

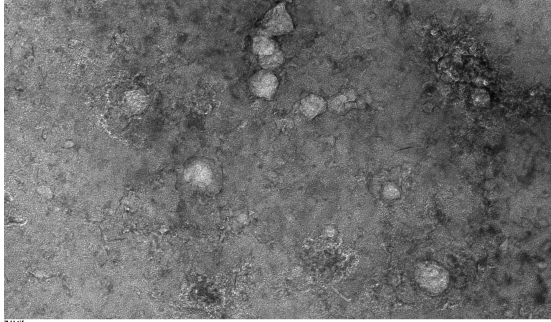
Supporting Information

Addressing MISEV Guidance Using Targeted LC-MS/MS: A Method for the Detection and Quantification of Extracellular Vesicle-Enriched and Contaminant Protein Markers from Blood

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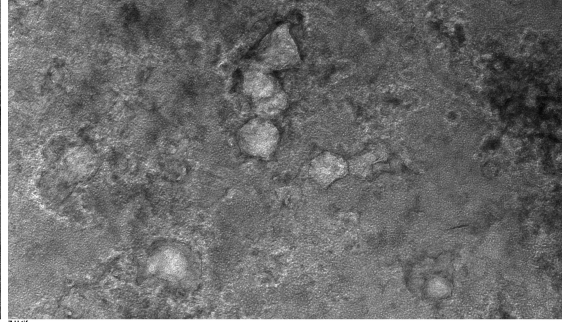
qEV35



Z.U.HJ
Cal: 0.219644 nm/pix

100 nm
Direct Mag: 49000 x
Adelaide Microscopy

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Gamma: 1.00, No Sharpening, Normal Contrast

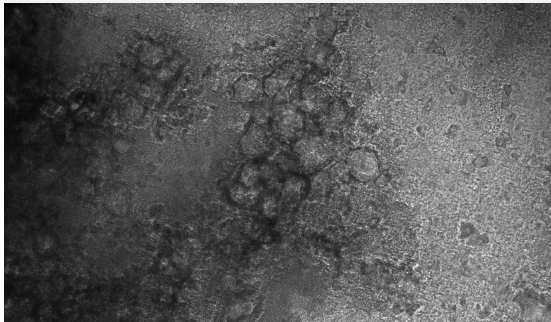


Z.U.HJ
Cal: 0.168273 nm/pix

100 nm
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Adelaide Microscopy

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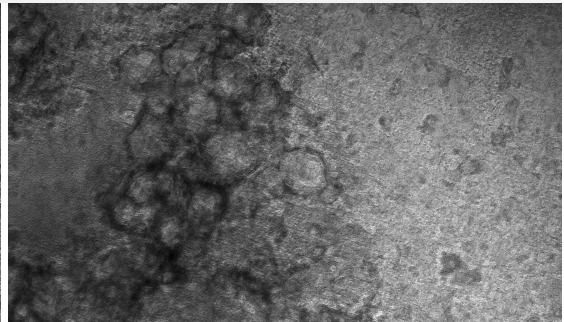
qEV70



Z.U.HJ
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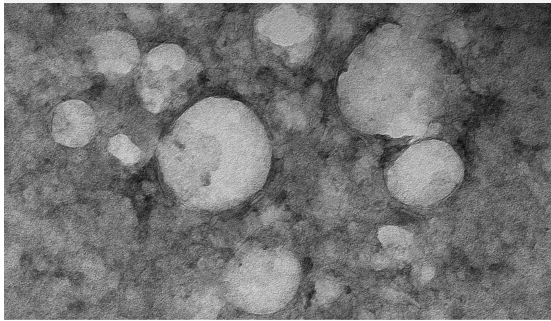


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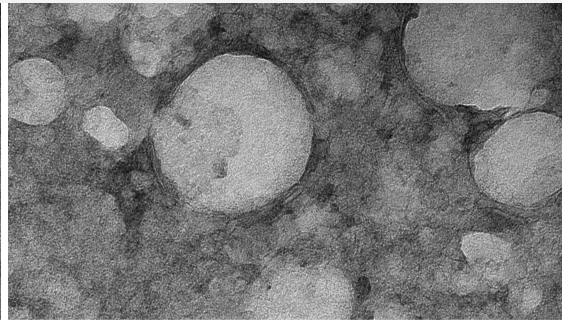
ExoQuick



Z.U.HJ
Cal: 0.219644 nm/pix

100 nm
Direct Mag: 49000 x
Adelaide Microscopy

Camera: MANOSPR115, Exposure: 1200 (ms) x 1 std. frames, Gain: 1, Bin: 1
Gamma: 1.00, No Sharpening, Normal Contrast



