**Supplementary Fig. 6.** Anti-SMASH predicted BGCs (lasso peptides) derived from strain BRMEA1\(^T\). (A) Identification of secondary metabolite regions using a 'relaxed' strictness. (B) Comparative analysis of type I polyketide synthase metabolite cluster blaster against the National Center for Biotechnology Information (NCBI) database. (C) Gene annotation with Pfam-based GC terms. Rows highlighted in dark pink denote core biosynthetic genes; rows highlighted in light pink show additional biosynthetic genes.