

- inactive
- Actinomyces chiangmaiensis
- Actinoplanes atraurantiacus
- Agreia bicolorata
- Allokutzneria albata
- Arenimonas oryzae
- Arthrobacter globiformis
- Auraticoccus monumenti
- Bacteriovorax sp. BAL6 X
- Brachybacterium alimentarium
- Brachybacterium paraconglomeratum
- Brevibacterium antiquum
- Brevibacterium mcbrellneri
- Cellulomonas cellasea
- Chlamydiales bacterium SCGC AG-110-P3
- Chloroflexi bacterium RBG 13 51 36
- Chromatococcus halotolerans
- Citricoccus muralis
- Dermabacter hominis
- Dermatophilus congolensis
- Enhygromyxa salina
- Geodermatophilus obscurus
- Granulosicoccus antarcticus
- Halobacteriovorax marinus
- Humibacillus sp. DSM 29435
- Intrasporangiaceae bacterium URHB0013
- Isoptericola variabilis
- Kocuria flava
- Kocuria varians
- Kytococcus sedentarius
- Legionella brunensis
- Leifsonia sp. Leaf325
- Luteimonas tolerans
- Microlunatus phosphovorus
- Micromonospora avicenniae
- Micromonospora peucetia
- Mobilicoccus massiliensis
- Mycolicibacterium obuense
- Nakamurella multipartita
- Nesterenkonia alba
- Okibacterium fritillariae
- Paenibacillus polysaccharolyticus
- Plantibacter sp. VKM Ac-1784
- Pseudoxanthomonas
- Psychromonas aquimarina
- Rothia mucilaginosa
- Sandaracinus amylolyticus
- Sanguibacter keddiei
- Sediminhabitans luteus
- Silanimonas lenta
- Subtercola sp. Z020
- Thermomonas haemolytica
- Virgibacillus salinus
- Wenzhouxiangella sediminis
- Xanthomonas translucens
- XC

