

**Table S1. Antimicrobial Potential.** Gene clusters associated with the synthesis of nonribosomal peptide-synthetases (NRPS), polyketide synthases (PKSs) and ribosomally synthesized and post-translationally modified peptides (RiPPs) identified in the genomes of the *Bacillus* isolates AB6, CG1, X1 and Z1.

Strain	Source	Gene Cluster	Antibiotic <sup>#</sup>	Similarity (%) <sup>*</sup>	Reference
<i>B. sonorensis</i> AB6	Honey	NRPS	Lichenysin	100	(61)
		NRPS	Bacitracin	88	(62)
		NRPS	Bacillibactin, paenibactin, griseobactin	46	(63)
		RiPP, Lantipeptide	Cytolysin	40	(64)
		NRPS	Fengycin	30	(65)
		RiPP, Lantipeptide	Streptin	20	(66)
		NRPS-PKS	Plipastatin	23	(67)
		NRPS	Plipastatin- Fengycin	15	(65, 67)
		RiPP, Lasso peptide	ND	-	(68)
<i>B. licheniformis</i> CG1	Honey	NRPS	Lichenysin	100	(61)
		RiPP, Lantipeptide	Lichenicidin	100	(57)
		NRPS	Bacillibactin, griseobactin	46	(63)
<i>B. sonorensis</i> X1	Royal Jelly	NRPS	Lichenysin	100	(61)
		NRPS	Bacitracin	88	(62)
		NRPS	Bacillibactin, paenibactin, griseobactin	46	(63)
		NRPS	Fengycin	30	(65)
		NRPS-PKS	Plipastatin	23	(67)
		NRPS	Plipastatin- Fengycin	23	(65, 67)
		RiPP, Lantipeptide	Streptin	20	(66)
		RiPP, Lasso peptide	ND	-	(68)
<i>B. subtilis</i> Z1	Honey	NRPS	Fengycin	100	(65)
		RiPP, sactipeptide	Sporulation killing factor, skfA	100	(69)
		RiPP, sactipeptide	Subtilosin A	100	(70)
		NRPS-PKS	Bacillaene	92	(71)
		NRPS	Bacillibactin	92	(63)
		NRPS	Bacilysin	85	(72)
		NRPS	Surfactin	78	(73)
		RiPP, glyocin	Sublancin	66	(74)

#ND, not described; \*Percentage of similar genes found in other clusters

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