

Supplementary Materials: A Dual-Reporter Platform for Screening Tumor-Targeted Extracellular Vesicles

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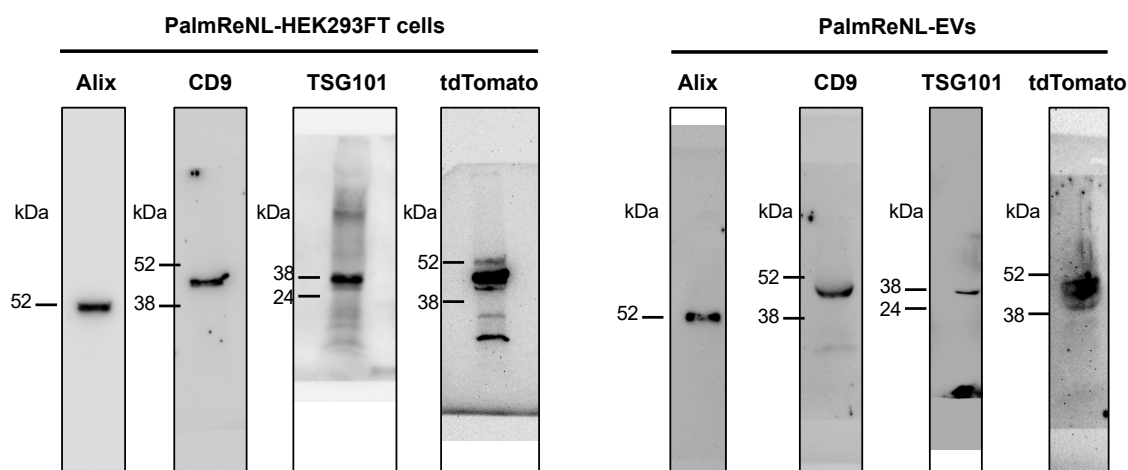
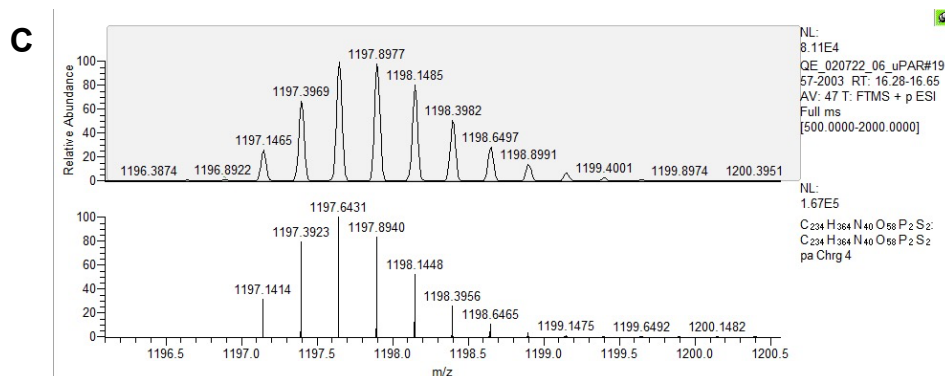
