

- inactive
- Actinomycetospora chiangmaiensis
- Actinoplanes atraurantiacus
- Agreia bicolorata
- Agrococcus casei
- Allokutzneria albata
- Arenimonas oryzae
- Arthrobacter globiformis
- Auraticoccus monumenti
- Bacillus sp. 7586-K
- Bacteriovorax sp. BAL6 X
- Brachybacterium alimentarium
- Brevibacterium mcbrellneri
- Cellulomonas cellasea
- Chlamydiales bacterium SCGC AG-110-P3
- Chromatococcus halotolerans
- Citricoccus muralis
- Clavibacter michiganensis
- Clostridium combesii
- Dermatophilus congolensis
- Enhygromyxa salina
- Frankiales bacterium YIM 75000
- Geodermatophilus obscurus
- Granulosicoccus antarcticus
- Humibacillus sp. DSM 29435
- Intrasporangiaceae bacterium URHB0013
- Isoptericola variabilis
- Kineosphaera limosa
- Kocuria flava
- Kocuria varians
- Kytococcus sedentarius
- Leifsonia sp. Leaf325
- Luteimonas tolerans
- Microlunatus phosphovorius
- Micromonospora avicenniae
- Micromonospora peucetia
- Mobilicoccus massiliensis
- Mycolicibacterium obuense
- Nakamurella multipartita
- Nesterenkonia alba
- Nonomuraea candida
- Okibacterium fritillariae
- Paenibacillus polysaccharolyticus
- Pseudoxanthomonas
- Psychromonas aquimarina
- Rothia mucilaginosa
- Saccharothrix espanaensis
- Sandaracinus amylolyticus
- Sediminihabitans luteus
- Silanimonas lenta
- Streptomyces ambofaciens
- Streptomyces avermitilis
- Subtercola sp. Z020
- Thermomonas haemolytica
- Virgibacillus salinus
- Wenzhouxiangella sediminis
- Xanthomonas translucens

