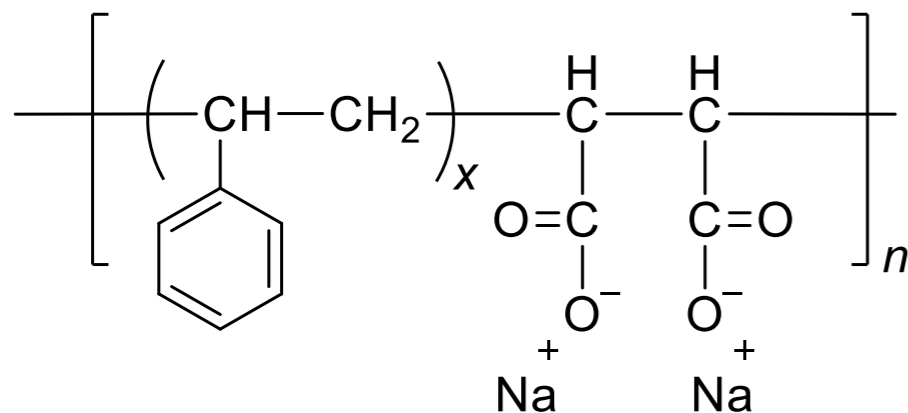


Berkeley, 24th March 2017

DIBMA: A Non-Aromatic Polymer for Solubilising Membrane Proteins

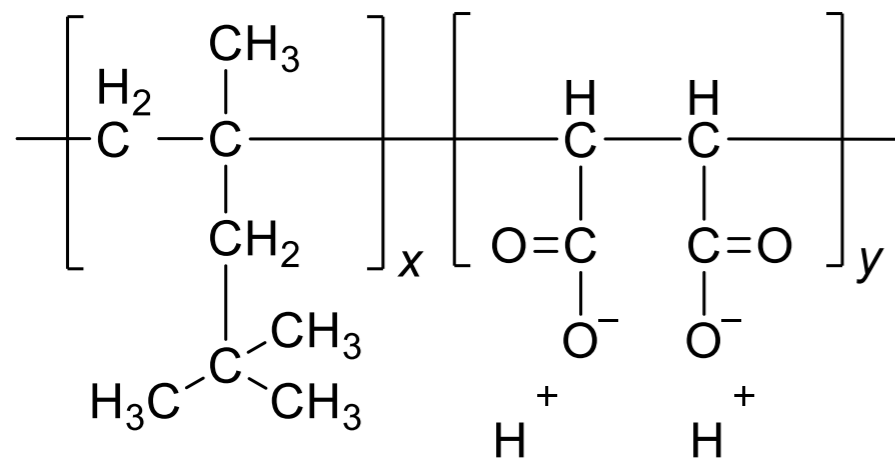
.....
Abraham Olusegun Oluwole, Bartholomäus Danielczak, Annette
Meister, Jonathan Oyebamiji Babalola, Carolyn Vargas, Sandro Keller

