

Supplementary Information

**Chloramphenicol-borate/boronate Complex for Controlling Infections
by Chloramphenicol-Resistant Bacteria**

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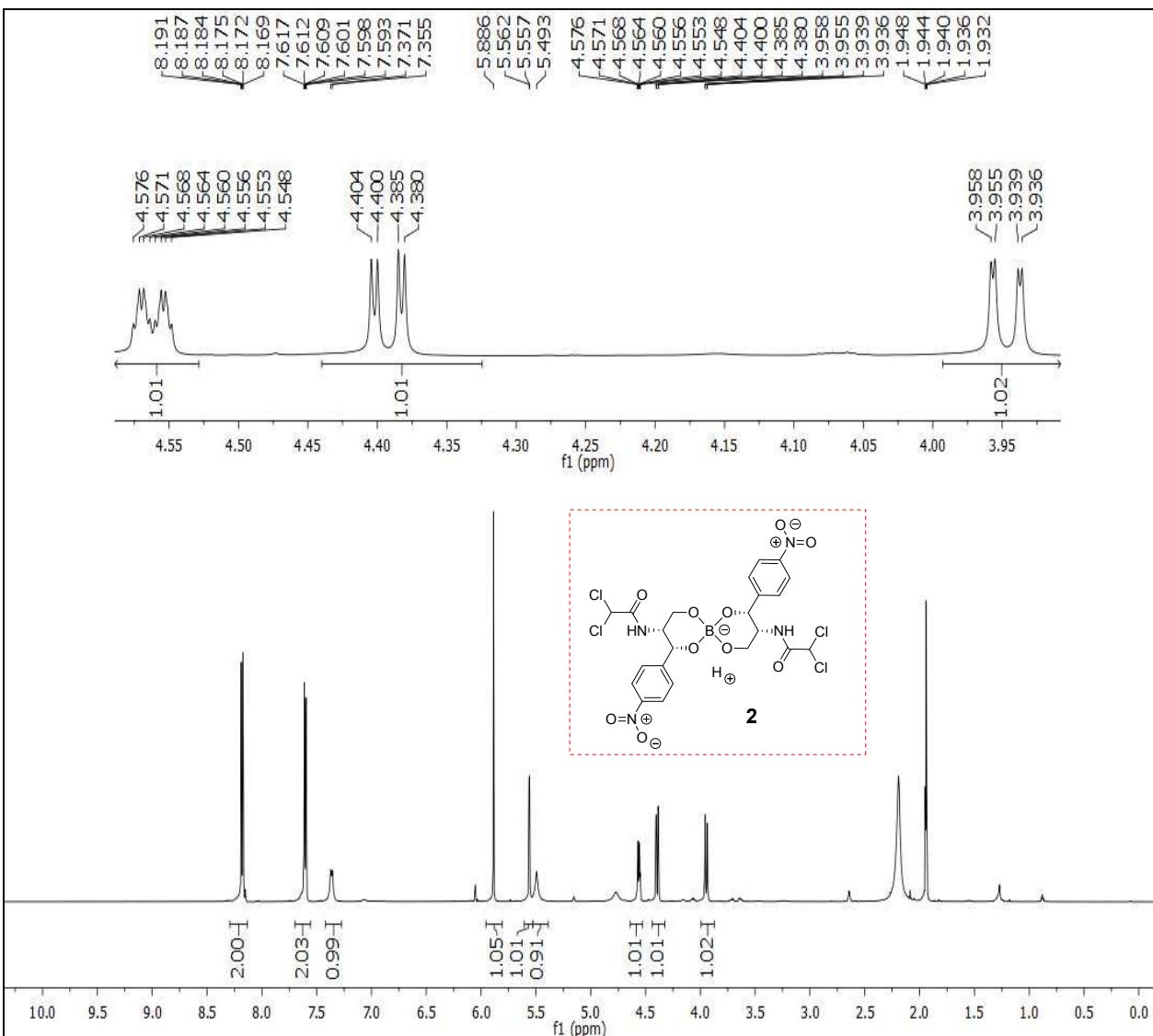


Figure S1. ^1H NMR spectrum of Chloramphenicol-borate ester (**2**) in CD_3CN , in 600 MHz.

