Supporting Information

Citrinin Is a Potential Quorum Sensing Inhibitor Against *Pseudomonas aeruginosa*

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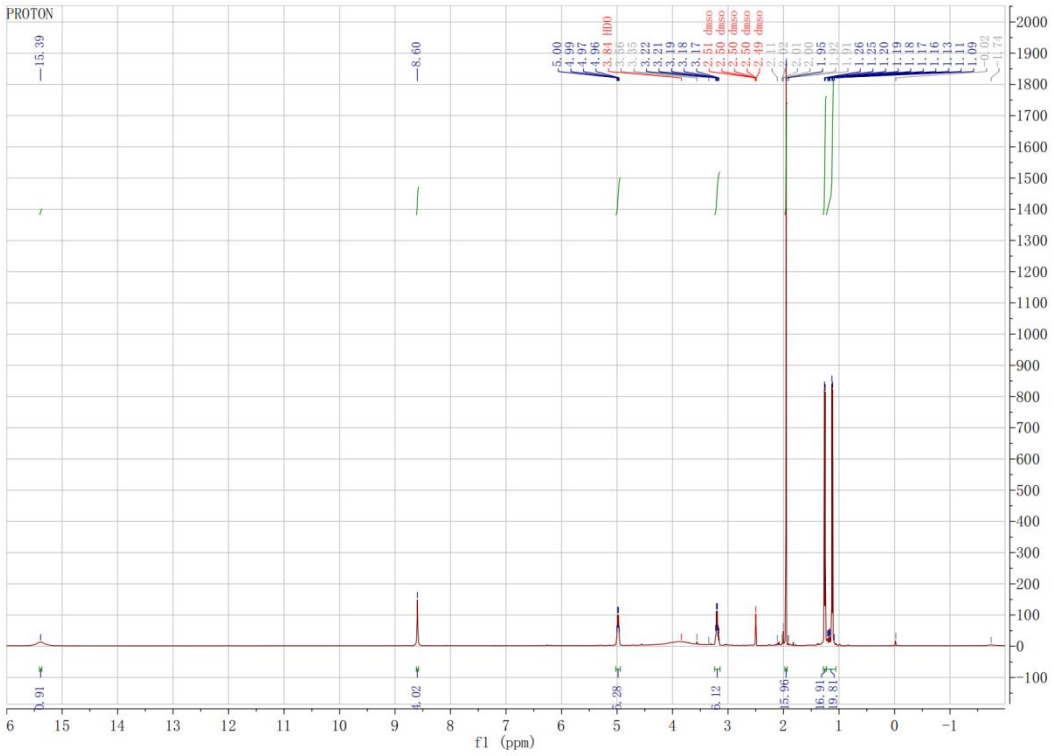

Figure S1. ^{1}H-NMR spectroscopy of the anti-QS compound (Solvent: DMSO-d6). ^{1}H NMR (500M Hz, DMSO-d_6) δ 15.39 (s, 1H), 8.60(s, 1H), 4.98(q, *J*=6.7Hz, 1H), 3.20(q, *J*=7.2Hz, 1H), 1.95(s, 4H), 1.26(d, *J*=6.7Hz, 4H), 1.12(d, *J*=7.2Hz, 1H).

