

## **Supplemental Information**

### **Optimized Cholesterol-siRNA Chemistry Improves Productive Loading onto Extracellular Vesicles**

**Reka Agnes Haraszti, Rachael Miller, Marie-Cecile Didiot, Annabelle Biscans, Julia F. Alterman, Matthew R. Hassler, Loic Roux, Dimas Echeverria, Ellen Sapp, Marian DiFiglia, Neil Aronin, and Anastasia Khvorova**

**Optimized cholesterol-siRNA chemistry improves productive loading onto extracellular vesicles**

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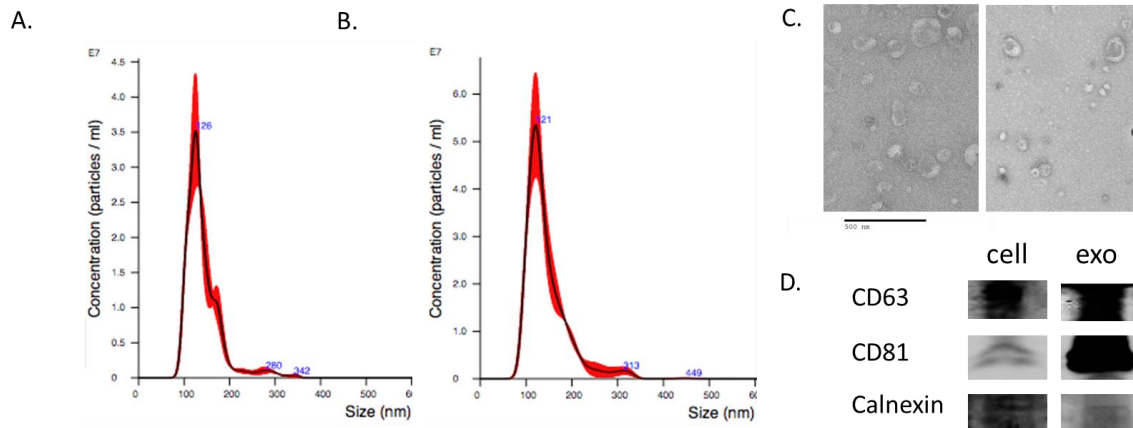
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Keywords: Extracellular vesicles; siRNA; oligonucleotides; nanovesicles; chemical modification; RNA therapy