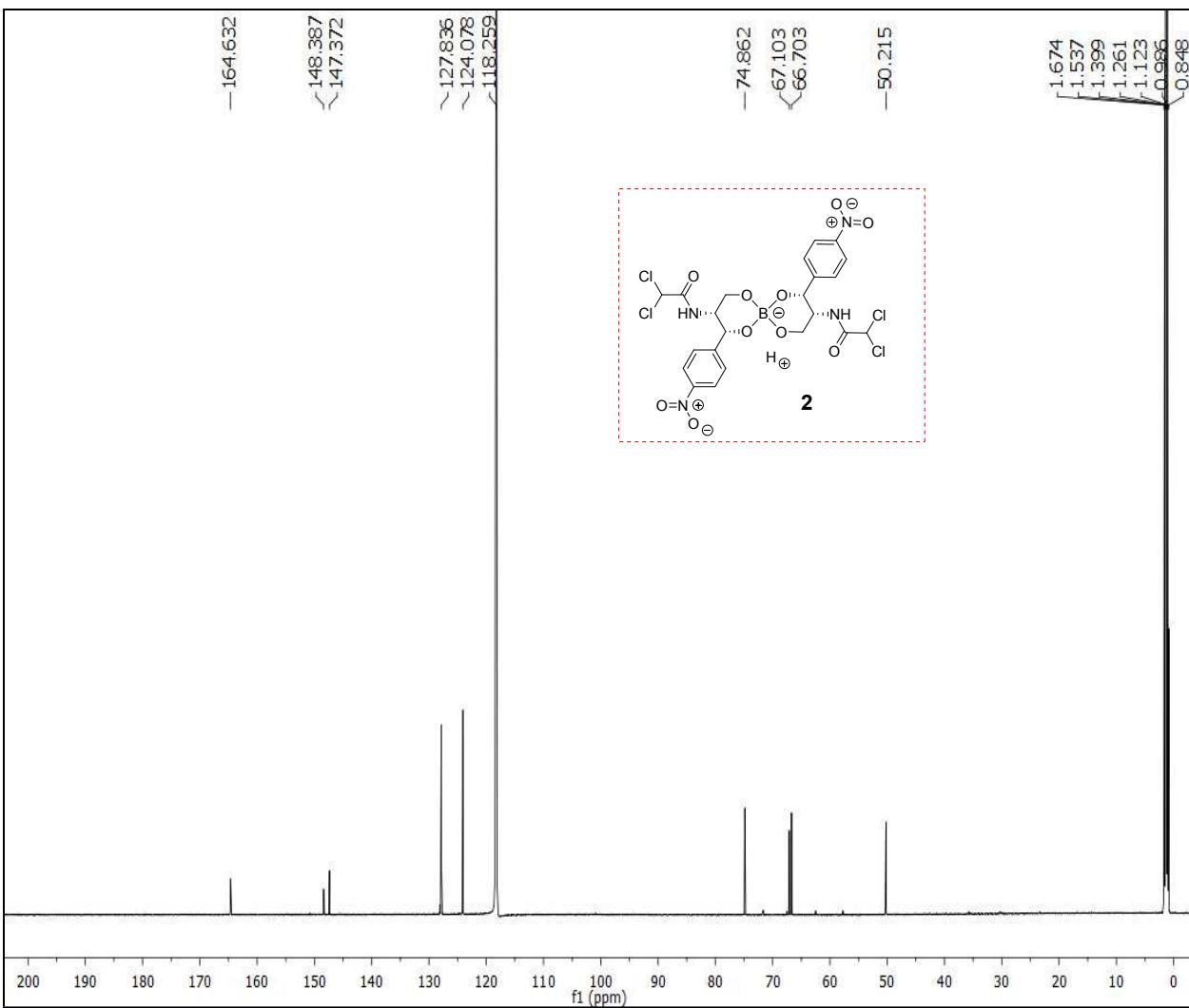


Figure S2. ^{13}C NMR spectrum of Chloramphenicol-borate ester (**2**) in CD_3CN , in 150 MHz.