Aromatic vs. aliphatic copolymers



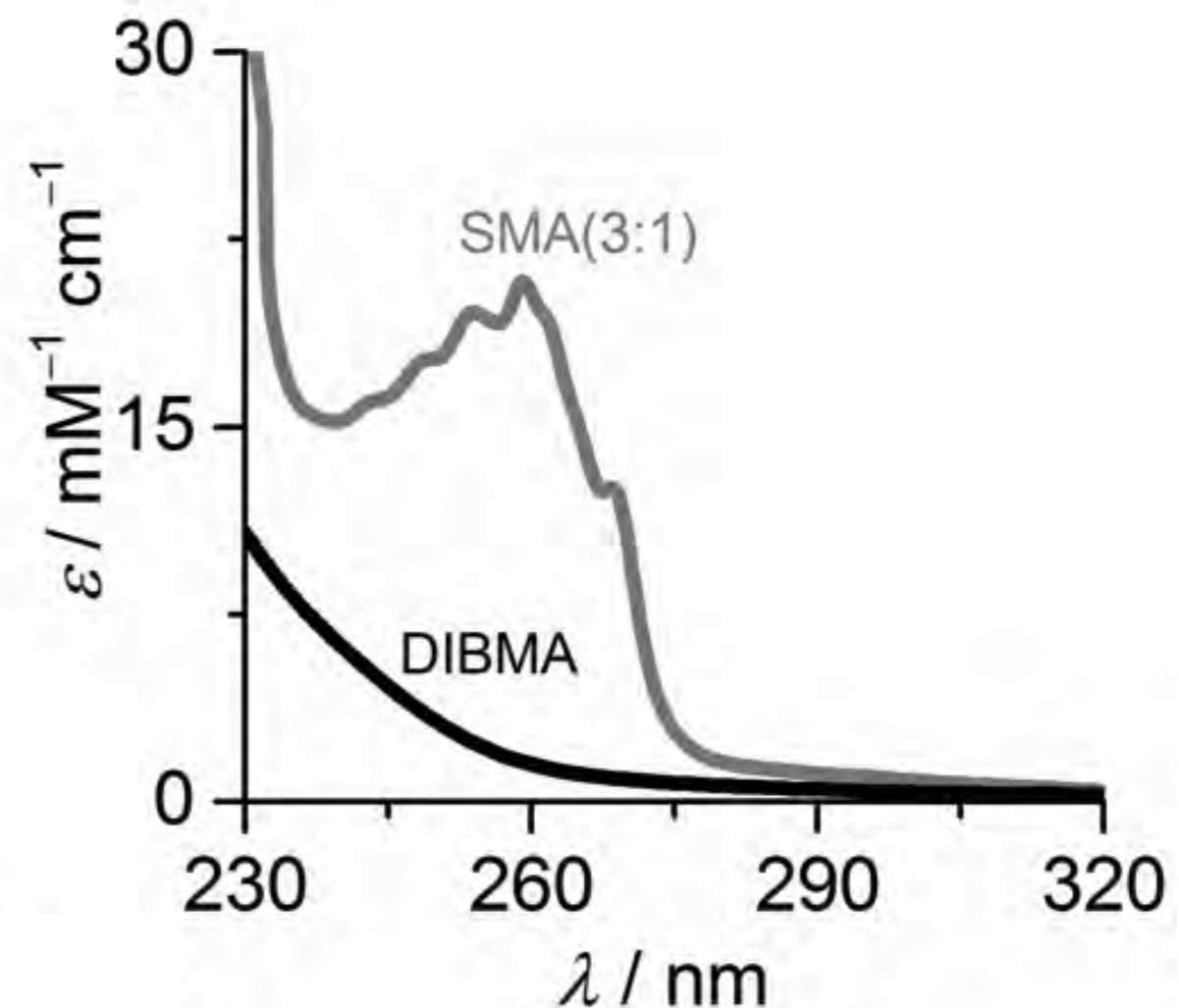
SMA: poly(styrene-co-maleic acid)

