

**Supplementary Information for:**

**Soluble Regions of GlpG Influence Protein-Lipid Interactions and Lipid Distribution**

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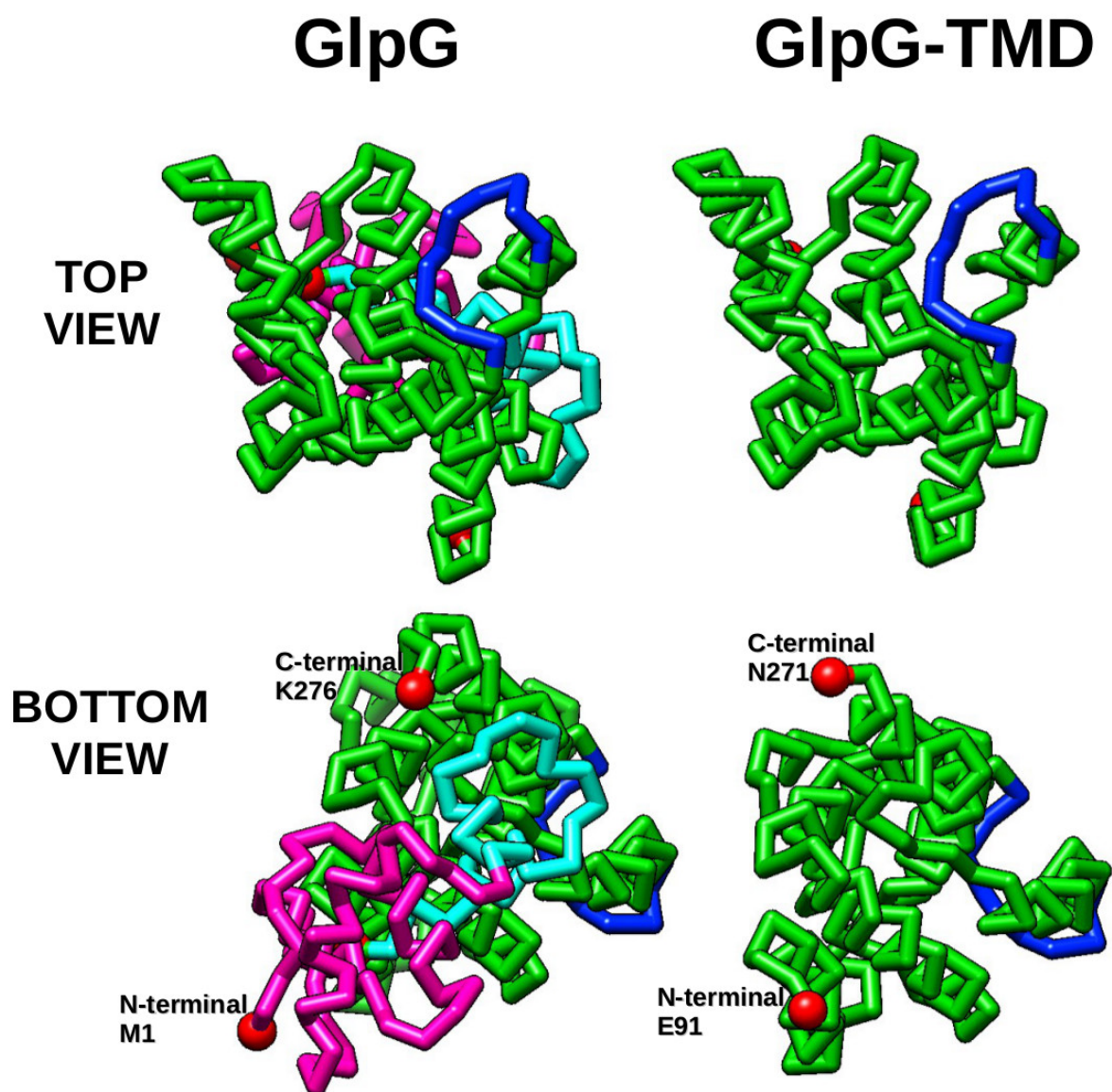
e-mail: [tidow@chemie.uni-hamburg.de](mailto:tidow@chemie.uni-hamburg.de)

