

Quantitative Analysis of Cargo Density in Single-extracellular Vesicles by Imaging

Taketoshi Kajimoto* and Shun-ichi Nakamura*

Department of Biochemistry and Molecular Biology, Division of Biochemistry, Kobe University Graduate School of Medicine, Kobe, Japan

*For correspondence: tkajimot@med.kobe-u.ac.jp; snakamur@kobe-u.ac.jp

[Abstract] Function of extracellular vesicles such as exosomes and microvesicles is determined by their wide ranges of cargoes inside them. Even in the pure exosomes or microvesicles the cargo contents are very heterogeneous. To understand this heterogeneous nature of extracellular vesicles, we need information of the vesicles, which will give us some parameters including vesicle size, number and cargo content of each vesicle. Here, we describe a new method to quantify cargo density in single-extracellular vesicles. Staining of extracellular vesicles in a membrane lipid content-proportionate manner and immobilization of extracellular vesicles onto glass substrate allow us to obtain cargo density information of single-extracellular vesicles. This protocol will be useful to analyze the effects of various drugs or genetic manipulation on vesicle generation and maturation including cargo sorting into heterogeneous extracellular vesicles.

Keywords: Extracellular vesicle, Exosome, Microvesicle, Cargo content, Cargo sorting, Single-extracellular vesicle, Imaging

[Background] Extracellular vesicles are small membrane-bound vesicles released from a variety of cells, comprising exosomes and microvesicles based on the current knowledge of their biogenesis (van Niel *et al.*, 2018). Extracellular vesicles contain various kinds of cargoes including cytosolic proteins, membrane proteins, lipids, microRNAs, and mRNAs. The cargoes are highly heterogeneous in their contents and variety depending on cell types. Extracellular vesicles play a role in erythrocyte maturation, antigen presentation, tumor metastasis, or prion disease and Alzheimer's disease propagation, through cell-to-cell communication by carrying their cargoes from the donor to the recipient cells (Maas *et al.*, 2017).

Exosomes are formed by fusion of multivesicular endosomes (MVEs) with the plasma membranes. On the other hand, microvesicles are formed by shedding of the plasma membranes. Endosomal sorting complexes required for transport (ESCRT) is known to be critical for the generation of lysosomal MVEs. It is reported that ceramide is important for the generation of exosomal MVEs (Trajkovic *et al.*, 2008). The size of exosomes ranges from 30 nm to 100 nm, whereas that of microvesicles is from 50 nm to 1,000 nm (Karpman *et al.*, 2017; van Niel *et al.*, 2018). The mechanism of cargo sorting into extracellular vesicles, especially microvesicles, is still mysterious. For understanding the heterogeneous nature of extracellular vesicles, it is important to get more information on cargo density as well as the number and the size of single extracellular vesicles.

Conventionally, we can quantitate the amount of cargo in total extracellular vesicles obtained from

culture media using western blot analysis. Since extracellular vesicles are highly heterogeneous as mentioned above, we need to get more precise figures of extracellular vesicles for the understanding of vesicle generation and maturation including the cargo sorting. We can solve this issue with detecting “cargo density of single-extracellular vesicles”. Here, we provide details on our new method for getting information of each extracellular vesicle with a cargo density. This protocol provides a useful readout of cargo content in each extracellular vesicle, and promote to unravel detailed molecular mechanism of the cargo sorting and accelerate clinical application of extracellular vesicles.

