Electronic Supplementary Material (ESI) for New Journal of Chemistry
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A new surfactin-C_{17} produced by *Bacillus subtilis* TD7 with a low critical micelle concentration and high biological activities

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Table S1 16S rDNA gene sequencing of *Bacillus subtilis* TD7.

GenBank accession number	16S rDNA gene sequence
Bacillus subtilis BAB-7106	ACGATGCGTAGCCGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACG GCCCAGACTCCTACGGGAGGCAGCAGTAGGGAATCTTCCGCAATGGACGAAAG TCTGACGGAGCAACGCCGCGTGAGTGATGAAGGTTTTCGGATCGTAAAGCTCTG TTGTTAGGGAAGAACAAGTACCGTTCGAATAGGGCGGTACCTTGACGGTACCTA ACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGTGGC AAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGCAGGCGGTTTCTTAAGTC TGATGTGAAAGCCCCCGGCTCAACCGGGGAGGGTCATTGGAAACTGGGGAACT TGAGTGCAGAAGAGGAGAGTGGAATTCCACGTGTAGCGGTGAAATGCGTAGAG ATGTGGAGGAACACCAGTGGCGAAGGCGACTCTCTGGTCTGTAACTGACGCTGA GGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCG TAAACGATGAGTGCTAAGTGTTAGGGGGTTTCCGCCCCTTAGTGCTGCAGCTAAC GCATTAAGCACTCCGCCTGGGGAGTACGGTCGCAAGACTGAAACTCAAATGAAT TGACGGA

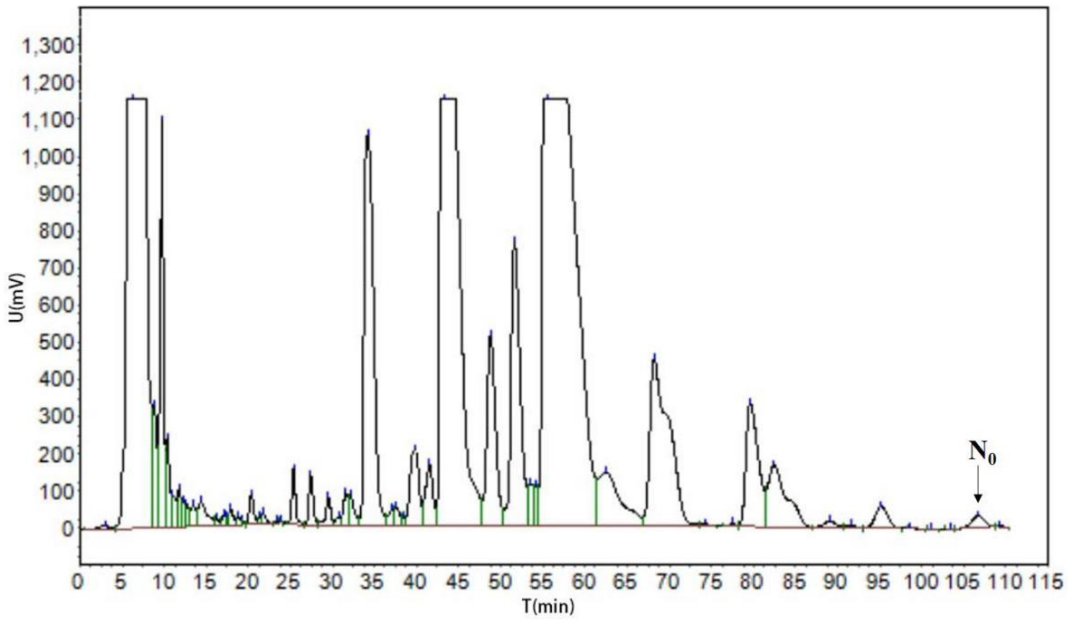

Figure. S1 RP-HPLC of lipopeptide and the peak of target substance (N_0).

