



Computational Indicator Approach for Assessment of Nanotoxicity of Two-Dimensional Nanomaterials

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1. SIMULATION DETAILS

Calculation of the free energy of lipid extraction

Constant velocity steered molecular dynamic (SMD) [1] simulations were performed to estimate free energy change during the single and a group of 2 and 3 lipids complete extraction from POPC bilayer into the bulk water solution.

Initial dimensions of the simulation box were $60 \times 60 \times 154 \text{ \AA}$. The model membrane was composed of 106 lipids (53 in each leaflet). The membrane was oriented in XY plane and its center of mass was restrained to the plane $Z = 0$ by the stiff spring with a constant $30000 \text{ kJ}/(\text{mol} \cdot \text{\AA}^2)$. Membrane model was generated using CHARMM-GUI v2.0 (<http://www.charmm-gui.org>) [2] and was parameterized with CHARMM36 force field [3]. Total number of atoms was 52 396, including 12 708 water molecules and 34 Na^+ and 34 Cl^- ions. The periodic boundary conditions were set along all the directions. System was modelled as isothermal-isobaric (NpT) ensemble at $T = 310 \text{ K}$ and $p = 1 \text{ atm}$ with a Nose–Hoover algorithm [4,5]. Dimensions of simulation box were changeable to provide target pressure. X and Y dimensions were coupled, while Z was changing independently. Long-range electrostatics was computed with particle-particle particle-mesh (PPPM) algorithm [6,7].

In order to estimate free energy change (ΔG) of the system during the N lipids extraction, the PMF was calculated as the sum of the works done by N virtual springs with constant of $2000 \text{ kJ}/(\text{mol} \cdot \text{\AA}^2)$, with one end of which attached to the center of mass of one from N lipids in the group and the other end moved at a constant velocity $0.1 \text{ \AA}/\text{ns}$ away from membrane and along Z. The X and Y components of the virtual force were zero.

The molecular dynamics models and other simulations-related data can be found at [8,9].

2. SUPPLEMENTAL RESULT DETAILS

Table S1. Quantitative parameters of ΔG profiles estimated for head and tail moieties of the POPC lipid interacting with the two-dimensional nanomaterials. M – point of local minimum, B – energy barrier, ΔG – an estimation of the free energy change, SD – standard deviation of ΔG (error estimation), ξ – distance between nanosheet and adsorbate molecule center of mass.

Nanomaterial	Lipid part	Min./Barrier	ΔG , kJ/mol	SD, kJ/mol	ξ , Å
GN	head	M	-14.2	1.62	4.75
	tails	M	-102.69	10.92	4.13
GON	head	M	-19.17	4.08	4.84
	tails	M	-29.64	14.04	5.21
Mg/Al-LDH	head	M4	-0.3	0.19	12.64
		M3	-0.61	0.28	10.17
		B2	1.1	0.37	9.06
		M2	-2.29	0.61	8.08
		B1	5.09	1.08	6.84
		M1	-3.47	1.41	6.17
	tails	(M)	0	0.12	(inf)
Aloohene	head	M'	-3.78	1.23	6.32
		M	-5.16	1.26	5.76
	tails	(M)	-0.029	0.046	(16.04)
BNN (± 0.5 e)	head	M	-15.61	6.55	4.82
	tails	M	-117.51	3.63	4.02
BNN (± 1.05 e)	head	M	-20.2	2.71	4.66
	tails	M	-115.35	11.91	4.02

