Meet Jess Your Complete Protein Analysis Solution



biotechne

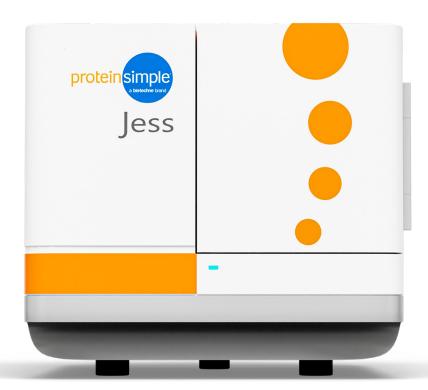
Meet Jess

Protein analysis comes with many challenges—labor-intensive protocols increase time to result and multiple hands-on steps increase user error and data variability. At best, you end up with Western blot-like, semi-quantitative results when what you really need, and deserve, is highly reproducible immunoassay quantitation.

Meet Jess, your protein analysis problem solver Jess automates the protein separation and immunodetection of traditional Western blotting, eliminating many of the tedious, error-prone steps. Just load your samples and reagents into the microplate, and Jess does the rest. She separates your protein by size and precisely manages antibody additions, incubations, washes and even the detection steps. Come back to fully analyzed

and quantitated results in just 3 hours. With Jess's chemiluminescent detection you'll get picogram-level sensitivity, letting you maximize the data you get from your sample. Go further with multiplexing—her fluorescent detection gets all the information you need in one shot. Best of all, you get reliable and reproducible results with target normalization to the amount of protein loaded.

Analysis is a breeze. Want to identify whether a protein is present or absent? Jess gives you the qualitative Western blot data you are used to seeing. Even better, she'll quantitate the results for you too. With just a few clicks you'll be analyzing immunoassay-like standard curves and precisely quantifying your protein. Jess, she's like Western blot meets ELISA in one.





How Can Jess Help You?



Reproducible

Jess precisely controls sample loading, incubations and washes; she eliminates the inconsistencies and user-dependent variability that can be introduced during traditional Westerns. She delivers intra-assay CVs <15%, giving you the consistency you need to be confident in your data.



Off High Throughput

With her 13 and 25 sample capillary cartridges, you'll get the throughput you need, with minimum hands-on time. Use Jess's fluorescent capabilities and multiplex your proteins for even higher throughput.



With Jess, it's pipette, run and done! Simply load your sample, antibodies and reagents into the plate, insert your plate and cartridge into Jess and press start. In just 3 hours of hands-free runtime, you can be analyzing data for your next publication or grant.



Quantitative

With Jess, protein quantification is a breeze. At the conclusion of your run, use the lane view option to compare band intensity or dive deep for fully quantitative analysis of protein size and concentration. With a few clicks, you'll be analyzing immunoassay-like standard curves and precisely quantifying your protein.

Why Settle? Do Protein Analysis Your Way, Quickly.

Jess Gives You **Four Different Ways** to Analyze Proteins

1 Fluorescence Detection

Why bother **stripping and reprobing**? Maximize your time and sample, and get the information you need in one shot, by multiplexing.

2 Chemiluminescence Detection

Working with low abundance targets or precious samples? **Chemiluminescent detection** gives you picogram-level sensitivity, letting you maximize the data you get from your sample.

3 Protein Normalization

Jess gives you an easy way to see if your samples contain a consistent protein load: either use her proprietary fluorescent protein normalization reagent or total protein reagent in a RePlex assay to measure proteins immobilized in the same capillary as your immunoassay. Best of all, Jess's multiple detection capabilities enable two-color protein detection for multiplexing and chemiluminescent detection, on top of protein normalization.

4 Blot Imaging

Still doing traditional Westerns? Snap! Get the picture with **Jess's blot imaging system**.

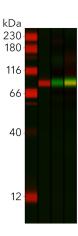


Figure 1.

Jess's fluorescent detection capabilities enable two-color protein detection for multiplexing. Detection of Stat3 (green) and phospho-Stat3 (red) in K562 cell lysate. Lane 1: ladder, Lane 2: phospho-Stat3; Lane 3: Stat 3; Lane 4: phospho-Stat3/Stat3.

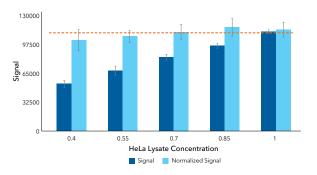


Figure 2.

With fluorescent protein normalization, you can take your protein load comparison and transform your data to effectively normalize your samples, increasing your confidence in your data interpretation.

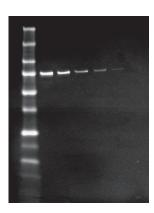
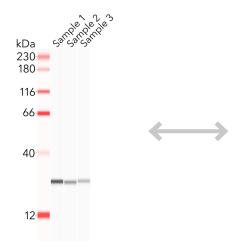


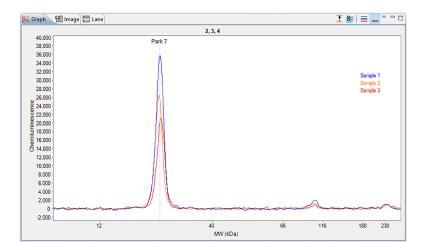
Figure 3.

