

79/4/1 Rhodobacter\_sphaeroides\_2.4.1\_assembly  
79/10/1 Escherichia\_coli\_K-12\_MG1655\_assembly  
79/13/1 Shewanella\_amazonensis\_SB2B\_assembly  
34837/65/4 Campylobacter\_jejuni.assembly  
34837/66/3 Caulobacter\_vibrioides.assembly  
34837/67/2 Coxiella\_burnetii.assembly  
34837/69/1 Escherichia\_coli\_K-12.assembly  
34837/70/1 Escherichia\_coli\_Sakai\_assembly  
34837/100/3 Staphylococcus\_aureus\_assembly  
34837/101/2 Salmonella\_enterica\_assembly  
34837/102/1 Shigella\_flexneri\_assembly  
43623/9/1 Klebsiella\_pneumoniae\_assembly  
43623/24/1 Mycobacterium\_tuberculosis\_pneumoniae  
43623/39/1 Acinetobacter\_pitii\_assembly  
34837/60/1 Escherichia\_coli\_K-12\_MG1655

- 79/10/1 Escherichia\_coli\_K-12\_MG1655\_assembly

34837/61/1 Rhodobacter\_sphaeroides\_2.4.1

- 79/4/1 Rhodobacter\_sphaeroides\_2.4.1

34837/62/1 Shewanella\_amazonensis\_SB2B

- 79/13/1 Shewanella\_amazonensis\_SB2B\_assembly

34837/107/3 Klebsiella\_pneumoniae

- 43623/9/3 Klebsiella\_pneumoniae\_assembly

34837/108/2 Mycobacterium\_tuberculosis

- 43623/24/2 Mycobacterium\_tuberculosis\_pneumoniae

34837/109/2 Acinetobacter\_pitii

- 43623/39/2 Acinetobacter\_pitii\_assembly

**34837/71/1 Some\_refseq\_assemblies**

- 34837/65/4 Campylobacter\_jejuni.assembly
- 34837/70/1 Escherichia\_coli\_Sakai\_assembly
- 34837/69/1 Escherichia\_coli\_K-12.assembly

**34837/105/1 S\_assemblies**

- 34837/70/1 Escherichia\_coli\_Sakai\_assembly
- 34837/101/2 Salmonella\_enterica\_assembly
- 34837/102/1 Shigella\_flexneri\_assembly
- 34837/100/3 Staphylococcus\_aureus\_assembly

**34837/73/1 Some\_genomes**

- 34837/60/1 Escherichia\_coli\_K-12\_MG1655
  - 79/10/1 Escherichia\_coli\_K-12\_MG1655\_assembly
- 34837/61/1 Rhodobacter\_sphaeroides\_2.4.1
  - 79/4/1 Rhodobacter\_sphaeroides\_2.4.1

**34837/75/1 Escherichia\_genome\_set**

- 34837/60/1 Escherichia\_coli\_K-12\_MG1655
  - 79/10/1 Escherichia\_coli\_K-12\_MG1655\_assembly

**34837/110/1 AMK\_genomes**

- 34837/107/3 Klebsiella\_pneumoniae
  - 43623/9/3 Klebsiella\_pneumoniae\_assembly
- 34837/108/2 Mycobacterium\_tuberculosis
  - 43623/24/2 Mycobacterium\_tuberculosis\_pneumoniae
- 34837/109/2 Acinetobacter\_pitii
  - 43623/39/2 Acinetobacter\_pitii\_assembly

34837/2/1 SURF\_B\_MetaBAT2\_CheckM

- Bin.012.fasta Bin.018.fasta Bin.027.fasta Bin.034.fasta Bin.038.fasta  
Bin.039.fasta Bin.040.fasta Bin.044.fasta

34837/16/1 SURF\_B\_MaxBin2\_CheckM

- Bin.008.fasta, Bin.009.fasta, Bin.012.fasta

34837/46/1 small\_arctic\_metabat

- bin.001.fasta, bin.002.fasta, bin.003.fasta

34837/77/2 capybyaraGut\_MaxBin2\_CheckM

- Bin.002.fasta Bin.003.fasta Bin.011.fasta Bin.012.fasta Bin.013.fasta  
Bin.015.fasta Bin.019.fasta Bin.020.fasta Bin.021.fasta Bin.023.fasta  
Bin.028.fasta Bin.030.fasta Bin.031.fasta Bin.033.fasta