

- inactive
- Actinomyces chiangmaiensis
- Actinoplanes atractionticus
- Agreia bicolorata
- Agrococcus casei
- Allokutzneria albata
- Anaerocolumna xylanovorans
- Arenimonas oryzae
- Arthrobacter globiformis
- Auraticoccus monumenti
- Bacillus sp. 7586-K
- Bacteriovorax sp. BAL6 X
- Bacteroides luti
- Brachybacterium alimentarium
- Brachybacterium paraconglomeratum
- Brevibacterium antiquum
- Brevibacterium mcbrellneri
- Cellulomonas cellasea
- Cephalotococcus capnophilus
- Chlamydiales bacterium SCGC AG-110-P3
- Chloroflexi bacterium RBG 13 51 36
- Chromatococcus halotolerans
- Citricoccus muralis
- Clavibacter michiganensis
- Clostridium combesii
- Dermabacter hominis
- Dermatophilus congolensis
- Desulfobacter curvatus
- Desulfobulbus elongatus
- Enhygromyxa salina
- Frankiales bacterium YIM 75000
- Geodermatophilus obscurus
- Granulosicoccus antarcticus
- Halobacteriovorax marinus
- Humibacillus sp. DSM 29435
- Intrasporangiaceae bacterium URHB0013
- Isoptericola variabilis
- Kineospira limosa
- Kocuria flava
- Kocuria varians
- Kytococcus sedentarius
- Legionella brunensis
- Leifsonia sp. Leaf325
- Luteimonas tolerans
- Micrococcus phosphovorus
- Micromonospora avicenniae
- Micromonospora peucetia
- Mobilicoccus massiliensis
- Mycobacterium obuense
- Nakamurella multipartita
- Nesterenkonia alba
- Nonomuraea candida
- Okibacterium fritillariae
- Paenibacillus polysaccharolyticus
- Pelobacter propionicus DSM 2379
- Plantibacter sp. VKM Ac-1784
- Pseudoxanthomonas
- Psychromonas aquimarina
- Rothia mucilaginosa
- Saccharothrix espanaensis
- Sandaracinus amylolyticus
- Sanguibacter keddiei
- Sedimentihabitans luteus
- Shewanella putrefaciens
- Silanimonas lenta
- Streptococcus pneumoniae
- Streptomyces ambofaciens
- Streptomyces avermitilis
- Subtercola sp. Z020
- Thermomonas haemolytica
- Virgibacillus salinus
- Wenzhouxiangella sediminis
- Xanthomonas translucens
- XC

