## Toll-like receptor 2 promiscuity is responsible for the immunostimulatory activity of nucleic acid nanocarriers

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Supplemental Figures S1-S9. Docking poses of lipopolyamines bound to <u>TLR2.</u> Representations generated by LigPlot+ software.



Figure S1. RPR 128506 docking to TLR2/TLR1 with one chains per TLR. The binding energy of this complex is -31.4 kcal/mol.



**Figure S2**. **RPR 120535 docking to TLR2/TLR1 with one chain in each TLR.** The binding energy of this complex is -32.2 kcal/mol. This is the best pose for this compound. The head group mediates five hydrogen bonds with TLR2 and one with TLR1.



Figure S3. RPR 120525 docking to TLR2/TLR1 with one chain in each TLR. The binding energy of this complex is -32.5 kcal/mol.



Figure S4. RPR 128506 docking to TLR2/TLR1 with both chains in TLR2. The binding energy of this complex is -31.9 kcal/mol.



Figure S5. RPR 120535 docking to TLR2/TLR1 with both chains in TLR2. The binding energy of this complex is -31.1 kcal/mol). A hydrogen bond to TLR1 Gln316' that was found using Chimera is not depicted by LigPlot+.



Figure S6. RPR 120525 docking to TLR2/TLR1 with both chains in TLR2. The binding energy of this complex is -32.1 kcal/mol.



**Figure S7. RPR 128506 docking to TLR2/TLR6.** The binding energy of this complex is -29.9 kcal/mol. Additional hydrogen bonds to TLR2 Tyr326 and TLR6 Ser320", Thr322" and Tyr 323" were found using Chimera but is not depicted by LigPlot+.



**Figure S8. RPR 120535 docking to TLR2/TLR6.** The binding energy of this complex is -28.5 kcal/mol. An additional polar contact with TLR6 Ser320 was found using Chimera (right panel) but is not depicted by LigPlot+ (left panel).



**Figure S9. RPR 120525 docking to TLR2/TLR6.** The binding energy of this complex is -31.7 kcal/mol.



<u>Supplemental Figure S10:</u> RPR 120525 is a weaker agonist then lipopeptides. Primed THP1 cells were incubated for 5h with serum free medium alone (control) or with the indicate amount of (A)  $Pam_2CSK_4$ , (B)  $Pam_3CSK_4$  or (C) RPR 120525 cationic lipids in serum free medium, then cell supernatants were collected and TNF- $\alpha$  was quantified using ELISA assay following manufacturer's instructions. Each point represents the mean ± standard deviation of three replicate values (n=3). The experiment is representative of at least 3 replicates.



<u>Supplemental Figure S11</u>. RPR 208484 with a single unsaturated alkene chain 18:1 ( $\Delta$ 9-Cis) signals as strongly as its saturated counterpart RPR 120535 and TLR2/TLR1 ligand Pam<sub>3</sub>CSK<sub>4</sub>. Raw-blue cells were incubated for 5h with serum free medium alone (C) or with the indicate amount of cationic lipids in  $\mu$ M in serum free medium, then cell supernatants were retired and replaced with complete medium. Supernatants were collected after further 17h of incubation and the NF-kB activation was quantified using Quanti-blue assay following manufacturer's instructions. Each bar represents the mean + standard deviation of three replicate values (n=3). The experiment is representative of at least 3 replicates.



<u>Supplemental Figure S12</u>. Chemical structure of lipofectamine transfection reagent component DOSPA.