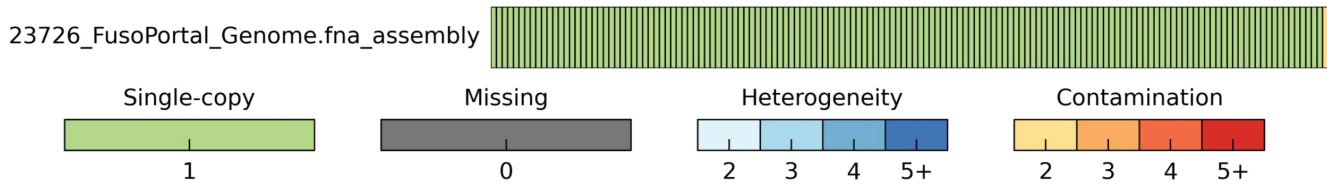
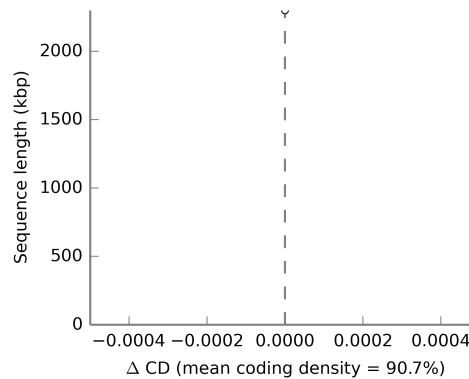
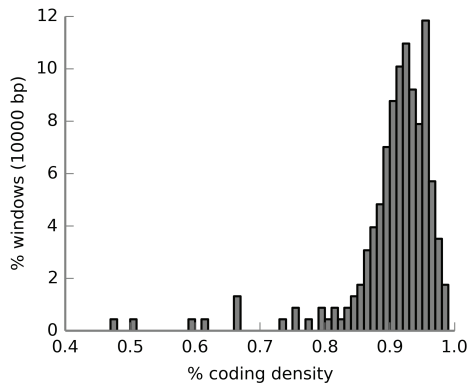
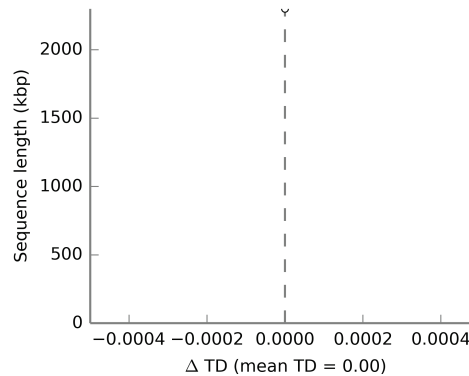
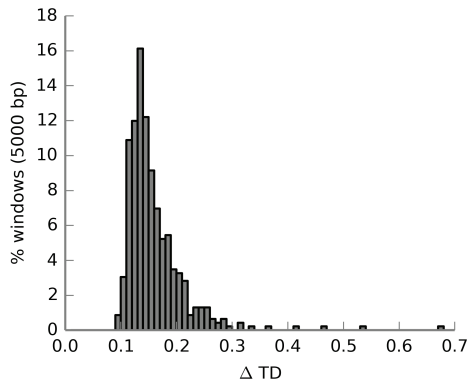
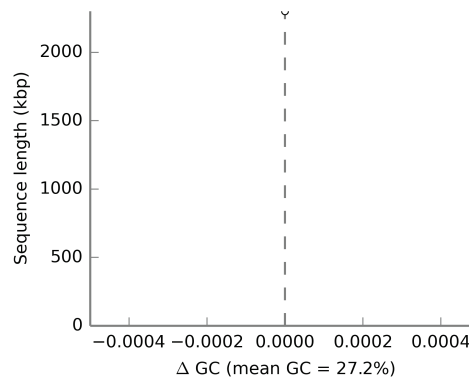
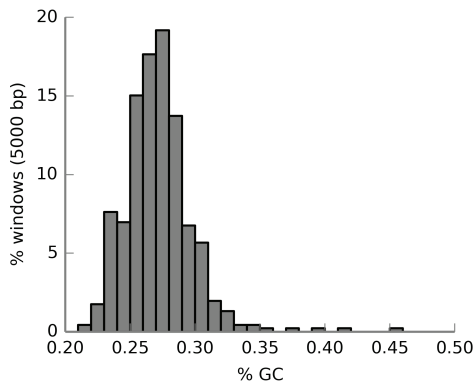