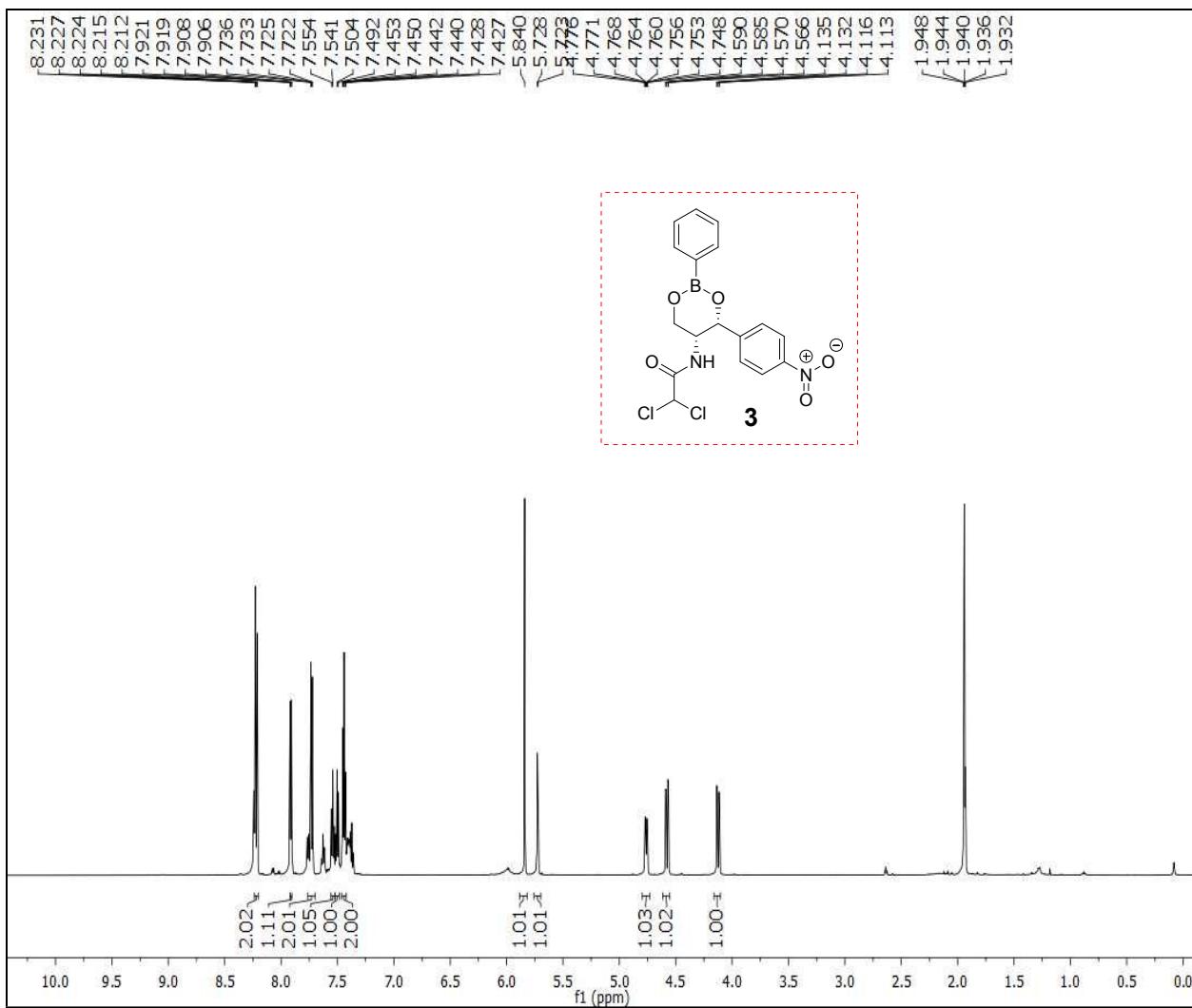


Figure S3. ^1H NMR spectrum of Chloramphenicol-phenyl boronate (**3**) in CD_3CN , in 600 MHz.

