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Supplemental Material

The ‘SElection End points in Communities of bacTeria’ (SELECT) Method: A Novel Experimental Assay to Facilitate Risk Assessment of Selection for Antimicrobial Resistance in the Environment

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Figure S2. Predicted no effect concentrations for resistance ($PNEC^R$, logged) determined using the SELECT (blue triangle) and qPCR methods (pink circle). Error bars represent the test concentrations directly above and directly below the NOECs used to calculate the $PNEC^R$ s. All SELECT $PNEC^R$ s were determined by taking the no observed effect concentrations and applying an assessment factor of 10. For all qPCR $PNEC^R$ s, the *intI1* gene target is presented. QPCR $PNEC^R$ s for azithromycin, clarithromycin, erythromycin and ciprofloxacin were taken from Stanton *et al.* (2020).

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Figure S5. Logged SELECT predicted no effect concentrations for resistance ($PNEC^R$ s) were determined for four antibiotics using wastewater treatment plant (WWTP) A influent (2018) samples using Iso-sensitest broth or artificial sewage, at 20 °C or 37 °C. Bars represent the test concentrations directly above and below the no observed effect concentrations used to determine the $PNEC^R$.

References

Additional File- Excel Document

Table S1. All primers and gBlocks used in this study for targeting 16S rRNA, *CTX-M* group, *ermF*, *qnrS* and *intI1* genes.

Gene target	Forward primer (5' to 3')	Reverse primer (5' to 3')	Product size (bp)	gBlock sequence and length (bp)	References
16S rRNA	CGGTGAAT ACGTCYC GG	GGWTACC TTGTTACG ACT	142	ACGGTGAATACGTTCCCG GGCCTTGTACACACCGCC CGTCACACCATGGGAGTG GGTTGCAAAGAAGTAGG TAGCTTAACCTTCGGGAG GGCGCTTACCACTTTGTG ATTCATGACTGGGGTGAA GTCGTAACAAGGTAACCG - 144	(Suzuki et al. 2000) (Murray et al. 2018)
<i>CTX-M</i> group	ATGTGCAG YACCAGTA ARGTKATG GC	ATCACKC GGRTCG CCXGGR AT	~300	GATGTGCAGCACCAGTAA AGTGATGGCCGCGGCCG CGGTGCTGAAGAAAAGTG AAAGCGAACCGAATCTGT TAAATCAGCGAGTTGAGA TCAAAAAATCTGACCTTGT TAACTATAATCCGATTGCG GAAAAGCACGTCAATGGG ACGATGTCAGTGGCTGAG CTTAGCGCGCCGCGCT ACAGTACAGCGATAACGT GGCGATGAATAAGCTGAT TGCTCACGTTGGCGCC CGGCTAGCGTCACCGCG TTCGCCCCGACAGCTGGG AGACGAAACGTTCCGTC TCGACCGTACCGAGCCG ACGTTAAACACCGCCATT	(Birkett et al. 2007) (Murray et al. 2018)

				CCGGGCGATCCGCGTGA TA - 338	
<i>ermF</i>	TCTGGGAG GTTCCATT GTCCT	ACTTTCAG GACCTAC CTCATAGA		TCTGATGCCCGAAA TGTTCAAGTTGTCC GTTGTGATTTTAGG AATTTTGCAGTTCC GAATTTCTTTCAA AGTGGTGTCAAATA TTCTTATGGCATT CTTCCGATATTTTC AAAATCTGATGTTT GAGAGTCTTGGAA ATTTTCTGGGAGGT TCCATTGTCCTTCA ATTAGAACCTACAC AAAAGTTATTTTCGA GGAAGCTTTACAAT CCATATACCGTTTT CTATCATACTTTTT TTGATTTGAACTT GTCTATGAGGTAG GTCCTGAAAGTTT CTTGCCACCGCCA -294	(Stanton et al. 2020)
<i>qnrS</i>	CGACGTGC TAACTTGC GTGA	GGCATTGT TGGAACT TGCA		TTCGACGTGCTAACT TGCGTGATACGACAT TCGTCAACTGCAAG TTCATTGAACAGGGT GATATCGAAGGCTG CCACTTTGATGTCCG CAGATCTTCGTGAT GCAAGTTTCCAACA ATGCCAACTT - 125	(Colomer- Lluch et al. 2014) (Stanton et al. 2020)

