

Complete genome sequence and epigenetic profile of *Bacillus velezensis* UCMB5140 used for plant and crop protection in comparison with other plant-associated *Bacillus* strains

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Supplementary material

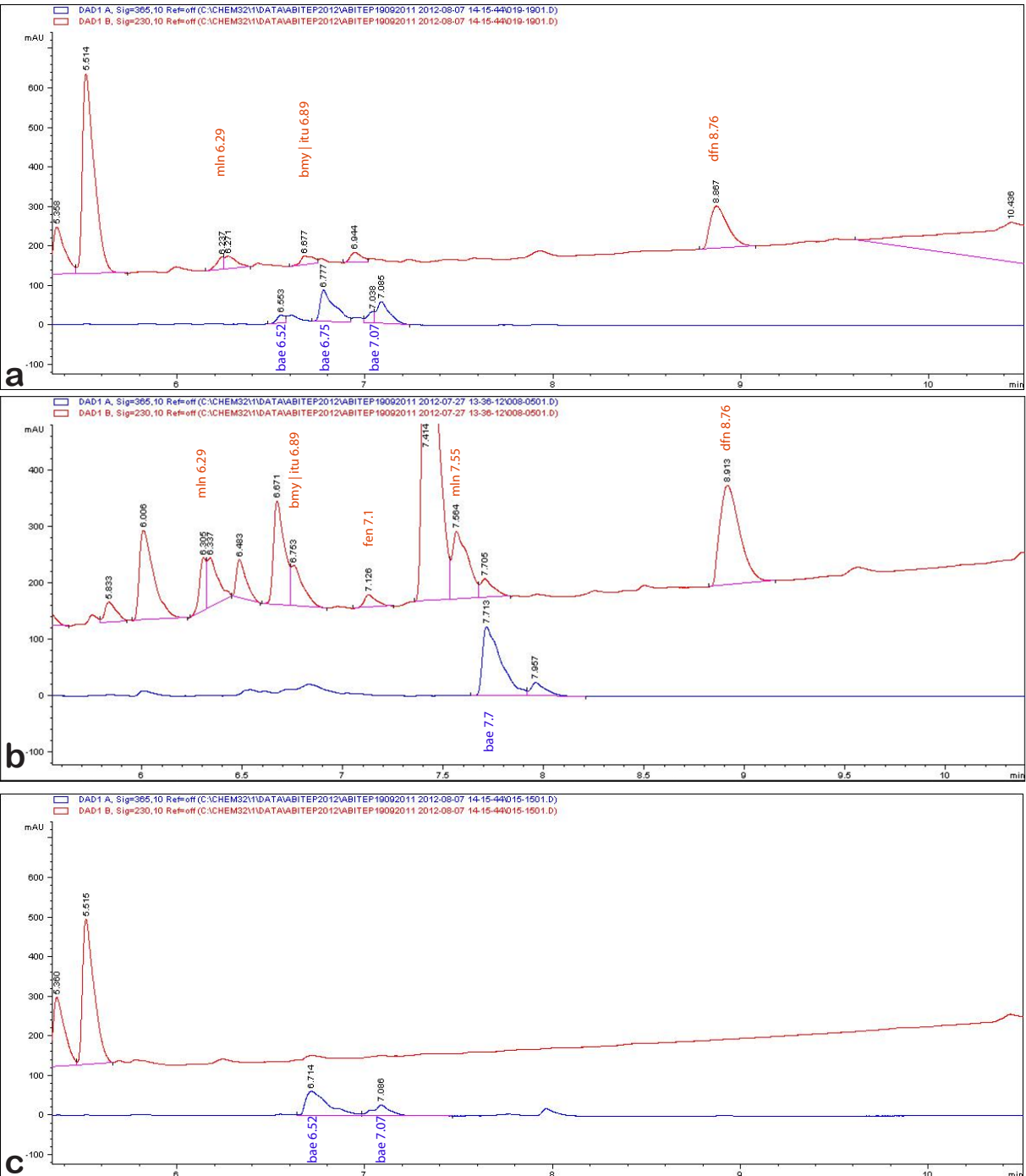


Fig. S1 Characteristic peak areas retrieved from HPLC chromatograms in culture media after 24 h of cultivation of a) UCMB5140; b) UCMB5113; and c) UCMB5021. Product abbreviations: bae – bacillaene; bmy – bacillomycin D; dfn – difficidin; fen – fengycin; itu – iturin A; mln – macrolactin D

