



- z_GCF_001433875.1 *Lactobacillus delbrueckii* subsp. *delbrueckii* DSM 20074 = JCM 1012 strain=DSM 20074, ASM143387v1
z_GCF_013394085.1 *Lactobacillus fermentum* strain=DSM 20052, ASM1339408v1
z_GCA_019120335.1 *Candidatus Enterococcus stercoripullorum*, ASM1912033v1
z_GCF_001436315.1 *Lactobacillus aviarius* subsp. *aviarius* DSM 20655 strain=DSM 20655, ASM143631v1
z_GCF_000407565.1 *Enterococcus cecorum* DSM 20682 = ATCC 43198 strain=ATCC 43198, Ente_ceco_ATCC43198_V1
z_GCF_001654615.1 *Lactobacillus aviarius* strain=UMNLAv11, ASM165461v1
z_GCA_019120355.1 *Candidatus Ligilactobacillus faecavium*, ASM1912035v1
z_GCA_004556255.1 *Lactobacillus delbrueckii*, ASM455625v1
z_GCF_002706375.1 *Lactobacillus amylovorus* DSM 20531 strain=DSM 20531, ASM270637v1
z_GCF_000959525.1 *Burkholderia multivorans* ATCC BAA-247 strain=BAA-247, ASM95952v1
z_GCF_013409425.1 *Nocardioides perillae* strain=DSM 24552, ASM1340942v1
z_GCF_000317125.1 *Chroococcidiopsis thermalis* PCC 7203 strain=PCC 7203, ASM31712v1
z_GCF_000817735.3 *Scytonema millei* VB511283 strain=VB511283, ASM81773v3
z_GCF_015291885.1 *Pseudomonas stutzeri* strain=ATCC 14405, ASM1529188v1
z_GCA_003963435.1 *Bradyrhizobiaceae* bacterium, ASM396343v1
z_GCF_000219605.1 *Pseudomonas stutzeri* strain=CGMCC 1.1803, ASM21960v1
z_GCF_018137945.1 *Brevibacterium* sp. W7.2 strain=W7.2, ASM1813794v1
z_GCF_001298265.1 *Bosea vaviloviae* strain=SD260, ASM129826v1
z_GCF_000308295.2 *Afiopia birgiae* 34632 strain=34632, Ab_34632
z_GCF_900169275.1 *Brevibacterium casei* CIP 102111 strain=CIP 102111, Assembly_BC102111
z_GCF_001419795.1 *Streptomyces niveiscabiei* strain=NRRL B-24457, ASM141979v1
z_GCF_002901845.1 *Staphylococcus hominis* subsp. *hominis* strain=NCTC 11320, ASM290184v1
z_GCF_003730015.1 *Chryseobacterium hominis* strain=DSM 22165, ASM373001v1
z_GCF_000368045.1 *Acinetobacter johnsonii* CIP 64.6 strain=CIP 64.6, Acin_john_CIP_64_6_V1
z_GCF_003860525.1 *Chryseobacterium* sp. F5649 strain=F5649, ASM386052v1
z_GCF_006716815.1 *Microbacterium lacticum* strain=DSM 20427, ASM671681v1
z_GCF_000225825.1 *Brachybacterium squillarum* M-6-3 strain=M-6-3, ASM22582v2
z_GCF_016907455.1 *Brachybacterium muris* strain=DSM 15460, ASM1690745v1
z_GCF_010211765.1 *Pseudoxanthomonas kaohsiungensis* strain=DSM 17583, ASM1021176v1
z_GCF_015707995.1 *Microbacterium* sp. 2C strain=2C, ASM1570799v1
z_GCA_016791045.1 *Methyloversatilis* sp., ASM1679104v1
z_GCA_005240065.1 *Gammaproteobacteria* bacterium, ASM524006v1
z_GCF_001570945.1 *Janibacter anophelis* NBRC 107843 strain=NBRC 107843, ASM157094v1
z_GCF_001517405.1 *Erwinia gerundensis*, EM595
z_GCF_005503355.1 *Sphingomonas* sp. 1F27F7B strain=1F27F7B, ASM550335v1
z_GCF_004367745.1 *Herbaspirillum huttiense* strain=NFYY 53159, ASM436774v1
z_GCF_003851545.1 *Ralstonia* sp. SET104 strain=SET104, ASM385154v1
z_GCF_000023205.1 *Micrococcus luteus* NCTC 2665 strain=NCTC 2665 (Fleming 2665), ASM2320v1
z_GCF_000368765.1 *Acinetobacter junii* CIP 64.5 strain=CIP 64.5, Acin_juni_CIP_64_5_V1
z_GCF_003030305.1 *Cutibacterium acnes* strain=ATCC 6919, ASM303030v1
z_GCA_900539805.1 uncultured *Phyllobacterium* sp., UMGS170
z_GCF_000785515.1 *Streptococcus salivarius* strain=NCTC 8618, ASM78551v1
z_GCF_000187465.1 *Streptococcus infantis* ATCC 700779 strain=ATCC 700779, ASM18746v1
z_GCF_000223335.1 *Streptococcus infantis* X strain=X, ASM22333v2
z_GCF_002287065.1 *Paracoccus salipaludis* strain=WN007, ASM228706v1
z_GCA_002863225.1 *Mycobacterium* sp., ASM286322v1
z_GCF_900110815.1 *Aliicoccus persicus* strain=IBRC-M10081, IMG-taxon 2654588195 annotated assembly
z_GCF_004136555.1 *Kocuria carniphila* strain=CCM 132, ASM413655v1
z_GCF_007641235.1 *Corynebacterium sanguinis* strain=CCUG 58655, ASM764123v1
z_GCF_016907795.1 *Kocuria palustris* strain=TAGA27, ASM1690779v1

Taxon