$M_w = 10$ kg/mol

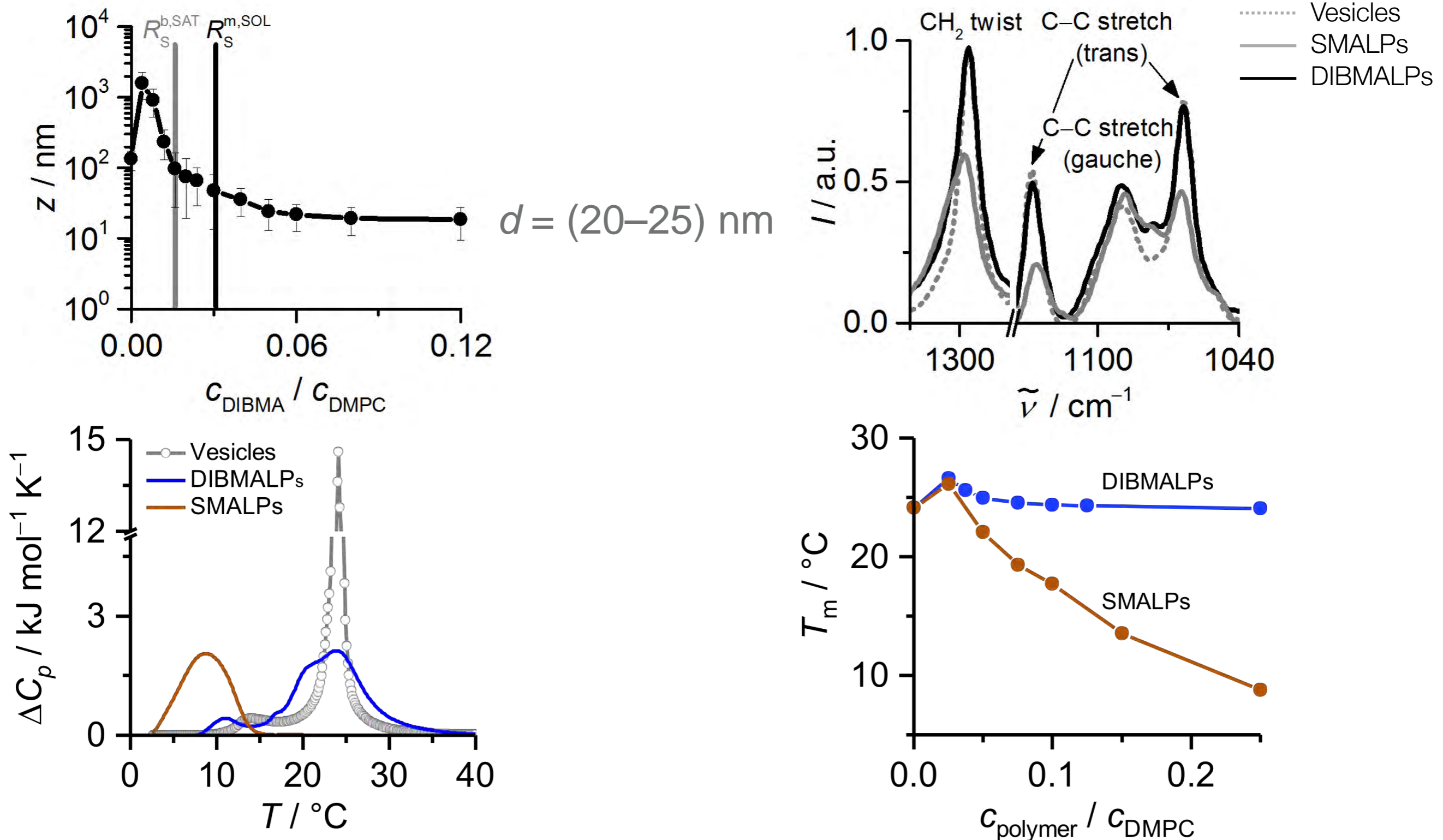


DIBMA: poly(diisobutylene-*alt*-maleic acid)

