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Phase separation in lipid-based nanoparticles: exploring the nano-bio interface

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Appendix I

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Protocol for *in situ* formation of gold nanoparticles in phase-separated liposomes

Herein, a protocol to achieve *in situ* formation of gold nanoparticles (AuNPs) inside the phase-separated PAP3 liposome core, is introduced. The protocol is adapted and modified from a previously described method.¹

Materials

Tris(hydroxymethyl)aminomethane (Tris), Trisodium Citrate Dihydrate ($\text{HOC}(\text{COONa})(\text{CH}_2\text{COONa})_2 \cdot 2\text{H}_2\text{O}$), Gold(III) chloride trihydrate ($\text{HAuCl}_4 \cdot 3\text{H}_2\text{O}$) and 1,2-distearoyl-sn-glycero-3-phosphatidylcholine (DSPC) were purchased from Sigma Aldrich. DOaG was synthesized as described in Chapter 2 and in reference ². $\text{HAuCl}_4 \cdot 3\text{H}_2\text{O}$ was stored in a dark place.

Method

1. Tris buffer was prepared at 10 mM concentration and pH was adjusted to 7.4 with 1M HCl.
2. Using a volumetric flask (50 mL), an aqueous stock solution of 50.1 mM $\text{HAuCl}_4 \cdot 3\text{H}_2\text{O}$ was prepared immediately after opening the manufacturer's bottle (using all the content from the bottle). From this stock, a stock of 5 mM was prepared.
3. Using a volumetric flask (50 mL), an aqueous stock solution of 20.4 mM trisodium citrate dihydrate was prepared.
4. Individual lipids as stock solutions of DSPC and DOaG (10 mM) in chloroform, were combined in a glass vial to 1:1 molar ratio and dried to a thin film, first under N_2 stream, then >1 h under vacuum.
5. Lipid films were redissolved in 50 μL absolute ethanol with gentle vortexing if necessary, to a total lipid concentration of 50 mM.
6. Using the stock solutions prepared in steps 2 and 3 an aqueous solution of HAuCl_4 : Sodium Citrate was prepared at a 1 : 4.08 ratio (Turkevich solution).³ Two concentrations of HAuCl_4 have been successfully used: 1.75 mM or 2.5 mM. Turkevich solution is prepared fresh every time and is used immediately after preparation to ensure prevention of premature AuNP formation prior to liposome formation.
7. Using a (pre-heated) glass micro-syringe (Hamilton, syringe series 700, volume 50), 35.7 μL of the ethanolic lipid solution (warm after submersion in a water bath of 50 °C for 5 sec) was rapidly injected in a glass vial containing 2.5 mL of Turkevich solution (1:71 v/v; EtOH:Turkevich solution) submerged in a 50 °C water bath (freshly made and submerged only for ~2-3 min before injection), under constant vigorous stirring (650 rpm – stirring bar dimensions: 12 x 4 x 4 cm), to form large unilamellar vesicles.
8. Liposomal solution (2.5 mL) was immediately passed through a size exclusion chromatography column (Illustra™ NAP25, GE Healthcare, Thermofischer Scientific)

equilibrated with Tris buffer 10 mM pH = 7.4, to replace the unencapsulated Turkevich solution with buffer.