Z.U.HJ
Cal: 0.168273 nm/pix

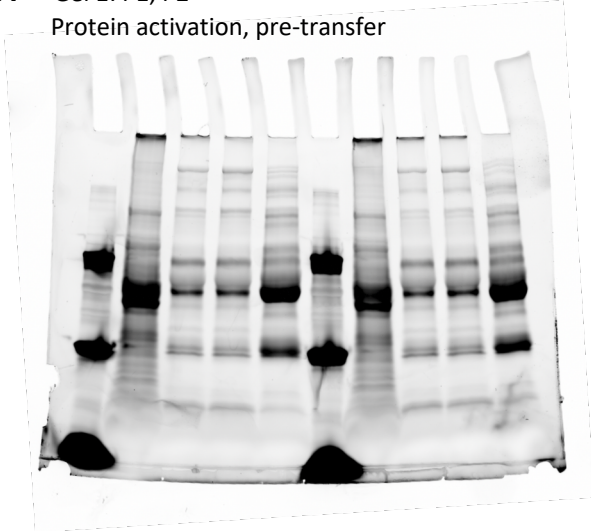
100 nm
Direct Mag: 68000 x
Adelaide Microscopy

Camera: MANOSPR115, Exposure: 1200 (ms) x 1 std. frames, Gain: 1, Bin: 1
Gamma: 1.00, No Sharpening, Normal Contrast

Figure S1. Unedited TEM images at 49,000 x magnification (left panels) and 68,000 x magnification (right panels).