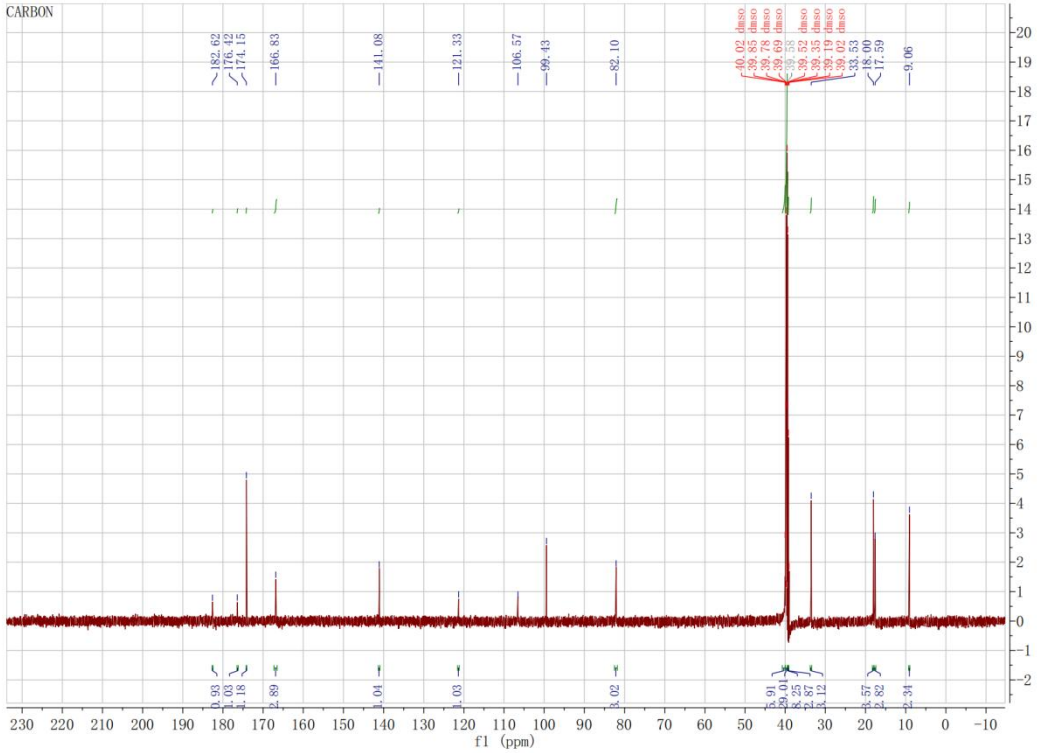

Figure S2. ^{13}C-NMR spectroscopy of the anti-QS compound (Solvent: DMSO-d6). ^{13}C NMR (126 MHz, DMSO) δ 182.62, 176.42, 174.16, 166.83, 141.08, 121.33, 106.57, 99.43, 82.10, 33.53, 18.00, 17.59, 9.06.

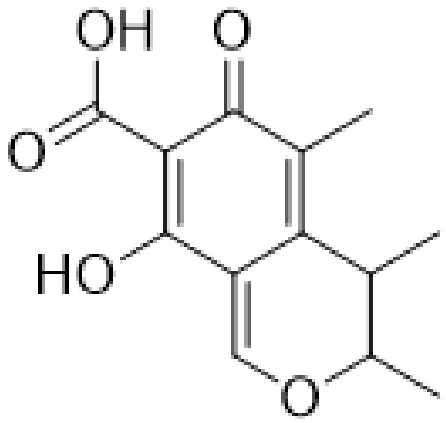

Figure S3. Chemical structure of citrinin.