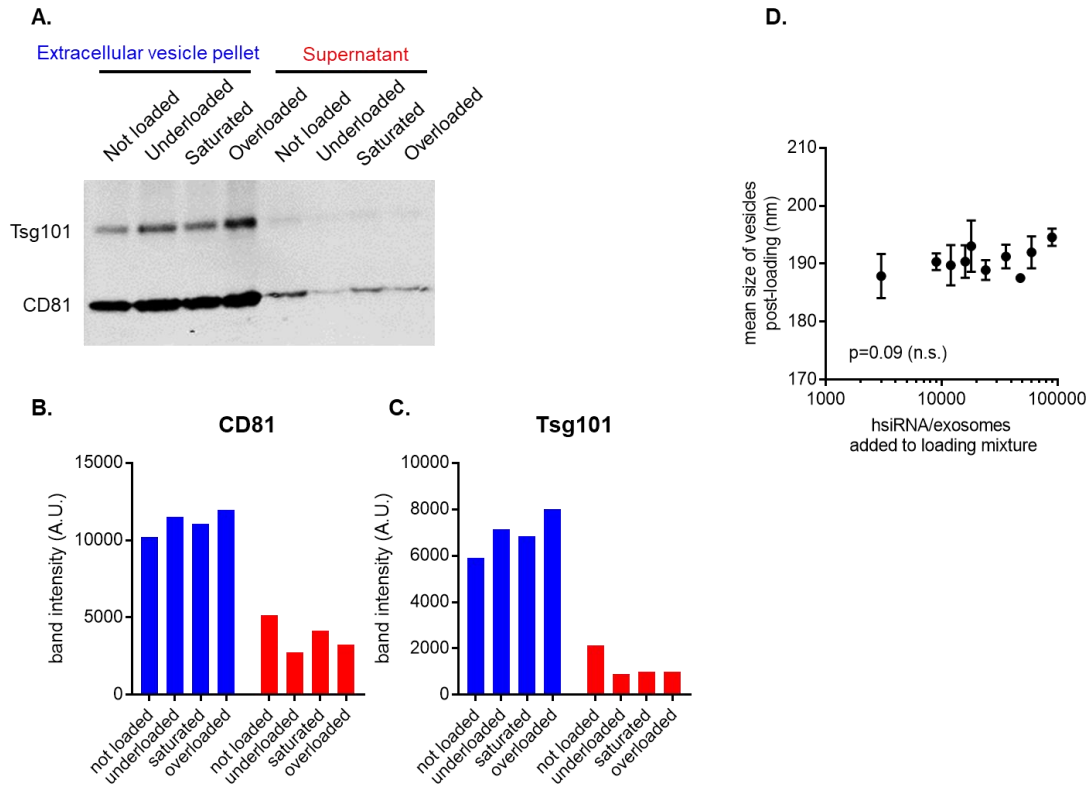
## Supplementary Information



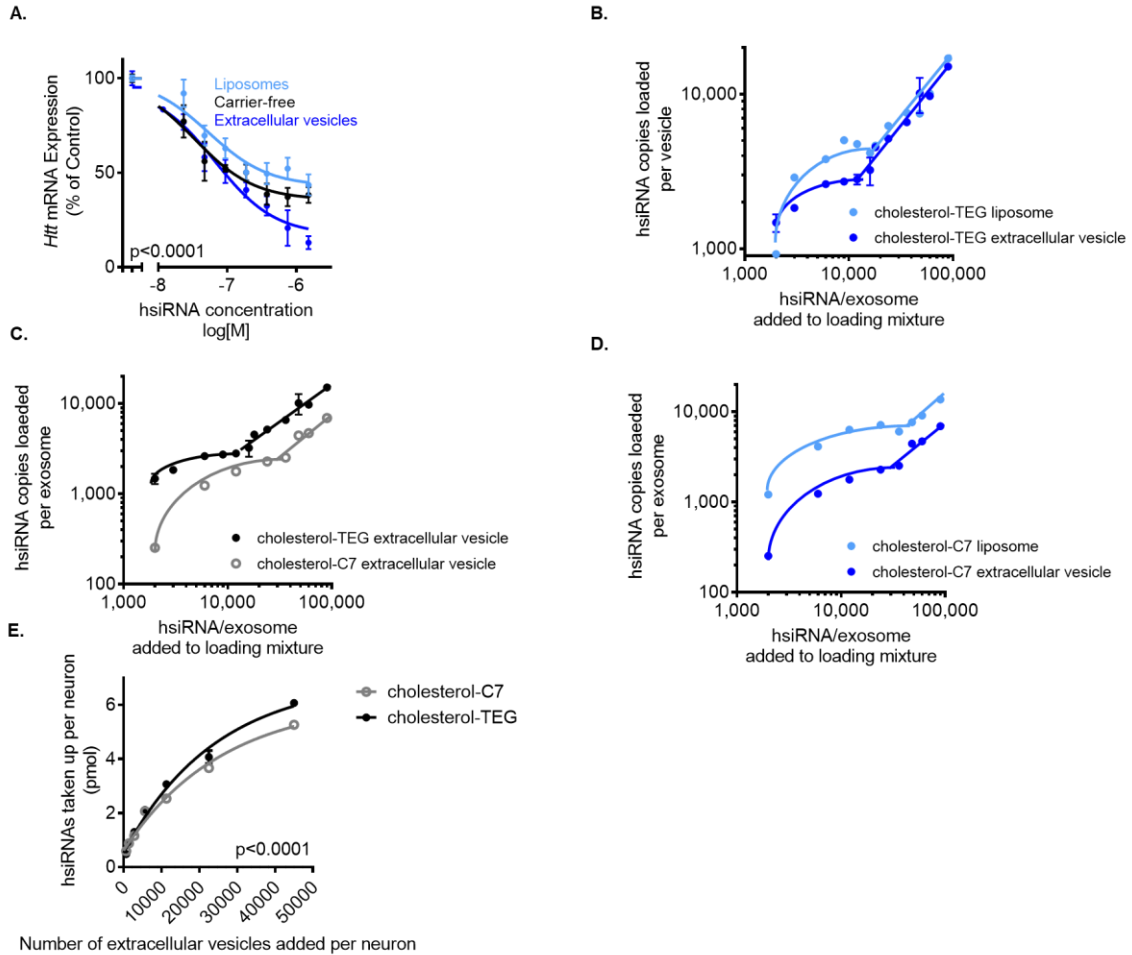
### Supplementary Figure 1. Characterization of liposomes and umbilical cord,