CheckM genome analysis run on KBase



| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|--|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 23726_FusoPortal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 148 | 1 | 0 | 0 | 0 | 100.0 | 0.161 |

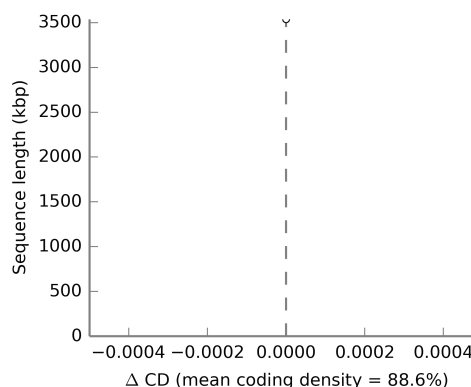
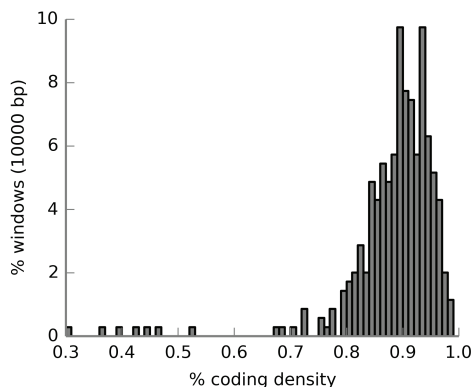
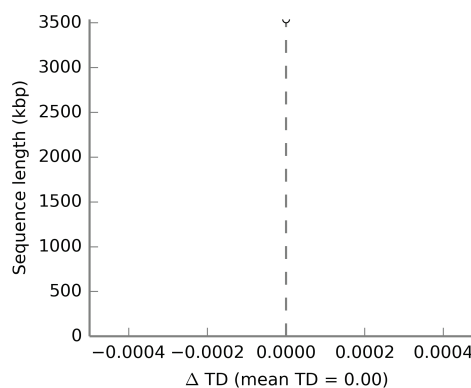
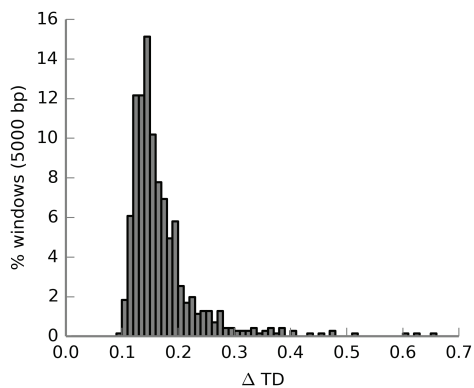
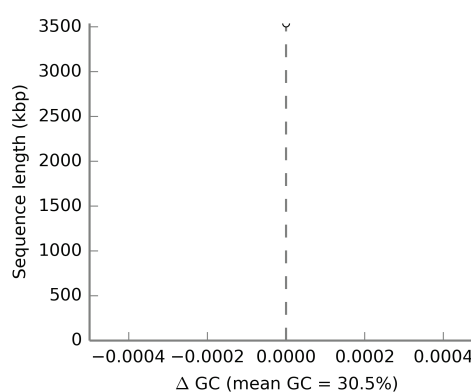
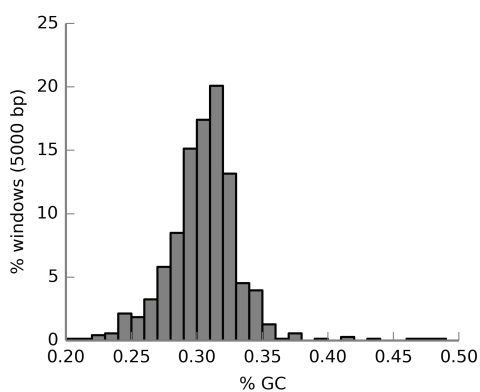


CheckM genome analysis run on KBase

49185_FusoPortal_Genome.fna_assembly

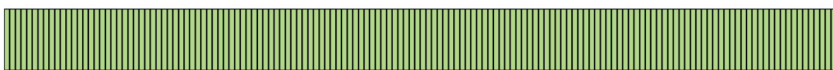


| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|--|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 49185_FusoPortal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 147 | 2 | 0 | 0 | 0 | 100.0 | 2.247 |

