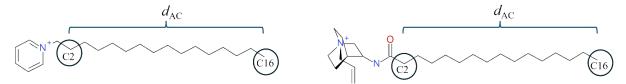
## **Supporting Information**

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). <sup>1</sup>			

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.<sup>1</sup>

-	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, <b>J</b> 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 QC<sub>16</sub>-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.

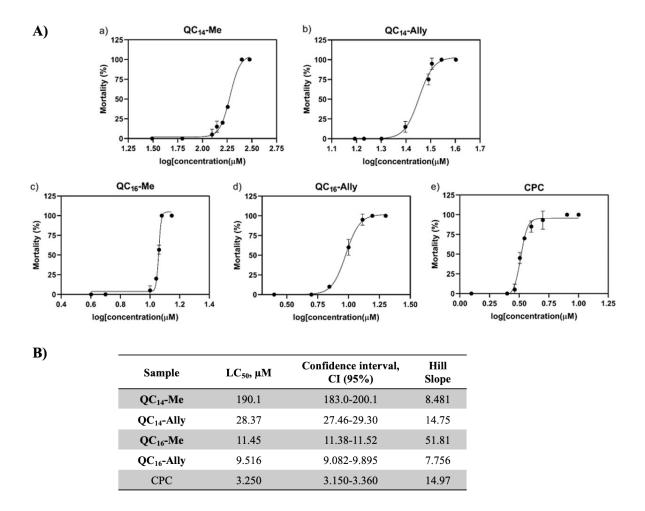


Figure S2. A) Concentration-response curves used for the calculation of the *Danio rerio* mortality after 120 hours of exposure to: a)  $QC_{14}$ -Me, b)  $QC_{14}$ -Ally, c)  $QC_{16}$ -Me, d)  $QC_{16}$ - Ally and e) CPC. B) Dose-inhibition results obtained after 120 hours of *Danio rerio* exposure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

<sup>&</sup>lt;sup>1</sup>F. 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