$M_w = 12$ kg/mol



Mild effects on membrane lipid order



Compatibility with divalent cations

$c_{\text{Ca}^{2+}} / \text{mM}$

0 1 2 3 4 5



$c_{\text{Ca}^{2+}} / \text{mM}$

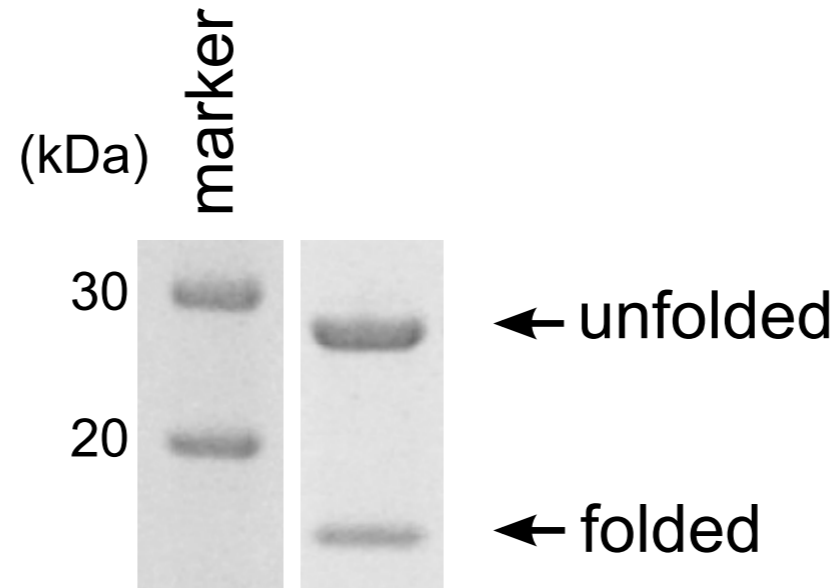
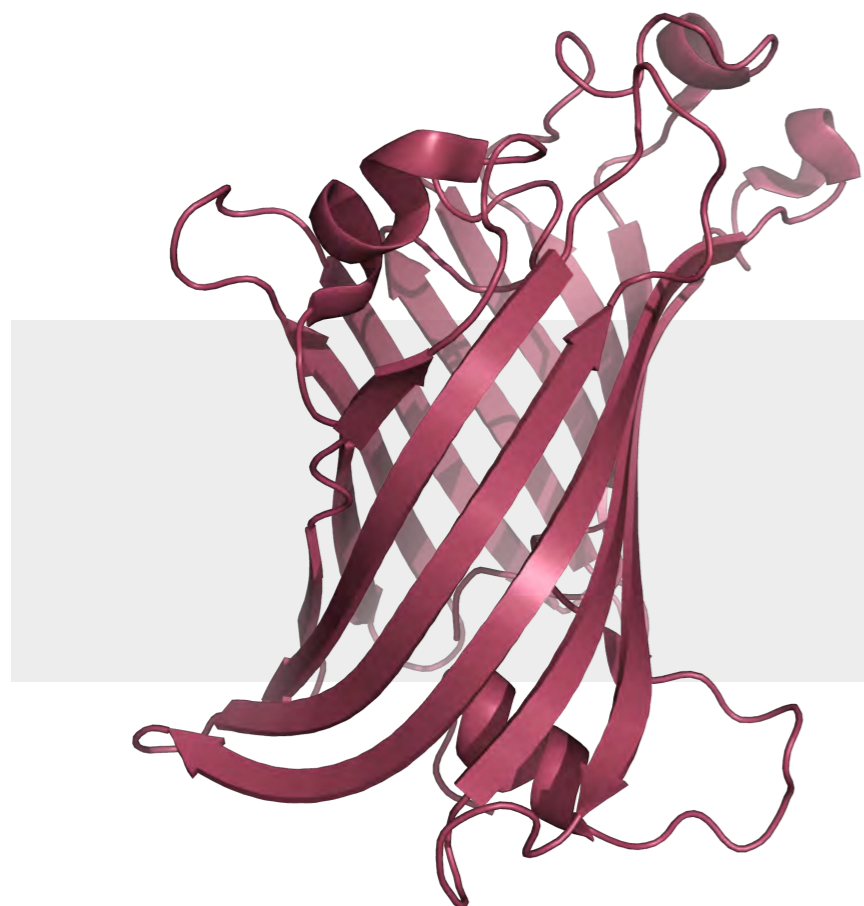
0 5 10 15 20 25 30 35 40 50



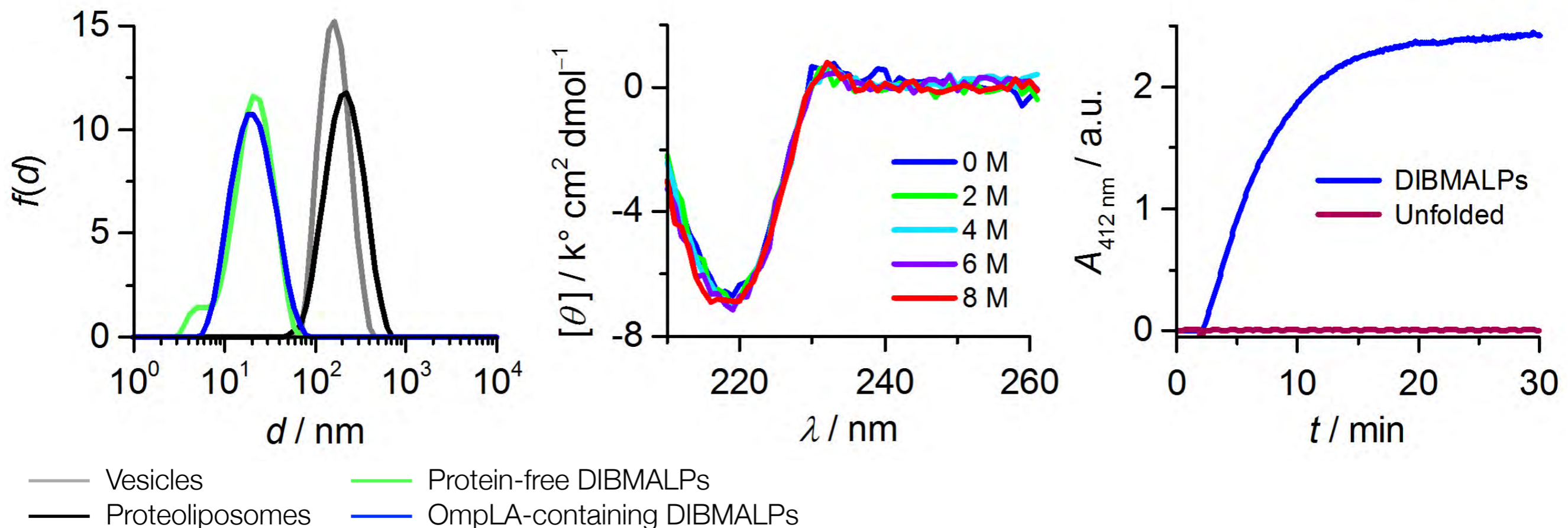
Model protein

Outer membrane phospholipase A (OmpLA):