Materials and Reagents

1. 10 µl, 200 µl, or 1,000 µl tips (Fukae-Kasei, Watson, catalog numbers: 123R-254CS, 123R-755CS, 110-706C)
2. 200 µl gel loading tips (Quality Scientific Plastics, catalog number: 010-Q)
3. 5 ml or 10 ml pipettes (Corning, catalog numbers: 4487, 4488)
4. 50 ml conical centrifuge tubes (BD, Falcon, catalog number: 352070)
5. 1.5 ml microcentrifuge tubes (Ina Optika, BIO-BIK, catalog number: CF-0150)
6. 35 mm or 100 mm culture dish (Corning, catalog numbers: 430165, 430167)
7. 6-well plate (Corning, catalog number: 3516)
8. 35 mm glass bottom culture dish (MatTek, catalog number: P35G-0-14-C)
9. 0.22 µm syringe filters (Millipore, catalog number: SLGVJ13SL)
10. Kimwipes (Nippon Paper Crecias, catalog number: 62011)
11. Aluminum foil (Mitsubishi Aluminum, Nippaku foil)
12. Ultracentrifuge thick wall polycarbonate tubes (Beckman Coulter, catalog number: 349622)
13. HeLa cells (RIKEN cell bank, catalog number: RCB0007)
14. MDA-MB-231 cells (ATCC, catalog number: HTB-26)
15. Dulbecco's modified Eagle medium (DMEM), high glucose, with L-Glutamine and Phenol Red (FUJIFILM Wako Pure Chemical, catalog number: 044-29765)
16. Fetal bovine serum (FBS) (Biowest, catalog number: S1820)
17. Penicillin-streptomycin Mixed Solution (P/S) (NACALAI TESQUE, catalog number: 09367-34)
18. Trypsin-EDTA solution with phenol red (NACALAI TESQUE, catalog number: 32778-34)
19. FuGENE HD transfection reagent (Promega, catalog number: E2311)
20. Opti-MEM, reduced serum medium (Thermo Fisher Scientific, Invitrogen™, catalog number: 31985070)
21. Extracellular vesicles-free fetal bovine serum (Thermo Fisher Scientific, Gibco™, catalog number: A2720803)
22. Total exosome isolation reagent (from cell culture media) (Thermo Fisher Scientific, Invitrogen™, catalog number: 4478359)
23. Paraformaldehyde (PFA) powder (Sigma-Aldrich, catalog number: P6148)
24. Normal goat serum (Jackson Immuno Research, catalog number: 005-000-121)

25. Lipophilic Tracer, DiD (Ex. 644 nm; Em. 665 nm) (Thermo Fisher Scientific, Invitrogen™, catalog number: D7757)
26. Biotinyl Cap PE (PE-biotin) (Avanti Polar Lipids, catalog number: 870277)
27. Biotin-conjugated bovine serum albumin (BSA-biotin) (Sigma-Aldrich, catalog number: A8549)
28. Bovine serum albumin (BSA) (Sigma-Aldrich, catalog number: A8022)
29. Streptavidin (Thermo Fisher Scientific, Invitrogen, catalog number: 434301)
30. Antibody of cargo (e.g., anti-cMET antibody; Cell Signaling, catalog number: 3127S)
31. Secondary antibody (e.g., anti-mouse IgG Alexa Fluor 594; Cell Signaling, catalog number: 8890S)
32. Plasmid DNA of cargo (e.g., constructing plasmid DNA of GFP- or mCherry-fused cargo protein)
33. Sodium chloride (NaCl) (NACALAI TESQUE, catalog number: 31319-45)
34. Potassium chloride (KCl) (Wako, catalog number: 163-03545)
35. di-Sodium hydrogenphosphate (Na₂HPO₄) (NACALAI TESQUE, catalog number: 31723-35)
36. Potassium dihydrogenphosphate (KH₂PO₄) (NACALAI TESQUE, catalog number: 28721-55)
37. NaOH (NACALAI TESQUE, catalog number: 31511-05)
38. Phosphate-buffered saline (PBS) (see Recipes)
39. 4% PFA (see Recipes)

