SUPPORTING INFORMATION

Investigating non-specific binding to chemically engineered sensor surfaces using liposomes as models

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1) Cyclic voltammograms of self-assembled monolayer on Gold





Figure S1. Cyclic voltammograms before and after the formation of self-assembled monolayers on gold electrodes. The solutions are 5 mM $Fe(CN)_6^{3-}$ in HEPES buffer. Scan rate 100 mV/s.

2) Fitting parameters of binding isotherms

Table S1. Parameters for the interaction of untagged liposomes with self-assembled monolayers

 on gold obtained by binding isotherms.

Fit model	Surface	$\Delta n_{D, max}^{a)} / RIU$	<i>К</i> _{<i>L</i>^{b)} / L·pmol⁻¹}	<i>К_Dс) /</i> pmol·L ⁻¹	h	R ²
simple	-COOH	0.0104±0.0013	0.00074 ± 0.00012	1400±200		0.9991
	-OH	0.0094 ± 0.0009	0.0032±0.0005	310±60		0.9924
	-CH ₃	0.0013±0.0002	0.012±0.006	80±40		0.8501
	-NH ₂	0.019±0.002	0.0061±0.0018	160±50		0.9730
	gold	0.0134±0.0010	0.00271±0.0003	370±40		0.9968
Fit model	Surface	$\Delta n_{D, max}^{a)} / RIU$	Κ _L ^{b)} / (L·pmol ⁻¹) ^h	<i>К_D^{с)} /</i> (pmol·L ⁻¹) ^h	h	R ²
Fit model	Surface -COOH	$\Delta n_{D, max}^{a)} / RIU$ 0.014±0.010	$\frac{K_L^{b}}{(L \cdot pmol^{-1})^h}$ 0.0007±0.0003	K _D ^{c)} / (pmol·L ⁻¹) ^h 1400±600	h 0.95±0.08	R ² 0.9989
Fit model	Surface -COOH -OH	$\Delta n_{D, max}^{a)} / RIU$ 0.014±0.010 0.026±0.006	$\frac{K_L^{b}}{(L \cdot pmol^{-1})^h}$ 0.0007±0.0003 0.0036±0.0006	$\frac{K_{D}^{c)} / (pmol \cdot L^{-1})^{h}}{1400 \pm 600}$ 280 ± 50	h 0.95±0.08 0.71±0.02	R ² 0.9989 0.9998
Fit model	Surface -COOH -OH -CH ₃	$\Delta n_{D, max}^{a)} / RIU$ 0.014±0.010 0.026±0.006	$\frac{K_L^{b}}{(L \cdot pmol^{-1})^h}$ 0.0007±0.0003 0.0036±0.0006 -	$\frac{K_{D}^{c)} / (pmol \cdot L^{-1})^{h}}{1400 \pm 600}$ 280 ± 50 -	<i>h</i> 0.95±0.08 0.71±0.02 -	R ² 0.9989 0.9998 -
Fit model	Surface -COOH -OH -CH ₃ -NH ₂	$\Delta n_{D, max}^{a)} / RIU$ 0.014±0.010 0.026±0.006 - 0.0148±0.0008	$ K_L^{b)} / (L \cdot pmol^{-1})^h $ 0.0007±0.0003 0.0036±0.0006 - 0.0006±0.0005	$\frac{K_{D}^{c)} / (pmol \cdot L^{-1})^{h}}{1400 \pm 600}$ 280 ± 50 - 2000 ± 1400	<i>h</i> 0.95±0.08 0.71±0.02 - 1.62±0.19	R ² 0.9989 0.9998 - 0.9941

^{a)}refractive index change at maximum surface loading, ^{b)}Langmuir equilibrium constant, ^{c)}Dissociation constant $K_D=1/K_L$