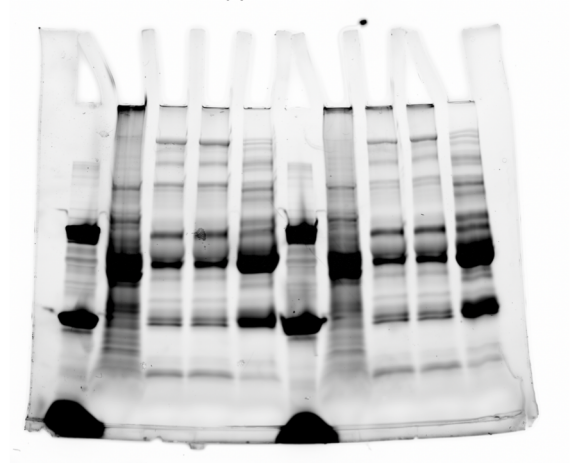
A Gel 1: P1, P2

Protein activation, pre-transfer



B Gel 2: P3, P4

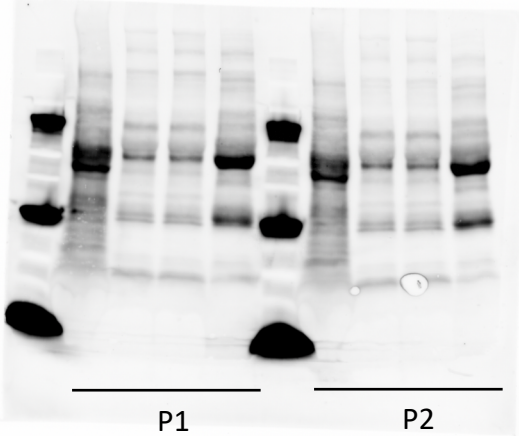
Protein activation, pre-transfer



C Blot 1: P1, P2

Protein transferred to PVDF membrane

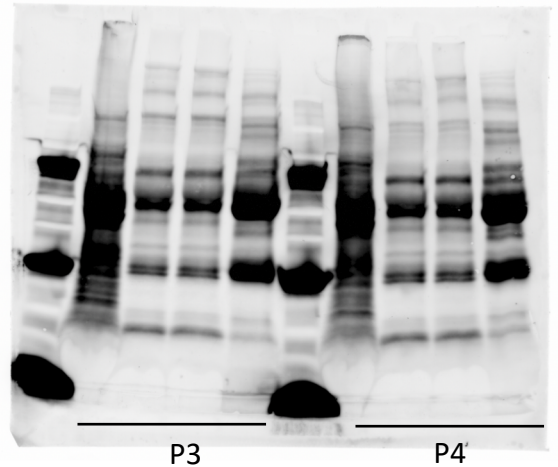
1 2 3 4 5 1 2 3 4 5



D Blot 1: P3, P4

Protein transferred to PVDF membrane

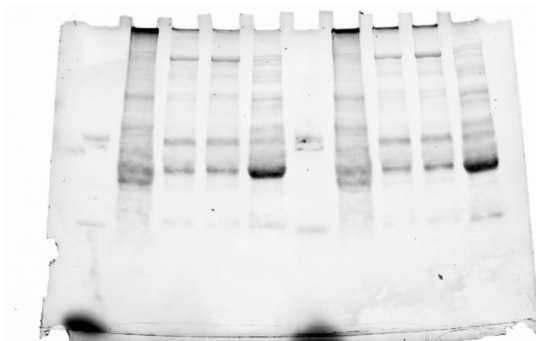
1 2 3 4 5 1 2 3 4 5



Lane 1= MW marker, Lane 2=HLM, Lane 3=qEV35, Lane 4=qEV70, Lane5=ExoQuick

E Gel 1: P1, P2

Post-transfer, 7min mixed MW



F Gel 2: P3, P4

Post-transfer, 7min mixed MW

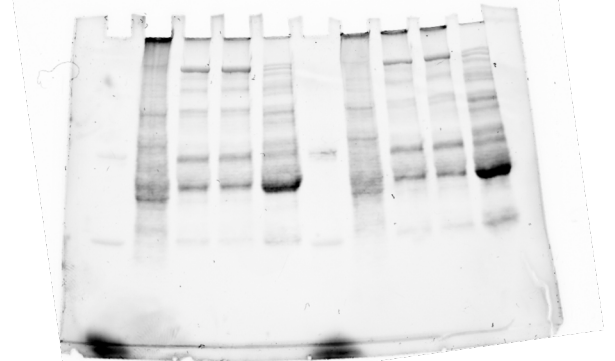


Figure S2. Full images of immunoblot gels showing total protein activation pre-transfer (**A-B**) and post-transfer (**E-F**) onto PVDF membranes (**C-D**). No stain, 35 μ g of lysed HLM or EV protein loaded to each well.

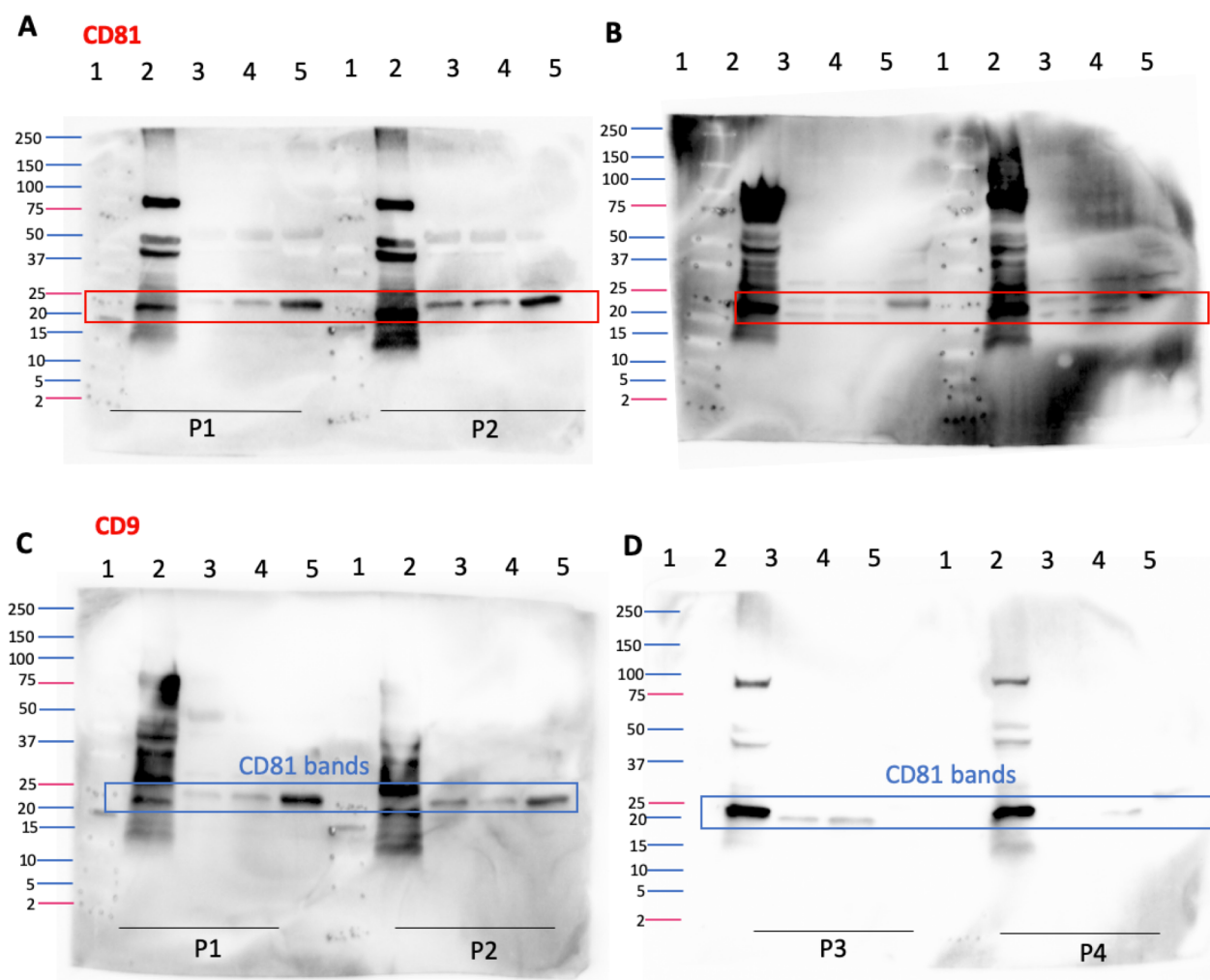


Figure S3. Full images of immunoblots the expression of EV-associated tetraspanins: CD81 (**A-B**) and CD9 (**C-D**) on PVDF. No CD9 band observed, unsuccessful stripping of CD81 antibody. Auto-exposure, 35 μ g of lysed HLM or EV protein loaded to each well. Lane 1= MW marker, Lane 2=HLM, Lane 3=qEV35, Lane 4=qEV70, Lane 5=ExoQuick

CD81 predicted band 25kDa, observed band 22kDa

CD9 predicted band 25kDa, reported 22kDa on manufacturers website. Not observed.

