

Supplementary Materials: A Genomic and Proteomic Approach to Identify and Quantify the Expressed *Bacillus thuringiensis* Proteins in the Supernatant and Parasporal Crystal

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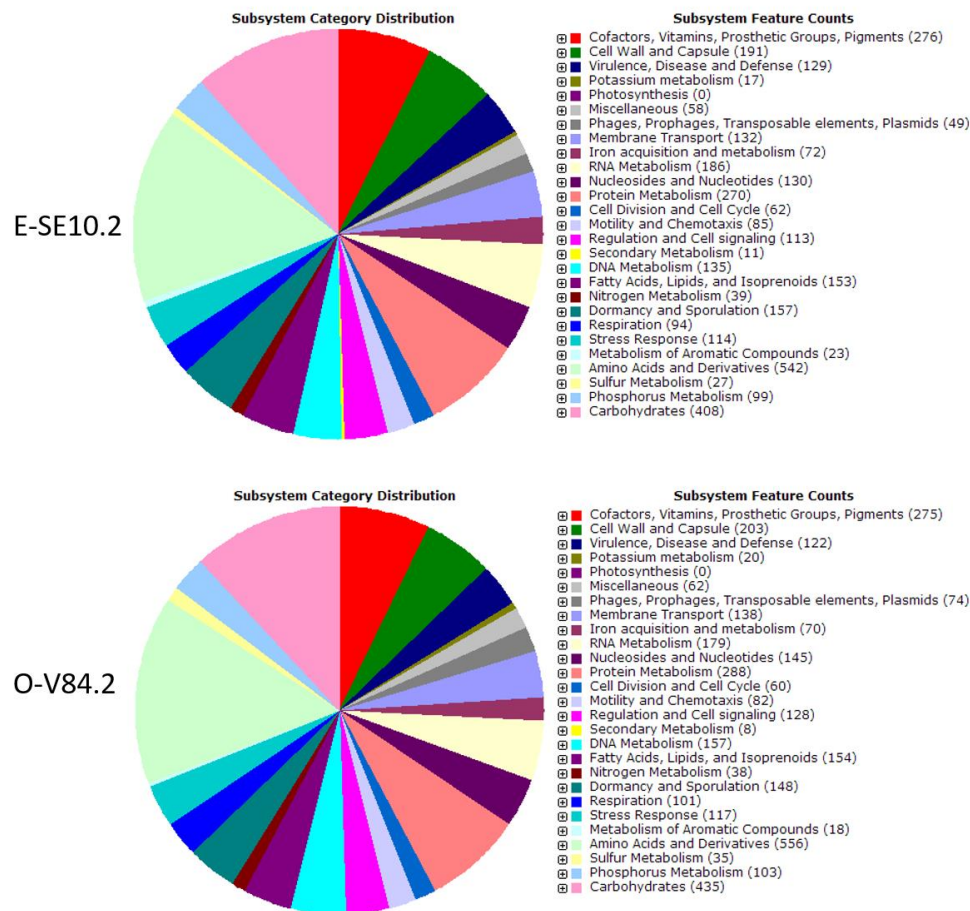


Figure S1. Subsystem category distribution of the *Bt* isolates E-SE10.2 and O-V84.2 by the genome annotation based on the Rast server. The right pie chart indicates the percentage or predicted encoding genes associated to at least one subsystem.