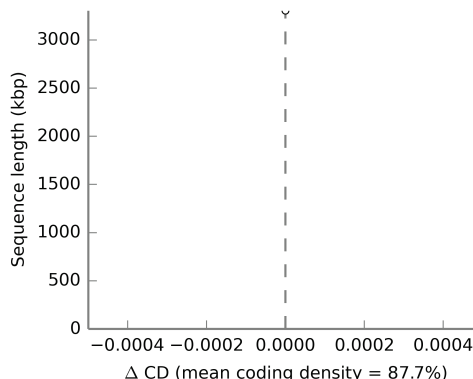
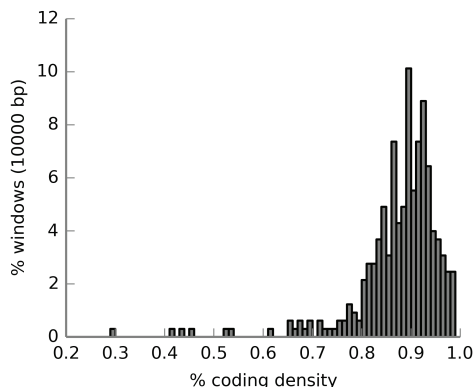
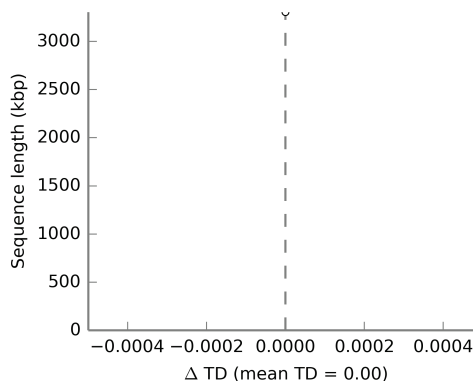
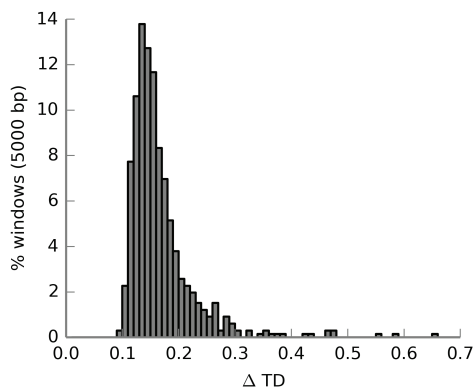
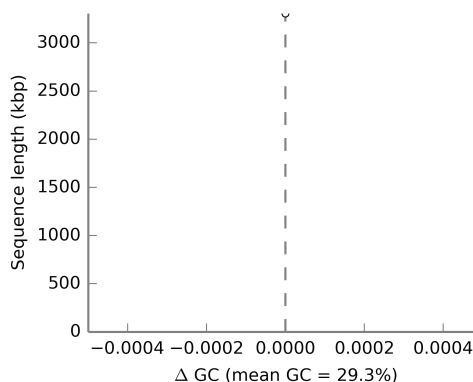
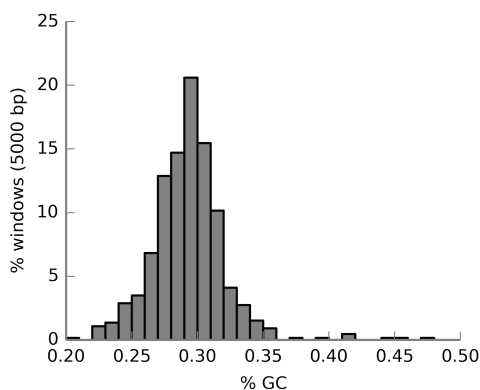


CheckM genome analysis run on KBase

27725_FusoPortal_Genome.fna_assembly

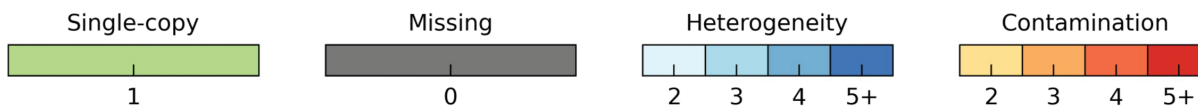


| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|--|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 27725_FusoPortal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 148 | 1 | 0 | 0 | 0 | 100.0 | 1.124 |