<i>intl1</i>	GCCTTGAT GTTACCCG AGAG	GATCGGT CGAATGC GTGT		GGCCTTGATGTTACCC GAGAGCTTGGCACCC AGCCTGCGCGAGCAG CTGTGCGGTGCACGG GCATGGTGGCTGAAG GACCAGGCCGAGGG CCGCAGCGGCGTTG CGCTTCCCGACGCC CTTGAGCGGAAGTA TCCGCGCGCCGGG CATTCTGGCCGTG GTTCTGGGTTTTTG CGCAGCACACGCA TTCGACCGATCC - 198	(Barraud et al. 2010) (Stanton et al. 2020)
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For Excel Tables S1, S2, S3 & S4 see attached excel sheet.

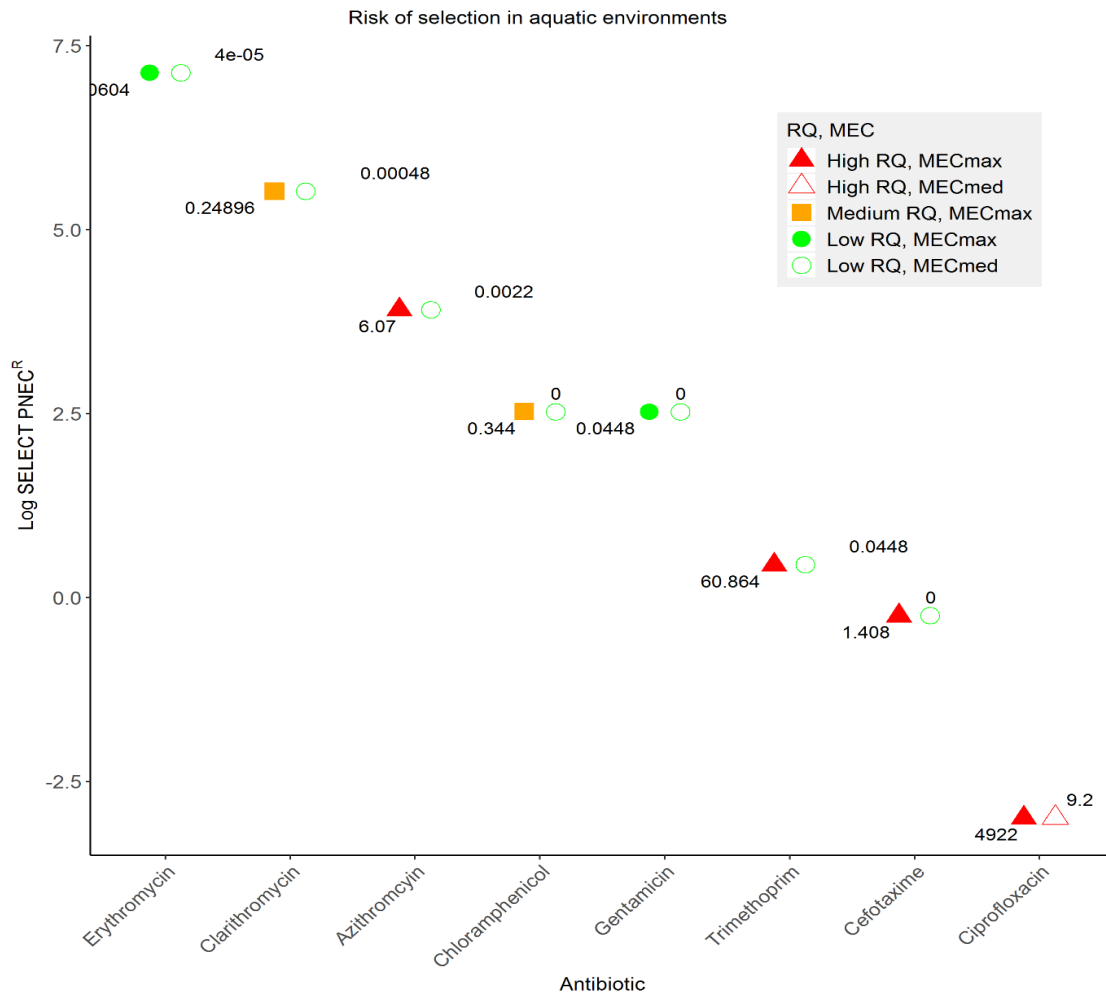


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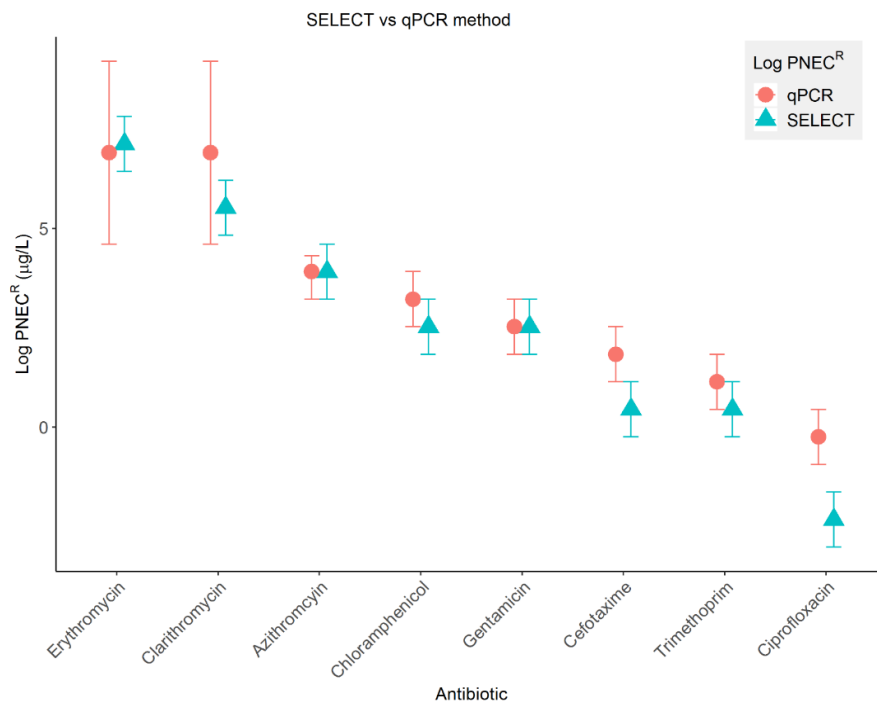


Figure S2. Predicted no effect concentrations for resistance (PNEC^Rs, logged) determined using the SELECT (blue triangle) and qPCR methods (pink circle). Error bars represent the test concentrations directly above and directly below the NOECs used to calculate the PNEC^Rs. All SELECT PNEC^Rs were determined by taking the no observed effect concentrations and applying an assessment factor of 10. For all qPCR PNEC^Rs, the *intI1* gene target is presented. QPCR PNEC^Rs for azithromycin, clarithromycin, erythromycin and ciprofloxacin were taken from Stanton *et al.* (2020).

