

HIGH THROUGHPUT

DETERGENT SCREENING

FOR MEMBRANE PROTEINS

VERSION 5.2 Isabel Moraes, PhD, Principal Research Scientist at National Physical Laboratory (NPL)
Amy Danson, PhD, Higher Research Scientist at National Physical Laboratory (NPL)
Riccardo Marabini, PhD, Senior R&D and Application Scientist at Fida Biosystems

- 12 detergents screened in 3 hours (in triplicates)
- Minimal sample consumption (1,5 uL for entire screen)
- No need for purification directly in crude matrix
- Walk-away automation
- Multiple readouts in a single run (size, peak area, polydispersity index)



INTRODUCTION

More than 50% of the present-day drugs, and an increasing number of drug discovery and development programs target proteins situated in a membrane. Membrane proteins are challenging to express, maintain in-solution, purify and stabilize after extraction from cells. As a result, most of the membrane proteins undergo detergent screening to identify the detergents that solubilize the highest amount of protein from the insoluble fraction of cell lysate. Effective detergents offer high degree of solubilization as well as stabilization of membrane proteins.

In this Application Note, Fida 1 is used for high throughput detergent screening to assess the degree of solubilization and polydispersity of a GFP-tagged membrane protein directly in crude matrix, using only 40nL of sample per measurement. Moreover, the FIDA detergent screening is compared with Fluorescence Size- Exclusion Chromatography (FSEC). Contrary to standard inhibiting drugs, "degraders" trigger an endogenous cell mechanism, targeted protein degradation, which captures and "slays" unwanted or damaged proteins via the ubiquitin- proteasome system.



FIDA MEASURING PRINCIPLE

Flow-Induced-Dispersion-Analysis (FIDA) is a capillary-based microfluidic method, exploiting that flow rates in the center of the capillary are faster than at the edges of the capillary. The resulting radial concentration gradients at the front and the tail of the sample zone result in the diffusion of your chosen indicator. The distinct dispersion profiles determine the shape of the generated curves and enables a "first principle" biophysical measurement of size (Figure 1). The FIDA measurement of size is broadly applicable for studying biomolecular stability, interactions, etc.

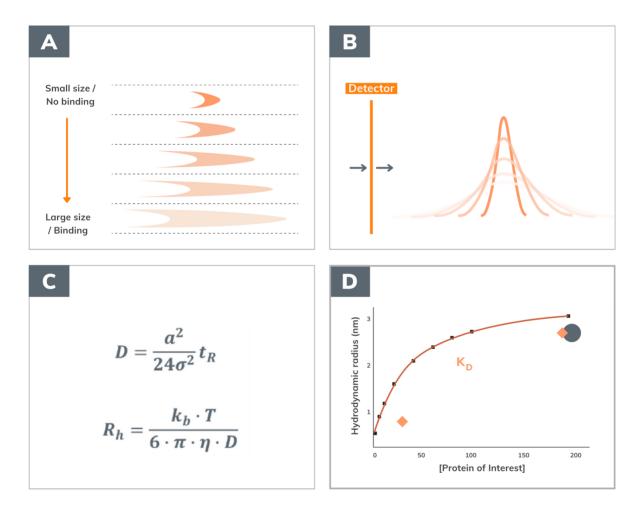


Figure 1: FIDA measuring principle.

- A. Dispersion profiles occurring in capillary. The shape of the profile depends on the size of the fluorescent molecule or on the size of the complexes it forms with its binding partners,
- B. Corresponding detector readout,
- C- D. Calculation of diffusivity, D, and hydrodynamic radius, Rh, is used to measure the affinity, KD, of binding partners.



MATERIAL & METHODS

The intensity of the fluorescence signal is representative of the solubilized fluorophore (GFP) concentration and is directly proportional to the peak area (Figure 1B). Therefore, the peak area can be used to assess the solubilization performance of detergents, enabling an extremely simple setup with no labeling requirements and only a few nL of sample. In the experiment shown in Figure 2, GFP-fused Lactose Permease (GFP-LacY) was the indicator, solubilized in 12 different detergents to assess their performance. The capillary was equilibrated with the buffer containing a specific detergent, e.g. CHAPS and GFP-LacY solubilized in CHAPS injected as indicator (40nL) at 0.1 mg/ml.

