1 SUPPLEMENTAL MATERIAL

- 2
- 3

Table S1. Goodness of fit of the Gompertz's model and magnitude of in vivo growth parameters for diverse strains of Enterococci under different incubation conditions. The following modified Gompertz's equation was used:

 $y = B + A * e^{-e^{(2.781*(\frac{A}{5})*(0.04-x)+1)}}$; where y is the bacterial count (in log₁₀ CFU/g) obtained at time x (in hours), B (basal) is the 6 initial number of bacteria after induction of infection, A (from Auctus, the Latin word for "growth") is the difference between the 7 maximal and minimal predicted bacterial count, and S (slope) is the growth rate. Model fitting (SigmaPlot v12.3, Systat software 8 inc., USA) was evaluated by the adjusted coefficient of determination (adj. R^2), the standard error of the estimate ($S_{v|x}$), normality 9 (Shapiro-Wilk test), homoscedasticity (constant variance test) and non-multicollinearity (defined as a variance inflation factor 10 <10). Comparison of the magnitude of the parameters between experimental groups was done by curve fitting analysis (CFA; 11 Prism 6.04) [23, 24]. Other abbreviation used in the table are: IIS for in vivo inoculum size (log₁₀ CFU/g) measured in non-12 treated mice at 0 h post-infection; NT and CV for normality and constant variance in terms of yes (passed) or no (failed); VIF: 13 highest variance inflation factor reported; ΔG_{26-2} is the net growth (mean) in non-treated mice; IVP for invalid parameter because 14 NLR failed any of the presumptions normality or homoscedasticity; ** when it was not possible to fit Gompertz' equation to data. 15 the closest equation was exponential decay (parameters not shown). Normal status means that non-neutropenic mice were 16 used. All the parameters are presented with their respective standard errors (±SE). 17

Strains	Conditions	Status	IIS	Adj.R ²	S _{y x}	NT	CV	VIF	Basal	Slope	Auctus	ΔG_{26-2}
<i>E. faecalis</i> ATCC 29212	Aerobiosis	Neutropenic	5	0.55	0.66	No	No	6	IVP	IVP	IVP	0.01
	Aerobiosis+mucin	Neutropenic	5	0.95	0.29	Yes	No	7	IVP	IVP	IVP	2.07
	Anaerobiosis	Neutropenic	7	0.96	0.14	Yes	No	6	IVP	IVP	IVP	0.84
		Neutropenic	6	0.72	0.52	No	No	4	IVP	IVP	IVP	0.91
		Neutropenic	5	0.65	0.52	No	No	6	IVP	IVP	IVP	0.46
		Normal	4	**	**	**	**	**	**	**	**	**
	Anaerobiosis+mucin	Neutropenic	5	0.99	0.20	Yes	Yes	4	4.81 ± 0.06	0.62 ± 0.03	4.04 ± 0.07	2.64
		Neutropenic	4	0.99	0.13	Yes	Yes	4	3.86 ± 0.09	0.68 ± 0.07	4.94 ± 0.11	3.64
		Normal	4	0.99	0.10	Yes	Yes	2	3.93 ± 0.07	0.74 ± 0.05	4.09 ± 0.09	2.38
E. faecium ATCC 19434	Aerobiosis	Neutropenic	6	0.57	0.35	Yes	No	4	IVP	IVP	IVP	0.19
	Aerobiosis+mucin	Neutropenic	6	0.97	0.21	Yes	Yes	4	5.94 ± 0.08	0.27 ± 0.02	3.26 ± 0.11	2.54
	Anaerobiosis	Neutropenic	6	0.71	0.52	Yes	No	4	IVP	IVP	IVP	0.66
	Anaerobiosis+mucin	Neutropenic	6	0.98	0.18	Yes	Yes	3	6.17 ± 0.06	0.30 ± 0.02	3.12 ± 0.09	2.40
		Neutropenic	7	0.94	0.23	Yes	Yes	5	7.00 ± 0.06	0.37 ± 0.10	2.25 ± 0.19	1.62
<i>E. faecalis</i> ATCC 51299	Anaerobiosis+mucin	Neutropenic	5	0.94	0.31	Yes	Yes	4	4.73 ± 0.16	0.73 ± 0.07	3.04 ± 0.19	1.65
<i>E. faecium</i> ATCC 51559	Anaerobiosis+mucin	Neutropenic	5	0.99	0.10	Yes	Yes	4	5.66 ± 0.06	0.66 ± 0.04	3.09 ± 0.07	1.72

Figure S1. Dissemination of *E. faecium* ATCC 19434 to bloodstream and spleen in neutropenic animals 26 hours after thigh infection with inocula prepared using anaerobiosis+mucin (black bar, N=7 mice), aerobiosis+mucin (light gray bar, N=8) or aerobiosis (dark gray bar, N=7). Data are representative of two independent experiments. The P value of each post-hoc comparison (Dunn for Kruskal-Wallis and Holm-Sidak after ANOVA) is represented above the bars by asterisks (*= P≤0.05, **= P≤0.01 and ***= P≤0.001) or by NS (P>0.05). In spleen, the anaerobiosis+mucin group had higher dissemination than aerobiosis (Δ +1.8 log₁₀ CFU/g, P<0.001) and aerobiosis+mucin (Δ +0.9 log₁₀ CFU/g, P=0.022). In blood, the anaerobiosis+mucin group again was associated with greater dissemination than aerobiosis (Δ +1.6 log₁₀ CFU/g, P≤0.01) and aerobiosis+mucin (Δ +0.1 log₁₀ CFU/g, P≤0.01). Bacteremia was demonstrated in 14% (1 of 7), 63% (5 of 8), and 100% (7 of 7) of the mice allocated to aerobiosis, aerobiosis+mucin and anaerobiosis+mucin, respectively.