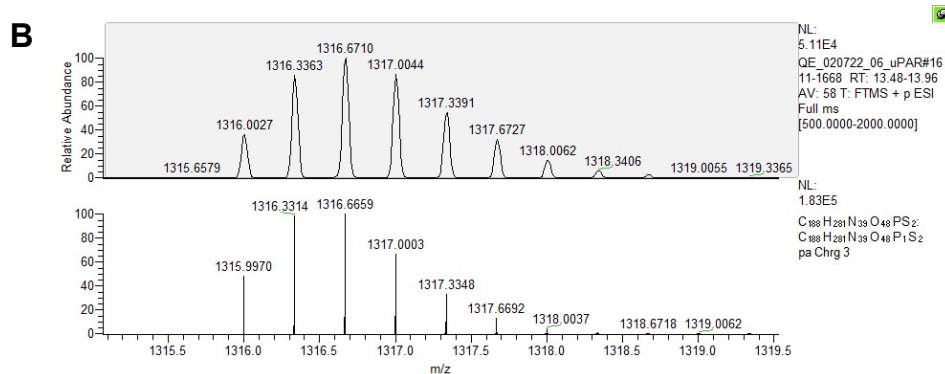
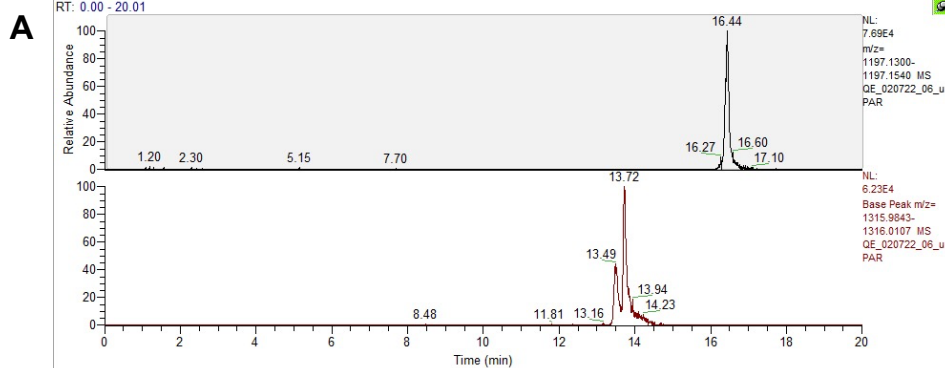
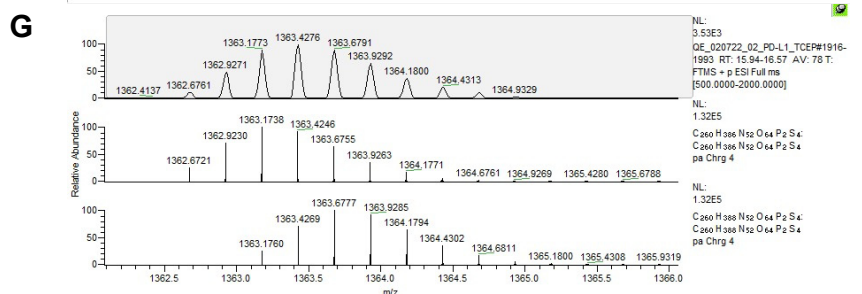
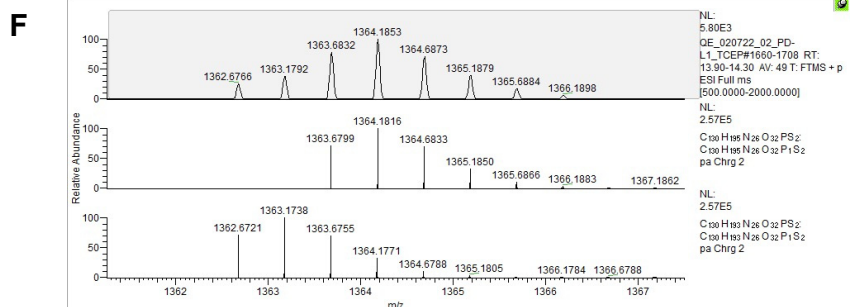
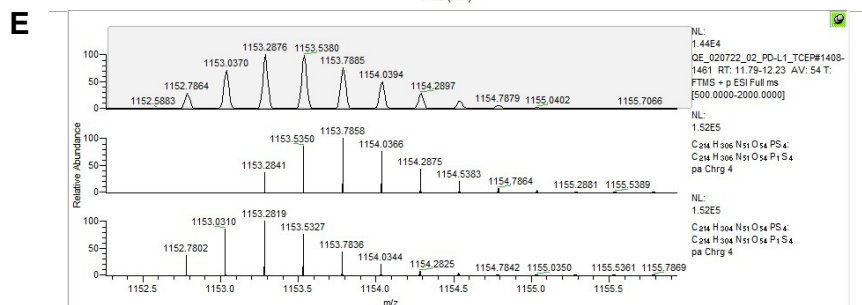
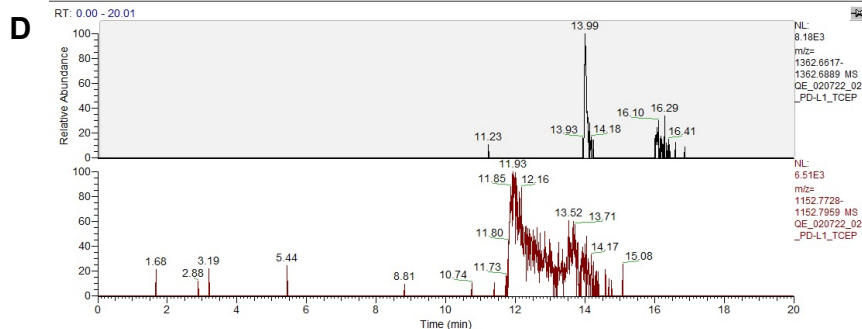


