

Effect of stepwise exposure to high-level erythromycin on anaerobic digestion

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Table S1. qPCR primer sequences for antibiotic resistance genes analysis.

Resistance mechanism	gene	5' to 3'	Reference
esterase gene	<i>ereA</i>	F: CCTGTGGTACGGAGAACATTGATGT R: ACCGCATTCGCTTGCTT	
	<i>ereB</i>	F: GCTTTATTCAGGAGGCAGGAAT R: TTTAAATGCCACAGCACAGAAC	
phosphorylase gene	<i>mphA</i>	F: CTGACGCGCTCCGTGTT R: GGTGGTGCATGGCGATCT	
	<i>-01</i>	F: CGCAGCGCTTGATCTTAG R: TTACTGCATCCATACGCTGCTT	
methylase gene	<i>mphB</i>	F: TTGAGAAGGGATTGCGAAAAG R:	
	<i>ermA</i>	ATATCCATCTCCACCATTAAATAGTAAACC F: TAAAGGGCATTAAACGACGAAA	[1-3]
efflux pump gene	<i>ermB</i>	R: TTTATACCTCTGTTGTTAGGGATTGAA	
	<i>ermC</i>	F: TTTGAAATCGGCTCAGGAAAA R: ATGGTCTATTCAATGGCAGTTACG	
	<i>ermX</i>	F: GCTCAGTGGTCCCCATGGT R: ATCCCCCGTCAACGTTT	
	<i>mefA</i>	F: CCGTAGCATTGGAACAGCTTT R: AACGGAGTATAAGAGTGCTGCAA	
	<i>mefE</i>	F: CGTATTGGGTGCTGTGATTG R: TATGCACAGGCCTTCCATTA	

References

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