- integral membrane enzyme
- differential migration on SDS-PAGE (folded vs. unfolded)



Solubilisation of OmpLA in DLPC



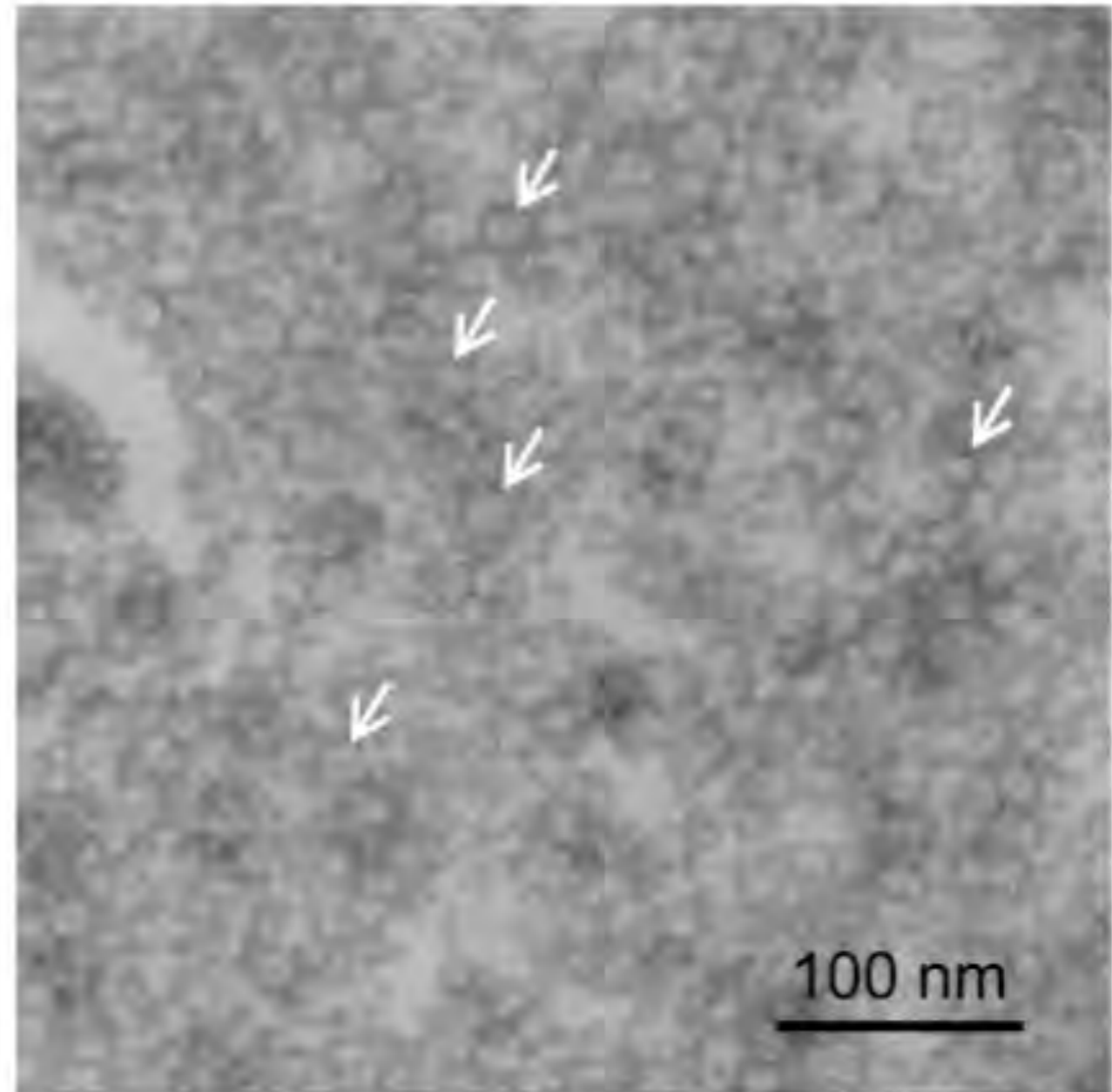