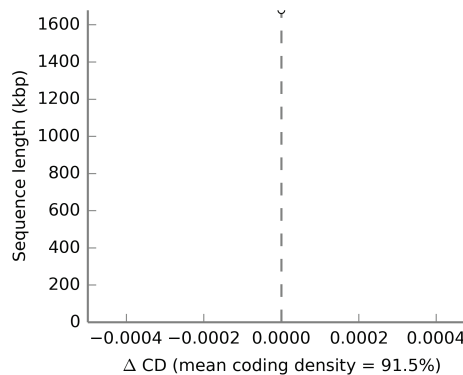
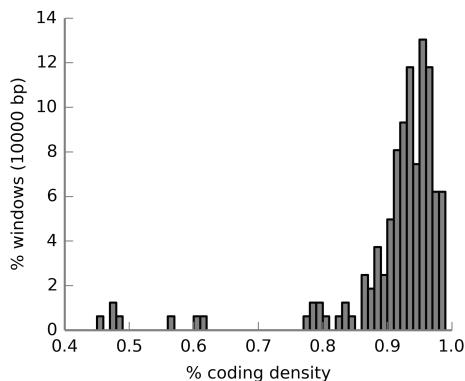
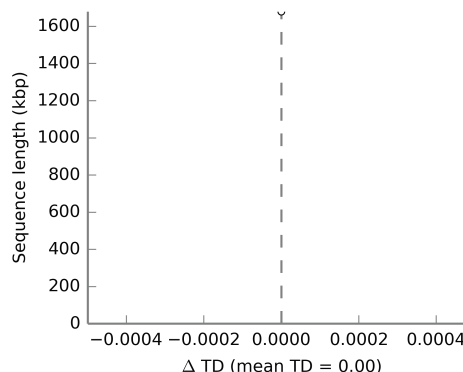
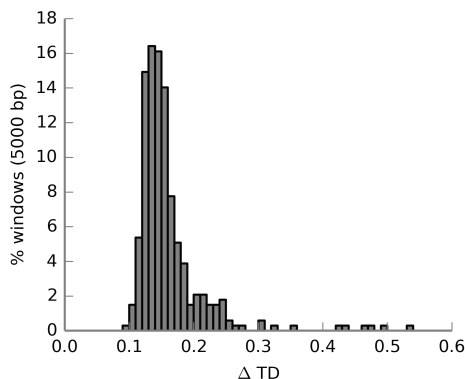
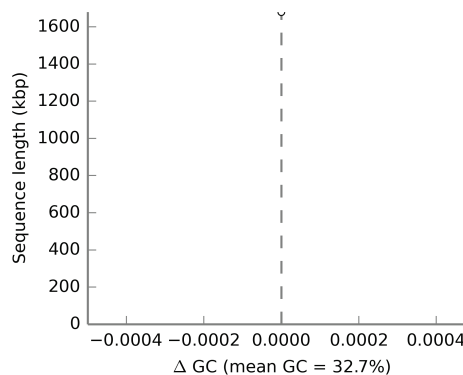
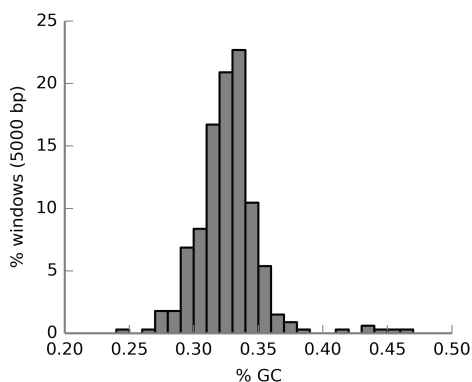


CheckM genome analysis run on KBase

25563_Fusoportal_Genome.fna_assembly



| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|--|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 25563_Fusoportal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 149 | 0 | 0 | 0 | 0 | 100.0 | 0.0 |