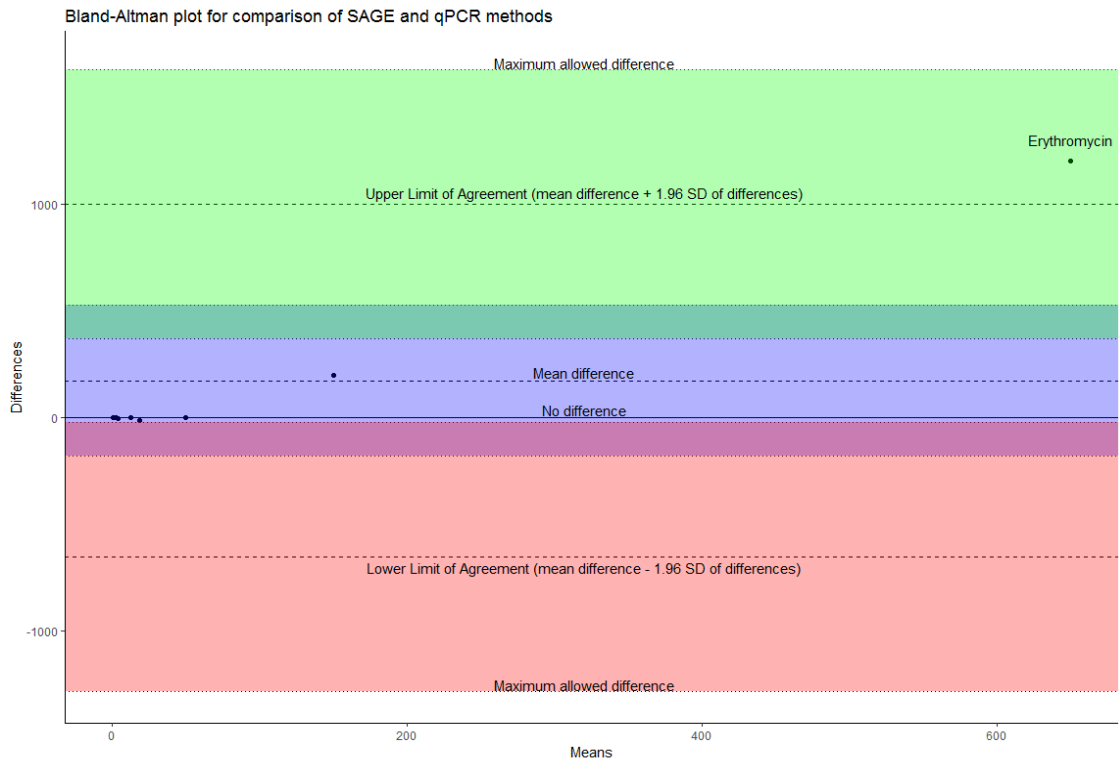


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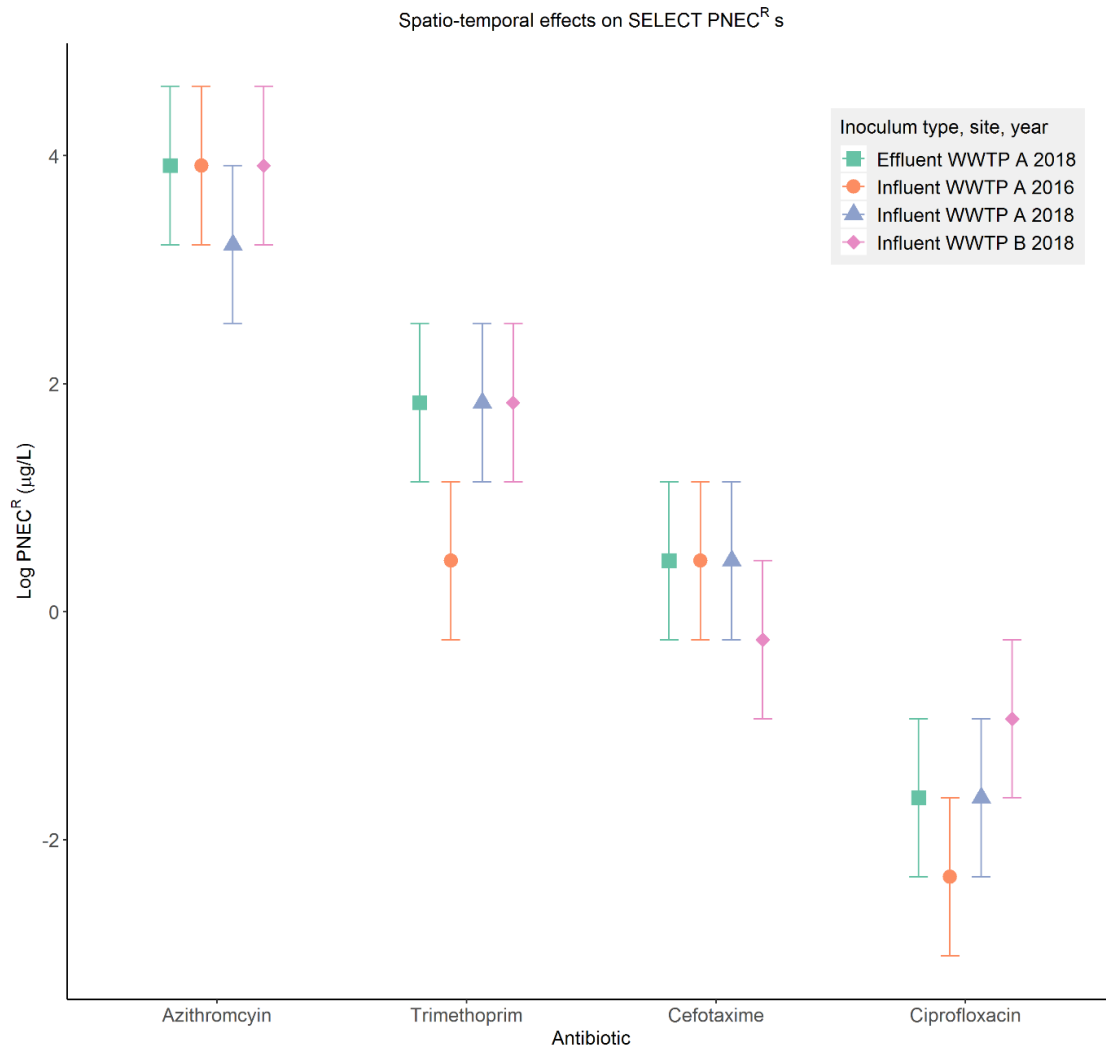