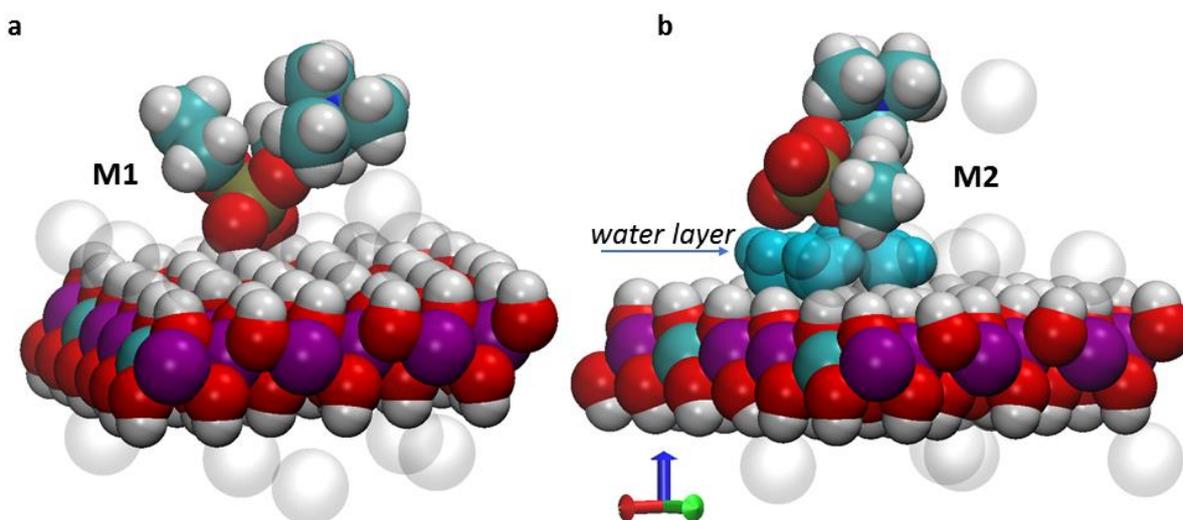


Figure S1. Configurations of head part of POPC lipid adsorbed at local minima M1 (a) $\xi = 6.17$ Å and M2 (b) $\xi = 8.08$ Å on the surface of Mg/Al-LDH. Water is not shown, except several molecules, which form single-molecule-thin water layer between LDH surface and PO_4 -group of a lipid head. Colors: aluminum – cyan, magnesium – purple, carbon – cyan, oxygen – red, hydrogen – white, phosphorus – golden, nitrogen – blue, chlorine ions – transparent, water – light blue.

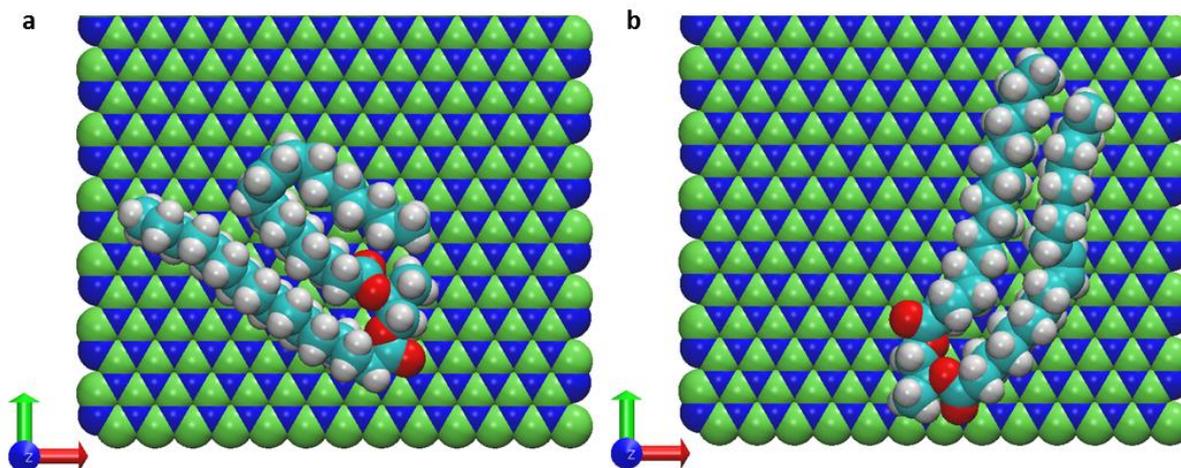


Figure S2. Typical configurations of tails part of POPC lipid adsorbed on the surface of BNN (near point of local minimum M, $\xi = 4.02 \text{ \AA}$). Colors: boron – green, nitrogen – blue, carbon – cyan, oxygen – red, hydrogen – white.

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