Fit model	Surface	$\Delta n_{D, max}^{a)} / RIU$	<i>К</i> _L ^{b)} / L∙pmol ⁻¹	<i>К</i> _D ^{с)} / pmol·L ⁻¹	h	R ²
simple	-COOH	0.0052±0.0019	0.0024±0.0012	400±200		0.9830
	-OH	0.0143±0.0009	0.0030±0.0003	330±30		0.9992
	-CH ₃	0.0018 ± 0.0005	0.02±0.013	50±30		0.7174
	-NH ₂	0.018±0.002	0.014±0.004	70±20		0.9791
	gold	0.022±0.012	0.0010±0.0006	1000±600		0.9930
Fit model	Surface	$\Delta n_{D, max}^{a)} / RIU$	<i>K</i> _L ^{b)} / (L·pmol ⁻¹) ^h	$K_D^{c)} / (\text{pmol·L}^{-1})^h$	h	R ²
Fit model	Surface -COOH	$\Delta n_{D, max}^{a)} / \text{RIU}$ 0.01±0.03	$\frac{K_L^{\rm b)}}{(L \cdot pmol^{-1})^h}$ 0.002±0.004	$K_{D}^{c)} / (pmol \cdot L^{-1})^{h}$ 500±1000	h 0.9±0.3	R ² 0.9780
Fit model extended	Surface -COOH -OH	$\Delta n_{D, max}^{a)} / RIU$ 0.01±0.03 0.020±0.005	$\frac{K_L^{b)} / (L \cdot pmol^{-1})^h}{0.002 \pm 0.004}$ 0.0029 \pm 0.0004	K_D^{c} / (pmol·L ⁻¹) ^h 500±1000 340±50	<i>h</i> 0.9±0.3 0.91±0.04	R ² 0.9780 0.9995
Fit model extended	Surface -COOH -OH -CH ₃	$\Delta n_{D, max}^{a)} / RIU$ 0.01±0.03 0.020±0.005 -	$\frac{K_{L^{b}}}{(L \cdot pmol^{-1})^{h}}$ 0.002±0.004 0.0029±0.0004 -	K_{D}^{c} / (pmol·L ⁻¹) ^h 500±1000 340±50 -	<i>h</i> 0.9±0.3 0.91±0.04 -	R ² 0.9780 0.9995 -
Fit model extended	Surface -COOH -OH -CH ₃ -NH ₂	$\Delta n_{D, max}^{a)} / RIU$ 0.01±0.03 0.020±0.005 - 0.0142±0.0004	$\frac{K_{L^{b}}}{(L \cdot pmol^{-1})^{h}}$ 0.002±0.004 0.0029±0.0004 - 0.0028±0.0009	<i>K_D</i> ^{c)} / (pmol ·L ⁻¹) ^{<i>h</i>} 500±1000 340±50 - 360±110	<i>h</i> 0.9±0.3 0.91±0.04 - 1.57±0.10	R ² 0.9780 0.9995 - 0.9983

Table S2. Parameters for the interaction of liposomes with N-glutaryl-DPPE tag with self-assembled monolayers on gold obtained by simple and extended Langmuir fits.

^{a)}refractive index change at maximum surface loading, ^{b)}Langmuir equilibrium constant, ^{c)}Dissociation constant $K_D=1/K_L$

3) SPR Simulation

SPR simulations were performed with the software Winspall (Freeware from MPIP Mainz, Germany). A 640 nm light source and a triangular prism (50 °) were chosen in best resemblance of the experimental setup. The following layer setup was used for approximation of a closest packed liposome monolayer:

Layer	Thickness / nm	εX - real	εX - imaginary
Glass	Infinite (starting layer)	2.59	0
Chromium ¹	3	-6.901	28.818
Gold ²	50	-12.555	1.1464
Thiol	1.5	1.847	0
Lipid bilayer	4.8	2.25	0
300 mM NaCl	200	1.786	0
Lipid bilayer	4.8	2.25	0
Water	Infinite (ending layer)	1.778	0



Figure S2. Change of the minimum of the SPR angle induced by anionic liposomes (300 mM NaCl encapsulant, without N-glutaryl-DPPE tag) on a $-NH_2$ SAM modified gold surface simulated for maximum surface loading (left) and obtained from experiment at 100 μ M phospholipid concentration (right).



Figure S3. Van 't Hoff plot of the K_D values obtained from the Langmuir fits of the temperature dependent interactions of anionic liposomes (300 mM NaCl encapsulant, no tag) with -OH SAM modified gold surfaces at varying phospholipid concentrations at different temperatures.

References

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