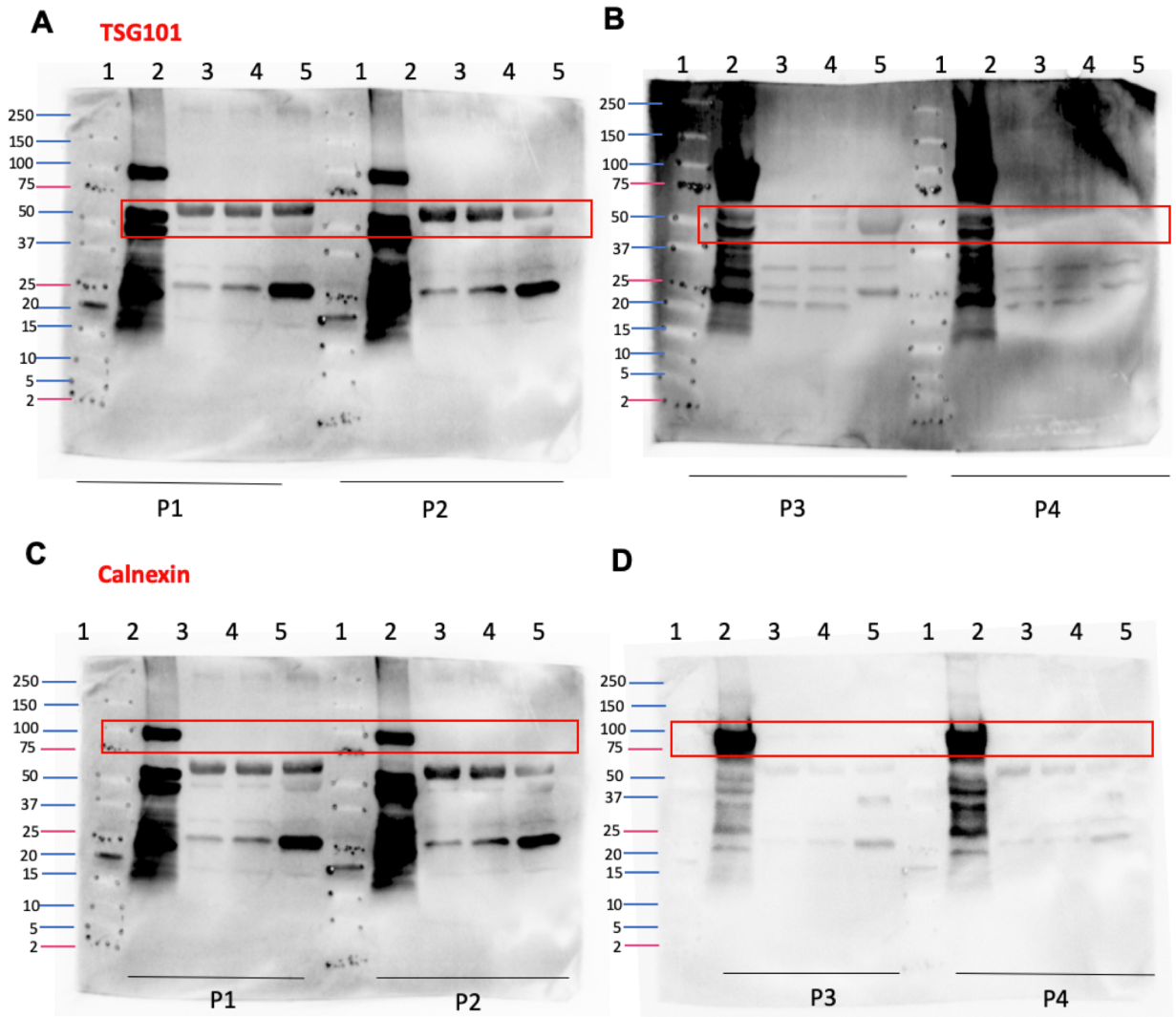


Figure S4. Full images of immunoblots the expression of EV-associated TSG101 (**A-B**) and (**A-B**) and ER marker Calnexin (**C-D**) on PVDF. Auto exposure, 35µg of lysed HLM or EV protein loaded to each well. Lane 1= MW marker, Lane 2=HLM, Lane 3=qEV35, Lane 4=qEV70, Lane5=ExoQuick.

