Supporting Information

Table S1.	Phospholipid	constituents	of the	model	Staphylococcus	Aureus	(left	column)
together wi	ith their assigned	ed 4-letter coo	des (rig	ht colun	nn). ¹			

Lipids	Lipid code
PG (a15:0/a15:0)	FFPG
PG (i15:0/a15:0)	IFPG
PG (16:0/a15:0)	PFPG
PG (i17:0/a15:0)	JFPG
PG (a17:0/a15:0)	ZFPG
PG (i17:0/i15:0)	JIPG
PG (a17:0/i15:0)	ZIPG
PG (18:0/a15:0)	SFPG
PG (i19:0/a15:0)	VFPG
PG (a19:0/a15:0)	TFPG
PG (a19:0/i15:0)	TIPG
PG (20:0/a15:0)	XFPG
PG (18:1/i15:0)	OIPG
PG (18:1/a15:0)	OFPG
LPG (i17:0/a15:0)	JFGK
LPG (a17:0/a15:0)	ZFGK
LPG (18:0/a15:0)	SFGK
LPG (a19:0/a15:0)	TFGK
CL (a17:0/a15:0)	ZFCL

Table S2. Letter Codes representing Fatty Acyl Tails which are not present in CHARMM-GUI Membrane Builder module.¹

-	Lipid tail	Code	Explanation
	a15:0	F	anteisopentadecanoyl, Fifteen for pentadecane
	i15:0	Ι	Isopentadecanoyl
	a17:0	Ζ	anteisomargaroyl, Z for g
	i17:0	J	isomargaroyl, J for g
	a19:0	Т	anTeisononadecanoyl
	i19:0	V	isononadecanoyl, V is after T and resembles an iso branch
	20:0	Х	arachidoyl, X for ch

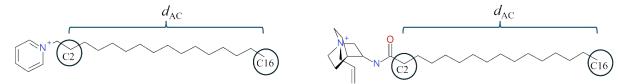


Figure S1. CPC (left) and QC₁₆-Ally molecules. Atoms used to define the distance d_{AC} (distance between the C2 and C16 atoms in the aliphatic chain) are denoted with the blue ellipses.

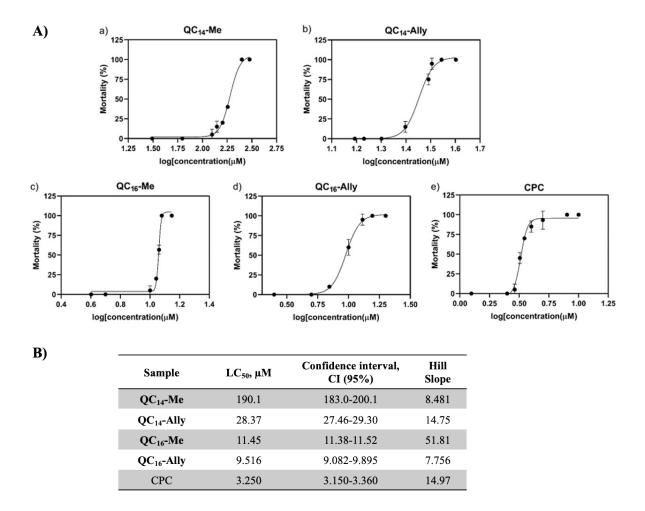


Figure S2. A) Concentration-response curves used for the calculation of the *Danio rerio* mortality after 120 hours of exposure to: a) QC_{14} -Me, b) QC_{14} -Ally, c) QC_{16} -Me, d) QC_{16} - Ally and e) CPC. B) Dose-inhibition results obtained after 120 hours of *Danio rerio* exposure to tested samples.

¹F. Joodaki, L. M. Martin, and M. L. Greenfield, Generation and Computational Characterization of a Complex Staphylococcus aureus Lipid Bilayer, Langmuir 38 (2022), pp. 9481-9499