Supplementary Table S1. Primers used in this study

Bacteria	Primer	Sequence (5′-3′)	Size	Reference
E. coli	16S F	GTTAATACCTTTGCTCATTGA	340	[45]
	16S R	ACCAGGGTATCTAATCCTGTT		
	ChuA-F	GACGAACCAACGGTCAGGAT	279	
	ChuA-R	TGCCGCCAGTACCAAAGACA		
	YjaA-F	TGAAGTGTCAGGAGACGCTG	211	[46]
	YjaA-R	ATGGAGAATGCGTTCCTCAAC		
	TspE4C2.1-F	GAGTAATGTCGGGGCATTCA	152	
	TspE4C2.1-R	CGCGCCAACAAAGTATTACG		
	clbB-F	GCGCATCCTCAAGAGTAAATA	280	[25]
	clbB-R	GCGCTCTATGCTCATCAACC		
	CNF-1s	GGGGGAAGTACAGAAGAATTA	1112	[27]
	CNF-1as	TTGCCGTCCACTCTCACCAGT		
	CDT-s1	GAAAGTAAATGGAATATAAATGTCCG	555	[47]
	CDT-IIIasa	TTTGTGTCGGTGCAGCAGGGAAAA		
	cif-int-s	AACAGATGGCAACAGACTGG	383	[13]
	cif-int-as	AGTCAATGCTTTATGCGTCAT		
E. coli-EPEC	eae-F	TCAATGCAGTTCCGTTATCAGTT	482	[48]
	eae-R	GTA AAGTCCGTTACCCCAACCTG		
	bfp-F	GGA AGTCAAATTCATGGGGGTAT	300	
	bfp-R	GGAATCAGACGCAGACTGGTAGT		
E. coli-DAEC	daaE -F	GAACGT TGGTTAATGTGGGGTAA	542	[48]
	daaE -R	TATTCACCGGTCGGTTATCAGT		
Klebsiella	16S F	ACTCCTACGGGAGGCAGCAGT	383	[49]
	16S R	TATTACCGCGGCTGCTGGC		
	clbB-F	GCGCATCCTCAAGAGTAAATA	280	
	clbB-R	GCGCTCTATGCTCATCAACC		
Enterococcus faecalis	16S F	CCCTTATTGTTAGTTGCCATCATT	144	[50]
	16S R	ACTCGTTGTACTTCCCATTGT		

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