TSG101 predicted band ~50kDa, observed double band at 48-55kDa
 Calnexin 5 predicted band at ~90kDa, observed at ~95kDa

Albumin in Serum vs EV lysates

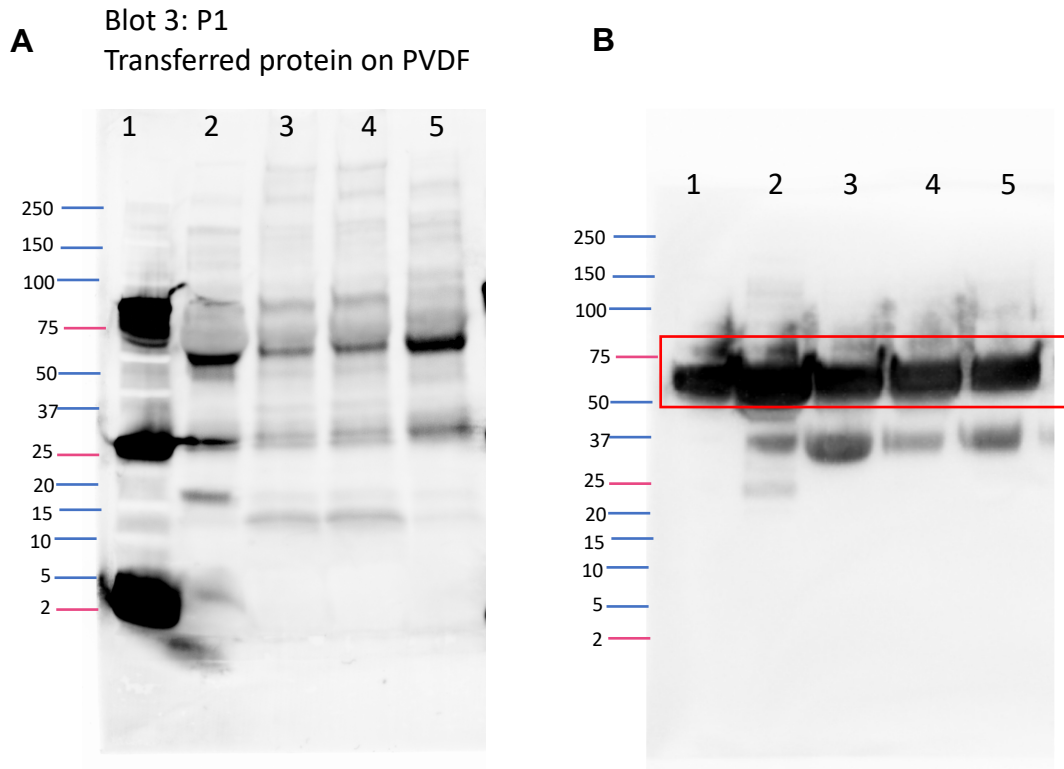


Figure S5. (A) Full images protein on PVDF/stain-free; **(B)** Full image of immunoblot showing Albumin expression at ~ 70 kDa. Lane 1=MW marker, Lane 2=HLM, Lane 3=qEV35, Lane 4=qEV70, Lane5=ExoQuick.; Albumin protein leaked from Lane 2 into Lane 1, therefore, Lane 1 and Lane 2 are considered to be one well in terms of Albumin protein expression. EV protein and matched serum samples from P1 were used; $35\mu\text{g}$ of lysed serum or lysed EV protein loaded to each well.

