

BGSC	Original	GenBank
1A1	Bacillus subtilis subsp. subtilis 168	AL009126
1A4	Bacillus subtilis subsp. subtilis QB928 KIT-2	CP003783
1A96	Bacillus subtilis subsp. subtilis JH642	CM000489
28A4	Bacillus vallismortis DV1-F-3 (=NRRL B-14890)	AFSH00000000
28A5	Bacillus Mojavensis RO-H-1	AFSI00000000
29A3	Bacillus sp. NRRL B-14911	AAOX01000000
2A11 ^T	Bacillus subtilis subsp. spizizenii TU-B-10T (= NRRL B-23049T)	CP002905
2A12	Bacillus subtilis subsp. spizizenii DV1-B-1 (= NRRL B-23054)	AFSG00000000
2A13	Bacillus subtilis subsp. spizizenii NRRL B-765 (= ATCC 6633 = NRS-231)	ADGS00000000
2A9	Bacillus subtilis subsp. spizizenii NRRL B-14472 (= ATCC 23059 = W23)	CP002183
3A1 ^T	Bacillus subtilis subsp. subtilis Marburg NCIB3610T	CM000488
3A26	Bacillus subtilis AUSI98	AFSF00000000
3A27	Bacillus subtilis RO-NN-1 (= NRRL B-14823)	CP002906
3A35	Bacillus subtilis BSn5	CP002468
4A11 ^T	Bacillus thuringiensis subsp. thuringiensis HD735T	CM000753
4B4	Bacillus thuringiensis subsp. finitimus YBT-020	CP002508 , CP002509 , CP002510
4D4	Bacillus thuringiensis subsp. kurstaki HD73	CP004069 - CP004076
4E4	Bacillus thuringiensis subsp. sotto T04001 (= HD6)	CM000749
4Q12	Bacillus thuringiensis subsp. israelensis HD522 (= OVR60) (= ATCC 35646)	AAJM00000000
4Y1	Bacillus thuringiensis subsp. tochiensis HD868 (117-72)	CM000746
4AJ1	Bacillus thuringiensis subsp. monterrey GM-33 (= T28001)	CM000752
4AW1	Bacillus thuringiensis subsp. andalucensis T37001 (= EA10192)	CM000754
4BA1	Bacillus thuringiensis subsp. pondicheriensis T20A001 (= B23)	CM000755
4BD1	Bacillus thuringiensis subsp. huazhongensis T40001 (= YBT 978)	CM000756
4CC1	Bacillus thuringiensis subsp. pulsiensis T65001 (= NARC Bt17)	CM000757
4CG1	Bacillus thuringiensis subsp. chinensis CT-43	CP001907 - CP001917
4XX6	Bacillus thuringiensis BMB171	CP001903 , CP001904
5A36 ^T	Bacillus licheniformis ATCC 14580T	AE017333 , CP000002
6A15	Bacillus cereus ATCC 10987 (=NRS 248)	AE017194 , AE017195
6A35	Bacillus cereus AH621	CM000719
6A54	Bacillus cereus m1293	CM000714
6A56	Bacillus cereus m1550 (=277/2)	CM000722
6A5 ^T	Bacillus cereus ATCC 14579T	AE016877 , AE016878
6A65	Bacillus cereus R_3098/03	CM000720
7A16	Bacillus megaterium QM B1551	CP001983 - CP001990
8A3 ^T	Bacillus pumilus NRS-272T (= ATCC 7061T)	ABRX01000000
9A21	Geobacillus thermoleovorans 10	U Oklahoma
9A24	Geobacillus sp. C56-T2	PRJNA41699
10A6 ^T	Bacillus amyloliquefaciens subsp. plantarum FZB42T	CP000560
11A3	Bacillus atrophaeus 1942	CP002207
11A4	Bacillus atrophaeus Camp Detrick, clone 1	AEFP00000000
11A5	Bacillus atrophaeus Camp Detrick, clone 2	AEFQ00000000
35A1	Paenibacillus sp. JDR-2	CP001656
36A2	Paenibacillus lautus Y412.MC10	CP001793
94A5	Geobacillus thermodenitrificans G11MC16	ABVH00000000
95A3	Geobacillus thermoglucosidasius Y4.1MC1	CP002293 , CP002294
95A4	Geobacillus thermoglucosidasius C56-YS93	CP002835 , CP002836

96A11	Geobacillus thermoleovorans Y4.12MC52	CP002442 , CP002443
96A12	Geobacillus thermoleovorans Y4.12MC61	CP001794 , CP001795
96A13	Geobacillus thermoleovorans C56-T3	CP002050
96A14	Geobacillus thermoleovorans GHH01	CP004008
99A2	Geobacillus teobii WCH70	CP001638 , CP001639 , CP001640