

Table S1. Genomic features of strains NKC19-3^T, NKC19-16^T, and DSM25609^T. All data were from this study unless stated otherwise.

Strains: 1, NKC19-3^T; 2, NKC19-16^T; 3, *Virgibacillus natechei* DSM25609^T.

Properties	1	2	3
BioProject	PRJNA726327	PRJNA726329	PRJNA894318
BioSample	SAMN18927773	SAMN18927775	SAMN31452450
Accession No.	JAGYHC000000000	CP074373	CP110224
Sequencing method	PacBio RSII	PacBio RSII	PacBio RSII
Assembly method	HGAP v. 3.0	HGAP v. 3.0	HGAP v. 3.0
Genome size (bp)	4,054,365	3,829,764	3,867,590
Contig	2	1	1
Genome coverage	122.3x	343.7x	322.6x
CDSs (total)	3,785	3,726	3,776
rRNA genes	88	86	85
tRNA genes	65	63	62
G + C content (mol%)	38.9	39.5	38.4

CDSs, coding sequence