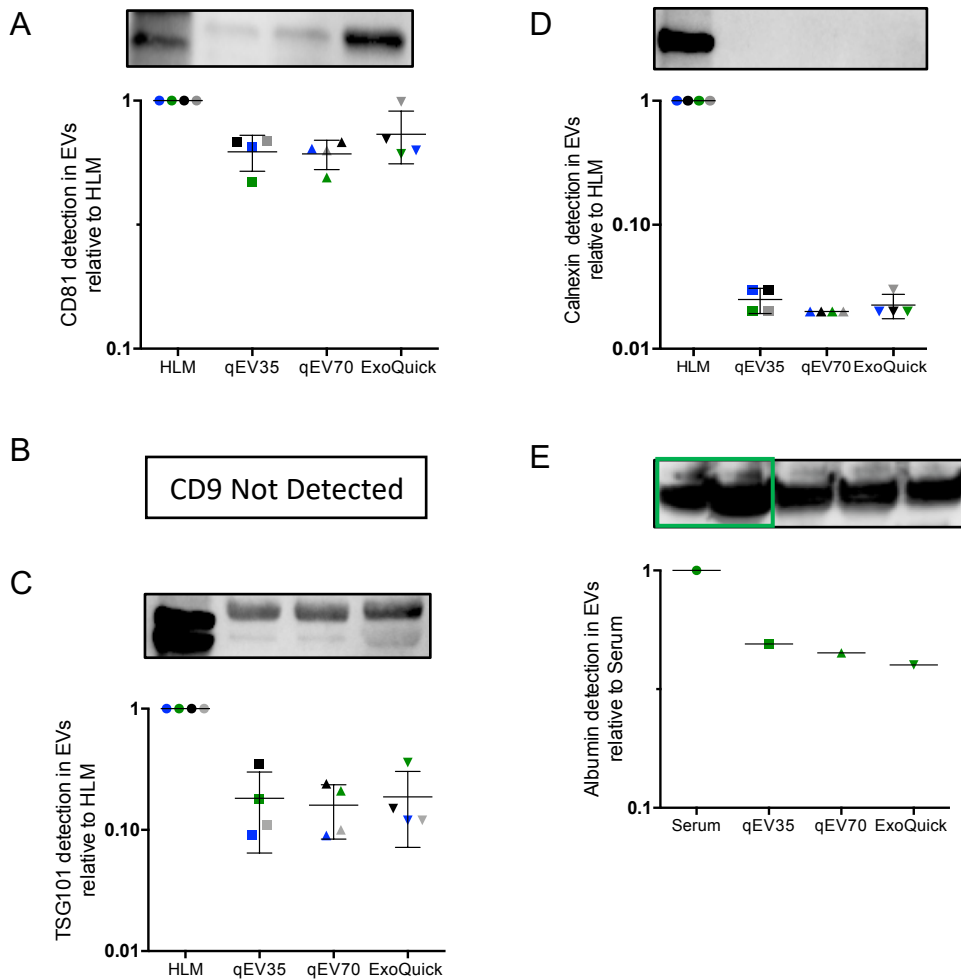
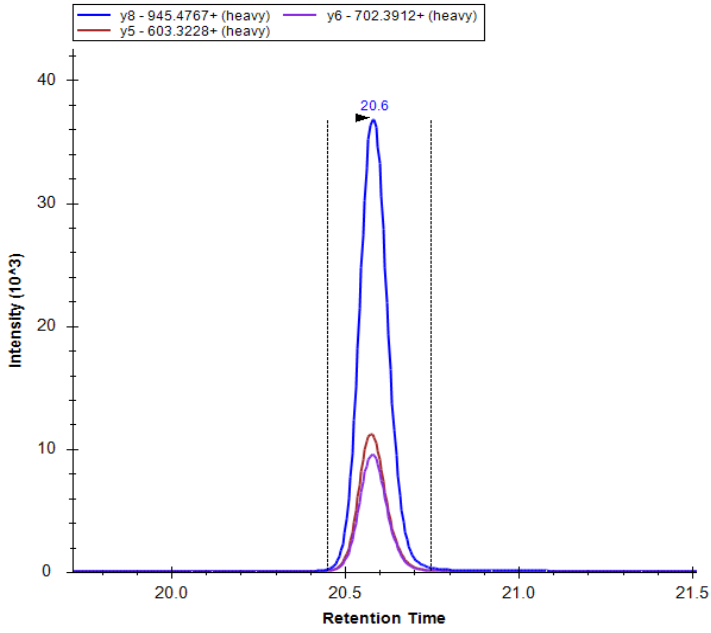


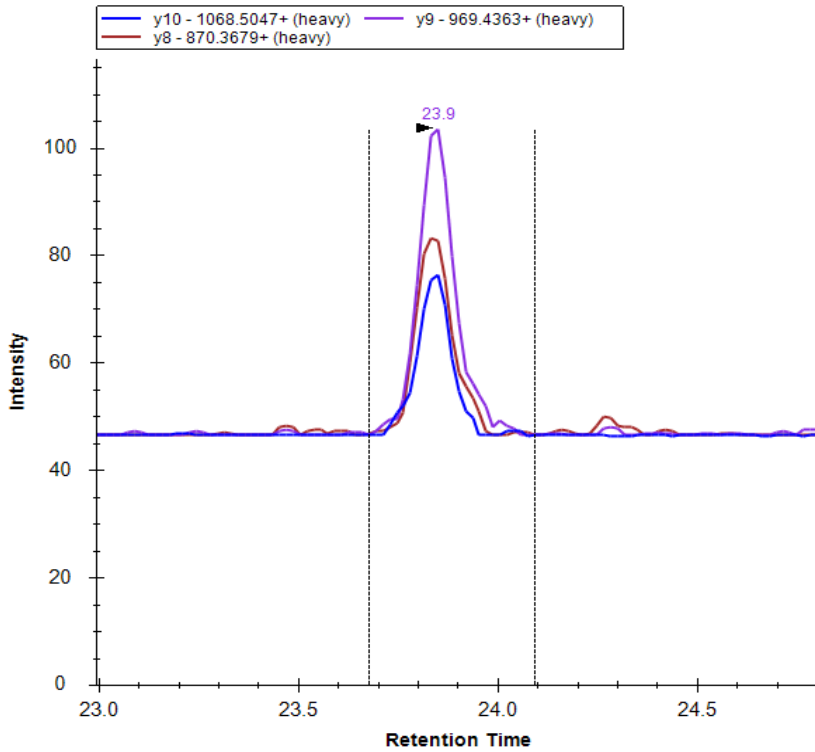
Figure S6. Relative protein quantification by ImageJ. ImageJ obtained values for non-EV controls (HLM) (A-D) and serum (E) were set to one and values obtained for corresponding EV lanes were expressed as fold change relative to non-EV control. The main focus was to compare protein expression between EVs obtained by three isolation methods, however, non-EV controls were included.