9. Liposomes encapsulating Turkevich solution in their core were then transferred in an eppendorf tube (1.5 mL) and placed in a thermomixer with a stable temperature at 75 °C for 10 min to ensure initiation of gold reduction from Au⁺³ to Au⁰ and subsequent nucleation and growth of AuNPs.
10. Liposomes containing formed AuNPs were transferred to a dialysis tube (Float-a-lyzer® G2, 1 mL capacity, 1000 kDa MWCO, Spectrum labs) and dialyzed against Tris buffer 10 mM pH = 7.4 overnight at 4 °C, to ensure complete removal of ethanol.
11. The hydrodynamic diameter and polydispersity index (PDI) of liposomes containing AuNPs were characterized by Dynamic Light Scattering (DLS) (Malvern Zetasizer Nano ZS). DLS measurements were carried out at room temperature in 10 mM Tris buffer (pH = 7.4) at a total lipid concentration of approx. 100 µM. Reported DLS measurements are the average of three measurements.
12. Liposomes containing AuNPs (3 µL) were applied to a freshly glow-discharged carbon 200 mesh Cu grid (Lacey carbon film, Electron Microscopy Sciences, Aurion, The Netherlands). Grids were blotted for 3 sec at 99% humidity in a Vitrobot plunge-freezer (FEI Vitrobot™ Mark III, Thermo Fisher Scientific). Cryo-Transmission Electron Microscopy (cryo-TEM) images were collected on a Talos L120C (NeCEN, Leiden University) or a TITAN (Eindhoven University of Technology) operating respectively at 120 kV or 300kV. In the case of Talos, images were recorded manually at a nominal magnification of 17500x or 36000x yielding respectively a pixel size of 5.87 or 2.89 ångström (Å) at the specimen. In the case of TITAN, images were recorded manually at a nominal magnification of 24000x or 30000x yielding a pixel size of 3.87 or 2.81 ångström (Å) at the specimen, respectively.
13. Cryo-electron tomography (cryo-ET) was performed on a Talos L120C (NeCEN, Leiden University) operating at 120kV. Tomographic tilt series acquisition was performed with Tomo4 software from Thermo Fisher Scientific with a total electron dose of less than 100 e⁻/nm². Alignment and reconstruction of the series were performed using IMOD.⁴
14. Liposomes containing AuNPs were concentrated up to 5 mM total lipid concentration (relevant for *in vivo* use) with a Vivaspin® 2 centrifugal concentrator (2 mL volume, 300k MWCO). Note: to prevent liposomes from aggregating, centrifugal forces were not used. Briefly, liposomes were transferred in the concentrator which was let in an upright position. Solvent was slowly removed by gravity and by occasional shaking.

