

variable

•	Actinomycetospora chiangmaiensis	•	Chloroflexi bacterium RBG 13 51 36	•	Kocuria varians	•	Pseudoxanthomonas
•	Actinoplanes atraurantiacus	•	Chromatocurvus halotolerans	•	Kytococcus sedentarius	•	Psychromonas aquimarina
•	Agreia bicolorata	•	Citricoccus muralis	•	Legionella brunensis	•	Rothia mucilaginosa
•	Agrococcus casei	•	Clavibacter michiganensis	•	Leifsonia sp. Leaf325	•	Saccharothrix espanaensis
•	Allokutzneria albata	•	Clostridium combesii	•	Luteimonas tolerans	•	Sandaracinus amylolyticus
•	Anaerocolumna xylanovorans	•	Dermabacter hominis	•	Microlunatus phosphovorus	•	Sanguibacter keddieii
•	Arenimonas oryziterrae	•	Dermatophilus congolensis	•	Micromonospora avicenniae	•	Sediminihabitans luteus
•	Arthrobacter globiformis	•	Desulfobacter curvatus	•	Micromonospora peucetia	•	Shewanella putrefaciens
•	Auraticoccus monumenti	•	Desulfobulbus elongatus	•	Mobilicoccus massiliensis	•	Silanimonas lenta
•	Bacillus sp. 7586-K	•	Enhygromyxa salina	•	Mycolicibacterium obuense	•	Streptococcus pneumoniae
•	Bacteriovorax sp. BAL6 X	•	Frankiales bacterium YIM 75000	•	Nakamurella multipartita	•	Streptomyces ambofaciens
•	Bacteroides luti	•	Geodermatophilus obscurus	•	Nesterenkonia alba	•	Streptomyces avermitilis
•	Brachybacterium alimentarium	•	Granulosicoccus antarcticus	•	Nonomuraea candida	•	Subtercola sp. Z020
•	Brachybacterium paraconglomeratum	•	Halobacteriovorax marinus	•	Okibacterium fritillariae	•	Thermomonas haemolytica
•	Brevibacterium antiquum	•	Humibacillus sp. DSM 29435	•	Paenibacillus polysaccharolyticus	•	Virgibacillus salinus
•	Brevibacterium mcbrellneri	•	Intrasporangiaceae bacterium URHB0013	•	Pelobacter propionicus DSM 2379	•	Wenzhouxiangella sediminis
•	Cellulomonas cellasea	•	Isoptericola variabilis	•	Pet28_1	•	Xanthomonas translucens
•	Cephaloticoccus capnophilus	•	Kineosphaera limosa	•	Pet28_2	•	XC
•	Chlamydiales bacterium SCGC AG-110-P3	•	Kocuria flava	•	Plantibacter sp. VKM Ac-1784		