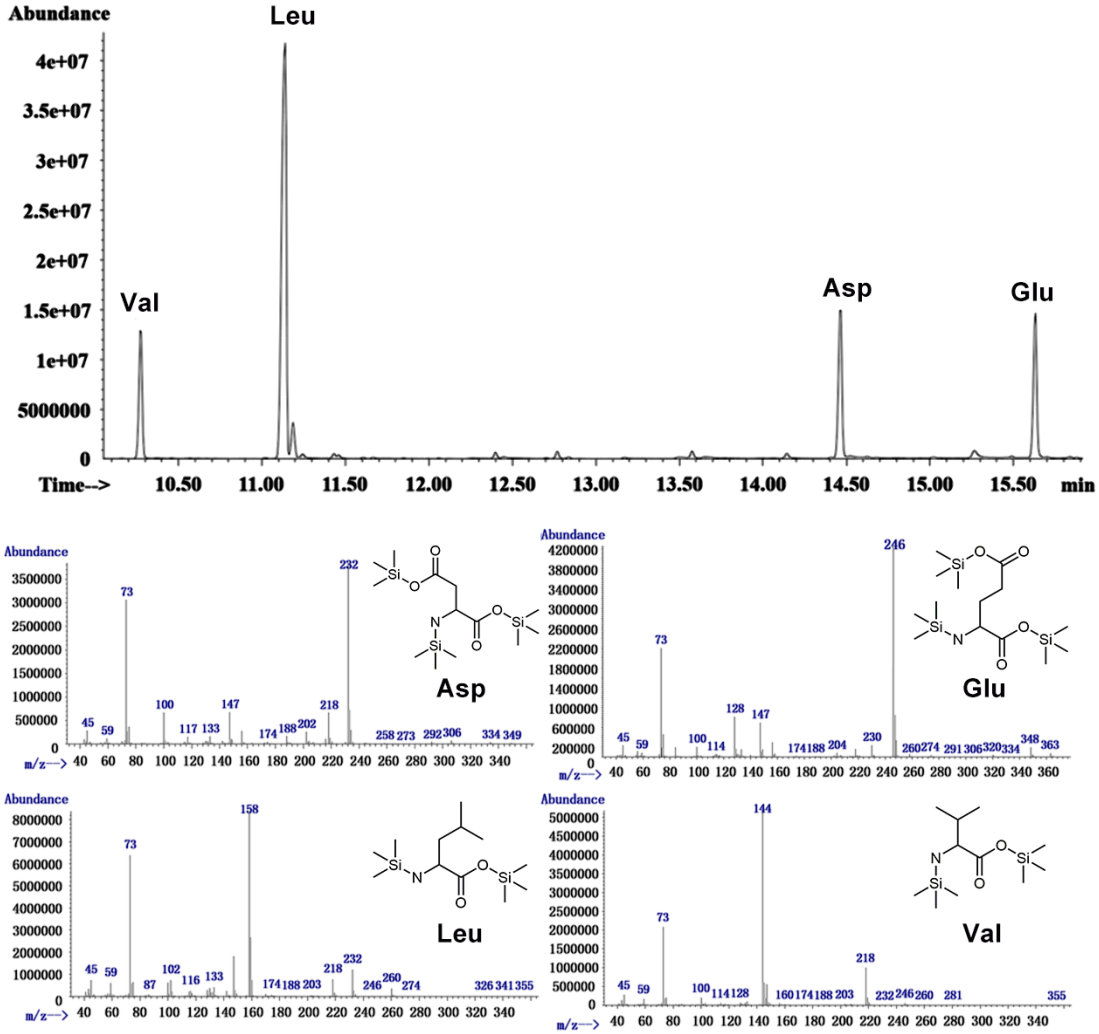

Figure. S2 The total ion and mass chromatograms of the amino acids in the N_0 after trimethylsilylation.

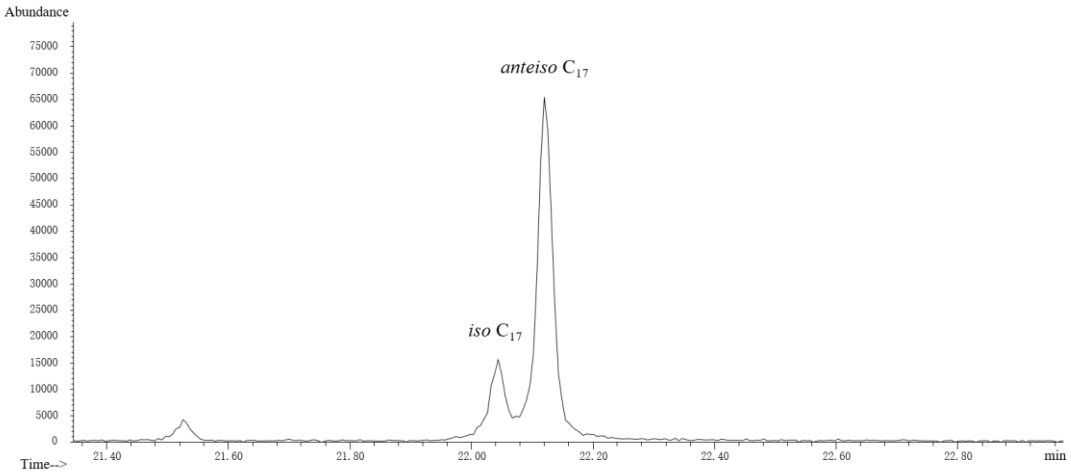

Figure. S3 The extracted ion chromatogram of the fraction of N_0 (*m/z* = 233).