Characterization

Size (nm)	PDI
157.7	0.301

Cryo-TEM images

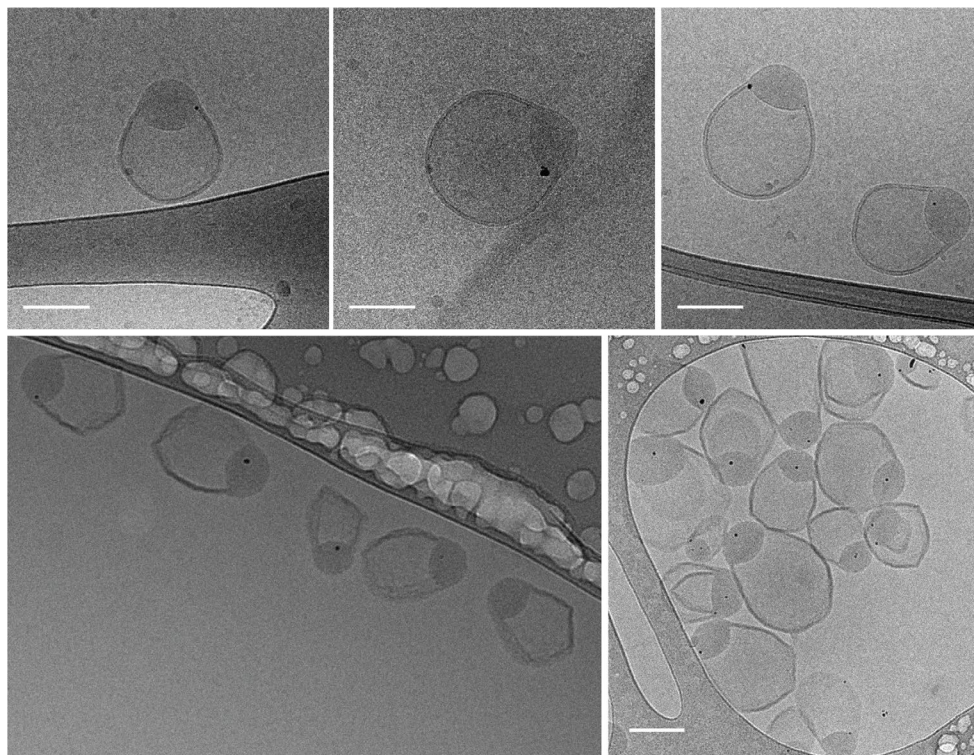


Figure 1: Cryo-TEM images depicting phase-separated PAP3 liposomes encapsulating AuNPs (black, electron dense dots) after *in situ* formation (Turkevich solution used: 1.75 mM HAuCl₄; 7.14 mM Sodium Citrate). Average AuNP size = 9.95 nm based on quantification (N = 103). Scale bars = 100 nm. Sample is free of unencapsulated AuNPs.

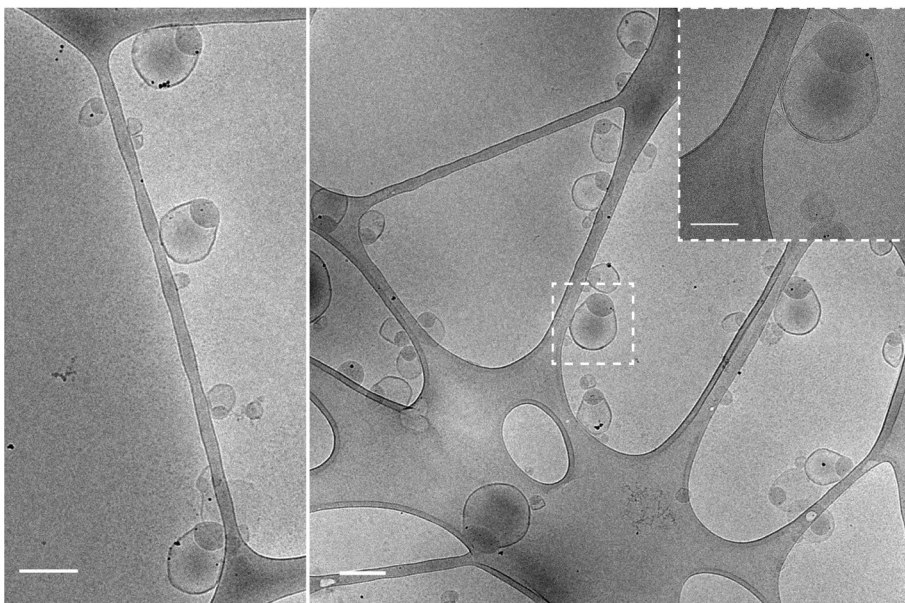


Figure 2: Cryo-TEM images depicting phase-separated PAP3 liposomes encapsulating AuNPs (black, electron dense dots) after *in situ* formation (Turkevich solution used: 2.5 mM HAuCl₄; 10.2 mM Sodium Citrate). Scale bars = 200 nm for low magnification images and 100 nm for insert. Sample is free of unencapsulated AuNPs.