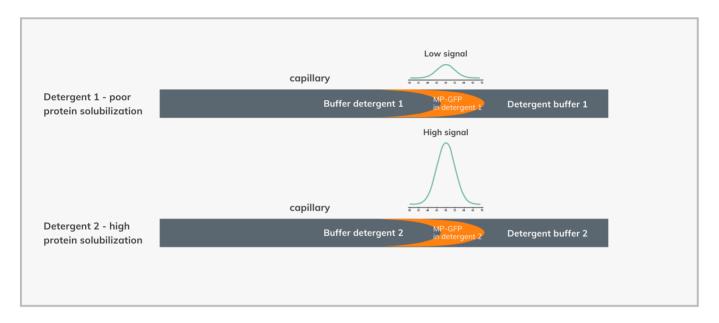


Figure 2. Schematic representation of the experimental conditions and raw signal in a GFP-tagged membrane protein detergent screening. MP stands for Membrane Protein.

Instrument:

Fida 1 with 480 nm LED fluorescence detection. Fidabio standard capillary (i.d.: 75 μ m, LT: 100 cm, Leff: 84 cm). The sample format was Fida glass vials for all the tests.

Indicator:

The indicator was the crude lysate containing the solubilized GFP-tagged protein..

Analyte

The analyte is just the buffer containing the corresponding detergent used to solubilize the protein of interest.

FIDA experiments:

20µL of crude membrane solubilized with a specific detergent was loaded into a Fidabio glass insert. A total of 12 samples, each with a different detergent, were loaded into the Fida 1 system, and the sample tray was kept at 5°C for the duration of the whole measurements. The washing buffers and the capillary chamber were at 25°C. The method shown below was used for the GFP-LacY experiments:



Rinse 1 M NaOH 45 3500 Pre-equilibration MilliQ 75 3500 Analyte Detergent A 20 3500 Indicator GFP-LacY in det. A 10 50 Analyte Detergent A 180 400			Time(s)	Pressure(mbar)
Analyte Detergent A 20 3500 Indicator GFP-LacY in det. A 10 50	Rinse	1 M NaOH	45	3500
Indicator GFP-LacY in det. A 10 50	Pre-equilibration	MilliQ	75	3500
	Analyte	Detergent A	20	3500
Analyte Detergent A 180 400	Indicator	GFP-LacY in det. A	10	50
	Analyte	Detergent A	180	400

Membrane protein sample preparation:

The cells overexpressing the protein of interest were lysed, centrifuged and the insoluble fraction resuspended in PBS, aliquoted and frozen. Afterwards, a specific detergent buffer was added to a crude membrane aliquot to solubilize the protein of interest. The samples were ultracentrifuged at 100,000 x g for 45 mins at 4 0C. The supernatant containing the solubilized membrane protein was diluted to 0.1 mg/ml of total protein concentration, determined with Bradford assay. The diluted sample was measured with the Fida 1 instrument and FSEC.

Buffer composition: 1X PBS, 150 mM NaCl + detergent Final detergent concentrations: CHAPS 2%; CHAPSO 2%; CYMAL-6 1%; C12E8 1%; DDM 1%; FC12 1%; LDAO 1%: LMNG 1%; OG 2%; UDM 1%; Control: No Detergent.

RESULTS

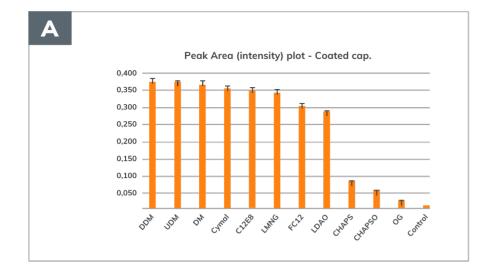
Flow-Induced Dispersion Analysis

The GFP-LacY was solubilized in 11 different detergents plus a negative control without any detergent. Figure 3A shows the histograms of the peak area for all detergents, suggesting that DDM and UDM most effectively solubilize GFP- LacY. The raw signals used to calculate the peak areas are overlayed in Figure 3B. Fida 1 also measures the Polydispersity Index (PDI) based on the peak shape. Polydisperse samples have higher PDI values, meaning that the indicator exists in a distribution of different sizes. In contrast, when the PDI is below 0.05, the sample is considered monodisperse.

Among the detergents screened in this assay, CHAPS, CHAPSO and OG have relatively high PDI values, while the rest of the detergents have similar values, close to 0, indicating monodisperse samples (Figure 3C).

These data show that Fida 1 can accurately screen for different detergents in unpurified samples, using only 40 nL per measurement, assessing 12 conditions in triplicates in just 3 hours. Most importantly, the volume used to load the instrument - $20\mu L$ - is fully recoverable and can be used for downstream analysis eliminating batch-to batch differences.