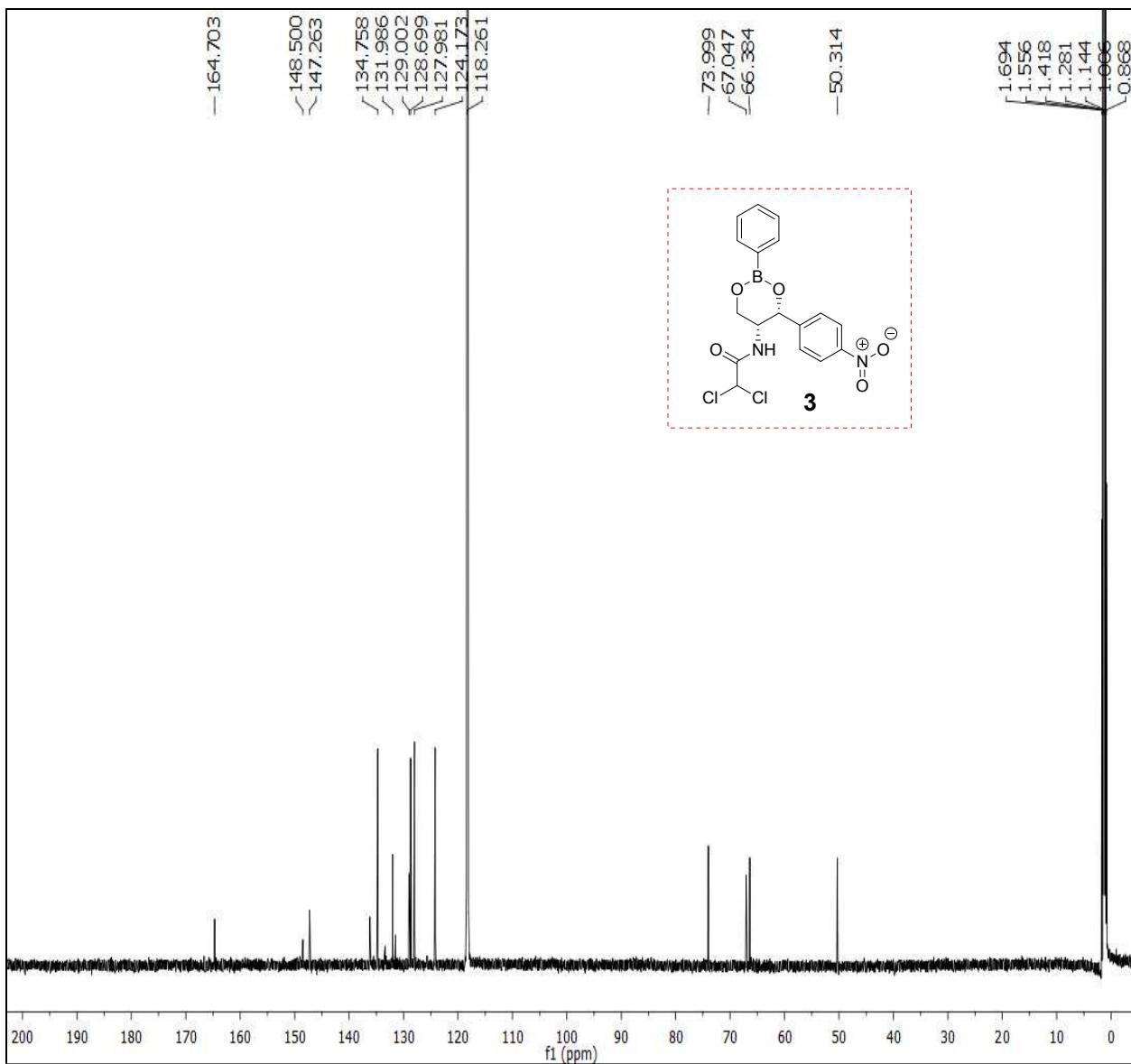


Figure S4. ^{13}C NMR spectrum of Chloramphenicol-phenyl boronate ester (**3**) in CD_3CN , in 150 MHz.

