Supporting Information

Bacterial Aggregation Triggered by Fibril Forming Tryptophan-Rich Sequences: Effects of Peptide Side Chain and Membrane Phospholipids

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Figure S1. Changes of the bacterial growth turbidity upon incubation with the 14 days old HHC nanofibrils. The experiments were performed once in triplicate.

Figures S2–S9. Representative TEM and FE-SEM micrographs of control *S. aureus* (ATCC 12600) and the cells exposed to HHC-10 and ^{4Har}HHC-10 at ½ MIC and 2 MIC, as well as the bacteria treated with the peptide nanofibrils. The bacterial aggregates are marked by orange arrows. Yellow arrows highlight nanostructure of different morphologies including but not limited to peptide aggregates formed over the course of bacterial aggregation. Membrane disruption (deformation) and degradation of cell wall septum are, respectively, presented by green and cyan arrows. The colorful square insets were magnified in the images outlined by the same color.

Figures S10–S17. Representative TEM and FE-SEM micrographs of control *E. coli* (ATCC 10536) and cells exposed to HHC-10 and ^{4Har}HHC-10 at ¹/₂ MIC and 2 MIC, as well as the bacteria treated with the peptide nanofibrils. Orange arrows highlight the aggregates of bacteria. Yellow arrows point to nanostructure of different morphologies including but not limited to peptide aggregates formed over the course of the cell aggregation. Pink arrows mark the inside cell protein aggregates that likely result from the interruption of bacterial proteostasis. Membrane buds (red), blebs (blue), thinning (cyan), and disruption and separation of bacterial outer and cytoplasmic membranes (green) are presented by the noted colors. The colorful square insets were magnified in the images outlined by the same color.

Figure S18. Fluorescence emission spectra for the membrane (SUVs) binding of HHC peptides.

Figure S19. CD spectra of ^{4Har}HHC-10 upon interaction with the SUVs. The peptide concentration was 200 μ M at $c_P/c_L = 0.04$, where the optical artifacts due to the liposomal particulates were minimal. The peptide and lipid were incubated at 37 °C for 30 min prior to the measurement.

Figure S20. FRET emission spectra for the interaction of POPE/POPG [3/1 (mol/mol)] (A) single/double labelled LUVs and (B) single labelled SUVs with HHC-10 and ^{4Har}HHC-10 (λ_{ex} = 460 nm). The spectra were registered for the fluorescence emission of NBD at 480–650 nm and Rhod at 580–650 nm.

Figures S21–S29. FE-SEM and TEM of ultrastructure images of anionic SUVs alone and in the presence of HHC peptides ($c_P/c_L = 0.08$). Red arrows highlight the exemplar free (and/or control) SUVs of different phospholipid composition. Yellow arrows point to the aggregates of lipid vesicles (SUVs) in complexation with the HHC peptides. The micrographs show different morphologies and aggregate nanostructures of POPC/POPG liposomes in comparison with those made of POPE/POPG [3/1 (mol/mol)].





S. aureus Control



HHC-10 @ 1/2 MIC



HHC-10 @ 2 MIC



HHC-10 @ 128 xM



^{4Har}HHC-10 @ ¹/₂ MIC



^{4Har}HHC-10 @ 2 MIC



$^{4\text{Har}}\text{HHC-10}$ @ 128 ${}^{\text{\tiny \mbox{M}}}\text{M}$





HHC-10 nanofibers ^{4Har}HHC-10 nanofibers



E. coli Control



HHC-10 @ 1/2 MIC



HHC-10 @ 2 MIC



HHC-10 @ 128 \propto M



^{4Har}HHC-10 @ ¹/₂ MIC



^{4Har}HHC-10 @ 2 MIC



$^{4\text{Har}}\text{HHC-10}$ @ 128 ${\propto}\text{M}$





HHC-10 nanofibers ^{4Har}HHC-10 nanofibers





Figure S19





SEM HV: 7.00 kV View field: 6.191 μm SEM MAG: 35.00 kx WD: 9.863 m Det: SE _____ 1 μm

POPE/POPG [3/1 (mol/mol)]



View field: 5.417 µm Det: SE SEM MAG: 40.00 kx

POPC/POPG [1/3 (mol/mol)]



HHC-10:POPE/POPG [3/1 (mol/mol)]



^{4Har}HHC-10:POPE/POPG [3/1 (mol/mol)]



^{4Har}HHC-10:POPC/POPG [1/3 (mol/mol)]



POPE/POPG [3/1 (mol/mol)]



POPC/POPG [3/1 (mol/mol)]



^{4Har}HHC-10:POPE/POPG [3/1 (mol/mol)]



^{4Har}HHC-10:POPC/POPG [3/1 (mol/mol)]