Equipment

1. Glass vial (Maruemu, catalog number: 5-115-02)
2. Forceps
3. Pipettes (Gilson, models: PIPETMAN P10, P20, P200, P1000)
4. Pipettor (Drummond Scientific, model: Pipet-Aid XP)
5. Polypropylene microtube storage rack, 80 positions
6. Test tube rack (Sanwakaken, catalog number: SS30-10)
7. Clean hood (NK System, model: VH-1300BH-2A)
8. CO₂ incubator (Forma Scientific, model: 3158)
9. Confocal laser scanning microscope (Zeiss, model: LSM 510 Meta)
10. Transmitted Light Microscope (Olympus, model: IMT-2)
11. Ultracentrifuge (Beckman Coulter, model: Optima TL)
12. Ultracentrifuge rotor (Beckman Coulter, model: TLA-100.3)
13. Refrigerated microcentrifuge (Tomy Seiko, model: MRX-151)
14. Vortex mixer (Scientific Industries, model: Vortex-Genie 2)
15. Refrigerator, 4 °C
16. Freezer, -20 °C
17. Cold room, 4 °C

Software

1. LSM510 v4.2 operating software (Zeiss)
2. ImageJ (<https://imagej.nih.gov/ij/>)
3. Excel for Mac 2011 (Microsoft)

Procedure

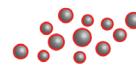
A. Overall procedure

Overall procedure for analyzing cargo density of single-extracellular vesicles is shown in Figure 1.

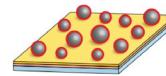
Production and purification of extracellular vesicles from cell culture supernatants or tissues.



Labeling of extracellular vesicle membrane with biotin-DOPE and lipophilic carbocyanine fluorescent dye.



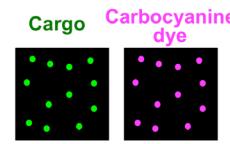
Immobilization of the labeled extracellular vesicles on avidin-coated glass surface.



Immunofluorescence staining of cargo of interest in the immobilized extracellular vesicles with specific antibody as like immunostaining of cells.



Acquisition of fluorescent images of cargo and lipophilic carbocyanine dye of each extracellular vesicle using fluorescence microscope.



Quantification of cargo and lipophilic carbocyanine dye of each extracellular vesicle using ImageJ software and calculation of cargo density using Excel software.

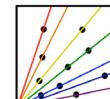


Figure 1. Overall procedure for detecting cargo density in single-extracellular vesicles

B. Production and purification of extracellular vesicles

1. Cell culture and transfection for producing released extracellular vesicles
 - a. HeLa cells or MDA-MB-231 cells are plated onto 35 mm culture dish or 6-well plate and maintained in Dulbecco's modified Eagle medium (DMEM) containing 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin (P/S) at 37 °C in 5% CO₂.
 - b. When cells are at 70% confluence, remove the medium and wash the cells 2 times each with 2 ml of 37 °C DMEM without FBS and P/S.
 - c. Replace medium with 1.2 ml of 37 °C DMEM containing 10% extracellular vesicles-free fetal bovine serum and 1% P/S.
 - d. Incubate at 37 °C in 5% CO₂ for 24 to 48 h.

Note: In case of analyzing the exogenous cargo (e.g., GFP- or mCherry-fused protein), transient transfection needs to be carried out using lipid-based transfection reagent (e.g., FuGENE HD) 24 to 48 h before the replacement with the extracellular vesicles-free medium.

2. Purification of extracellular vesicles

a. Flowchart of this part is shown in Figure 2.

