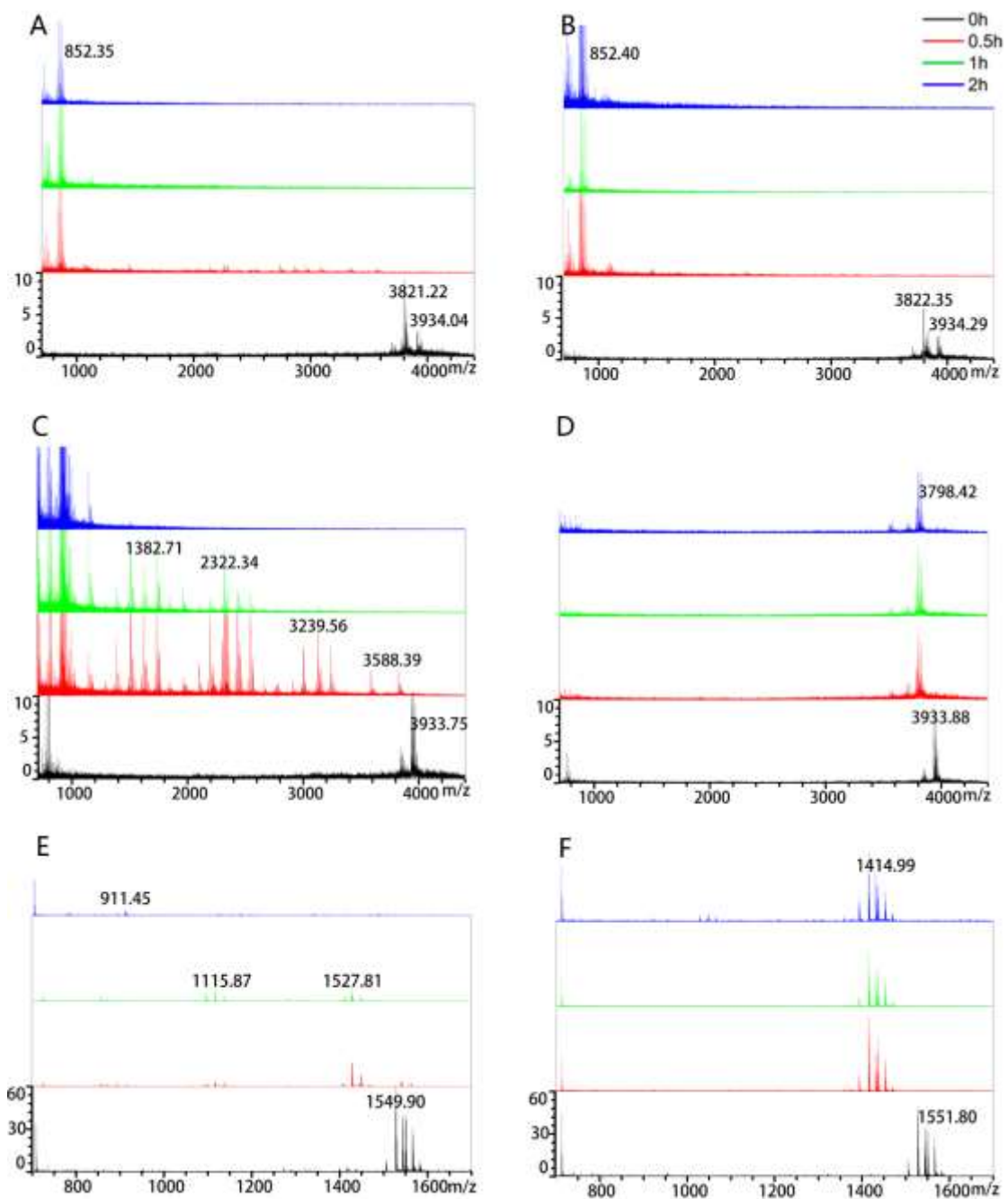


1 Figure S1



2

3 Figure S1. MALDI-TOF analysis of the degradation profiles of the immunotoxic
 4 peptides. The cell densities used for the incubations were OD₆₂₀ 1.2. The 33-mer (0.5
 5 mg/mL) gliadin peptide was incubated with *B. cereus* AFA01 from the medium of
 6 MCG-1 (A), SCB (B), BA (C) or LB (D) for 0, 0.5, 1 and 2 h. The 13-mer gliadin

- 7 peptide (0.25 mg/mL) was incubated with *B. cereus* AFA01 from the medium of BA (E)
- 8 or LB (F) for 0, 0.5, 1 and 2 h, respectively.

9 Table S1 Comparative genome statistics of th *B. cereus*

	CH	21155	AFA01
Genome Size (bp)	6098577	6258015	5843781
Chromosome No.	1	1	1
Plasmid No.	1	4	4
GC Content(%)	34.92	34.91	35.22
CDS No.	6063	6362	5925
tRNA No.	106	107	107
rRNA No.	42	42	42
Accession	SAMN17369856	SAMN17369855	SAMN17369857