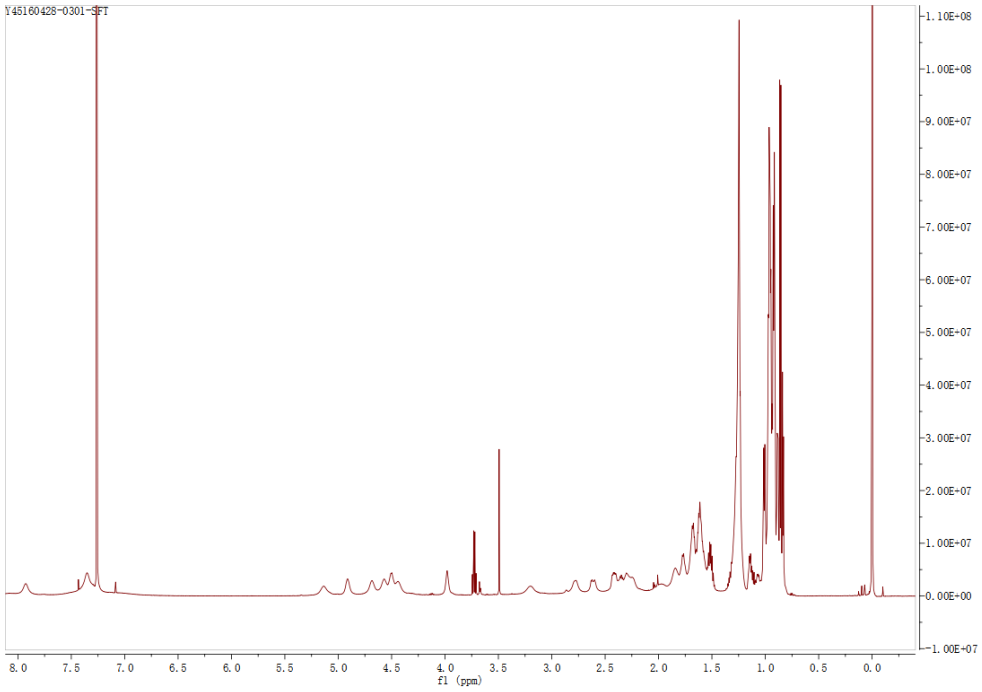

Figure. S4 ^{1}H NMR spectrum ($CDCl_3$) of N_0.

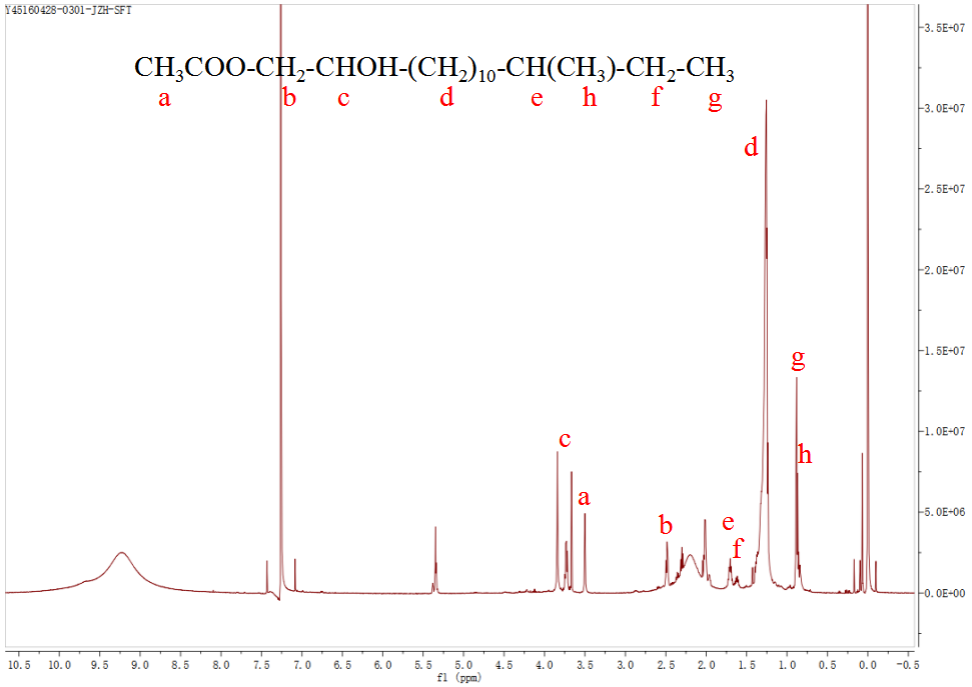

Figure. S5 ^{1}H NMR spectrum ($CDCl_3$) of fatty acid methyl ester.