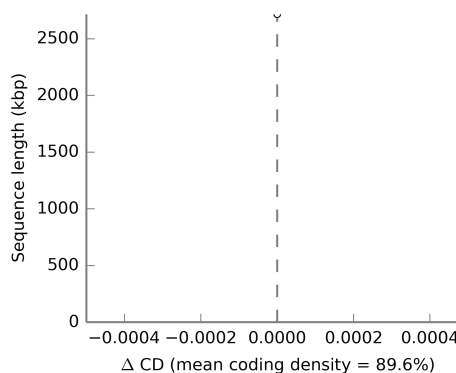
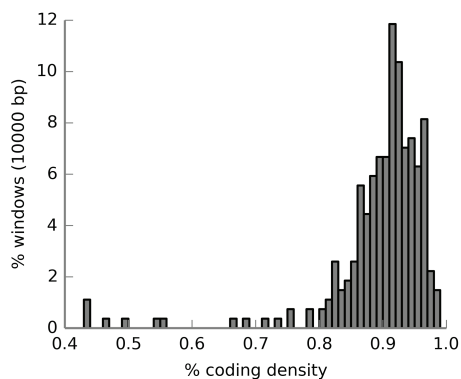
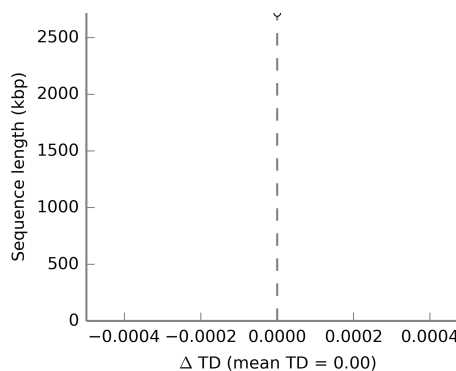
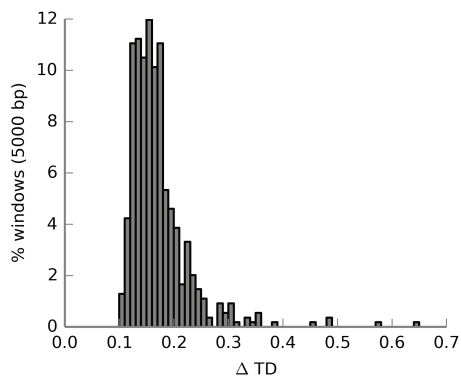
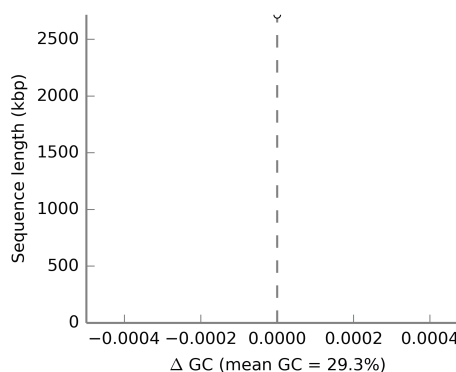
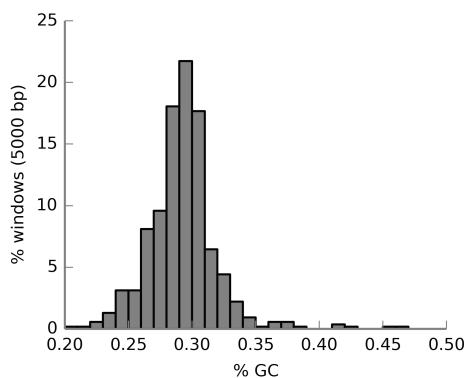


CheckM genome analysis run on KBase

9817_FusoPortal_Genome.fna_assembly

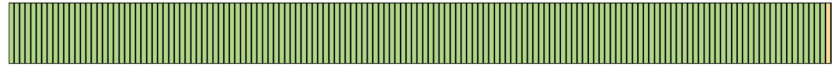


| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|-------------------------------------|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 9817_FusoPortal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 1 | 147 | 1 | 0 | 0 | 0 | 98.876 | 1.124 |

