

*Supporting Information for*

## **Effects of Molecular Weight and Orientation on the Membrane Permeation and Partitioning of Polycyclic Aromatic Hydrocarbons: A Computational Study**

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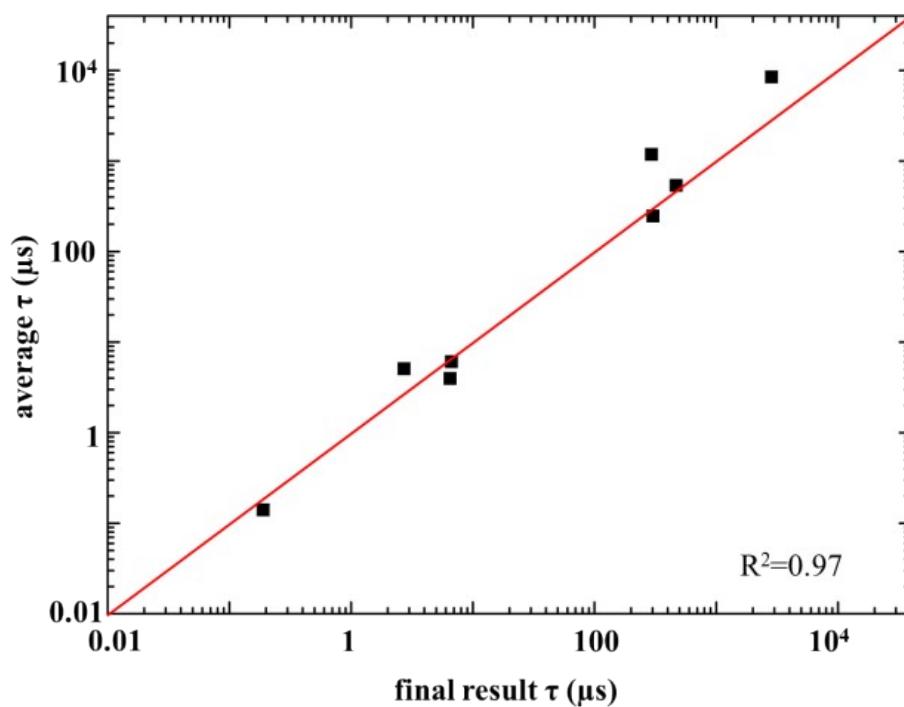
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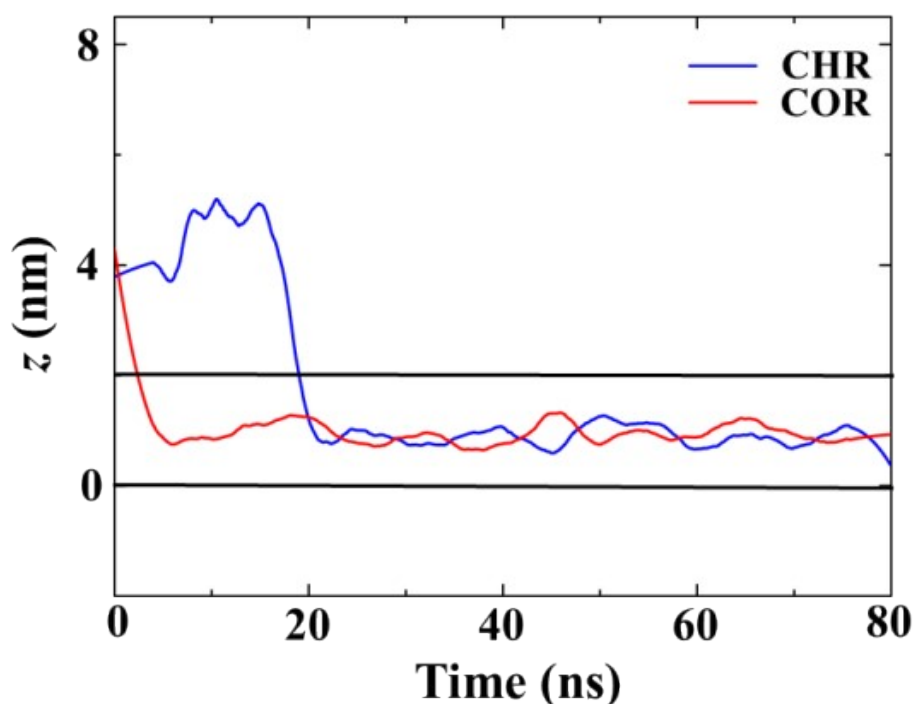
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**Table S1.** The cooperation between calculated parameters from 30-50 ns and the average value