С	
DETERGENT	PDI
DDM	0,0097
UDM	0,0097
DM	0,0097
Cymal	0,0098
C12E8	0,0097
LMNG	0,0098
FC12	0,0097
LDAO	0,0097
CHAPS	0,662
CHAPSO	0,685
OG	0,66
Control	NA

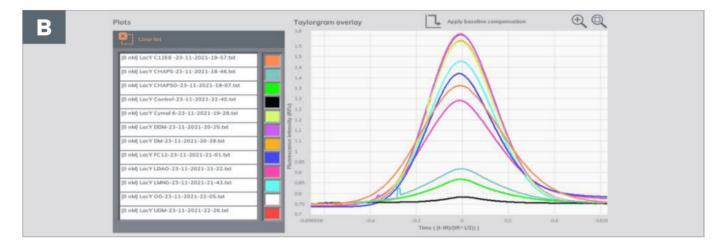


Figure 3.

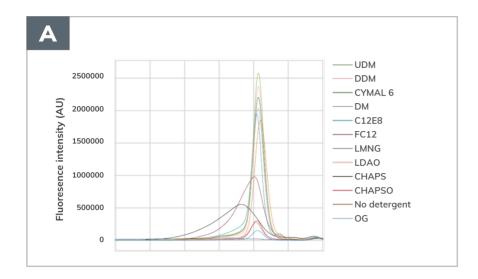
- **A**. FIDA detergent screening of 0.1mg/ml of GFP-LacY solubilized in different detergents. GFP-LacY peak area histograms for each detergent,
- B. Peak overlay for each of the GFP-LacY signals in different detergents, C. PolyDispersity Index (PDI) calculated for each GFP-LacY signal in different detergents

Fluorescence Size-Exclusion Chromatography

To validate the FIDA data for membrane protein detergent screening, the samples were also analyzed with Fluorescence Size-Exclusion Chromatography (FSEC), one of the standard methods for detergent screening. The signal from the FSEC is used to assess the level of solubilized protein in a detergent buffer, as well as the dispersity of the protein of interest. Figure 4A shows the overlapping FSEC chromatograms of all 12 conditions. UDM and DDM were shown to be the best detergents for solubilizing GFP-LacY confirming the results obtained with FIDA.

These data demonstrate that the Fida 1 system provides comparable results to the technologies used to date, with the striking advantages of using 10 to 100 times less sample volume, being much faster, and offering sample recovery opportunity as summarized in Table 1. Figure 4B compares detergent performance ranking with FSEC and FIDA based on signal intensity, while taking into consideration that the FIDA assay was run in triplicate while FSEC just one replica. The two rankings match very well, given that DDM and UDM have practically the same intensity in the two technologies.





В		
Ranking	FIDA	FSEC
1	DDM	DDM
2	UDM	UDM
3	DM	Cymal-6
4	Cymal	DM
5	C12E8	C12E8
6	LMNG	LMNG
7	FC12	FC12
8	LDAO	LDAO
9	CHAPS	CHAPS
10	CHAPSO	CHAPSO
11	OG	Control
12	Control	OG

Figure 4.

- A. Fluorescence Size-Exclusion Chromatography (FSEC) detergent screening of the samples of Figure 3.
- B. Screening ranking comparison between the FIDA assay and FSEC, ranking 1 was the best detergent that solubilized the most protein. ** The samples with DDM and UDM have practically the same level of solubilized GFP- LacY.

	FIDA	FSEC
Sample volume for 12 detergents	1.5uL (triplicates)	1440uL (singlets)
Exp. time for 12 detergents	<3h (triplicates)	20h (singlets)

Table 1: Comparison of experimental specifications between FIDA and FSEC detergent screening

CONCLUSIONS

The data demonstrate that the Fida 1 system offers a new approach to detergent screening that has significant advantages allowing essential data to be obtained faster using less material. The FIDA assay uses only 120nL of crude protein solution at 0.1 mg/ml for three replicates, and the whole volume leftover (\sim 19.8 μ L) is fully recovered, available for further experiments. The experimental time was less than 3h for 12 detergents in triplicates compared to \sim 20h for one replica in FSEC. In addition to FSEC and Fida 1 providing the same answers, Fida 1 gives aquantifiable PDI.

Altogether, the results of this work presents a new approach to characterize membrane proteins that allows users to reducing costs and time as well as to analyze protein expressed at low levels in a short time thus overcoming any stability issues.