### **Chapter 19**

### **Isolation of Plasma Membranes from the Nervous System** by Countercurrent Distribution in Aqueous Polymer **Two-Phase Systems**

#### Jens Schindler and Hans Gerd Nothwang

#### Summary

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The plasma membrane separates the cell-interior from the cell's environment. To maintain homeostatic 7 conditions and to enable transfer of information, the plasma membrane is equipped with a variety of 8 different proteins such as transporters, channels, and receptors. The kind and number of plasma 9 membrane proteins are a characteristic of each cell type. Owing to their location, plasma membrane 10 proteins also represent a plethora of drug targets. Their importance has entailed many studies aiming 11 at their proteomic identification and characterization. Therefore, protocols are required that enable 12 their purification in high purity and quantity. Here, we report a protocol, based on aqueous polymer 13 two-phase systems, which fulfils these demands. Furthermore, the protocol is time-saving and protects 14 protein structure and function.

Key words: Brain, Plasma membrane, Countercurrent distribution, Two-phase system, Enrichment 16

#### 1. Introduction

Plasma membrane (PM) proteins mediate signal transduction, 18 solute transport, secretion, and cell-cell contact. They are also 19 the central players in the propagation and transmission of action 20 potentials, which are the lingua franca in the nervous system. 21 Finally,  $\sim 70\%$  of all known drug targets act on them (1). PM 22 proteins are thus of prime interest in many areas of both basic and 23 biomedical research. However, their proteome analysis is rather 24 difficult, as they encompass only 0.4-2.5% of the total cellular 25 protein amount (2). This renders their identification difficult in 26 27 the bulk of other, more abundant proteins of the cytoskeleton,

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74 75 the energy metabolism, and alike. Furthermore, many properties of PMs such as density overlap with those of other membranous compartments, mainly the endoplasmic reticulum. Classical purification protocols of PM proteins are therefore rather cumbersome and material-consuming. This is for instance the case with the most popular method, consisting of a combination of differential and density gradient centrifugation steps (3, 4). Other methods are either quite expensive and contamination-prone (e.g. immunoprecipitation) or apply only to cultured cells such as surface labeling, which cannot be applied to bulky tissue.

Interest in novel, more efficient subcellular purification protocols has recently emerged from the impressive progress in the analytical part of proteomics. Current mass spectrometry can identify proteins at concentrations of less than 1 pM. This allows for detection of proteins such as the neuronal PSD-95, which is present at 300 copies/postsynaptic density (Chen et al., AU1 2005) in as few as  $2 \times 10^6$  neurons, based on 1,000 postsynaptic densities/cells. Hence, proteomics studies on functional or anatomically well-characterized small tissue samples or scarce biopsy material come into reach. However, protocols for the isolation of defined subcellular compartments did not keep pace, despite the recognized need to analyze the subproteome of the various compartments separately. This cellular dissection is mandatory to detect low abundant proteins and to identify compartment-specific post-translational modifications or significant changes in protein localization. Changes therein often underlie physiological and pathophysiological processes (5).

A highly selective and efficient method to separate membranes of different subcellular origin was developed more than 30 years ago by Albertsson and colleagues and was based on the use of aqueous polymer two-phase systems (6, 7). These systems often form, when aqueous solutions of two structural different water-soluble polymers are mixed above a defined concentration. Most often, poly(ethylene glycol) (PEG) and dextran are used as polymers, as they are cheap, require only moderate concentrations, separate easily, and preserve protein structure and function well. When mixed, phases will form, and the upper phase will be enriched in PEG, whereas the bottom phase will mainly contain dextran. Interestingly, the various cellular membranes have different affinities to partition in either of the two phases. PMs prefer the upper phase, whereas mitochondria partition rather to the bottom phase. This different behavior can be attributed to differences in hydrophobic and hydrophilic surface properties of membranes, most likely arising from differences in their phosholipid composition.

Membranes differ only subtly in their surface properties and the isolation of PMs by aqueous polymer two-phase systems cannot be achieved in a single step. One possibility to increase the

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purity of the PM fraction is multistep extraction procedures such 76 as countercurrent distribution (CD) (Fig. 1). CD is based on 77 the Nernst distribution law. Membranes are separated by this 78 method on the basis of their different solubilities in two immis-79 cible aqueous solutions of structurally different polymers. These 80 two phases, flowing into opposite directions, are brought into 81 contact, mixed, and allowed to separate. PMs preferentially parti-82 tion to the top phase and will reside there throughout the mul-83 tiple extractions. Intracellular membranes such as mitochondria 84 will preferentially partition to the bottom phase throughout the 85 procedure. We recently adapted this principle to the isolation of 86



Fig. 1. Scheme of countercurrent distribution. In CD experiments, the top-phase of the first two-phase system A is transferred to a fresh bottom-phase B and the bottom-phase of two-phase system A is re-extracted with a fresh top-phase. After six iterations, biomaterial is efficiently separated.

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87	PMs from the nervous system (8). The protocol is fast, easy to
88	perform, and yields up to 30% of the initial PMs with high purity.
89	Contaminations by endoplasmic reticulum and mitochondria,
90	the major contaminations in standard protocols, were low.

