

Figure S1. Distribution of tilt angles (τ), azimuthal rotation angles (ρ), and depths of insertion sampled by the N- (blue) or C- (orange) terminal residues of P1 (left) or P3 (right) in POPC/POPG during the 2- μ s MD simulations. Curves for *apo* and metalated piscidins are shown as dashed and solid lines, respectively.



Figure S2. Distribution of tilt angles, azimuthal rotation angles, and depths of insertion sampled by the N- (blue) or C- (orange) terminal residues of P1 (left) or P3 (right) in POPC/POPG/aldo-PC during the 2-µs MD simulations. Curves for *apo* and metalated piscidins are shown as dashed and solid lines, respectively.



Figure S3. Average hydrogen bonds per frame between PC and PG headgroups in (A) POPC/POPG and (B) POPC/POPG/aldo-PC bilayer.



Figure S4. Average hydrogen bonds per frame between piscidin and lipid headgroup constituents in (A) POPC/POPG and (B) POPC/POPG/aldo-PC bilayer.

Density functional theory calculations

Density functional theory (DFT) calculations were performed on the P1 ATCUN motif using Gaussian09¹ and the M06-2X² exchange correlation functional. Snapshots extracted from MD trajectories were stripped of solvent, truncated to Ile5, and capped with a methylamine group. Nickel was represented with

the Wachters-Hay all-electron basis set with all other atoms using the TZVP basis set.^{3,4} Geometry optimizations were performed using the integral equation formalism polarizable continuum model (IEF-PCM) with heptane as the solvent.⁵



Figure S5. DFT-optimized structures of representative snapshots of Phe1- and Phe2-Ni²⁺ close contacts. Relative energies are calculated with respect to C. (A) Phe1 forms a short interaction consistent with a cation- π interaction between Phe1 and the metal ion (+3.7 kcal/mol). The longer close contact between Phe2 and the metal ion in B is higher in energy (+12.9 kcal/mol) than the conformation in which the Phe2-Ni²⁺ close contact is supported by a CH- π interaction between Phe2 and Phe1 (0.0 kcal/mol).

References

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