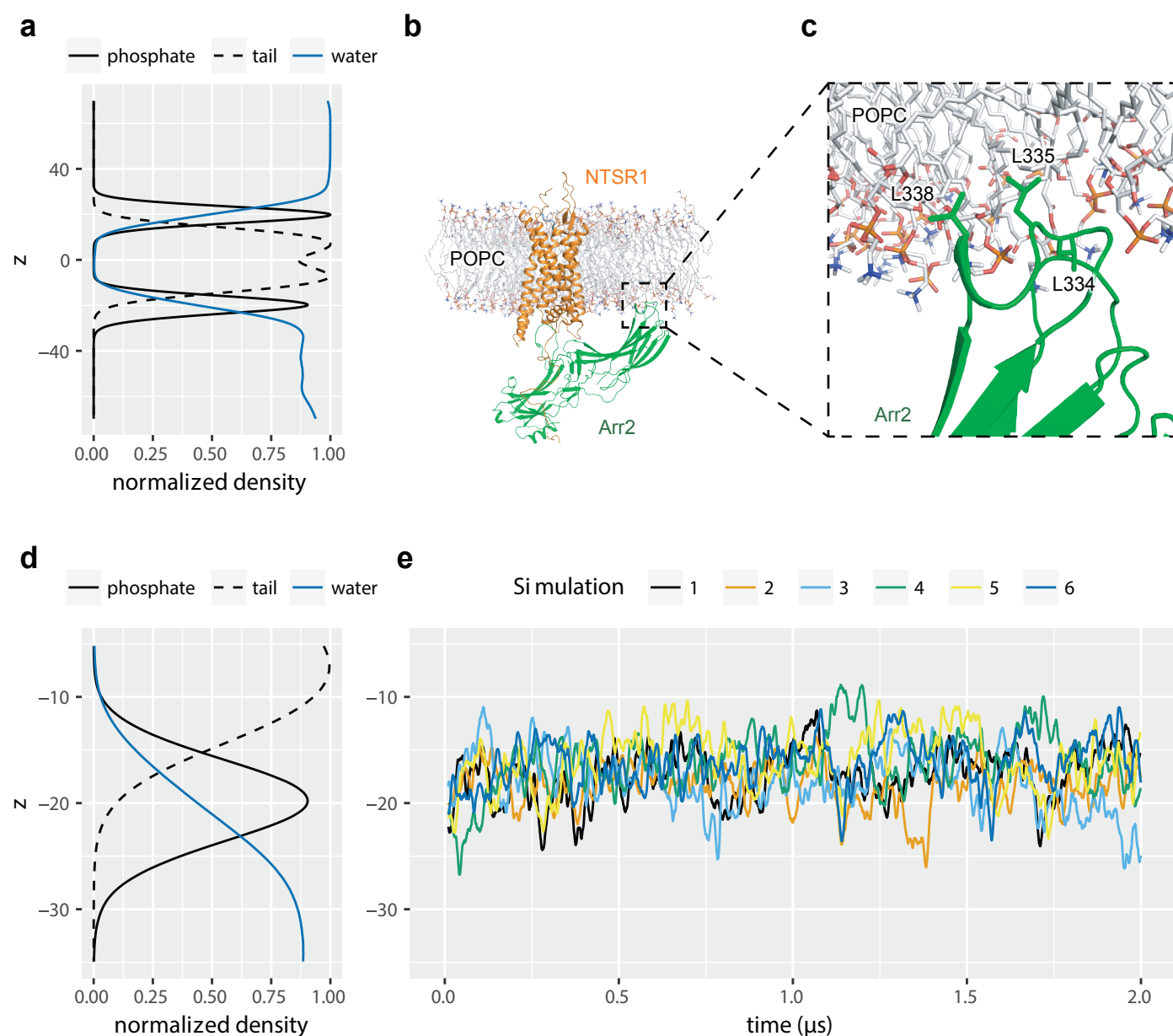


Supplementary information, Fig. S4



Supplementary information, Fig. S4| C-edge loop (L334 through L338) of Arr2 associates with the lipid bilayer

a-b, Normalized lipid and water solvent density and aligned system snapshot taken from simulation 1. c, Three leucine residues (L334, L335 and L338) on the arrestin C-edge loop provide a hydrophobic anchor in lipid bilayer. d, Zoomed-in normalized system density near the anchoring C-edge of arrestin. e, Average z-axis coordinates of heavy atoms of L334, L335 and L338 shown in (c).