#### Wharton's jelly derived extracellular vesicles

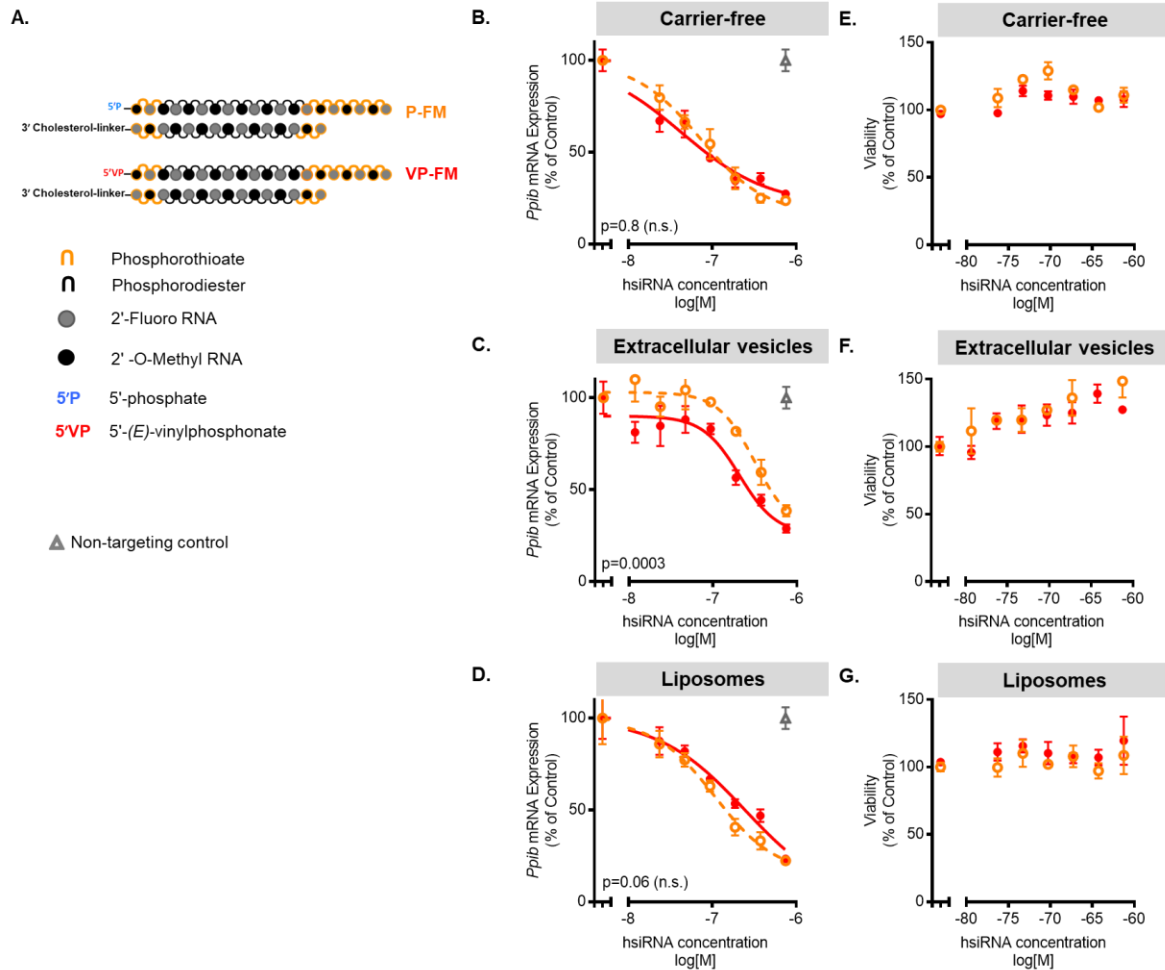
**A.** Nanoparticle Tracking Analysis shows homogenous extracellular vesicle size distribution with mean diameter  $141 \pm 40$  nm,  $n=3$  **B.** Nanoparticle Tracking Analysis of neutral liposomes, mean diameter  $144 \pm 47$  nm,  $n=3$  **C.** Transmission Electron Microscopy image of unloaded (left) and loaded (right) extracellular vesicles, size bar shows 500 nm. **D.** Western blot of positive and negative extracellular vesicle marker proteins.