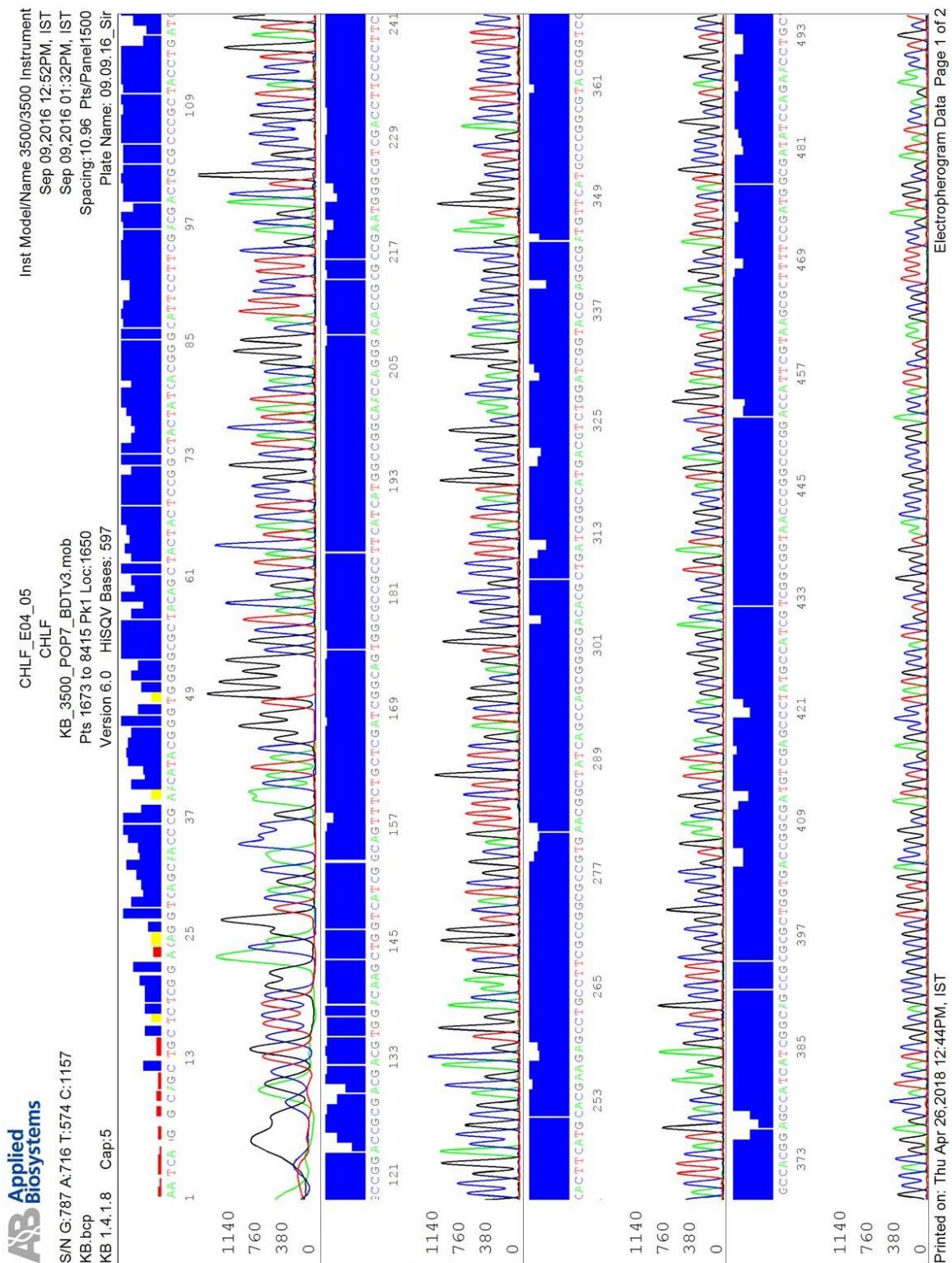


Figure S5. Forward primer based sequencing data of chloramphenicol acetyl transferase obtained from *P. aeruginosa* SS3.

