

## Bacteria in Bacillus genus

NCBI Accession	Project ID	Bacteria name
NC 000964	57675	Bacillus subtilis 168
NC 002570	57791	Bacillus halodurans C 125
NC 004722	57975	Bacillus cereus ATCC 14579
NC 003997	57909	Bacillus anthracis Ames
NC 003909	57673	Bacillus cereus ATCC 10987
NC 007530	58083	Bacillus anthracis Ames Ancestor
NC 005945	58091	Bacillus anthracis Sterne
NC 005957	58089	Bacillus thuringiensis serovar konkukian 97 27
NC 006270	58097	Bacillus licheniformis ATCC 14580
NC 006274	58103	Bacillus cereus E33L
NC 006322	58199	Bacillus licheniformis ATCC 14580
NC 006582	58237	Bacillus clausii KSM K16
NC 007103	58103	Bacillus cereus E33L
NC 004604	15862	Bacillus megaterium QM B1551
NC 004721	57975	Bacillus cereus ATCC 14579
NC 005707	57673	Bacillus cereus ATCC 10987
NC 007322	58083	Bacillus anthracis Ames Ancestor
NC 007323	58083	Bacillus anthracis Ames Ancestor
NC 006578	58089	Bacillus thuringiensis serovar konkukian 97 27
NC 007104	58103	Bacillus cereus E33L
NC 007105	58103	Bacillus cereus E33L
NC 007106	58103	Bacillus cereus E33L
NC 007107	58103	Bacillus cereus E33L
NC 008598	58795	Bacillus thuringiensis Al Hakam
NC 008600	58795	Bacillus thuringiensis Al Hakam
NC 009673	58317	Bacillus cytotoxicus NVH 391 98
NC 009674	58317	Bacillus cytotoxicus NVH 391 98
NC 009725	58271	Bacillus amyloliquefaciens FZB42
NC 009848	59017	Bacillus pumilus SAFR 032
NC 010008	15862	Bacillus megaterium QM B1551
NC 010009	15862	Bacillus megaterium QM B1551
NC 010010	15862	Bacillus megaterium QM B1551
NC 010180	58315	Bacillus weihenstephanensis KBAB4
NC 010181	58315	Bacillus weihenstephanensis KBAB4
NC 010182	58315	Bacillus weihenstephanensis KBAB4
NC 010183	58315	Bacillus weihenstephanensis KBAB4
NC 010184	58315	Bacillus weihenstephanensis KBAB4
NC 011654	58753	Bacillus cereus AH187
NC 011655	58753	Bacillus cereus AH187
NC 011656	58753	Bacillus cereus AH187
NC 011657	58753	Bacillus cereus AH187
NC 011658	58753	Bacillus cereus AH187
NC 011725	58757	Bacillus cereus B4264
NC 011771	58751	Bacillus cereus AH820
NC 011772	58759	Bacillus cereus G9842
NC 011773	58751	Bacillus cereus AH820
NC 011774	58759	Bacillus cereus G9842
NC 011775	58759	Bacillus cereus G9842
NC 011776	58751	Bacillus cereus AH820
NC 011777	58751	Bacillus cereus AH820
NC 011969	58529	Bacillus cereus Q1

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NC 011971	58529	Bacillus cereus Q1
NC 011973	58529	Bacillus cereus Q1
NC 012472	59299	Bacillus cereus 03BB102
NC 012473	59299	Bacillus cereus 03BB102
NC 012577	59303	Bacillus anthracis CDC 684
NC 012579	59303	Bacillus anthracis CDC 684
NC 012581	59303	Bacillus anthracis CDC 684
NC 012655	59385	Bacillus anthracis A0248
NC 012656	59385	Bacillus anthracis A0248
NC 012659	59385	Bacillus anthracis A0248
NC 013791	45847	Bacillus pseudofirmus OF4
NC 013792	45847	Bacillus pseudofirmus OF4
NC 013793	45847	Bacillus pseudofirmus OF4
NC 014019	15862	Bacillus megaterium QM B1551
NC 014023	15862	Bacillus megaterium QM B1551
NC 014025	15862	Bacillus megaterium QM B1551
NC 014031	15862	Bacillus megaterium QM B1551
NC 014103	48371	Bacillus megaterium DSM319
NC 014171	49135	Bacillus thuringiensis BMB171
NC 014172	49135	Bacillus thuringiensis BMB171
NC 014219	49513	Bacillus selenitireducens MLS10
NC 014331	50615	Bacillus cereus biovar anthracis CI
NC 014332	50615	Bacillus cereus biovar anthracis CI
NC 014333	50615	Bacillus cereus biovar anthracis CI
NC 014335	50615	Bacillus cereus biovar anthracis CI
NC 014479	51879	Bacillus subtilis spizizenii W23
NC 014551	53535	Bacillus amyloliquefaciens DSM 7
NC 014639	59887	Bacillus atrophaeus 1942
NC 014976	62463	Bacillus subtilis BSn5
NC 015634	68053	Bacillus coagulans 2 6
NC 016047	73967	Bacillus subtilis spizizenii TU B 10