**Supplementary Figure 2. Characterization of loaded extracellular vesicles** **A.** Western blot of unloaded, underloaded, saturated and overloaded extracellular vesicles (0, 3000, 12 000 and 90 000 RNA molecules per vesicle mixed in; 0, 1900, 3700 and 14300 RNA molecules per vesicle loaded). After incubation at 37°C for 1 hour, extracellular vesicle-hsiRNA mixture was centrifuged at 100,000 g for 1 hour to pellet loaded extracellular vesicles (blue) and remove non-loaded siRNA (red). SDS-PAGE was conducted after lysis in RIPA. **B.** CD81 signal was quantified using ImageJ Gel Analysis tool. **C.** Tsg101 signal was quantified using ImageJ Gel Analysis tool. **D.** Size (average diameter of particles in nm) of loaded extracellular vesicles measured *via* Nanoparticle Tracking Analysis. n=3, mean ± SD



**Supplementary Figure 3. Comparison of loaded extracellular vesicles *versus* loaded liposomes** **A.** Silencing activity of hsiRNA loaded onto either extracellular vesicles (dark blue) or liposomes (light blue) or delivered carrier-free (black) to primary neurons.  $n=3$ , mean  $\pm$  SEM **B-D.** Cholesterol-hsiRNA was loaded onto extracellular vesicles at different hsiRNA-to-extracellular vesicles ratios. The loading curve shows an initial saturation phase followed by a secondary linear phase. **E.** Guide strand accumulation of cholesterol-C7-hsiRNA and cholesterol-TEG-hsiRNA in neurons following delivery *via* extracellular vesicles.  $n=3$ , mean  $\pm$  SEM, two-way ANOVA. Guide strand accumulation measured by PNA hybridization assay.



**Supplementary Figure 4. Stabilization of 5'-phosphate is not toxic in and beneficial for EV-mediated delivery of siRNAs.** **A.** Scheme of chemically modified hsiRNAs. P-FM fully modified backbone with 5'-phosphate on guide strand, VP-FM fully modified backbone with 5'-(E)-vinylphosphonate on guide strand.  $n=3$ , mean  $\pm$  SEM **B-D.** HeLa cells were incubated for three days with cholesterol-hsiRNA variants with different 5' end modifications either alone (carrier-free), or loaded onto extracellular vesicles or liposomes, target *Ppib* mRNA silencing was measured, and silencing potency calculated (IC<sub>50</sub>).  $n=3$  Pairwise comparison of curves was conducted using two-way ANOVA. Significance is depicted in grey. **E-G.** Primary murine cortical neurons were incubated for one week with cholesterol-hsiRNA variants with different 5' end modifications either alone (carrier-free),

or loaded onto extracellular vesicles or liposomes, targeting *Huntingtin*. To measure cell viability, Alamar Blue® was added and incubated at 37°C for 12 hours, and fluorescence measured at 570 nm excitation, 585 nm emission. Signal is normalized to non-treated cells samples. n=3, mean  $\pm$  SD