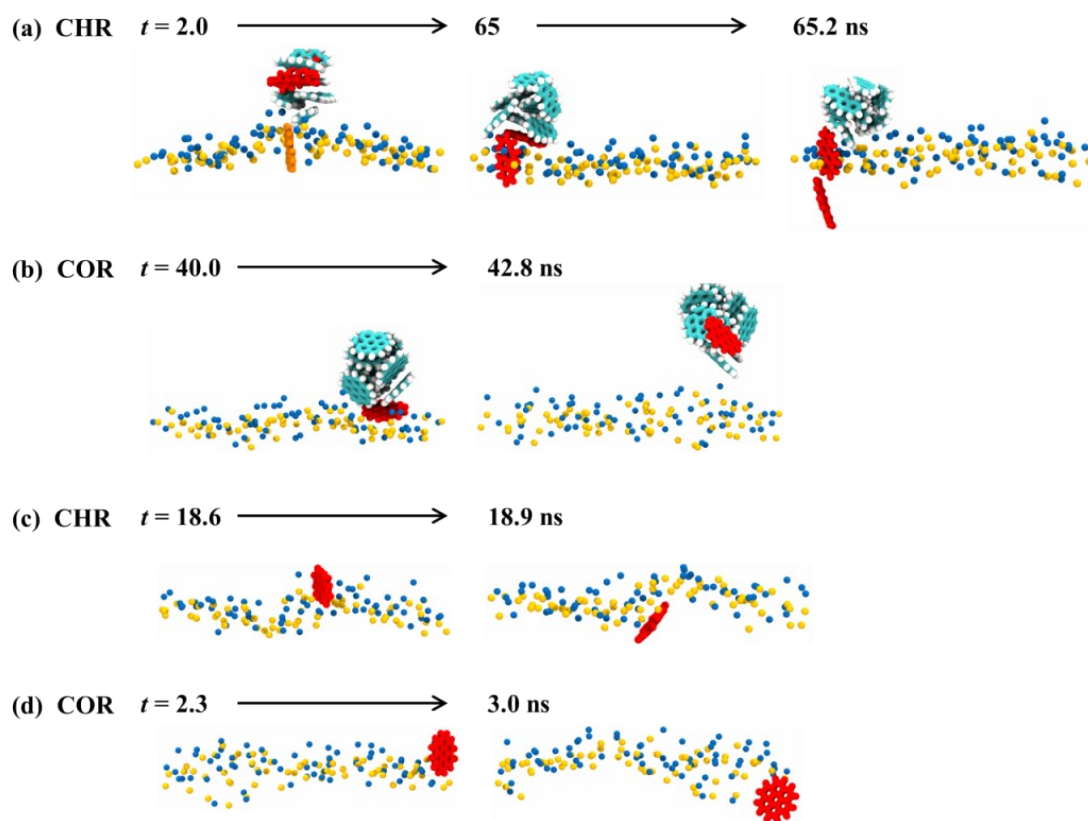
PAHs	$P$ (cm·s <sup>-1</sup> )					$\tau$ (ms)					$\log K_{LW}$				
	35-40	40-45	45-50	30-50	average	35-40	40-45	45-50	30-50	average	35-40	40-45	45-50	30-50	average
<b>NAP</b>	12.3	12.8	11.6	11.7	12.1	0.11	0.1	0.14	0.19	0.14	2.57	2.53	2.68	2.81	2.65
<b>PHE</b>	26.2	29.1	24.5	23.1	25.7	2.99	9.67	4.87	2.73	5.06	3.96	4.52	4.27	3.99	4.18
<b>ANT</b>	20.1	16.6	16.4	16.9	17.5	1.16	2.71	5.41	6.51	3.95	3.81	4.08	4.32	4.39	4.15
<b>PYR</b>	20.9	22.1	22.7	19.3	21.2	2.89	7.81	6.84	6.68	6.05	4.24	4.55	4.46	4.42	4.42
<b>CHR</b>	36.8	39.1	38.8	41.2	39.0	359.0	605.9	707	469.9	535	6.49	6.73	6.73	6.58	6.63
<b>BAP</b>	42.5	29.7	31.7	29.2	33.3	3339.3	613.5	475.7	294.8	1180	7.57	6.62	6.49	6.2	6.72
<b>DBA</b>	36.0	35.4	30.4	26.4	32.1	123.2	289.3	269.9	303.4	246	5.81	6.11	6.10	6.05	6.02
<b>COR</b>	35.2	33.1	35.4	32.7	34.1	16805	11878	2254	2858	8449	7.99	7.90	7.21	7.3	7.60



**Figure S1.** The fitting relationship between our final result of transmembrane time (30-50 ns) and average result (three simulation periods of 35-40 ns, 40-45 ns and 45-20 ns), the red line indicate a 1:1 agreement.



**Figure S2.** The time evolution of the  $z$  position of the center of mass of single CHR (blue line) and COR (red line). The horizontal lines at  $z = 0$  and  $z = 2$  nm represent the center and surface (which was substituted with the average position of the P) of lipid membrane, respectively.



**Figure S3.** The typical snapshots of the absorption process of (a) CHR, (b) COR. (c) single CHR, (d) single COR.

**Table S2.** Summary of all simulations performed

	solute	No. of solute	Duration (ns)
<b>Constrained MD</b>	NAP	1	50 × 32
	PHE	1	50 × 32
	ANT	1	50 × 32
	PYR	1	50 × 32
	CHR	1	50 × 32
	BAP	1	50 × 32
	DBA	1	50 × 32
	COR	1	50 × 32
<b>Unconstrained MD</b>	NAP	10	80

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PYR	10	80
CHR	10	80
COR	10	80
CHR	1	80
COR	1	80

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