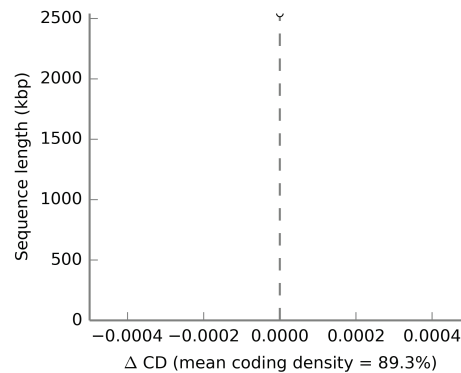
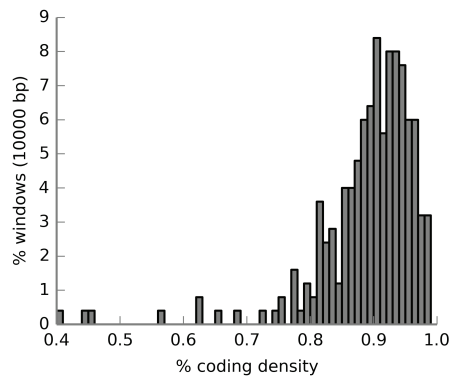
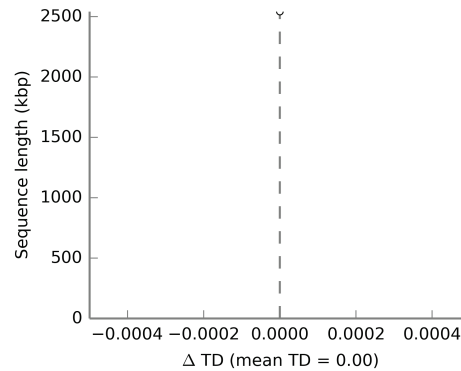
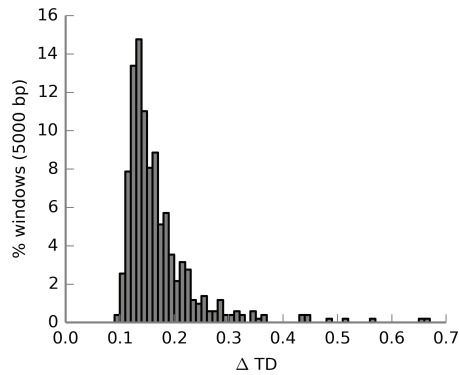
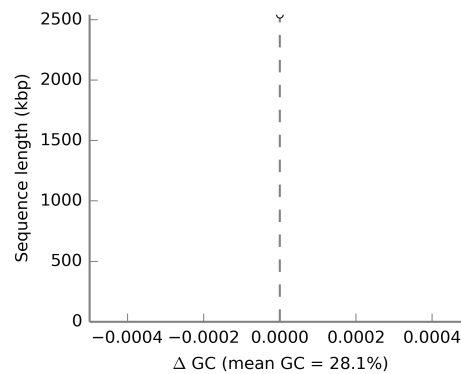
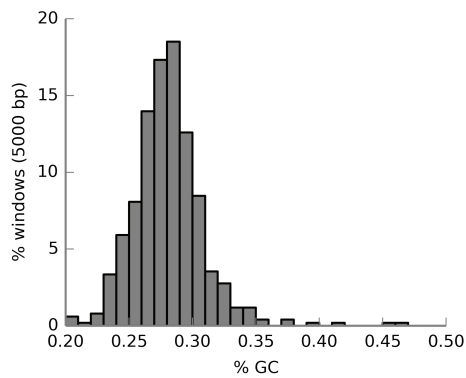


CheckM genome analysis run on KBase

2_1_31_Fusoportal_Genome.fna_assembly



| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|---|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 2_1_31_Fusoportal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 148 | 1 | 0 | 0 | 0 | 100.0 | 0.562 |

