| Reference Organism | Classification | RefSeq ID | Format | Year |
|---|----------------|-----------------|-----------------|------|
| Homo sapiens CHM1_1.1 | Human | GCF_000306695.2 | Chromosome | 2013 |
| Escherichia coli O157:H7 str. Sakai | Bacteria | GCF_000008865.2 | Complete Genome | 2018 |
| Mycoplasmopsis fermentans M64 | Bacteria | GCF_000186005.1 | Complete Genome | 2011 |
| Propionibacterium acidifaciens | Bacteria | GCF_003798325.1 | Complete Genome | 2018 |
| Staphylococcus aureus subsp. aureus NCTC 8325 | Bacteria | GCF_000013425.1 | Complete Genome | 2006 |
| Streptococcus pneumoniae | Bacteria | GCF_002076835.1 | Chromosome | 2017 |
| Malassezia furfur | Fungi | GCA_009938135.1 | Complete Genome | 2020 |
| Wickerhamomyces canadensis | Fungi | GCA_003707015.2 | Scaffold | 2018 |