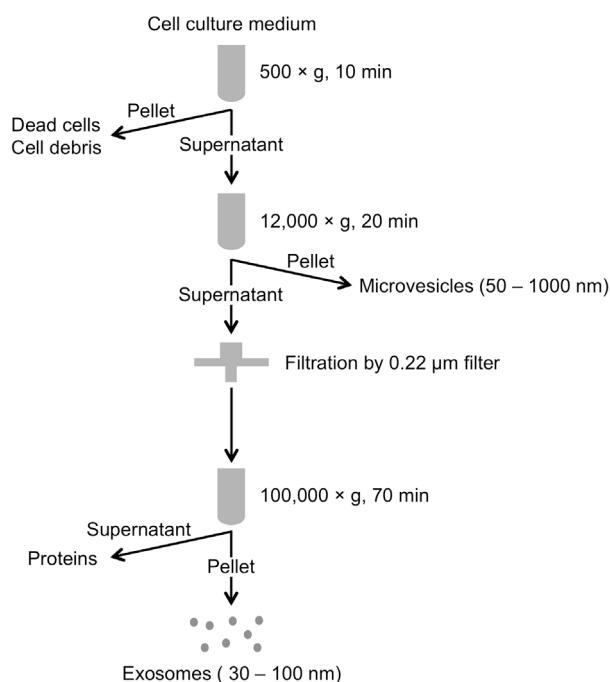


Figure 2. Flowchart of isolation of extracellular vesicles (exosomes and microvesicles) by differential centrifugation

- Cell culture media (1.2 ml) from 90%-95% confluent cells is collected in 1.5 ml microcentrifuge tube.
- Centrifuge at 500 \times g at 4 °C for 10 min to remove the cell debris using a refrigerated microcentrifuge.
- Collect all 1.2 ml supernatant and transfer to a new 1.5 ml microcentrifuge tube.
- Centrifuge at 12,000 \times g at 4 °C for 20 min using a refrigerated microcentrifuge to separate microvesicles and others.
- Collect the upper 1 ml of supernatant is directly purified by filtration through a 0.22 µm pore filter and transfer into the ultracentrifuge thick wall polycarbonate tube. Approximately 70 µl of sample is lost after the filtration.
- The pellet from Step B2e is diluted in 200 µl of phosphate-buffered saline (PBS) (Microvesicle-enriched fraction).
- Ultracentrifuge the filtrated supernatant using ultracentrifuge (Optima TL, rotor: TLA-100.3) at 100,000 \times g at 4 °C for 70 min.
- Remove supernatant (approximately 930 µl) carefully.

- j. Dilute the pellet in 200 μ l of PBS (Exosome-enriched fraction).

Note: Alternatively exosomes can be purified using total exosome isolation reagent according to the manufacturer's instructions.

C. Preparation of functionalized glass surface

1. Flowchart of this part is shown in Figure 3.

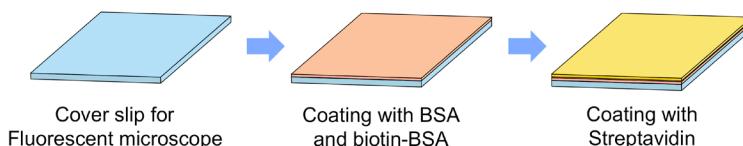


Figure 3. Flowchart of preparation of functionalized glass surface

2. Prepare BSA/BSA-biotin mixture with mixing 5 μ l of 10 M BSA and 5 μ l of 1 M BSA-biotin in 490 μ l of PBS.
3. Cover the glass surface of glass bottom 35 mm culture dish with 400 μ l of the BSA/BSA-biotin mixture.
4. Incubate at room temperature for 20-30 min.
5. Prepare 25 mM streptavidin solution by diluting 8 μ l of 1 M streptavidin in 312 μ l of PBS.
6. Remove excess BSA/BSA-biotin by gently rinsing the glass surface once with 1 ml of PBS.
7. Cover the glass surface with 300 μ l of the streptavidin solution.
8. Incubate at room temperature for 10 min.
9. Gently rinse the BSA/BSA-biotin/streptavidin-coated glass surface twice each with 1 ml of PBS.

D. Labeling of extracellular vesicles

1. Transfer the isolated extracellular vesicles (200 μ l total) into a glass vial.
2. Add 2 μ l of 10 mg/ml DOPE-biotin into the glass vial and suspend to mix.
3. Incubate at room temperature for 5 min.
4. Add 2 μ l of 1 mg/ml DiD into the glass vial and suspend to mix.

Note: Because DiD staining is the most critical step for calculating the cargo density, it is important to take exact amount of DiD.

5. Spin down the glass vial in a 50 ml conical tube at 200 $\times g$ at room temperature and take it out with forceps.
6. Then incubate at room temperature for 15 min in the dark (Figure 4).