#### 91 2. Materials

92	Owing to the strong influence of ions on membrane partitioning
93	in the two-phase systems, double distilled water should be used
94	throughout the experiments.
<ul> <li>95 2.1. Two-Phase</li> <li>96 Systems</li> <li>97</li> <li>98</li> </ul>	<ol> <li>Glass-Teflon homogenizer.</li> <li>Dextran stock solution: Dextran T500 (20%, w/w) (<i>see</i> Note 1).</li> <li>PEG stock solution: PEG 3350 (40%, w/w).</li> <li>Tris-H<sub>2</sub>SO<sub>4</sub>: Tris (200 mM), pH 7.8 adjusted with H<sub>2</sub>SO<sub>4</sub>.</li> </ol>

### 99 **3. Methods**

100	3.1. Two-Phase	All steps of the affinity two-phase partitioning protocol should be
101	Partitioning	performed at 4°C. Working at room temperature prevents phase
102		separation. The procedure is illustrated in Fig. 1. The numbers
103		in Fig. 1 correspond to the numbered two-phase systems given
104		in Table 1, and the letters refer to the phases as indicated in the
105		protocol given below.
106		1. Prepare seven two-phase systems with the compositions indi-
107		cated in Table 1, 1 day prior to use. Mix them by 20 inverta-
108		tions, vortexing for 10 s, another 20 invertations, and store

## Table 1Composition of two-phase systems

	Two-phase system "A" (g)	Two-phase systems "B–G" (g)
Dextran stock solution	1.035	1.035
PEG stock solution	0.518	0.518
Tris–H <sub>2</sub> SO <sub>4</sub>	0.750	0.750
Water	0.598	0.698

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the mixtures at 4°C overnight. Two-phase systems will form 109 overnight with the top phase enriched in PEG and the bottom 110 phase enriched in dextran. 111

- On the next day, remove all top phases from two-phase systems "B-G" and store them separately.
- 3. Homogenize 0.1 g brain tissue in two-phase system "A" using a114glass-Teflon homogenizer followed by 45 s of sonication. Centri-115fuge at  $700 \times g$  for 5 min to accelerate phase separation.116
- 4. Transfer the top phase (Top) of two-phase system "A" onto 117 the bottom-phase (Bot) of two-phase system "B" (TopA → BotB) 118 (*see* Note 2). Add an equal amount of fresh top-phase (stored 119 in step 2) onto bottom-phase "A" (I. in Fig. 1). Mix both 120 two-phase systems by 20 invertations, vortex for 10 s, then 121 mix again by another 20 invertations. Centrifuge at 700 × g 122 for 5 min to accelerate phase separation. 123
- 5. Transfer top phases in the following order (II. in **Fig. 1**): (1) 124 TopB  $\rightarrow$  BotC; (2) TopA  $\rightarrow$  BotB. Add an equal amount of 125 fresh top-phase (stored in step 2) onto bottom-phase "A." 126 Mix all two-phase systems by 20 invertations, vortexing for 10 127 s, and another 20 invertations. Centrifuge at 700 × g for 5 min 128 to accelerate phase separation. 129
- 6. Transfer top phases in the following order (III. in Fig. 1): 130
  (1) TopC → BotD; (2) TopB → BotC; (3) Top A → BotB. Add 131
  an equal amount of fresh top-phase (stored in step 2) onto 132
  bottom-phase "A." 133

Mix all two-phase systems by 20 invertations, vortexing for 13410 s, and another 20 invertations. Centrifuge at  $700 \times g$  for 5 min to accelerate phase separation. 136

- 7. Transfer top phases in the following order: (1) Top D  $\rightarrow$ BotE; (2) TopC  $\rightarrow$  BotD; (3) TopB  $\rightarrow$  BotC; (4) Top A  $\rightarrow$ BotB. Add an equal amount of fresh top-phase (stored in step 2) onto bottom-phase "A." Mix all two-phase systems by 20 invertations, vortexing for 10 s, and another 20 invertations. Centrifuge at 700 × g for 5 min to accelerate phase separation.
- 8. Transfer top phases in the following order: (1) TopE → BotF;
  (2) Top D → BotE; (3) TopC → BotD; (4) TopB → BotC; (5)
  Top A → BotB. Add an equal amount of fresh top-phase (stored in step 2) onto bottom-phase "A." Mix all two-phase systems by 20 invertations, vortexing for 10 s, and another 20 invertations.
  Centrifuge at 700 × g for 5 min to accelerate phase separation.
- 9. Transfer top phases in the following order: (1) TopF → BotG: 149
  (2) TopE → BotF; (3) Top D → BotE; (4) TopC → BotD; (5) 150
  TopB → BotC; (6) Top A → BotB. Add an equal amount of 151
  fresh top-phase (stored in step 2) onto bottom-phase "A." 152

Mix all two-phase systems by 20 invertations, vortexing for 15310 s, and another 20 invertations. Centrifuge at  $700 \times g$  for 5 154

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155 156 157	min to accelerate phase separation. After phase separation, you end up with seven two-phase systems (VII. in <b>Fig. 1</b> ). PMs are enriched in TopF and TopG.
158 1 159 160	0. PMs can be recovered from TopG or combined TopF + G ( <i>see</i> <b>Note 3</b> ) by diluting the phases 1:10 with water followed by ultracentrifugation at $1,50,000 \times g$ and $4^{\circ}$ C for 1 h.

(4)

#### 161 **4. Notes**

162	1. Dextran can contain up to 10% water and for that reason has to
163	be freeze-dried. For freeze-drying, dissolve dextran in distilled
164	water in a plastic dish with a large surface (e.g. Petri dish), freeze
165	it at -80°C, and dry it by sublimating the water under vacuum.
166	Store the freeze-dried dextran in closed plastic tubes sealed tightly
167	with parafilm at $-20^{\circ}$ C. Let it come to room temperature before
168	opening to protect it from humidity.
169	2. In two-phase systems, interphases are always considered as
170	part of the bottom phase.
171	3. The purity of PMs from TopG alone is slightly higher but
172	combining TopF and TopG nearly doubles the yield.

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# Author Queries

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