**Table S1** Quantification of the PG molecular species distribution found in the PG of strains grownin the thigh model. Multiple spectra were obtained using the methods outlined in Fig. 3, thepercentage of each peak was calculated and the percent distribution determined.

	% Total PG (Mean ± SD)					
PG Species	AH1263 (WT) Luria broth	AH1263 (WT) Thigh	JLB2 (∆ <i>fakA</i> ) Thigh	JLB31 (∆ <i>fakB1 ∆fakB</i> 2) Thigh	JLB27 (∆ <i>fakB1</i> ) Thigh	JLB30 (∆ <i>fakB2</i> ) Thigh
30:0-PG	21.71 ± 2.40	3.16 ± 0.61	15.24 ± 1.58	13.60 ± 1.31	3.11 ± 0.51	2.97 ± 0.18
31:0-PG	0.64 ± 0.11	19.29 ± 1.44	13.55 ± 0.34	11.90 ± 0.95	7.22 ± 0.80	25.66 ± 1.92
32:0-PG	55.29 ± 2.52	4.91 ± 0.64	19.77 ± 2.00	21.11 ± 2.06	5.91 ± 1.48	5.54 ± 0.32
33:1-PG	ND <sup>a</sup>	21.58 ± 1.26	ND	ND	26.78 ± 0.92	ND
33:0-PG	7.70 ± 0.66	23.46 ± 0.85	24.32 ± 0.97	24.52 ± 1.32	15.19 ± 1.10	37.02 ± 1.41
34:0-PG	11.58 ± 0.90	3.86 ± 1.32	9.95 ± 2.75	13.25 ± 2.43	8.36 ± 0.93	$3.60 \pm 0.36$
35:1-PG	ND	9.33 ± 0.91	ND	ND	21.76 ± 1.06	ND
35:0-PG	$3.08 \pm 0.43$	14.41 ± 0.98	17.16 ± 2.49	15.63 ± .91	11.67 ± 0.95	26.64 ± 1.27

<sup>a</sup>ND means not detected