Figure 4. Labeling of extracellular vesicles with DOPE-biotin and DiD in glass vial

E. Immobilization of extracellular vesicles on glass surface

1. The DOPE-biotin- and DiD-labeled extracellular vesicles are applied onto the functionalized glass surface using gel loading long tip (Figure 5).

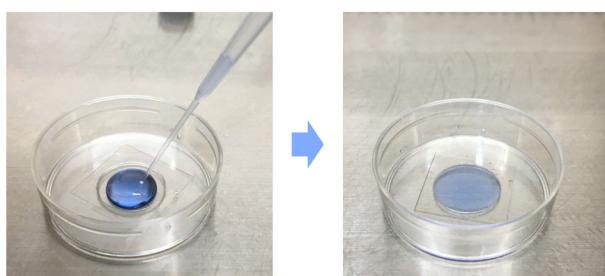


Figure 5. Immobilization of extracellular vesicles on functionalized glass surface.

DOPE-biotin- and DiD-labeled extracellular vesicle solution is applied onto functionalized glass surface and made it flat against the surface tension.

2. Leave to stand at 4 °C for 2-3 h in dark condition.
3. Rinse twice each with 2 ml of PBS.

Note: In case of analyzing the transiently expressed exogenous cargo (e.g., GFP- or mCherry-fused protein), skip to “(G) Imaging” immediately.

F. Immunofluorescence staining of endogenous cargo in extracellular vesicles

1. Fix immobilized extracellular vesicles with 4% PFA for 5 min in dark condition.
2. Rinse extracellular vesicles twice with 2 ml of PBS and pipette out all solution.
3. Add 150 µl of PBS containing 10% normal goat serum and incubate at room temperature for 30 min in dark condition.
4. Rinse extracellular vesicles once with 2 ml of PBS and pipette out all solution.
5. Add 150 µl of anti-cargo antibody (e.g., anti-cMet antibody) in PBS and incubate at room temperature for 60-90 min in dark condition.
6. Rinse extracellular vesicles once with PBS and pipette out all solution.
7. Add 150 µl of fluorescent-dye conjugated secondary antibody (e.g., Alexa 488 or Alexa 594) in PBS and incubate at room temperature for 30 min in dark condition.
8. Rinse extracellular vesicles twice each with 2 ml of PBS.

9. Store at 4 °C in dark condition, and observe in 2-3 days.

G. Imaging of fluorescence-labeled cargo and DiD using fluorescence microscope

Note: For this imaging part we will show about the exosome sample as a representative.

1. Turn on the confocal laser scanning microscope (Zeiss, LSM 510 Meta).
2. Open the Zeiss LSM510 v4.2 operating software.
3. Turn on the appropriate laser (e.g., Argon laser for excitation of GFP or Alexa 488, 543 nm wavelength He-Ne laser for excitation of mCherry or Alexa 594, 633 nm wavelength He-Ne laser for excitation of DiD).
4. Select 63x objective lens.
5. Set the glass bottom 35 mm culture dish with immobilized and cargo-labeled extracellular vesicles on the specimen holder.
6. Carefully find the glass surface using visible light under eyepiece viewing.
7. Change the observation status to laser scanning.
8. In the “Scan Control panel”, “Mode”, set the level of zoom to x1 (Default).
9. In the “Scan Control panel”, “Channels”, set the pinhole to “200”.
10. Adjust the focus with DiD channel using “Fast XY”.
11. Collect averaged images (e.g., number is selected to 4 for the average) of DiD and fluorescent cargo (e.g., GFP- or mCherry-fused cargo or immunostained cargo with anti-mouse IgG Alexa 488 or anti-mouse IgG Alexa 594 as secondary antibody) and save them as lsm format.
12. The acquired images of exosomes are shown in Figure 6 (in case of endogenous cargo) or in Kajimoto *et al.* (2013) or (2018) (in case of exogenous GFP- or mCherry-fused cargo).