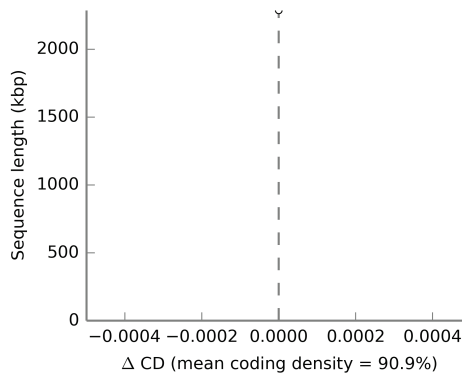
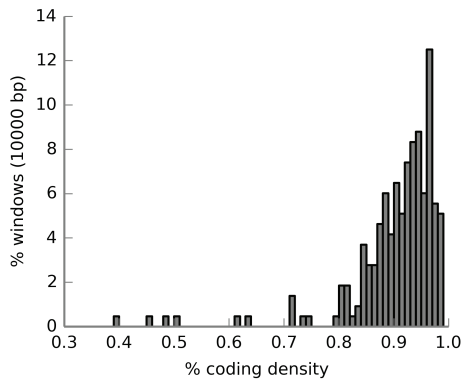
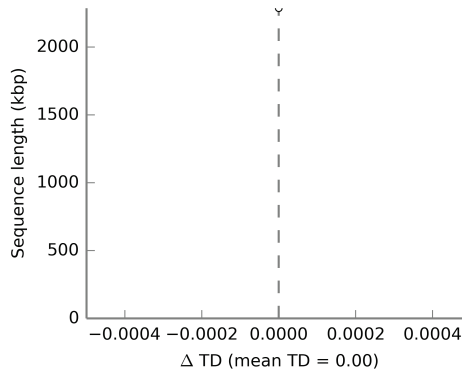
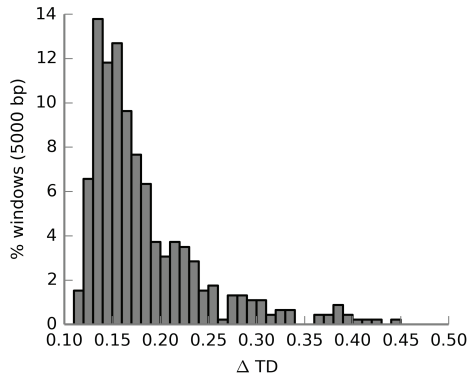
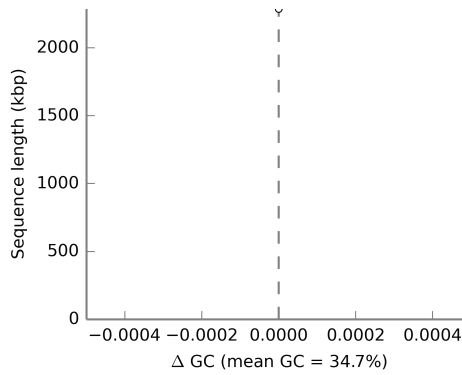
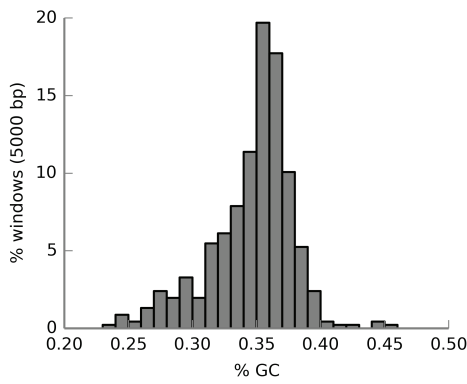


CheckM genome analysis run on KBase

1_1_36S_FusoPortal_Genome.fna_assembly

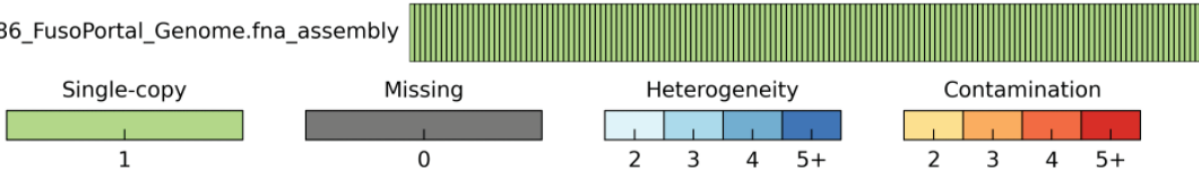


| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|--|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 1_1_36S_FusoPortal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 149 | 0 | 0 | 0 | 0 | 100.0 | 0.0 |



CheckM genome analysis run on KBase

25586_FusoPortal_Genome.fna_assembly



| Bin Name | Marker Lineage | # Genomes | # Markers | # Marker Sets | 0 | 1 | 2 | 3 | 4 | 5+ | Completeness | Contamination |
|--|----------------|-----------|-----------|---------------|---|-----|---|---|---|----|--------------|---------------|
| 25586_FusoPortal_Genome.fna_assembly | k_Bacteria | 174 | 149 | 89 | 0 | 149 | 0 | 0 | 0 | 0 | 100.0 | 0.0 |

