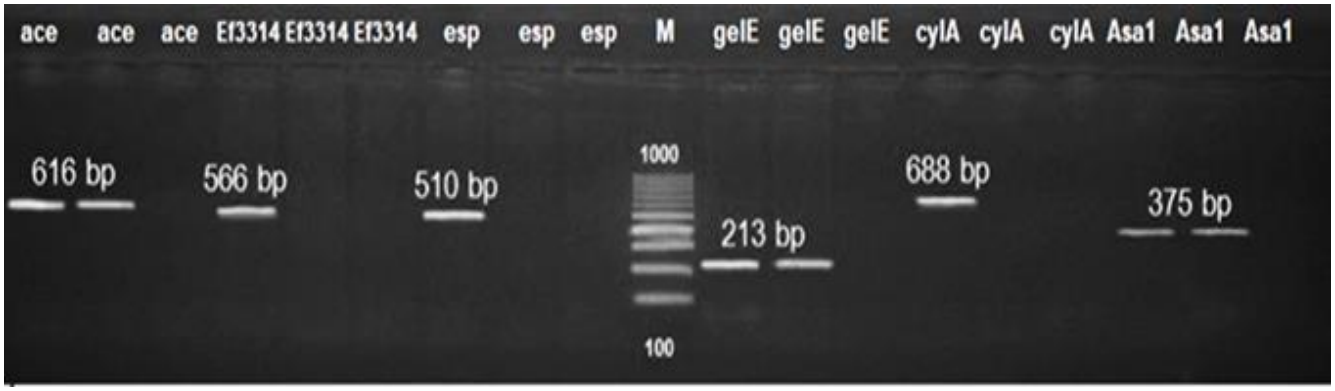
Supplementary Table S1. PCR primers and products for the detection of *E. faecalis* virulence genes.

Agent	Virulence factor	Oligonucleotide primers sequences (5'-3')	Product size (bp)	Reference
Enterococcus faecalis	*esp*	AGATTTCATCTTTGATTCTTGG	510 bp	**Vankerckhoven *et al.*, 2004**
		AATTGATTCTTTAGCATCTGG		
	gelE	TATGACAATGCTTTTTGGGAT	213 bp	
		AGATGCACCCGAAATAATATA		
	asa1	GCACGCTATTACGAACTATGA	375 bp	
		TAAGAAAGAACATCACCACGA		
	cylA	ACTCGGGGATTGATAGGC	688 bp	
		GCTGCTAAAGCTGCGCTT		
	ace	GGAATGACCGAGAACGATGGC	616 bp	**Creti *et al.*, 2004**
		GCTTGATGTTGGCCTGCTTCCG		
	EF3314	AGAGGGACGATCAGATGAAAAA	566 bp	
		ATTCCAATTGACGATTCACTTC		

Supplementary Table S2. Host range of *E. faecalis* bacteriophages

Bacterial sp.	Isolates sources	ϕZEF1	ϕZEF2
E. faecalis1	**Isolated by this study**	**+**	+
E. faecalis2	**Isolated by this study**	**+**	+
E .faecalis3	**Isolated by this study**	**-**	-
E. faecalis4	**Isolated by this study**	**+**	+
E .faecalis5	**Isolated by this study**	**+**	+
E .faecalis6	**Isolated by this study**	**-**	-
E. faecalis7	**Isolated by this study**	**+**	+
E. faecalis8	**Isolated by this study**	**+**	+
E. faecalis9	**Isolated by this study**	**+**	+
E. faecalis10	**Isolated by this study**	**+**	+
E. faecalis11	**Isolated by this study**	**-**	-
E. faecalis12	**Isolated by this study**	**+**	+
E. faecalis13	**Isolated by this study**	**+**	+
E. faecium	**Animal Health Research Institute, Dokki, Egypt.**	**-**	-
S. mutans	**Cairo MIRCEN reference (Gene bank accession number : Z95910)**	**-**	-
E. gallinarum	**Animal Health Research Institute, Dokki, Egypt.**	**-**	-
S. aureus	**Faculty of Science, Zagazig University, Egypt (Gene Bank accession number: KR270348).**	**-**	-
E. coli	**Department of Zoonoses, Faculty of Veterinary Medicine, Zagazig University.**	**-**	-
P. aeruginosa	**Faculty of Science, Zagazig University, Egypt (Gene Bank under accession number: LC514698).**	**-**	-



Supplementary Figure S1. Gel electrophoresis of PCR products of virulence genes amplified from *Enterococcus* isolate No. 4. PCR amplification of ace, EF3314, esp, gelE, cylA and asa1 genes genes at 616, 566, 510, 213, 688 and 375 bp, respectively. Presented from left to right: ace gene (lane 1, positive control, lane 2, positive isolate, lane 3, negative control); EF3314 gene (lane 4, positive control, lane 5, negative isolate, lane 6, negative control), esp gene (lane 7, positive control, lane 8 negative isolate, lane 9, negative control); lane 10, (M) 100 bp DNA Ladder; gelE gene (lane 11, positive control, lane 12, positive isolate, lane 13, negative control); cylA gene (Lane 14, positive control, lane 15, negative isolate, lane 16, negative control); asa1 gene (lane 17, positive control, lane 18, positive isolate, lane 19, negative control).