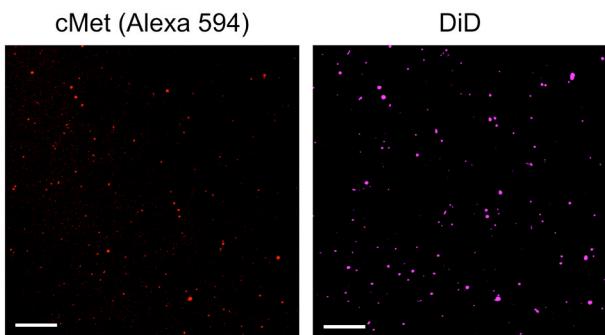


Figure 6. Fluorescent images of cargo (cMet-Alexa 594) and DiD in single-exosomes.
Exosomes released from MDA-MB-231 cells are stained with anti-cMet antibody with Alexa 594 secondary antibody and DiD. Scale bars, 10 μm.

Data analysis

1. Open the ImageJ software.
2. Go to “Adjust” → “Set Measurements”, and check “Area”, “Mean gray value”, and “Integrated Density”, then click “Ok” button.

3. Open the acquired images (lsm format file, here using the data of Figure 6 as example) with drag-and-drop to ImageJ icon (Figure 7A).
4. Select DiD image (Channel 2) (Figure 7B).
5. Set “Image” → “Type” → “8-bit” and confirm that the image is converted to greyscale (Figure 7C).
6. Set “Edit” → “Invert” → “No (against the question “Process all 3 images?”)” for inverting the image of DiD (Figure 7D).
7. Use “Image” → “Adjust” → “Threshold” to highlight the extracellular vesicles (Figure 7E). Basically using “Auto” would be working well. If necessary, adjust the threshold to the suitable level to count.
8. Go to “Analyze” → “Analyze Particles”. There are some options in the “Analyze Particles” window (Figure 7F). Basically use default setting (Size: 0-Infinity, Circularity: 0.00-1.00, Show: Nothing) and just check “Display results” and “Add to Manager”. If there are too many small noises or you want to exclude particles based on size, adjust the numbers of “Size”.

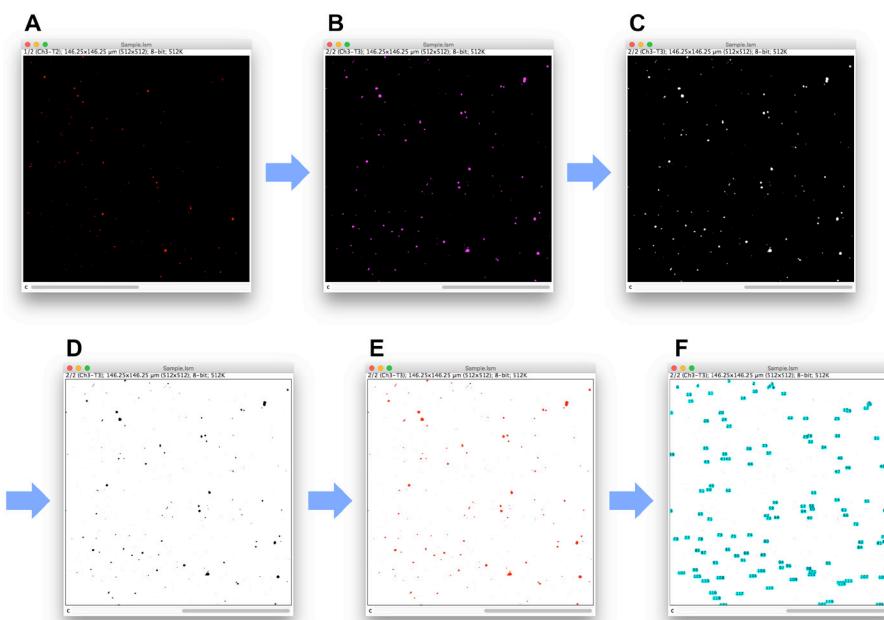


Figure 7. Conversion of acquired images of cargo (cMet-Alexa 594) and DiD to be calculated with ImageJ

9. In the “ROI Manager” window, go to “More” → “Multi Measure”, and just check “Measure All 2 Slices” and click “Ok” button.
10. Get “Area”, “Mean gray value (Mean)”, and “Integrated Density (IntDen)” information of cargo (in channel 1) and DiD (in channel 2) of each extracellular vesicle (Figure 8).

