Scoring of winning genomes

1848202.0

38570.0

1994919.0

2043417.0

2146411.0

2347830.0

1843816.0

24592.0

26218.0

1313189.0

1311241.0

1692182.0

1481180.0

1499231.0

1556788.0 27478.0

1473098.0

21079.0

19909.0

14396.0

2.9

2.8

10325.0

751055.0 8612.0

1.2

1.1

1.9

9560.0

9599.0

1.4

1.5

0.7

1.1

1.6

7299.0

1.6

1.7

9311.0

9165.0

2.2

5660.0

5087.0

6470.0

8312.0

1.8

8516.0

2.6

6710.0

7010.0

5998.0

4392.0

5364.0

5026.0

3934.0

5393.0

6265.0

7783.0

11196.0

5089.0

7204.0

3877.0

4800.0

5478.0

5325.0

3.2

3.5

3.6

3.5

4.5

4.5

18569.0

4.2

3.9

1304218.0

1129018.0

4.7 1091162.0

4.0

4.9

1219236.0

4.6

1272694.0 31725.0

> 6.0 1402200.0

5.2 1631781.0

5.1

53.6

51.6

7.7 2307451.0

0.6

Normalized Score

1704873.0

44.0

1632468.0

27112.0

24720.0

0.4

28.6

19667.0 26.7

23.1

0.2

7.2 1559283.0

1685327.0

14539.0

2.7

2.5

2.3

9804.0

2.8

3.4

19799.0

2.7

12982.0

18178.0

1491859.0

1337506.0 24093.0

1255568.0

1673572.0

1729210.0

1601184.0

1492058.0

1614882.0

1947794.0

1702685.0

1880205.0

2.3

10334.0

2.5

2.7

7419.0

18192.0

2303408.0

1643524.0

13379.0

14727.0

876888.0

43505.0

2395516.0

9020.bin.66.fa

8020.bin.35.fa

1001.bin.77.fa

8020.bin.44.fa

9020.bin.34.fa

9020.bin.71.fa -

9020.bin.82.fa -

9020.bin.29.fa -

9020.bin.92.fa -

9020.bin.5.fa -

1001.bin.37.fa -

1001.bin.63.fa -

9020.bin.85.fa -

1001.bin.47.fa

9020.bin.58.fa -

6009.bin.29.fa

1001.bin.61.fa -

1001.bin.48.fa

9020.bin.83.fa -

9020.bin.63.fa -

6009.bin.43.fa -

9020.bin.39.fa

6009.bin.39.fa -

1001.bin.38.fa -

6009.bin.79.fa -

9020.bin.10.fa -

9020.bin.21.fa -

1001.bin.83.fa -

6009.bin.84.fa -

1001.bin.23.fa -

6009.bin.70.fa -

9020.bin.91.fa -

1001.bin.42.fa -

6009.bin.36.fa -

9020.bin.48.fa -

⊕ E 0 8020.bin.39.fa − B

1001.bin.67.fa -

1001.bin.75.fa -

9020.bin.20.fa -

6009.bin.24.fa -

9020.bin.52.fa -

6009.bin.76.fa -

1001.bin.70.fa -

8020.bin.50.fa -

9020.bin.51.fa -

9020.bin.47.fa -

1001.bin.79.fa -

1001.bin.66.fa -

9020.bin.61.fa -

1001.bin.62.fa -

9020.bin.33.fa -

1001.bin.16.fa -

1001.bin.21.fa -

8020.bin.25.fa

6009.bin.51.fa -

9020.bin.94.fa -

8020.bin.59.fa -

1001.bin.27.fa -

6009.bin.50.fa -

6009.bin.15.fa -

9020.bin.64.fa

1001.bin.81.fa -

9020.bin.24.fa

9020.bin.31.fa -

6009.bin.3.fa -

9020.bin.30.fa

1001.bin.74.fa -

9020.bin.65.fa

9020.bin.60.fa -

1001.bin.45.fa

9020.bin.41.fa

0.0

0.6

6162.0

1.6

1.2

0.4

0.3

0.2

1.3

7057.0

1.6

4957.0

0.9

0.8

4943.0

12454.0

15288.0

18374.0

15911.0

99.9

101 4

98.6

100.6

100.0

95.8

97.0

94.8

93.6

94.6

69330.0 93.6

93.3

93.3 91.7

91.8

89.4 87.9

88.7

87.7

88.6

88.0 85.7

87.9

87.5

85.6

85.6

86.2

85.7

85.4

84.5

85.2

84.8

84.2

84.0

83.4

81.8

81.0

81.0

80.8

80.4

80.1

78.9

78.7

78.3

75.6

75.6

75.0

74.3

74.0

71.9

2831525.0

71.5

70.3

2779902.0

69.9

69.6

69.3

67.6

67.2

65.0

2527434.0

64.9

63.2

2399946.0

61.8

59.2

58.9

56.4

8.5 2177031.0

2505807.0

78.9

78.6

85.2

84.7

87.9

85.0

84.7

90.9

90.4

85.8

88.5

3532804.0

92.3

91.9

89.4

87.5

81.1

80.1

78.2

85.5

87.6

82.8

91.5

92.7

96.9

84.7

77.7

75.2

80.4

79.5

3065804.0

3075802.0

8.0

completeness contamination length N50

94.0

1.0

4106169.0

76022.0

81.7

2672539.0 50064.0

2487599.0

2642554.0

2211607.0

86.4

90.7

86.8

93.4

87.9

98.0

92.2

2720219.0

2391975.0

46085.0

2389656.0

2074703.0 41698.0

2176031.0

2192366.0

1869483.0

1658139.0

1664806.0

21773.0

19667.0

22256.0

90.3

94.3

2904012.0

99.3

97.0

96.3

420B005.0

4179240.0

66813.0