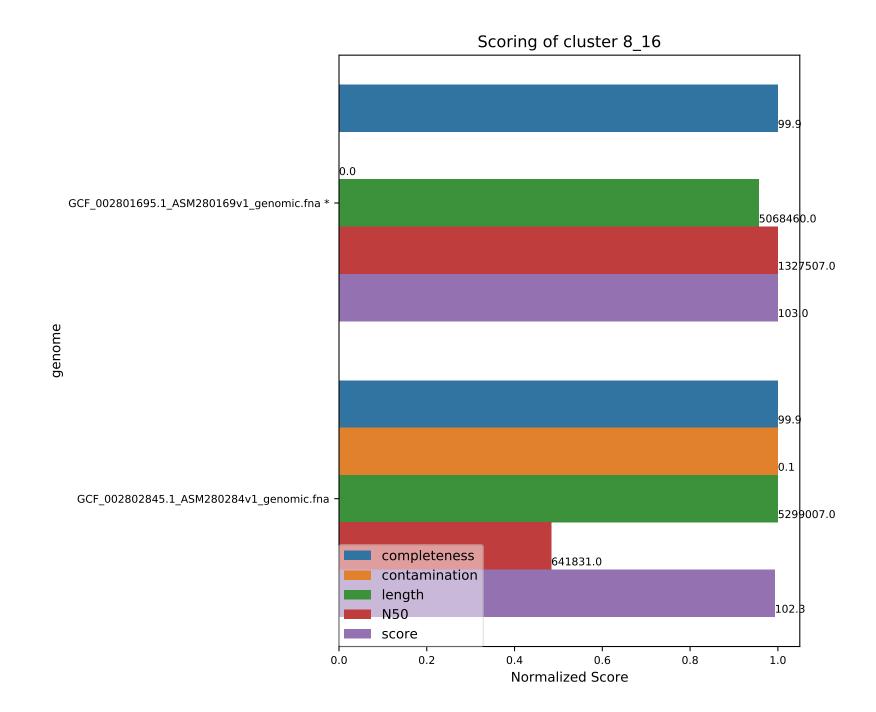












99.9

102.8

5100504.0

103.0 99.9

5100563.0

102.8

99.9

50827‡4.0

102.9 99.6

102.

5116077.0

103 2

1.0

3477186.0

8.0

0.6

Normalized Score

5229053.0

5237635.0

GCF_900140915.1_10702_1_57_genomic.fna

GCF_900141265.1_10702_1_86_genomic.fna

GCF_900141325.1_10702_1_92_genomic.fna -

GCF_900141395.1_11893_6_55_genomic.fna

GCF 900141435.1 11893 6 54 genomic.fna

GCF_900141525.1_12082_5_83_genomic.fna -

487685.0

650676.0

707369.0

completeness

contamination

0.2

0.4

length

score

■ N50

0.0

0.0

1210066.0

Scoring of cluster 8_17

979384.0

726114.0

410567.0

392489.0

435585.0

312980.0

296470.0

422343.0

360332.0

394721.0

1742956.0

501405.0

 ${\sf GCF_002800725.1_ASM280072v1_genomic.fna}$

GCF_002800765.1_ASM280076v1_genomic.fna

GCF_002802135.1_ASM280213v1_genomic.fna

GCF_017174855.1_ASM1717485v1_genomic.fna

 ${\sf GCF_017174975.1_ASM1717497v1_genomic.fna}$

GCF_017174995.1_ASM1717499v1_genomic.fna

 ${\sf GCF_017175215.1_ASM1717521v1_genomic.fna}$

GCF_017175945.1_ASM1717594v1_genomic.fna -

 ${\sf GCF_017176255.1_ASM1717625v1_genomic.fna}$

GCF_017176675.1_ASM1717667v1_genomic.fna

GCF_017183735.1_ASM1718373v1_genomic.fna

GCF_017183835.1_ASM1718383v1_genomic.fna *

 ${\sf GCF_017189355.1_ASM1718935v1_genomic.fna}$

 ${\sf GCF_017190695.1_ASM1719069v1_genomic.fna}$

GCF_900130685.1_11861_7_38_genomic.fna

GCF_900130695.1_11861_7_39_genomic.fna -

GCF_900130725.1_11861_7_66_genomic.fna -

99.9

5028267.0

102.9

5062964.0

102.9

5110690.0

102.7 ■99.9

5066539.0

99.6

5021378.0

102. 99.9

50858\$4.0

5042893.0

102.8 99.9

5097095.0

99.9

5083641.0

102.8

5096267.0

103.1 99.9

5098671.0 5098671.0 101.6

5107403.0 5107403.0 103 3

99.9

99.9

5073623.0 5073623.0 103 3

5074858.0 5074858.0

103 3

5066758.0

102.7

5066499.0

99.8 5075695.0