Channel 1					Channel 2						
cMet (Alexa 594)					DiD						
Results					Results						
Area	Mean	IntDen	RawIntDen	Ch	Area	Mean	IntDen	RawIntDen	Ch		
1	1.061	84.769	89.913	1102	1	123	1.061	37.615	39.988	489	2
2	0.326	0.250	0.082	1	124	0.326	101.500	33.126	406	2	
3	0.490	3.667	1.795	22	125	0.490	24.667	12.075	148	2	
4	0.653	19.750	12.891	158	1	126	0.653	139.875	91.303	1119	2
5	0.163	0	0	1	127	0.163	152	24.800	304	2	
6	0.163	24	3.916	48	1	128	0.163	152.500	24.885	305	2
7	0.082	0	0	1	129	0.082	105	8.567	105	2	
8	0.082	1	0.082	1	130	0.082	142	11.588	142	2	
9	1.469	9.278	13.626	167	1	131	1.469	49.944	73.350	899	2
10	0.163	0	0	1	132	0.163	134	21.000	268	2	
11	0.245	16	3.916	48	1	133	0.245	93.333	22.845	280	2
12	0.163	0	0	1	134	0.163	144	23.498	288	2	
13	0.326	4.750	1.550	19	1	135	0.326	102.750	33.534	311	2
14	0.326	8.250	2.692	33	1	136	0.326	77	25.130	308	2
15	1.387	178.059	246.975	3027	1	137	1.387	43	59.643	731	2
16	5.548	20.544	113.982	1397	1	138	5.548	27.794	154.206	1890	2
17	0.408	0.400	0.163	2	1	139	0.408	131.200	53.523	656	2
18	1.305	0.625	0.816	10	1	140	1.305	52.375	68.373	838	2
19	0.734	1	0.734	9	1	141	0.734	50.778	37.287	457	2
20	2.529	64.710	163.671	2006.000	1	142	2.529	26.774	67.720	830	2
21	0.326	11.500	3.753	46	1	143	0.326	41	13.381	164	2

Figure 8. Results of “Area”, “Mean gray value (Mean)”, and “Integrated Density (IntDen)” of cargo (cMet-Alexa 594) and DiD in each exosome quantified by ImageJ. The total number of exosomes analyzed in this image is 122.

11. Transfer the data sets to Excel by copy and paste.
12. Obtain “Mean gray value (Mean)” information of background of cargo and DiD channels respectively.
 - a. Open the acquired images again.
 - b. Select vesicle-free dark area with “Rectangular” or “Oval” tool.
 - c. Go to “Analyze” → “Measure”, and get “Area”, “Mean gray value (Mean)”, and “Integrated Density (IntDen)” information of both cargo and DiD channels.
 - d. Transfer the data sets to Excel with copy and paste.
13. Calculate background of cargo and DiD of each extracellular vesicle respectively.
Calculating formula: $\text{Background} = (\text{Mean gray value [Mean]} \text{ of background}) * (\text{Area} \text{ of extracellular vesicles})$
Note: If the calculating result becomes less than zero “0”, it would be fixed to zero “0”. This negative value means undetectable level of cargo or DiD.
14. Calculate cargo density of each extracellular vesicle.
Calculating formula:
$$\text{Cargo density} = [(\text{Integrated Density (IntDen)} \text{ of cargo}) - (\text{Background of cargo})] / [(\text{Integrated Density (IntDen)} \text{ of DiD}) - (\text{Background of DiD})]$$
15. Then make presentation diversely depending on the purpose of the study as in Kajimoto *et al.* (2013) or (2018).