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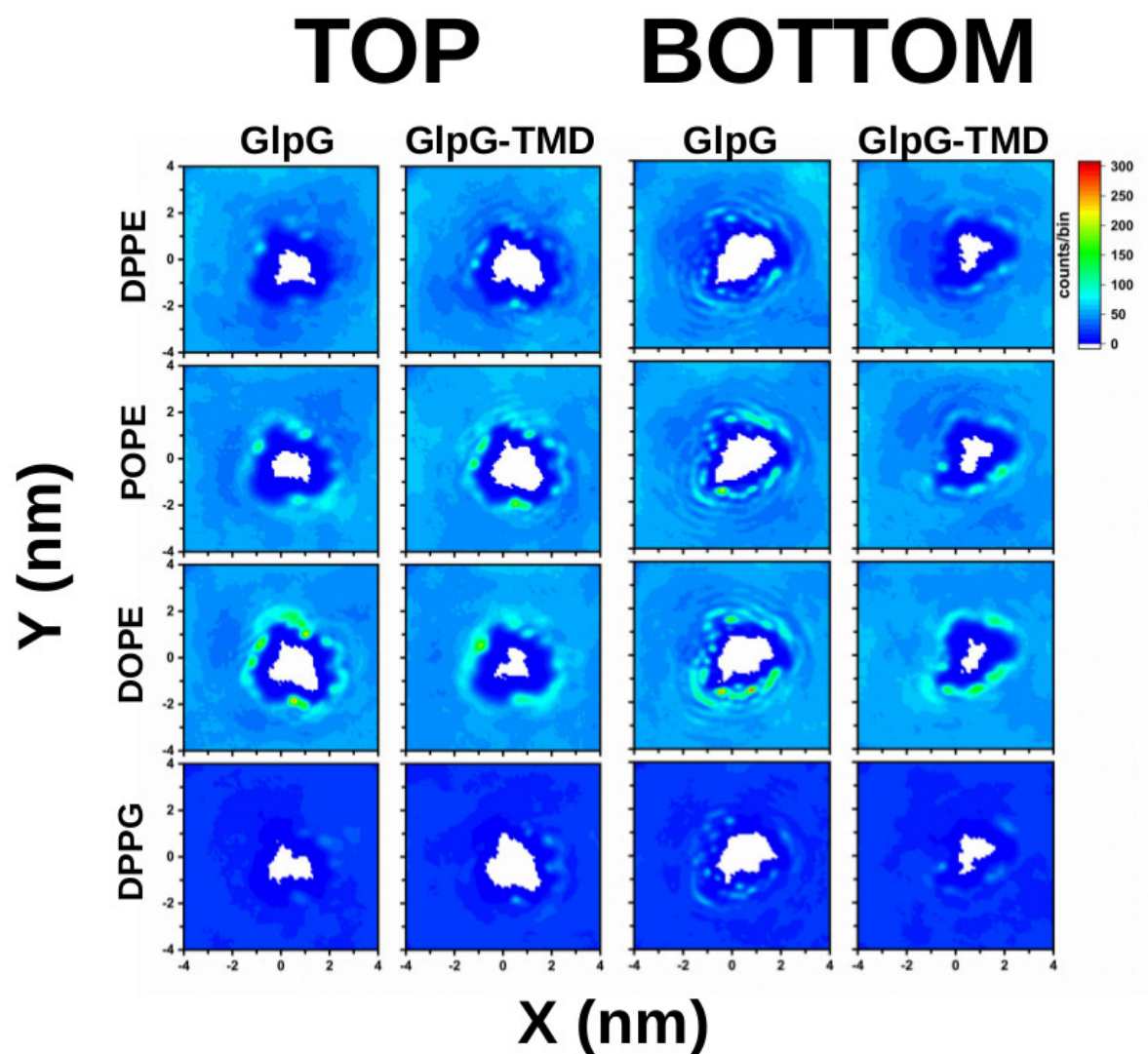
e-mail: [yasser.almeida.hernandez@chemie.uni-hamburg.de](mailto:yasser.almeida.hernandez@chemie.uni-hamburg.de)