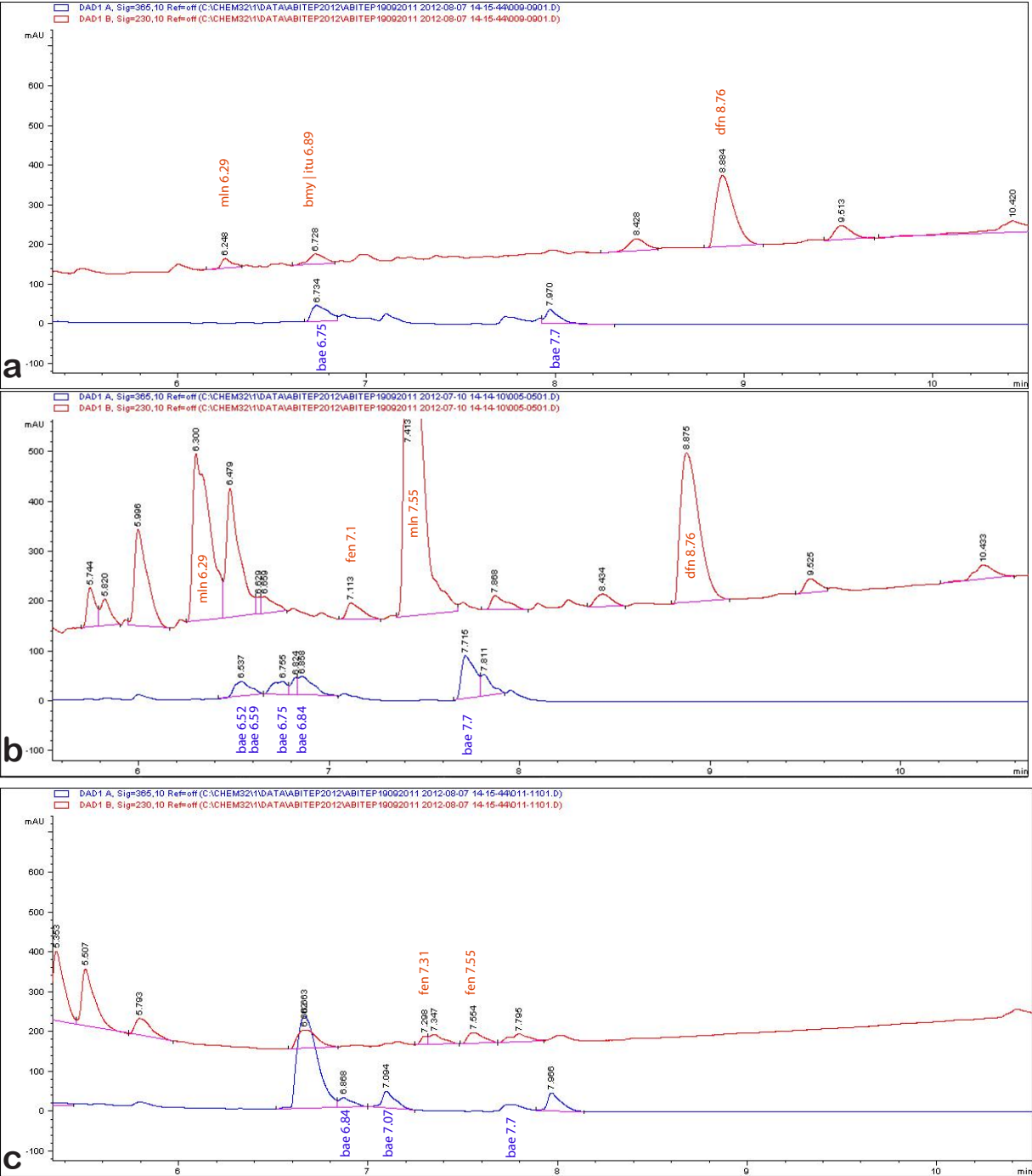
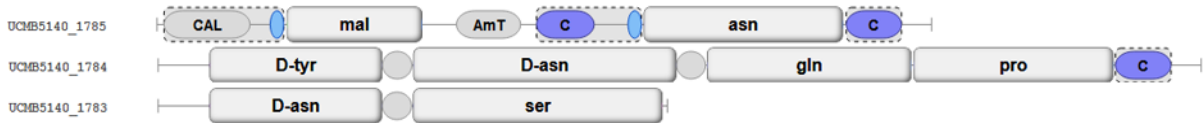


Fig. S2 Characteristic peak areas retrieved from HPLC chromatograms in culture media after 48 h of cultivation of a) UCMB5140; b) UCMB5113; and c) UCMB5021. Product abbreviations: bae – bacillaene; bmy – bacillomycin D; dfn – diffidin; fen – fengycin; itu – iturin A; mln – macrolactin D

UCMB 5140



UCMB 5113

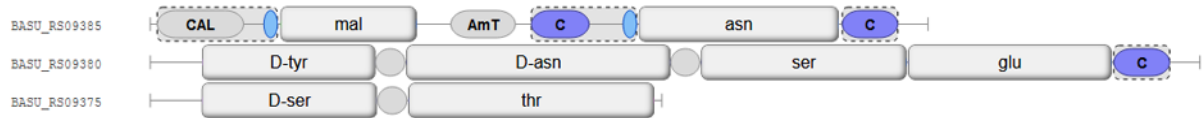


Fig. S3 Amino acid specific adenylation domains predicted by antiSMASH in NRPS gene clusters encoding iturin A in UCMB5140 and bacillomycin D in UCMB5113