Gene Targeted	Compound Name	Figure applicable	Strand	Sequence 5'-3'	Conjugate 5'	Conjugate 3'
<i>Huntingtin</i>	P-PM	Fig.3.	passenger	mC.mA.G.mU.A.A.A.G.A.G.A.mU.mU#mA#mA	Cy3	Cholesterol-TEG
			guide	PmU.fU.A.A.fU.fC.fU.fC.fU.fU.fU.A.fC.fU#G#A#U#A#U#A		
<i>Huntingtin</i>	VP-PM	Fig.3.	passenger	mC.mA.G.mU.A.A.A.G.A.G.A.mU.mU#mA#mA	Cy3	Cholesterol-TEG
			guide	VPmU.fU.A.A.fU.fC.fU.fC.fU.fU.fU.A.fC.fU#G#A#U#A#U#A		
<i>Huntingtin</i>	cholesterol-TEG hsiRNA P-FM	Fig.1-3.	passenger	fC#mA#fG.mU.fA.mA.fA.mG.fA.mG.fA.mU.fU#mA#fA	Cy3	Cholesterol-TEG
			guide	PmU#fU#mA.fA.mU.fC.mU.fC.mU.fU.mU.fA.mC#fU#mG#fA#mU#fA#mU#fA		
<i>Huntingtin</i>	VP-FM Endonuclease stable	Fig.3-4.	passenger	fC#mA#fG.mU.fA.mA.fA.mG.fA.mG.fA.mU.fU#mA#fA	Cy3	Cholesterol-TEG
			guide	VPmU#fU#mA.fA.mU.fC.mU.fC.mU.fU.mU.fA.mC#fU#mG#fA#mU#fA#mU#fA		
<i>Huntingtin</i>	cholesterol-C7	Fig.1.	passenger	fC#mA#fG.mU.fA.mA.fA.mG.fA.mG.fA.mU.fU#mA#fA	Cy3	Cholesterol-C7
			guide	PmU#fU#mA.fA.mU.fC.mU.fC.mU.fU.mU.fA.mC#fU#mG#fA#mU#fA#mU#fA		
<i>Huntingtin</i>	moderately endonuclease sensitive (DNA)	Fig.4.	passenger	fC#mA#fG.mU.A.mA.fA.mG.fA.mG.fA.mU.fU#mA#fA.dT.dT		Cholesterol-TEG
			guide	VPmU#fU#mA.fA.mU.fC.mU.fC.mU.fU.mU.fA.mC#fU#mG#fA#mU#fA#mU#fA		
<i>Huntingtin</i>	highly endonuclease sensitive (RNA)	Fig.4.	passenger	fC#mA#fG.mU.A.mA.fA.mG.fA.mG.fA.mU.fU#mA#fA.rU.rU		Cholesterol-TEG
			guide	VPmU#fU#mA.fA.mU.fC.mU.fC.mU.fU.mU.fA.mC#fU#mG#fA#mU#fA#mU#fA		
<i>PPIB</i>	P-FM	Supp.Fig.4.	passenger	fC#mA#fA.mA.fU.mU.fC.mC.fA.mU.fC.mG.fU#mG#fA	Cy3	Cholesterol-TEG
			guide	PmU#fC#mA.fC.mG.fA.mU.fG.mG.fA.mA.fU.mU#fU#mG#fC#mU#fG#mU#fU		
<i>PPIB</i>	VP-FM	Supp.Fig.4.	passenger	fC#mA#fA.mA.fU.mU.fC.mC.fA.mU.fC.mG.fU#mG#fA	Cy3	Cholesterol-TEG
			guide	vPmU#fC#mA.fC.mG.fA.mU.fG.mG.fA.mA.fU.mU#fU#mG#fC#mU#fG#mU#fU		

**Supplementary Table 1. Table describing hsiRNA sequences used in this study.** m = 2' -O-methyl; f = 2' -fluoro; # = phosphorothioate; P = phosphate; VP = 5' -(E)-vinylphosphonate; TEG = triethyl glycol; C7 = 2-aminobutyl-1-3-propanediol



P49184	DNSL1	HUMAN Deoxyribonuclease-1-like 1 OS=Homo sapiens GN=DNASE1L1 PE=1 SV=1	34 kDa
Q7KZF4	SND1	HUMAN Staphylococcal nuclease domain-containing protein 1 OS=Homo sapiens GN=SND1 PE=1 SV=1	102 kDa
O94919	ENDD1	HUMAN Endonuclease domain-containing 1 protein OS=Homo sapiens GN=ENDOD1 PE=1 SV=2	55 kDa
P13489	RINI	HUMAN Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2	50 kDa
P21589	SNTD	HUMAN 5'-nucleotidase OS=Homo sapiens GN=NT5E PE=1 SV=1	63 kDa
P22413	ENPP1	HUMAN Ectonucleotide pyrophosphatase/phosphodiesterase family member 1 OS=Homo sapiens GN=ENPP1 PE=1 SV=2	105 kDa
P09543	CN37	HUMAN Isoform CNPI of 2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Homo sapiens GN=CNP	45 kDa

**Supplementary Table 2. Nucleases detected in extracellular vesicles**