

Antisense RNA Knockdown and Genomic Deletion of wecD Sensitizes Escherichia coli to Bacteriophage T7 Infection

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SUPPLEMENTAL MATERIAL

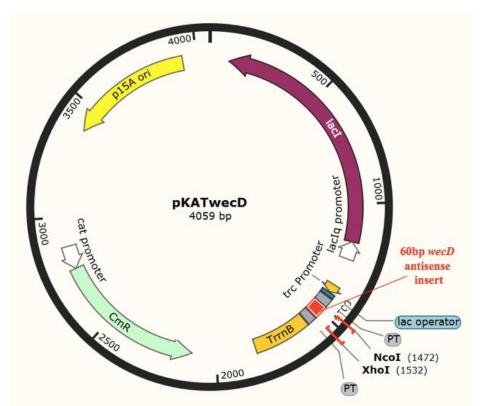


FIG. S1 pKATwecD plasmid map depicting the 60 base pair wecD antisense insert, restriction enzyme sites of NcoI and XhoI, and paired termini (PT).

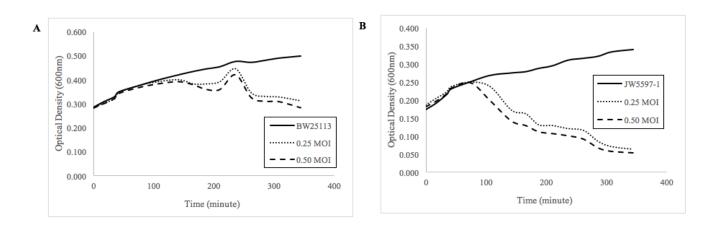


FIG. S2 The effect of *wecD* genomic deletion on bacteriophage T7 infectivity at 0.25 and 0.5 MOI. (A) Parent/WT strain BW25113. (B) *wecD* genomic deletion JW5597-1 strain. The optical density of cultures was measured at 600 nm for 330 minutes.

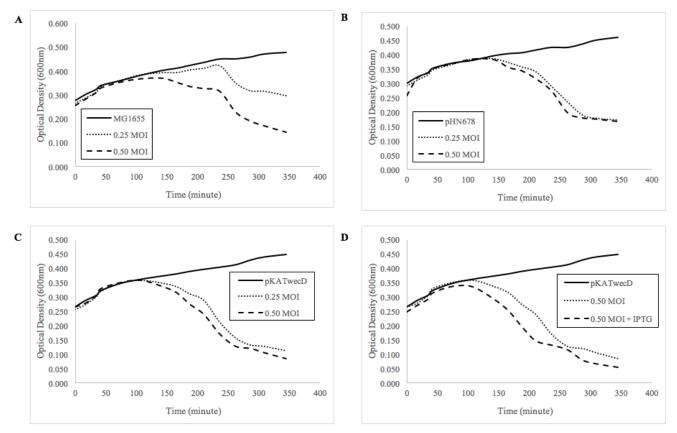


FIG. S3 The effect of *wecD* asRNA knockdown on bacteriophage T7 infectivity at 0.25 and 0.5 MOI. (A) WT MG1655 strain. (B) MG1655 transformed with pHN678 plasmid. (C) MG1655 transformed with pKAT*wecD* plasmid, without IPTG induction (D) MG1655 transformed with pKAT*wecD* plasmid, with 1 mM IPTG induction. The optical density of cultures was measured at 600 nm for 330 minutes.

Sequence of pKATwecD, regions around the wecD antisense wecDas60f sequence NNNNCTGTTGNNNTANTCNTCCGGCTCGTATAATGTGGTGGAATTGTGAGCGGATAACAATTTcagGaggAAtt CAT GGT GGC GCG GAC GGG CAC GAA GTA ACT CCA $AACCATGCAGTGGTGGTGGTGGTGGTGC\underline{CATGGTGgcGCGGACGGAGGAGGAAGTAACTCCAGCAGAGTATTG}\\ \underline{GCCGTAGCCAGTTTTCC}\\ \underline{TCGAGCACCACCACCACCACCACCACCACGGGTGGTTAATTCCTCCTACTAGTTTTGGN}$ GCA GAT ATT GGC CGT AGC CAG TTT TCC GGATGAGAGAAGATTTTCAGCCTGATACAGATTAAATCAGAACGCAGAAGCGGTCTGATAAAACAGAATTT GCCTGGCGGCAGTAGCNGGNNGGTCCCACCTGACCCCATGCCGAACTCAGAAGTGAAACGCCGTAGTGCC GATGGTAGTGTGGGGTCTCCCCATGCGAGAGTAGGGAACTGCCAGGCATCAAATAAAACGAAAGGCTCAG TCGAAAGACTGGGCCTTTCGTTTTATCTGAGGGGGAGGCAGAATAAATGATCATATCGTCAATTATTACCTC CACGGGGAGAGCCTGAGCAAACTGGCCTCAGGCATTTGAGAAGCACACGGTCACACTGCTTCCGGTAGTC AATAAACCGGTAAACCAGCAATAGACATAAGCGGCTATTTAACGACCCTGCCCTGAACCGACGACCGGGTC GAATTTGCTTTCGAATTTCTGCCATTCATCCGCTTATTATCACTTATTCAGGCGTAGCAACCAGGCGTTTAAG CGCCTTGCGTATAATATTTGCCCATCGNGAAANCGGGGGCNANAGTTGTCCATATTNNCCACGTTTAAATCA AACTGNGAAACTCACCCAGGGATTGGCTTGAANCGAAAACATATTCTCATTAACCCTTTAGGGAAATAGNN NGGTTTTCACCGTANACNNCACATCTTGNNAATATATGNGNAGAANTGCNAAATCGNCGTNGNNNTCNTTC AGNAGNNANGANNNGTTNNNGTTTGCTCATGGAAACGGNNGNANNAGGGNGGACNNNNTCCCNNATCN CNNNN Sequence ID: Query_170659 Length: 60 Identities Expect Gaps 60/60(100%) 0/60(0%) Plus/Plus 111 bits(60) 1e-29 Query CACGAAGTAACTCCAGCAGATATTGGCCGTAGCCAGTT Sbjct 1

FIG. S4 NCBI BLAST alignment results of pKATwecD sequences (around the wecD antisense region) and wecDas60f insert sequence.