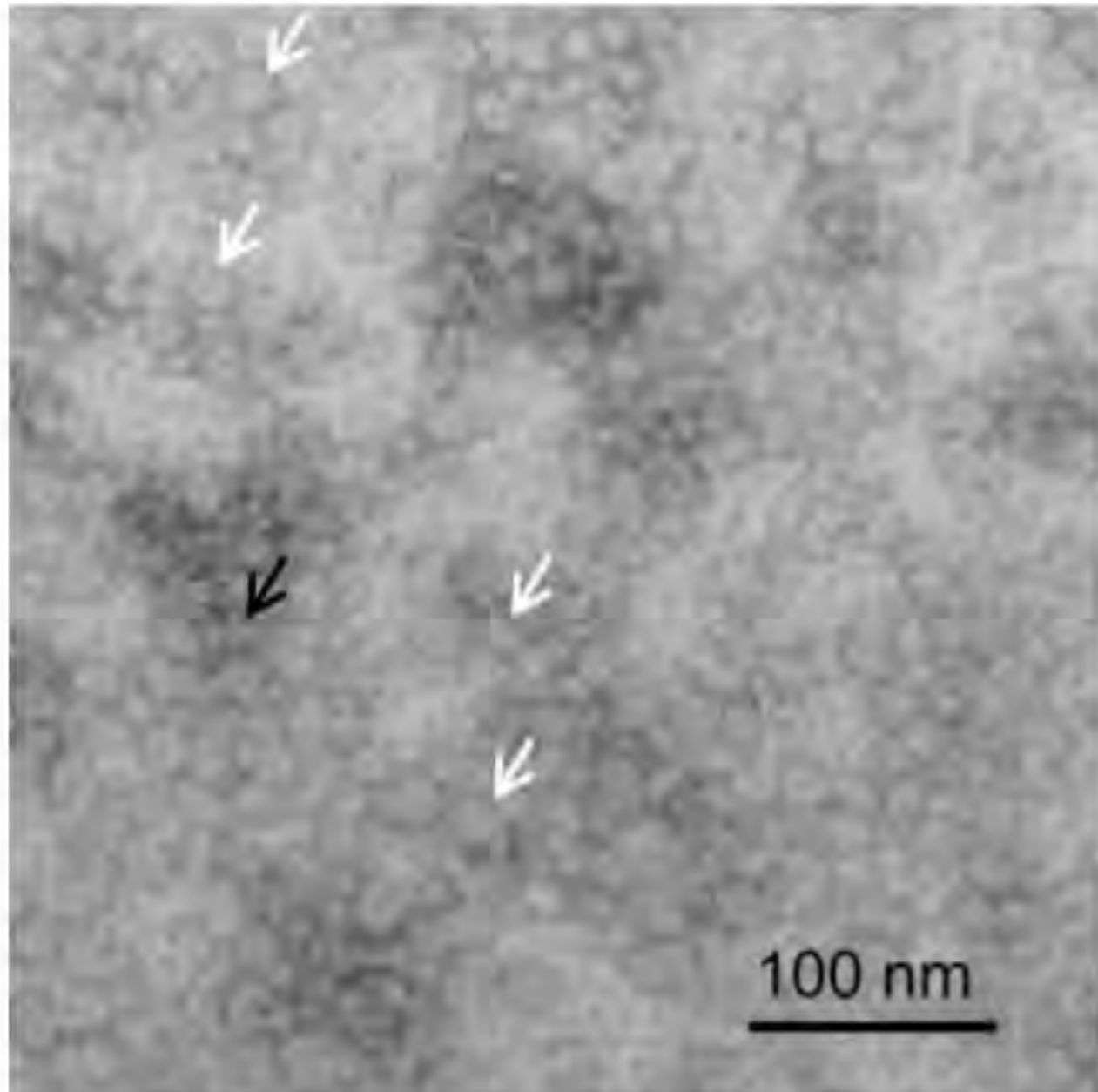
Nanodisc formation from proteoliposomes:

- as efficient as from protein-free DLPC vesicles
- OmpLA remains natively folded and enzymatically active