Figure S7. Representative chromatograms in Skyline. Stable isotope labelled peptides [ALB: 80 pmol/mL, CD81: 8 pmol/mL, CD9: 0.8 pmol/mL, CANX: 1.6 pmol/mL, TSG101: 1.6 pmol/mL] spiked into EV digests (qEV70) from healthy human serum.

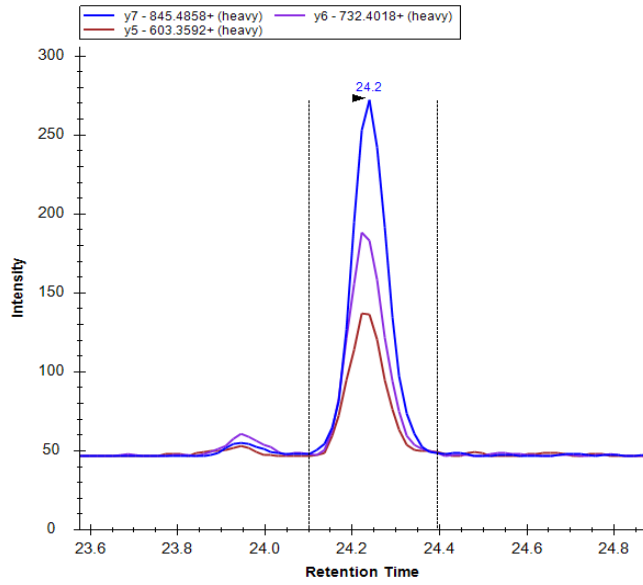
Albumin



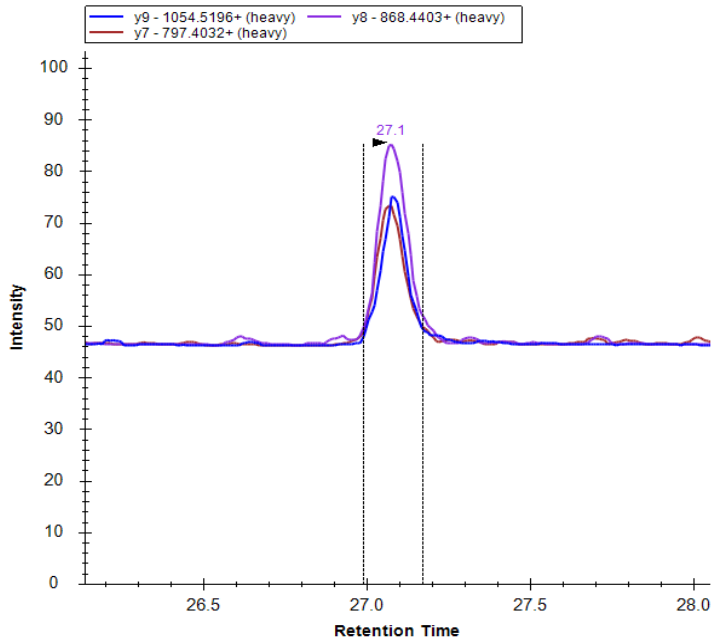
CD81



CD9



CANX



TSG101

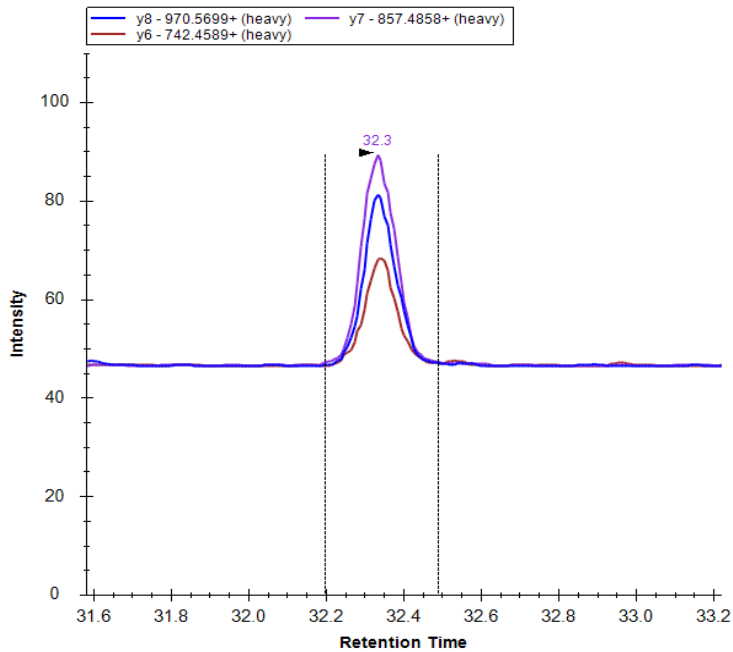


Figure S8. Comparison of sample consumption for immunoblots and LC-MS/MS injections in EV marker analytical workflow.