Notes

1. This protocol is applicable to all samples including adherent and non-adherent cells, plasma, and tissues. We used adherent cells as an example to describe the detailed protocol.
2. The amount of extracellular vesicles varies in different cell types. In most cases, small-scale culture in 35 mm dishes or 6-well plates is enough, but in some cases, a large-scale culture in

60 mm or 100 mm dishes is required.

3. If using the size- and density- dependent purification method for extracellular vesicles like differential centrifugation, it is important to use healthy cells for avoiding the contamination of apoptotic body or cell debris.
4. Detailed conditions of Procedures C, D, and E are basically according to the previously reported fluorescence-based assay to measure single liposomes (Hatzakis *et al.*, 2009).

Recipes

1. Phosphate-buffered saline (PBS)
 - a. Mix 1.37 M NaCl, 27 mM KCl, 100 mM Na₂HPO₄, and 18 mM KH₂PO₄ in 900 ml H₂O
 - b. Adjust to a final pH of 7.4
 - c. Add H₂O to 1,000 ml
 - d. Autoclave to store (This is 10x PBS stock)
 - e. Dilute 1:10 with H₂O to make 1x PBS
2. 4% PFA
 - a. Prepare 30 ml of H₂O in 50 ml conical tube and heat to 60 °C
 - b. Add 1.6 g of PFA powder in the 30 ml of H₂O, 60 °C in the fume hood and mix well
 - c. Add 4 µl of 10 N NaOH and mix well until the solution gets clear
 - d. Add 4 ml of 10x PBS
 - e. Add H₂O to 40 ml and mix
 - f. Cool, store at 4 °C in dark condition, and use in two weeks

Acknowledgments

This protocol was adapted from procedures published in Kajimoto *et al.* (2013) and (2018). This work was supported by grants from the JSPS KAKENHI, the Uehara Memorial Foundation, the Osaka Medical Research Foundation, and the Nakatani Foundation.

Competing interests

There are no conflicts of interest.

References

1. Hatzakis, N. S., Bhatia, V. K., Larsen, J., Madsen, K. L., Bolinger, P. Y., Kunding, A. H., Castillo, J., Gether, U., Hedegard, P. and Stamou, D. (2009). [How curved membranes recruit amphipathic helices and protein anchoring motifs](#). *Nat Chem Biol* 5(11): 835-841.
2. Kajimoto, T., Okada, T., Miya, S., Zhang, L. and Nakamura, S. (2013). [Ongoing activation of](#)

[sphingosine 1-phosphate receptors mediates maturation of exosomal multivesicular endosomes.](#) *Nat Commun* 4: 2712.

3. Kajimoto, T., Mohamed, N. N. I., Badawy, S. M. M., Matovelo, S. A., Hirase, M., Nakamura, S., Yoshida, D., Okada, T., Ijuin, T. and Nakamura, S. I. (2018). [Involvement of G \$\beta\$ \$\gamma\$ subunits of Gi protein coupled with S1P receptor on multivesicular endosomes in F-actin formation and cargo sorting into exosomes.](#) *J Biol Chem* 293(1): 245-253.
4. Karpman, D., Stahl, A. L. and Arvidsson, I. (2017). [Extracellular vesicles in renal disease.](#) *Nat Rev Nephrol* 13(9): 545-562.
5. Maas, S. L. N., Breakefield, X. O. and Weaver, A. M. (2017). [Extracellular vesicles: Unique intercellular delivery vehicles.](#) *Trends Cell Biol* 27(3): 172-188.
6. Trajkovic, K., Hsu, C., Chiantia, S., Rajendran, L., Wenzel, D., Wieland, F., Schwille, P., Brugger, B. and Simons, M. (2008). [Ceramide triggers budding of exosome vesicles into multivesicular endosomes.](#) *Science* 319(5867): 1244-1247.
7. van Niel, G., D'Angelo, G. and Raposo, G. (2018). [Shedding light on the cell biology of extracellular vesicles.](#) *Nat Rev Mol Cell Biol* 19(4): 213-228.