## **Supplemental Figures**



**Fig. S1. (**a) Confocal images of isolated GPMVs. Left: GFP labelled CD47. Right: Membrane marker DiIC18. The arrow indicates a typical vesicle selected for experiments. Scale bars 10µm (b) Anti-GST staining of SIRP $\alpha$  absorbed on a glass slide. The black region is a scratch with a pipette tip to estimate the background level. By florescence measurements and comparison to the known density of fluorophores on RBC incubated with free SIRP $\alpha$ , we estimate the SIRP $\alpha$  density to roughly [R<sub>0</sub>]  $\approx$ 4000 molecules/µm<sup>2</sup>. This estimate is consistent with E-selectin densities obtained by physisorption on glass slides under similar conditions(1) and in the same order of magnitude as SIRP $\alpha$  layers obtained on plastic surfaces (2). The density is below the maximum packing density of a SIRP $\alpha$  monolayer and we can expect that BSA, which is used for blocking uncovered glass surface, fill the gaps between individual SIRP $\alpha$  molecules. Scale bar 60µm. (c) CD47-GFP TIRF micrographs of the adhering GPMV segment on a SIRP $\alpha$  labeled glass surface. Micrographs were acquired on three different GPMVs during the first 5-20 minutes after initial GPMV adhesion. The ring-like CD47-GFP patterns equilibrated over time to a homogenous fluorescence signal. All further experiments were conducted on equilibrated, homogenous adhesion segments. Adhesion segments were typically about 10µm in diameter.



**Fig. S2** Probability distribution of membrane-substrate distance obtained from simulations of the coarse-grained CD47-SIRPα complex.



**Fig. S3.** (a) Binding of (water soluble) GST-SIRPα to RBC showed no significant effect of shift from pH=7.4 to pH=6 on K3d binding affinity. (b) Incubation of red blood cells (RBCs) together with GPMVs and fluorescent anti-CD47 in order to estimate the CD47 concentration on GPMV's. Arrow indicates RBC. Scale bar 10µm

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