Product Information Sheet

Fluidity One-M

Quantify and characterize any protein interaction – even in complex backgrounds, even with challenging targets

Molecular size, $K_{\rm D}$, concentration and stoichiometry from a single experiment





Explore interactions in solution under physiologically-relevant conditions

- Analyze samples in different backgrounds from simple buffers to cell lysates, serum and saliva
- Obtain simultaneous measures of $K_{\rm D}$ and concentration within 35 minutes
- Gain insights into interaction stoichiometry; distinguish on-target binding from off-target binding

Application examples

Characterize polyclonal antibodies

Track functional immune response in serum samples

Reveal therapeutic antibody / protein interaction mechanisms Characterize
disordered
proteins or
higher-order
complexes under
close to in vivo
conditions

Explore
aggregation
effects and
distinguish
between specific
and non-specific
binding

Take a closer look

- Uses microfluidic diffusional sizing (MDS) technology to measure changes in molecular size (hydrodynamic radius) as binding events occur
- Enables development of customized protocols to study a wide range of interactions typical run time 35 minutes for 24 datapoints to determine K_n
- Eliminates risk of binding artefacts or other surface constraints measure directly in solution no surface immobilization
- Minimizes consumption of precious samples 3.5 μ L per datapoint (application dependent), 50-80 μ L to determine K_D

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Specifications

System	
Application	Determine size, $K_{\rm D}$, concentration and stoichiometry in buffer systems or complex backgrounds such as serum or plasma
Technology	Microfluidic Diffusional Sizing (MDS)
Interaction analysis	
Run time	Typically within 35 min for 24 datapoints to determine $K_{\scriptscriptstyle D}$
Size range: hydrodynamic radius, $R_{\rm h}$	1 – 20 nm
Accuracy of size determination	± 10%
Reproducibility of size determination	CV < 10%
Working range molecular weight	1.4 kDa – 14 MDa
Sensitivity range (labeled HSA in PBS)	100 pM − 10 μ M Alexa Fluor TM 488 100 pM − 10 μ M Alexa Fluor TM 647
Typical sample consumption to determine protein $K_{\rm D}$	50 – 80 μL
Sample volume per datapoint	3.5 µL (application dependent)
Compatibility	Compatible with crude, complex backgrounds such as undiluted serum or plasma
	Compatible with aqueous and biological buffers including components such as TRIS, HEPES, PBS, NaCI, KCI, TWEEN, DMSO and DMF
Datapoints per run	Up to 24 datapoints per run
Fluorescent labels	Alexa Fluor™ 647 and equivalents, RFP/Cy5, Alexa Fluor™ 488 and equivalents, GFP/FITC, Fluidiphore labeling kit (fluidiphore rapid amine 503)
Data export	USB Mass Storage Device / Fluidity Cloud
Exported data file formats	CSV
Data output from Fluidity Cloud	Result tables, binding curves and affinity ($K_{\rm D}$), size ($R_{\rm h}$) of complex and labeled species
Consumables	
	Kits sufficient for 192 datapoints
Specifications	
Temperature control	25 °C (actively controlled)
Operating environment	15 °C to 30 °C
Power requirements	100 – 240 V AC, 50 – 60 Hz
Safety and EMC standards	Designed to comply with all relevant safety and EMC standards
Dimensions	
Dimensions (D x W x H; mm)	666 x 432 x 489 (Drawer Out)
Weight (kg)	35

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