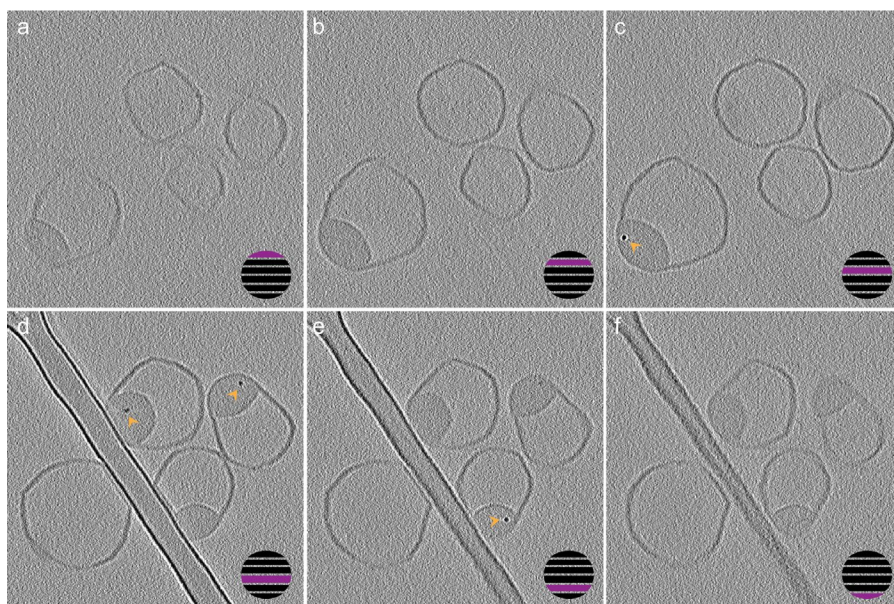


Figure 3. Cryo-electron tomography of PAP3 liposomes containing AuNPs. a-f) Representative slices of the tomogram showing AuNPs (orange arrows) reside in the inner

core of liposomes and show a preference for the liposome lipid droplet. Slice schematic (right bottom corner) shows which slice (in purple) corresponds to each image (a, f = top and bottom slices, b-e = middle slices). Turkevich solution used: 1.75 mM HAuCl₄; 7.14 mM Sodium Citrate.

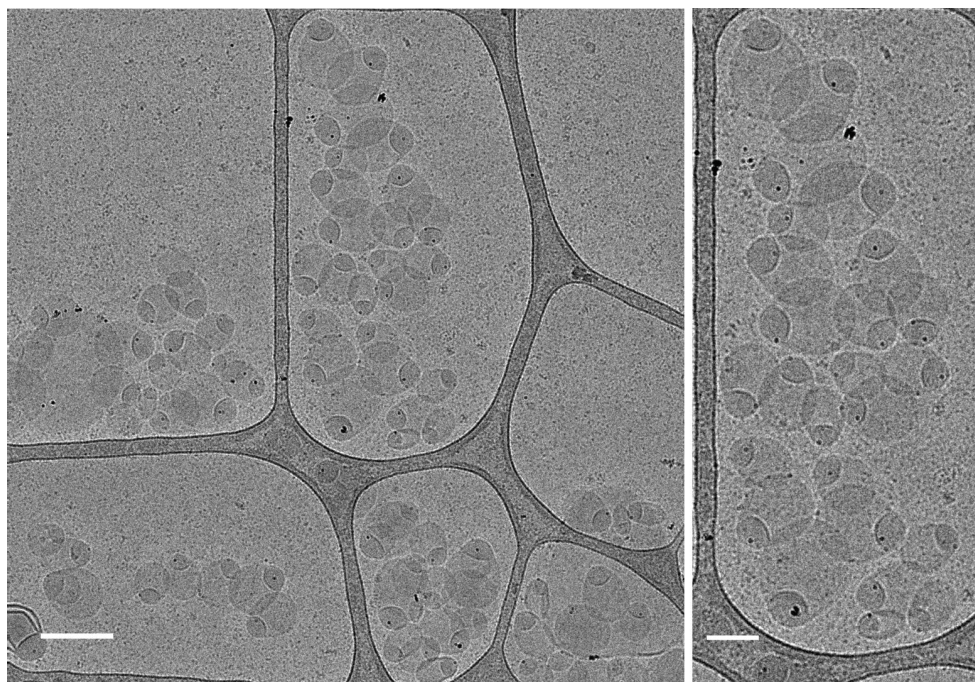


Figure 4. PAP3 liposomes containing AuNPs (black, electron dense dots) after incubation with human serum (1:1) for 10 min. Liposome morphology is preserved and AuNPs remain encapsulated (Turkevich solution used: 1.75 mM HAuCl₄; 7.14 mM Sodium Citrate). Average AuNP size = 9.62 nm based on quantification (N = 70). Scale bars = 200 nm for low magnification image and 100 nm for high magnification image.