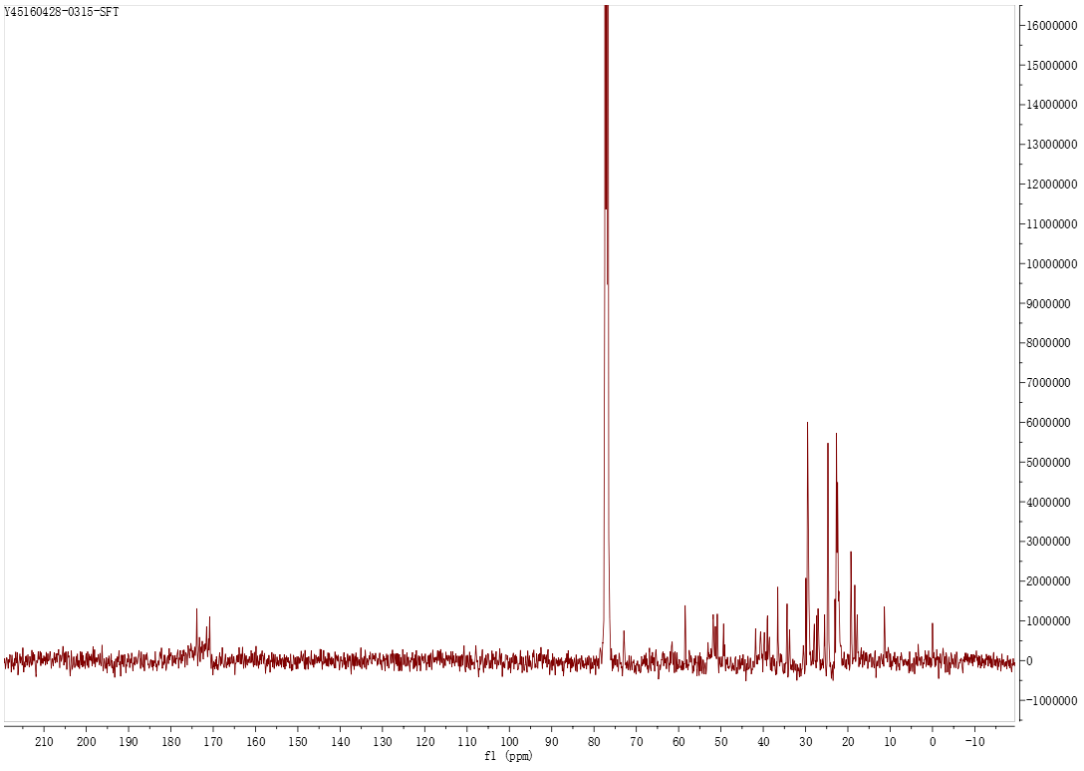

Figure. S6 ^{13}C NMR spectrum ($CDCl_3$) of N_0.

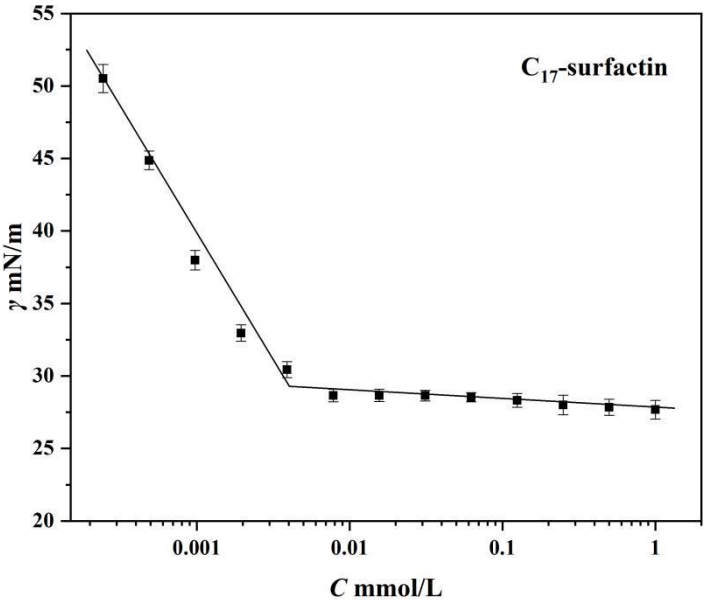

Figure. S7 Surface tension versus logarithm of concentrations value of surfactin-C_{17}.