Figure S1. Western blot analysis of exosome marker proteins in PalmReNL-HEK293FT cells and - EVs. Original images for Figure 1D.

DOPE-uPAR (VSNKYFSNIHWGC)



DOPE-PDL1 (YASYHCWCWRDPGRS)



DOPE-EGFR (KKKGGYHWYGYTPQNVI)

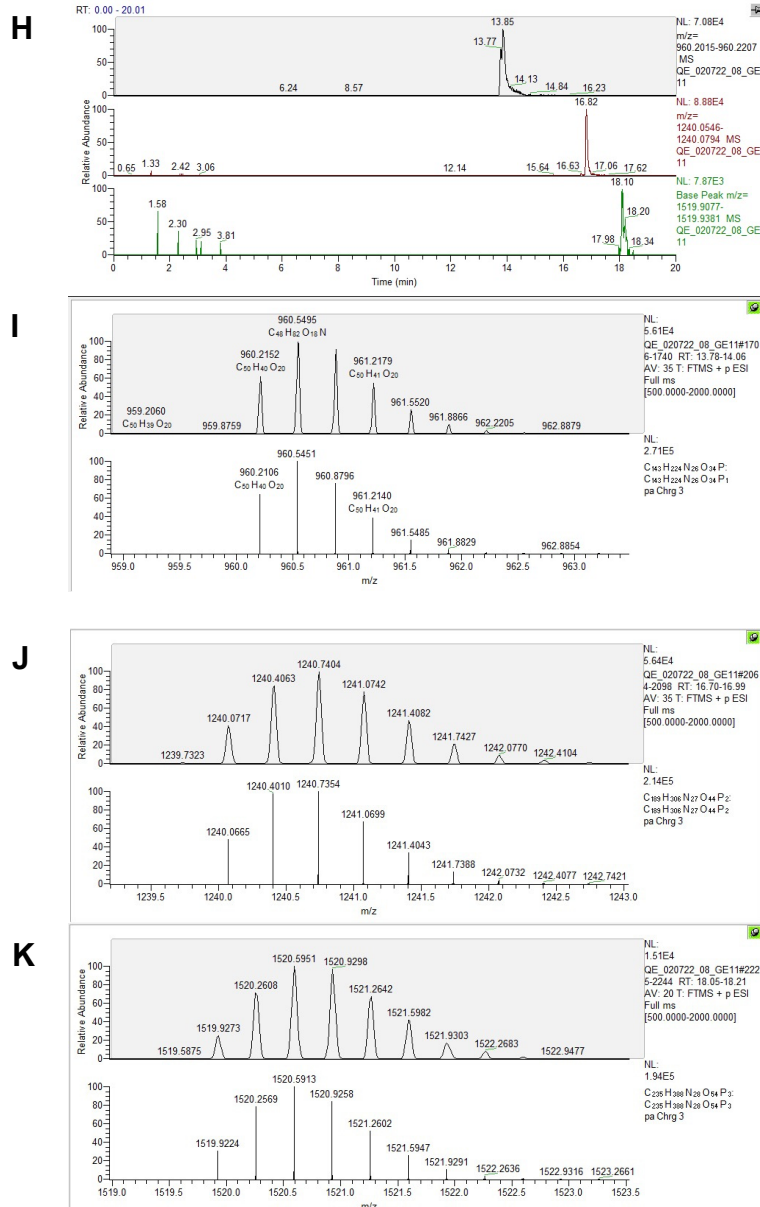


Figure S2. Characterization of DOPE-THPs by LC/MS. **A)** Extracted ion chromatograms for m/z 1315.9975 and 1197.142 (+/- 10 ppm). **B)** Mass spectrum at 13.48-13.96 min showing DOPE-uPAR-uPAR-DOPE conjugate with a disulfide linking two DOPE-uPARs. Isotope models are shown below the raw data. **C)** Mass spectrum at 16.28-16.65 min showing DOPE-uPAR-uPAR conjugate with a disulfide linking one DOPE-uPAR and one unmodified uPAR-binding peptide. Isotope models are shown below the raw data. **D)** Extracted ion chromatogram for m/z 1152.7844 and 1362.6753 (+/- 10 ppm). **E)** Mass spectrum at 11.79-12.23 min showing DOPE-PDL1-PD-L1 conjugate with one or two disulfides linking two PD-L1-binding peptides. Isotope models are shown below the raw data. **F)** Mass spectrum at 13.9-14.3 min showing DOPE-PDL1 conjugate with none or one internal disulfide linkage. Isotope models are shown below the raw data. **G)** Mass spectrum at 15.94-16.57 min showing DOPE-PDL1-PDL1-DOPE conjugate with one or two disulfides linking two DOPE-PDL1s. **H)** Extracted ion chromatograms for m/z 960.2111, 1240.067 and 1519.9929 (+/- 10 ppm). **I)** Mass spectrum at 13.78-14.06 min showing one EGFR-binding peptide with one DOPE attached. Isotope models are shown below the raw data. **J)** Mass spectrum at 16.70-16.99 min showing one EGFR-binding peptide with two DOPE molecules attached. Isotope models are shown below the raw data. **K)** Mass spectrum at 18.05-18.21 min showing one EGFR-binding peptide with three DOPE molecules attached. Isotope models are shown below the raw data.

