

- inactive
- Actinomycetospora chiangmaiensis
- Actinoplanes atraurantiacus
- Agreia bicolorata
- Agrococcus casei
- Allokutzneria albata
- Anaerocolumna xylanovorans
- Arenimonas oryzae
- Arthrobacter globiformis
- Bacillus sp. 7586-K
- Bacteriovorax sp. BAL6 X
- Bacteroides luti
- Brachybacterium alimentarium
- Brachybacterium paraconglomeratum
- Brevibacterium antiquum
- Brevibacterium mcbrellneri
- Cellulomonas cellasea
- Cephalotococcus capnophilus
- Chloroflexi bacterium RBG 13 51 36
- Chromatococcus halotolerans
- Citricoccus muralis
- Clavibacter michiganensis
- Clostridium combesii
- Desulfobacter curvatus
- Desulfobulbus elongatus
- Enhygromyxa salina
- Frankiales bacterium YIM 75000
- Geodermatophilus obscurus
- Granulosicoccus antarcticus
- Halobacteriovorax marinus
- Intrasporangiaceae bacterium URHB0013
- Isoptericola variabilis
- Kineospaera limosa
- Kocuria flava
- Kocuria varians
- Kytococcus sedentarius
- Leifsonia sp. Leaf325
- Luteimonas tolerans
- Microlunatus phosphovorus
- Micromonospora avicenniae
- Micromonospora peucetia
- Mobilicoccus massiliensis
- Mycolicibacterium obuense
- Nakamurella multipartita
- Nesterenkonia alba
- Okibacterium fritillariae
- Paenibacillus polysaccharolyticus
- Pelobacter propionicus DSM 2379
- Pet28\_1
- Plantibacter sp. VKM Ac-1784
- Pseudoxanthomonas
- Psychromonas aquimarina
- Rothia mucilaginosa
- Saccharothrix espanaensis
- Sandaracinus amylolyticus
- Sanguibacter keddii
- Sediminihabitans luteus
- Shewanella putrefaciens
- Silanimonas lenta
- Streptomyces ambofaciens
- Streptomyces avermitilis
- Subtercola sp. Z020
- Thermomonas haemolytica
- Xanthomonas translucens
- XC