Figure S1



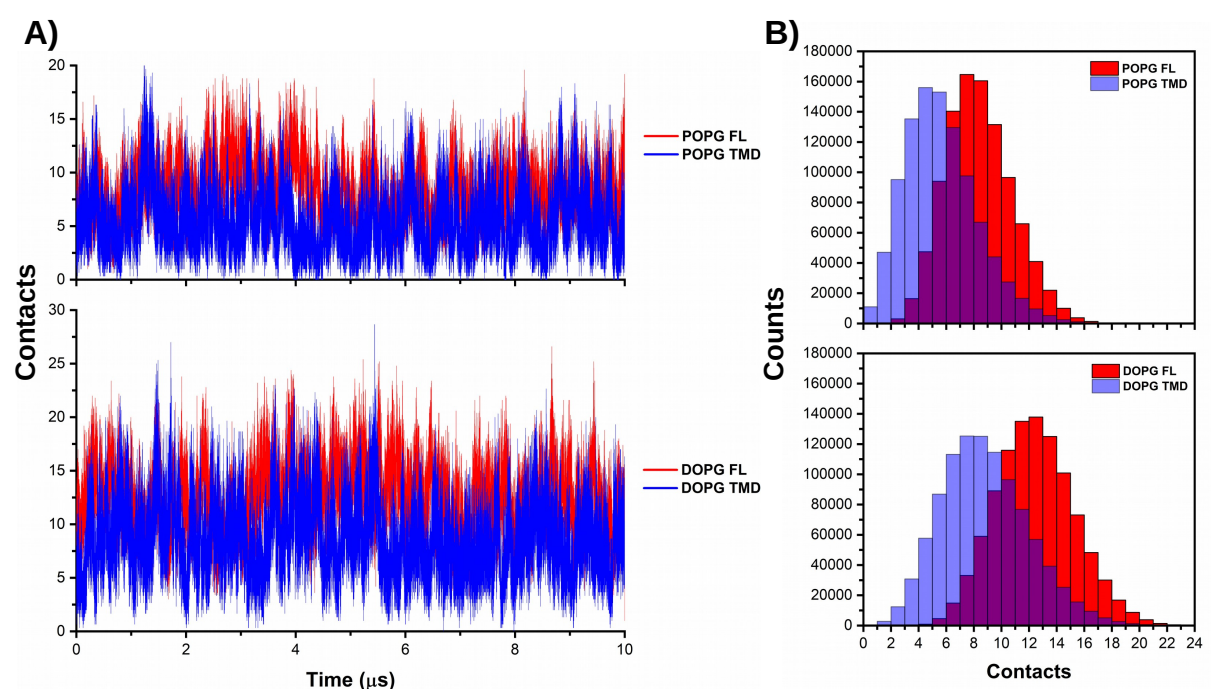
**Top and bottom views of GlpG and GlpG-TMD.** Magenta: CytoD, Cyan: Ln region, Green: TMD, Blue: L5 loop. These orientations have been used for the structural alignment in the 2D density maps (Figures 3, S2).

Figure S2



Averaged density maps of PE-based lipids and DPPG in the top and bottom leaflet for GlpG and GlpG-TMD.

**Figure S3**



**A) Averaged total number of contacts (both leaflets) of POPG and DOPG with GlpG and GlpG-TMD, along entire simulations. B) Distribution of the contacts along entire simulations.**