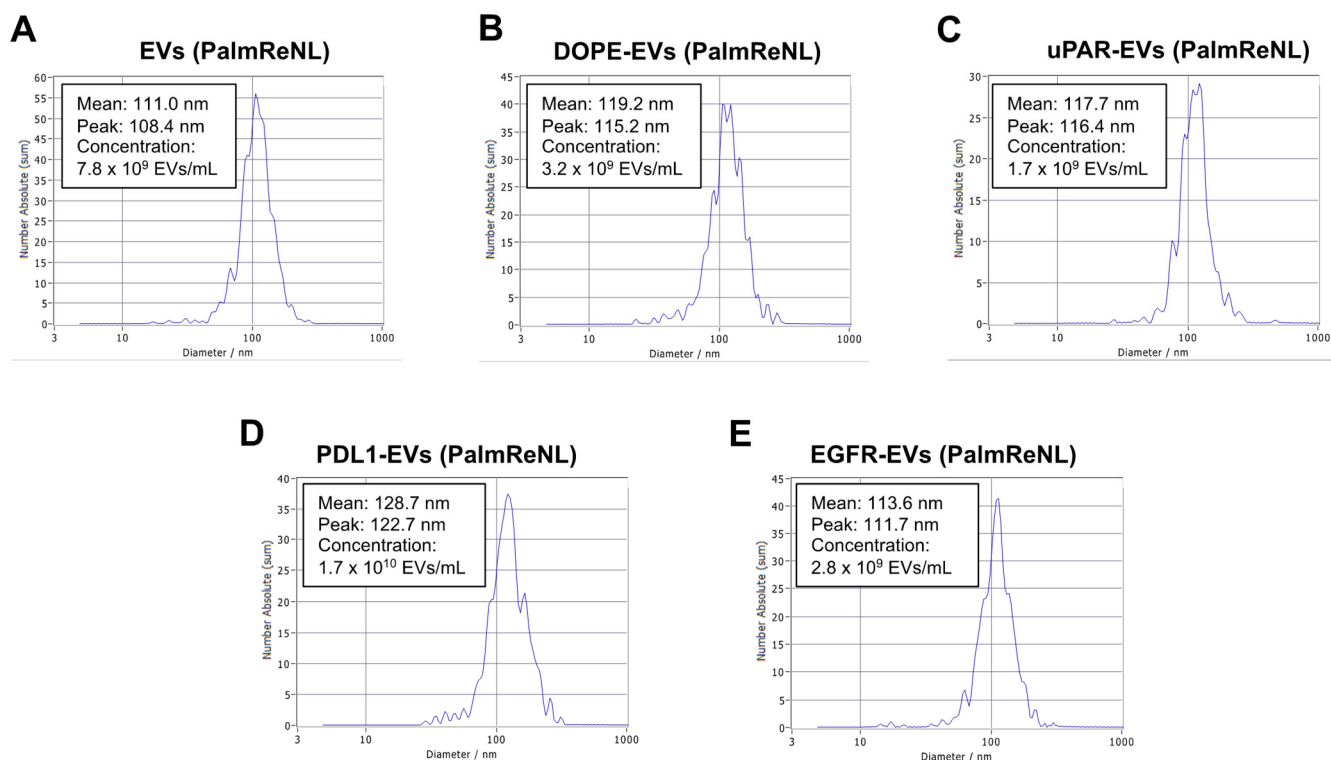


Figure S3. Characterization of PalmReNL-EVs engineered with THPs by nanoparticle tracking analysis (NTA). **A-B)** Unmodified and DOPE-only control PalmReNL-EVs. **C-E)** PalmReNL-EVs bearing uPAR-, PDL1-, and EGFR-binding peptides.

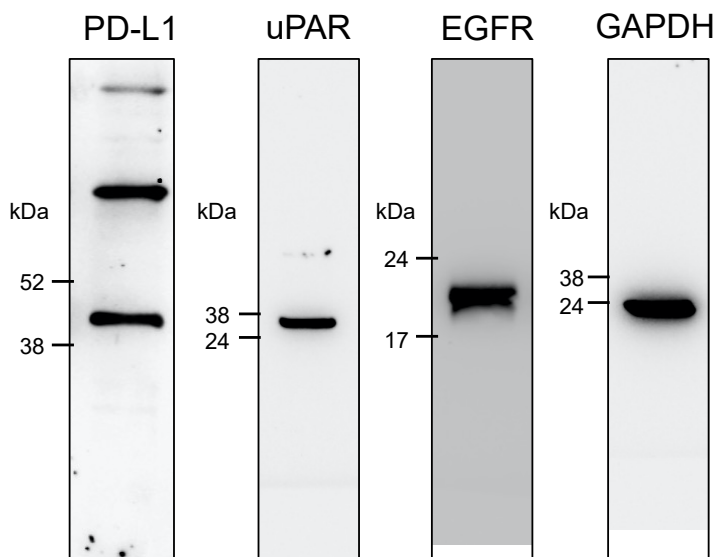


Figure S4. Western blot analysis of the target membrane protein expression in MDA-MB-231 cells under the non-reducing condition. GAPDH is a loading control. Original images for Figure 2D.