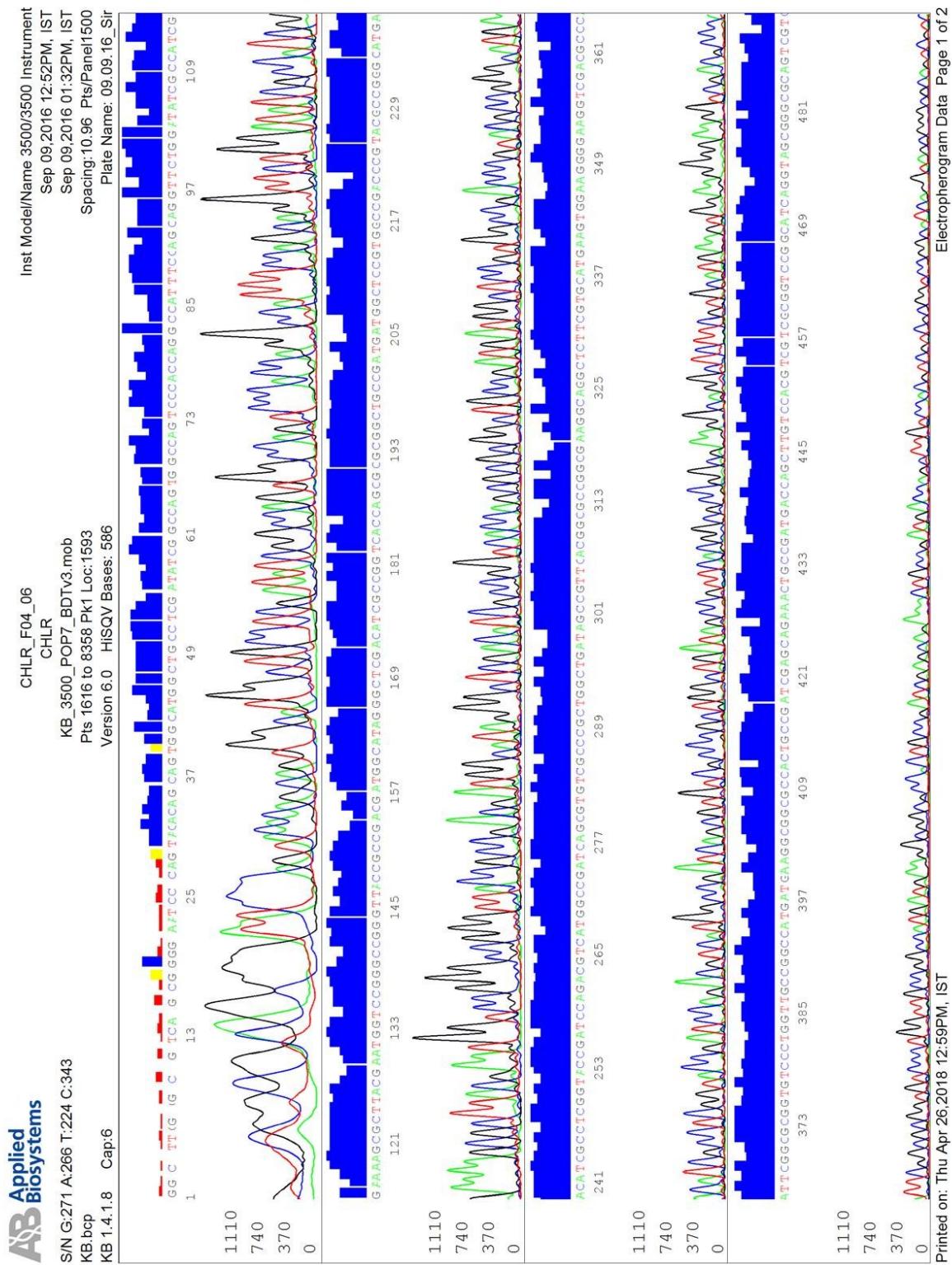


Figure S6. Reverse primer based sequencing data of chloramphenicol acetyl transferase obtained from *P. aeruginosa* SS3.