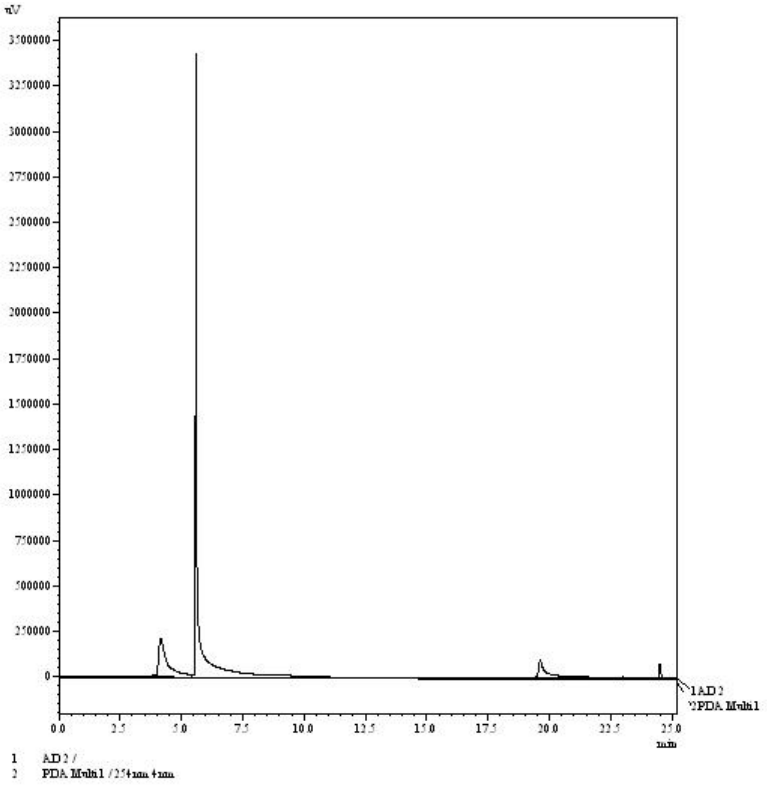

Figure S4. HPLC analysis of citrinin.

Table S1. Bacterial strains and plasmids.

Strain or plasmid	Relevant genotype	Source
Strains		
P. aeruginosa PAO1	Wild type	Schuster and Greenberg. (2007)
Chromobacterium violaceum CV026	Δ*cviI*, mini Tn5 mutant of CV31532	McLean. (2004)
Chromobacterium violaceum CV12472	Wild type	McLean. (2004)
E. coli MG4/pKDT17	*E. coli* DH5a harboring plasmid MG4/pKDT17	Hong et al. (2012)
E. coli pEAL08-2	*E. coli* DH5a harboring plasmid pEAL08-2	Cugini et al. (2007)
E. coli pDSY	*E. coli* DH5a harboring plasmid pDSY	Zhao et al. (2021)
Plasmids		
MG4/pKDT17	P*lasB-lacZ*, and P*lac-lasR* transcriptional fusion	Pearson et al. (1994)
pEAL08-2	P*pqsA-lacZ*, and P*lac-pqsR* transcriptional fusion	Cugini et al. (2007)
pDSY	P*rhlA-lacZ*, and P*lac-rhlR* transcriptional fusion	Zhao et al. (2021)

Table S2. The primer sequences for real-time RT-PCR.

primers	sequences (5′ to 3′)	primers	sequences (5′ to 3′)
rpsl (F)	GCAACTATCAACCAGCTGGTG	*rpsl* (R)	GCTGTGCTCTTGCAGGTTGTG
lasR (F)	CTGTGGATGCTCAAGGACTAC	*lasR* (R)	AACTGGTCTTGCCGATGG
lasI (F)	GGCTGGGACGTTAGTGTCAT	*lasI* (R)	AAAACCTGGGCTTCAGGAGT
lasB(F)	ACCAGAAGATCGGCAAGTAC	*lasB*(R)	GTTGACCTGCTTGTAGGTGTTG
rhlR (F)	CTGGGCTTCGATTACTACGC	*rhlR* (R)	CCCGTAGTTCTGCATCTGGT
rhlI (F)	GTAGCGGGTTTGCGGATG	*rhlI* (R)	CGGCATCAGGTCTTCATCG
rhlA (F)	GGCGATCGGCCATCT	*rhlA* (R)	AGCGAAGCCATGTGCTGAT
pqsR(F)	CTGATCTGCCGGTAATTGG	*pqsR*(R)	ATCGACGAGGAACTGAAGA
pqsA (F)	GACCGGCTGTATTCGATTC	*pqsA* (R)	GCTGAACCAGGGAAAGAAC
phzH (F)	TGCGCGAGTTCAGCCACCTG	*phzH* (R)	TCCGGGACATAGTCGGCGCA