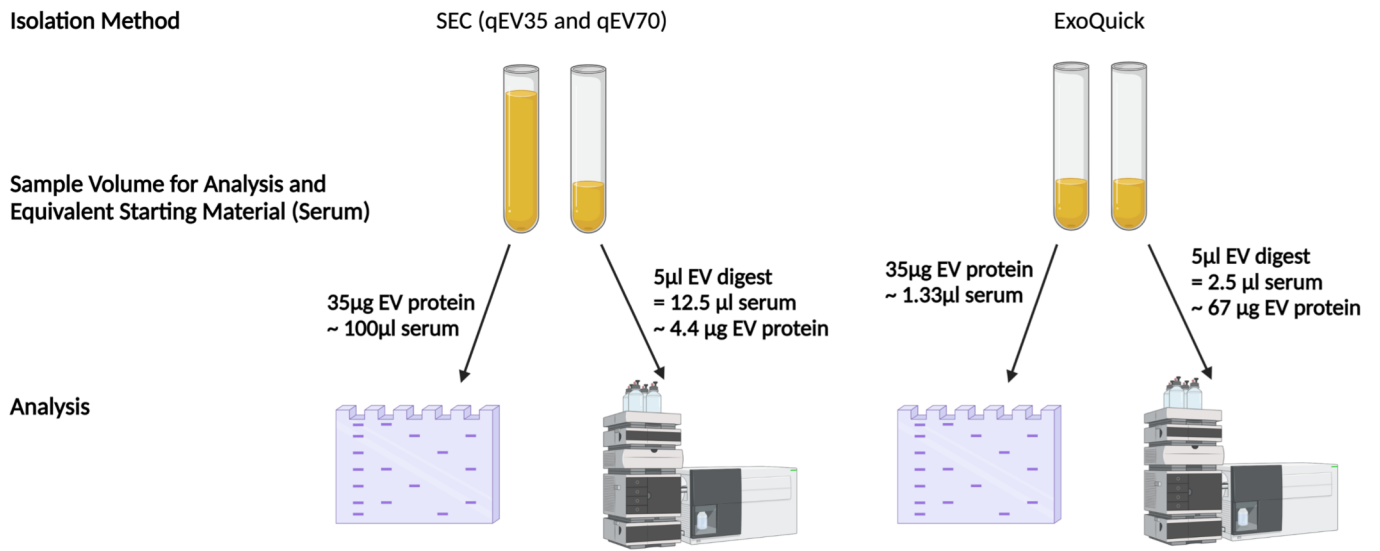


Table S1. Protein concentration of EV samples, measured by micro BCA, and comparison of sample consumptions for immunoblots and LC-MS/MS injections. Immunoblot protein loading = 35 µg. LC-MS/MS injection volume = 5 µl.

Method	Donor	Protein Concentration (µg/µl)	Immunoblots		LC-MS/MS	
			Volume EVs Loaded (µl)	Equivalent Serum Volume (µl)	Protein per Injection (µg)	Equivalent Serum Volume (µl)
qEV35	1	1.71	20.4	102.1	4.28	12.5
	2	1.68	20.8	104.2	4.20	
	3	1.54	22.7	113.6	3.85	
	4	1.41	24.8	124.1	3.53	
qEV70	1	1.92	18.2	91.1	4.80	12.5
	2	1.67	20.9	104.7	4.18	
	3	2.40	14.6	72.8	6.00	
	4	1.85	18.9	94.6	4.63	
ExoQuick	1	139.27	0.25	1.26	69.64	2.5
	2	116.42	0.30	1.50	58.21	
	3	113.22	0.31	1.55	56.61	
	4	168.33	0.21	1.04	84.17	

Table S2. Detection of EV markers in serum and plasma EVs from healthy donors and patients with NAFLD using qEV70 and ExoQuick.

Analyte	qEV70 EVs > LOD (%)			ExoQuick EVs > LOD (%)	
	Healthy Serum (n=9)	Healthy Plasma (n=5)	NAFLD Plasma (n=4)	Healthy Serum (n=9)	NAFLD Serum (n=5)
ALB	100	100	100	100	100
CD81	78	80	100	100	100
CD9	100	100	100	100	100
CANX	100	20	80	78	100
TSG101	89	80	100	100	100