Contig result of obtained sequences from *Pseudomonas aeruginosa* strain SS3, Chloramphenicol acetyl transferase

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CATGCACGAAGAGCCTGCCCTCGCCGGCGCGTGAACGGCTATCAGCCAGCGGGCACAGCTGATCG
GCCATGACGTCTGGATCGGTACCGAGGGATGTCATGCCCGCGTACGGGCGACGGAGCCATC
ATCGGCAGCCGCGCGCTGGTACCGCGATGTCGAGCCCTATGCCATCGTCGGCGGTAAACCGGCCG
GACCATTGTAAGCGCTTCCATGGCGATATCCAGAACCTGCTGGAAATGGCTGGTGGACTGGC
CACTGGCGATATCGAGGCAGCCATGCCACTGCTGTACTGGGATATCCCCGCTTGTACCAAGGC
ACTGGAACACAGCGCCAGGCCTCGGCCTGA
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Figure S7. Homology analysis of the obtained sequence of catB gene from *P. aeruginosa* EXR1.

Range	Score	Expect	Identities	Gaps	Strand
14837043 to 4837681	1166 bits(631)	0.0	638/641(99%)	2/641(0%)	Plus/Plus

Features: Chloramphenicol acetyltransferase

Query 10 ATGGGCAACTATTCGAGAGGCCATTAGGGGCAAGCTGCTCGAACAGGTAGCAACCCGAACAT 69
Sbjct 4837043 ATGGGCAACTATTCGAGAGGCCATTAGGGGCAAGCTGCTCGAACAGGTAGCAAC 4837102

Query 70 CCGAACATAACGGGTGGGGCGCTACAGCTACTACTCCGGCTACTATCACGGGATTCC 129
Sbjct 4837103 CCGAACATAACGGGTGGGGCGCTACAGCTACTACTCCGGCTACTATCACGGGATTCC 4837162

Query 130 GACGACTGCGCCCGCTACCTGATGCCGACCGCAGACGTGGACAGCTGGTATCGGC 189
Sbjct 4837163 GACGACTGCGCCCGCTACCTGATGCCGACCGCAGACGTGGACAGCTGGTATCGGC 4837222

Query 190 AGTTCTGCTCGATGGCACTGGGCCCGCTTACATCATGGCCGCAACCAAGGGACACCC 249
Sbjct 4837223 AGTTCTGCTCGATGGCACTGGGCCCGCTTACATCATGGCCGCAACCAAGGGACACCC 4837282

Query 250 GCGGAATGGCGCTGACCTTCCCTTCCACTTACATGACGAAAGAGCCTGCGCTCGCC 309
Sbjct 4837283 GCGGAATGGCGCTGACCTTCCCTTCCACTTACATGACGAAAGAGCCTGCGCTCGCC 4837342

Query 310 GCGCTGAAACGGCTATCAGCCAGCGGGCGACAGCTGATCGCCATGACGCTGGATCGG 369
Sbjct 4837343 GCGCTGAAACGGCTATCAGCCAGCGGGCGACAGCTGATCGCCATGACGCTGGATCGG 4837402

Query 370 ACCGAGGCATGTTCATGCCCGGTACGGGTGCCAACGGAGCCATCATCGGAGCCGC 429
Sbjct 4837403 ACCGAGGCATGTTCATGCCCGGTACGGGTGCCAACGGAGCCATCATCGGAGCCGC 4837462

Query 430 GCGCTGGTACCCGGCGATGTCGAACCCCTATGCCATGTCGGGGTAACCCGGCCGGACC 489
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Query 490 ATTCTGTAAGCGCTTCCGATGGGATATCCAGAACCTGCTGGAAATGGCTGGTGGGAC 549
Sbjct 4837523 ATTCTGTAAGCGCTTCCGATGGGATATCCAGAACCTGCTGGAAATGGCTGGTGGGAC 4837582