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List of Abbreviations

apoAI	apolipoprotein AI	DOTMA	dioctadecenyl-trimethylammonium propane
apoCII	apolipoprotein CII	dpf	days post fertilization
ApoE	apolipoprotein E	DSPC	distearoylphosphatidylcholine
ASOs	antisense oligonucleotides	EL	endothelial lipase
Asp	asparagine	EMA	European medicine agency
ATTR	transthyretin amyloidosis	EPC	phosphatidylcholine (egg)
BA	basilar artery	EPG	phosphatidylglycerol (egg)
BBB	blood brain barrier	EPR	enhanced permeation-retention
bECs	brain endothelial cells	ESI	electron spray ionization
CE	cholesterol ester	ET	electron tomography
CG	coarse grained	EV	extracellular vesicle
CHO	cholesterol	FDA	food and drug administration
CHT	caudal hematopoietic tissue	FFA	free fatty acid
COM	center of mass	FFT	fast Fourier transform
CSF	cerebrospinal fluid	GPIHBP1	glycosylphosphatidylinositol anchored high density lipoprotein binding protein 1
CT	computed tomography	GUVs	giant unilamellar vesicles
CtA	cerebral arteries	HDL	high-density lipoprotein
CV	caudal vein	H_{II}	hexagonal phase
DAG	diacylglycerol	His	histidine
DCM	dichloromethane	HL	hepatic lipase
DIPEA	diisopropylethylamine	hpi	hours post injection
DLS	dynamic light scattering	HRMS	high resolution mass spectrometry
DLV	dorsal longitudinal vein	HSPG	heparan sulfate proteoglycans
DMAP	dimethylaminopyridine	ID	injected dose
DMSO	dimethylsulfoxide	IV	intravenous
DOaG	dioleoylamidoglycerol	KC	Kupffer cells
DODAP	dioleoyldimethylammonium propane	L_d	liquid disordered
DOE	design of experiment	LDL	low-density lipoprotein
DOG	dioleoylglycerol	LDLr	LDL receptor
DOPC	dioleoylphosphatidylcholine	LNP	lipid nanoparticle
DOPS	dioleoylphosphatidylserine		

L_o	liquid ordered	PMBC	primordial midbrain channel
LPL	lipoprotein lipase	PMF	potential mean force
LR	lissamine rhodamine	POPC	palmitoyl-oleoyl-phosphocholine
LSECs	liver sinusoidal endothelial cells	RES	reticuloendothelial system
LUVs	large unilamellar vesicles	RNAi	RNA interference
MCeV	middle cerebral vein	rt	room temperature
MD	molecular dynamics	SASA	solvent-accessible surface area
MMcTA	mid.mesencephalic central artery	SEC	size exclusion chromatography
MPS	mononuclear phagocyte system	SECs	scavenging endothelial cells
mRNA	messenger RNA	Ser	serine
MS	mass spectrometry	siRNA	small interfering RNA
MsA	mesencephalic artery	SRB1	scavenger receptor B1
MsV	mesencephalic vein	TEM	transmission electron microscopy
NMR	nuclear magnetic resonance	TG	triglyceride (or triacylglycerol)
NN	neural network	TGL	triacylglycerol lipase
PBS	phosphate buffered saline	THF	tetrahydrofuran
PC	phosphatidylcholine	TLC	thin layer chromatography
PDI	polydispersity index	Trp	tryptophan
PEG	polyethylene glycol	US	umbrella sampling
PHBC	primordial hindbrain channel	VLDL	very-low density lipoprotein
PKC	protein kinase C	VLP	virus-like particle