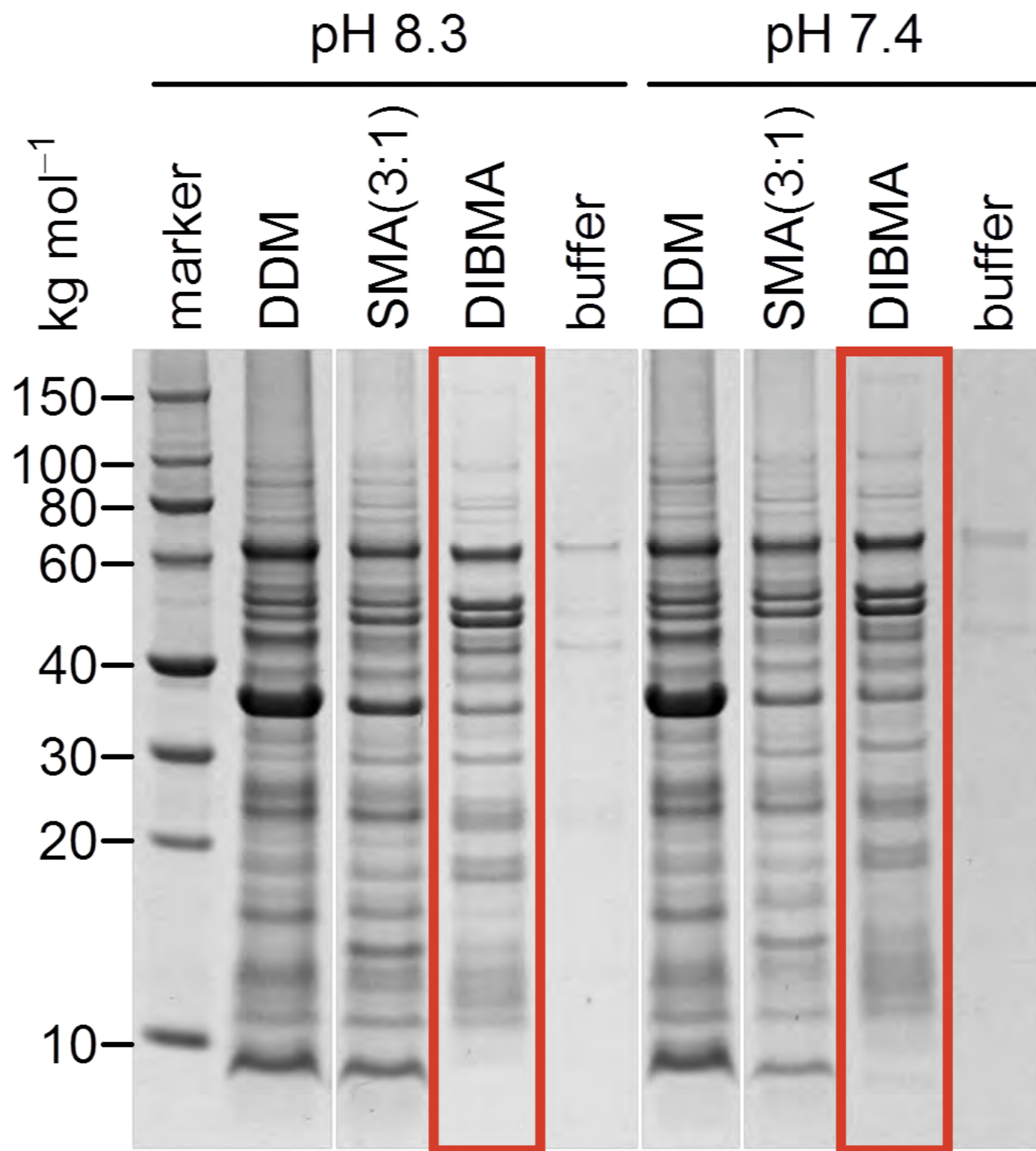
Transmission electron microscopy

Protein-free DLPC discs

OmpLA-containing discs



Solubilisation of *E. coli* membranes



Take-home messages

Aliphatic DIBMA is a promising alternative to SMA, as it is less perturbing to lipid order and compatible with UV spectroscopy and divalent cations.

Both SMA and DIBMA solubilise a wide range of membrane proteins directly from cellular membranes to form lipid-bilayer nanodiscs.

SMALPs rapidly exchange lipids and, thus, represent equilibrium rather than kinetically trapped structures (*Sci. Rep.*, DOI: 10.1038/srep45875).

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