

Experimental report

28/03/2024

Proposal: 8-02-983

Council: 10/2022

Title: The role of charge for interaction with bio-membranes: ζ -Synuclein and Synaptobrevin-2

Research area: Biology

This proposal is a new proposal

Main proposer: Irina APANASENKO

Experimental team: Irina APANASENKO
Margarita KRUTEVA
Ralf BIEHL
Benedetta Petra ROSI

Local contacts: Thomas SAERBECK
Philipp GUTFREUND
Nicolo PARACINI

Samples: h-d DMPC/DMPG mebrane + Synaptobrevin(1-96)
h-d DMPC/DMPG mebrane + Synaptobrevin(1-116)
h-d DMPC/DMPG mebrane + alpha synucleine

Instrument	Requested days	Allocated days	From	To
FIGARO	5	0		
D17	5	3	01/04/2023	04/04/2023

Abstract:

In our current research we focus on the interaction of two neuronal IDPs with bio-membranes: ζ -Synuclein and Synaptobrevin-2. We would like to investigate the interaction of both IDP with membranes of varying charge composition (fraction of negative DMPG in neutral DMPC) to examine the configuration in/at the membrane and in the adjacent solution. For ζ -Syn we want to identify the location of ζ -helices and the configuration of disordered regions. For Syb2 we will examine the disordered soluble region with and without the transmembrane region.

Experimental report – proposal no. 8-02-983

In this experiment, the configuration of intrinsically disordered protein alpha-synuclein (α -Syn) in/at differently charged lipid's membrane (fraction of negative DMPG in neutral DMPC) has been investigated.

For the experiment, 3 days of beamtime have been allocated at D17.

Sample	Protein concentration [mg/ml], total volume [ml]
DMPC lipid membrane + α -Syn	0.1, 15
25% of DMPG/DMPC lipid membrane + α -Syn	0.1, 15
50% DMPG/DMPC lipid membrane + α -Syn	0.1, 15

Table 1 List of samples measured at D17.

Data have been collected with the following setup parameters:

Angle	DAN	SAN	S2W	S3W	OPENING	TRS
Angle 1	2,8	0,8	1,5	0,75	0,85	0
Angle 2	7,6	3,2	6	3	4	0

Table 2 D17 setup parameters.

An average time of ≈ 3 hrs was required to measure empty membrane and then ≈ 5 hrs to measure membrane with the protein for each sample in three different contrasts (H₂O, D₂O, SMW (silicon-matched (38% D₂O/62% H₂O) water)). All the data have been collected at T=37°C.

DMPC and 25% DMPG/DMPC lipid bilayers were made by vesicle fusion method.

50% DMPG/DMPC lipid bilayer was made by the Langmuir-Blodgett trough method.

Preliminary results:

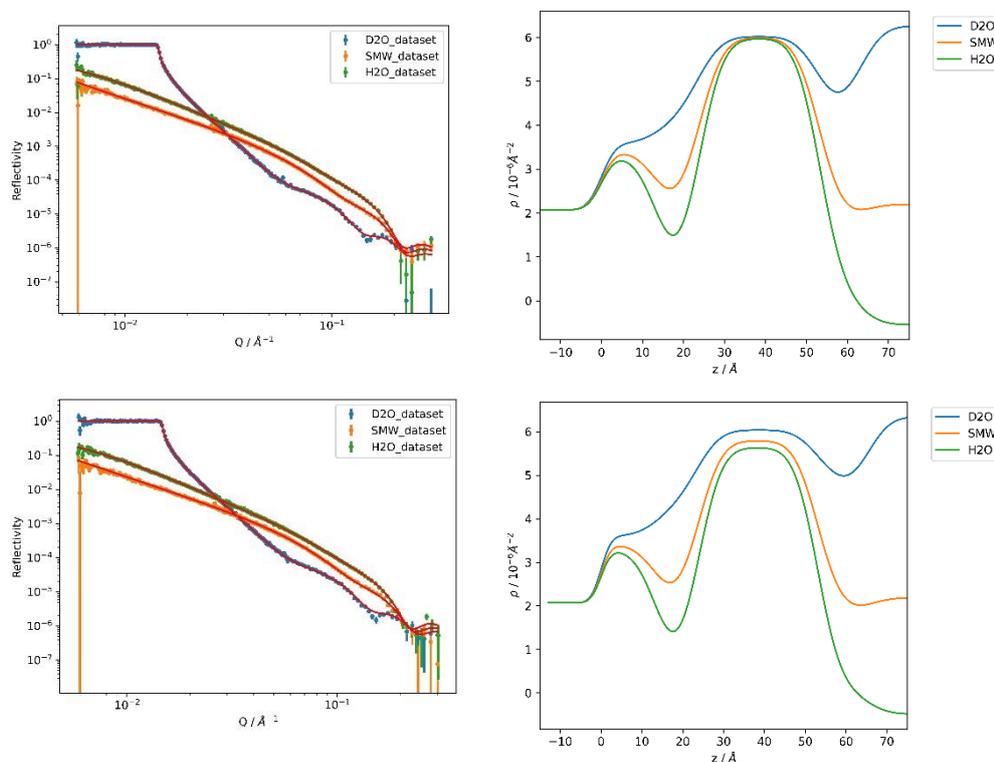


Figure 1 Neutron reflectometry profile and model data fits and the scattering length density profiles these fits describe for deposited DMPC bilayer without (top) and with α -Syn protein (bottom).

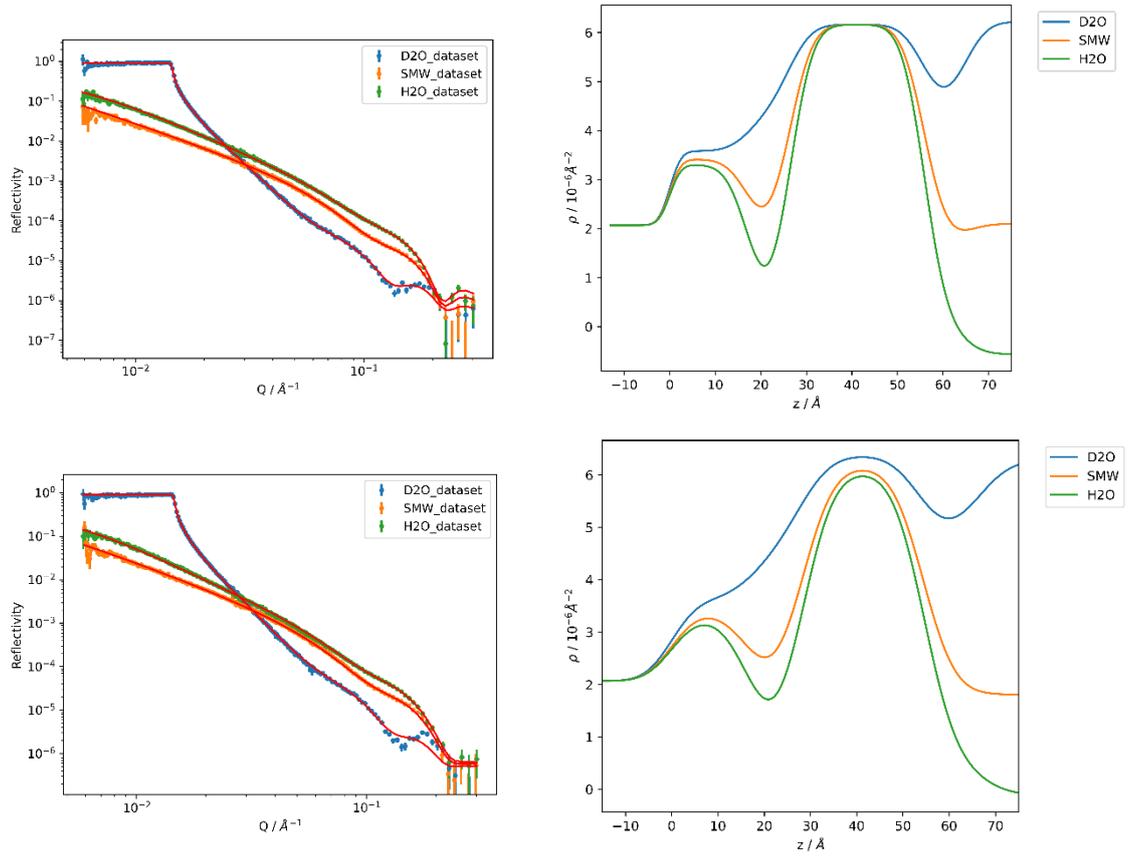


Figure 2 Neutron reflectometry profile and model data fits and the scattering length density profiles these fits describe for deposited 25% DMPG/DMPC bilayer without (top) and with α -Syn protein (bottom).