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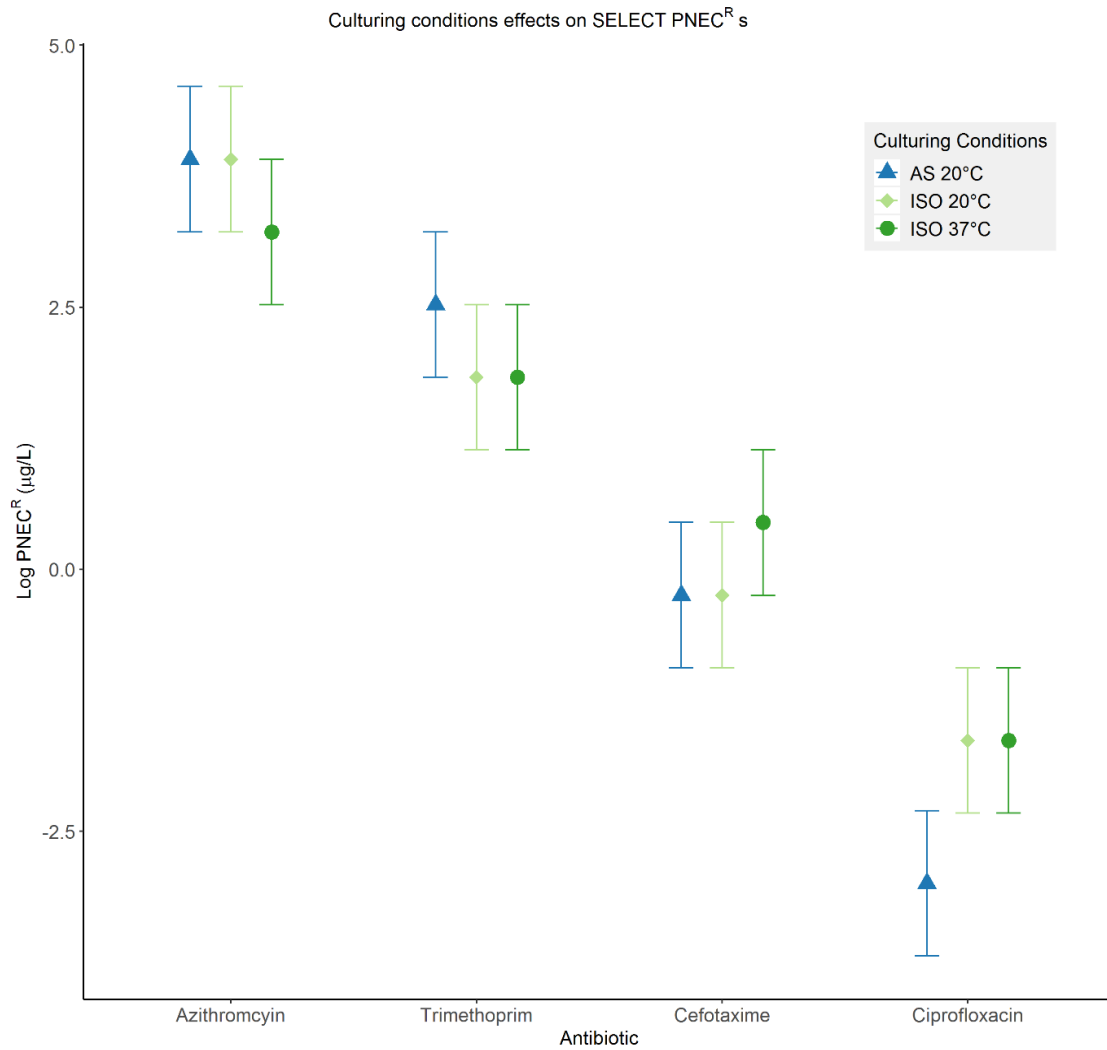


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