Jess's imaging system allows for imaging of traditional Western blotting membranes.

Stop, Analyze and Wow!

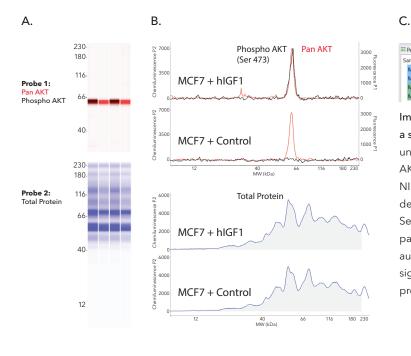
At the end of your run, use the lane view option to compare band identity or dive deep for fully quantitative analysis of protein size and concentration. Dive deeper to compare protein expression changes and analyze protein isoforms or size changes. Want to analyze expression changes between samples or compare runs? Jess's protein normalization will give you the confidence you need in your analysis.





Peaks Capillaries																
Sa	ample	Primary	Secondary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width	S/N	Baseline	Channel	^
	Hela Lysate	Park 7	GAM	2	1	Park 7	311	27	35508.8	316798	50.0	8.4	356.5	703.9	CHEMI	
	Hela Lysate	Park 7	GAM	3	1	Park 7	307	27	25799.8	228724	50.0	8.3	293.2	694.5	CHEMI	
	Hela Lysate	Park 7	GAM	4	1	Park 7	306	27	20867.5	193841	50.0	8.7	235.0	746.1	CHEMI	~

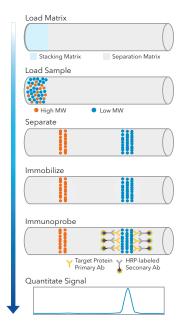
Automated Immunoassays and Total Protein Normalization **Equals Better Quantification**



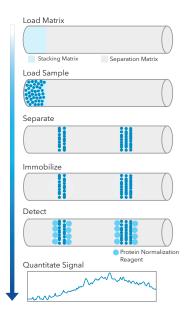
Immunoassay and total protein detection performed in a single capillary. AKT phosphorylation in MCF7 lysates untreated and activated with h-IGF1. Phospho-AKT and pan AKT were detected in Probe 1 using chemiluminescence and NIR fluorescence, respectively, while total protein signal was detected in Probe 2 (A). Example Graph views of pan AKT, AKT Ser473 phosphorylation and total protein signal for samples in panel A (B). Peaks Table in Compass for Simple Western shows automated normalization of phosphorylated and pan AKT signal to total protein signal demonstrates quantitation of target protein expression (C).

How Does Jess Work?

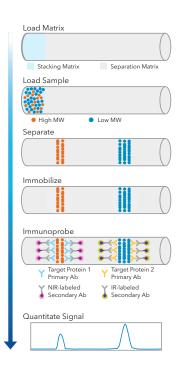
Chemiluminescent Detection



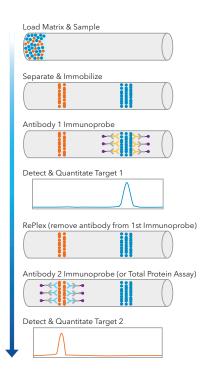
Fluorescent Protein Normalization



Fluorescent Detection



RePlex™ with Jess™



Specifications

	Total Protein Specification	Chemiluminescence Specification	Fluorescence Specification	Protein Normalization Specification						
Sample Required	0.3-1.2 µg	0.6-1.2 µg	2-4 μg	0.6-3.6 µg						
Volume Required	3 μL/well									
Size Range	Molecular weight (MW) ladder ranges from 2-440 kDa									
Sizing CV	<10%									
Intra-assay CV		<20%								
Inter-assay CV	<20%†	<20%*	<20%	<20%						
Resolution (± percent difference in MW)	± 15-20% for MW <20 kDa ± 10% for MW >20 kDa									
Quantitation CV	<20% (total pro	N/A								
Dyanamic Range	2-3 logs	3-4 logs	3-4 logs	1 log						
Sensitivity	ng	Low pg	High pg	ng						
Capillary	5 cm, 100 μm, 400 nL									
Runtime		<4 hours with immunoassay								
Samples Per Run	13 or 25									
Weight	23 kg									
Dimensions (Closed)	0.36 m H x 0.3 m W x 0.57 m D									
Dimensions (Open)	0.36 m H x 0.53 m W x 0.57 m D									
Power	US/CAN 120 V AC, 60 Hz, 4.2 amps Europe 240 AC, 50 Hz, 2.1 amps Japan 100 AC, 50/60 Hz, 5.0 amps									
Operating Temperature	18-24 °C									
Operating Humidity	20-60% relative, non-condensing									

^{*} Inter-assay CV is calculated with normalization to the system control protein

[†] Percent peak area

Where Science Intersects Innovation®

Bio-Techne* | R&D Systems™ Novus Biologicals™ Tocris Bioscience™ ProteinSimple™ ACD™ ExosomeDx™ Asuragen®

Contact Us

Global info@bio-techne.com bio-techne.com/find-us/distributors
North America TEL 800 343 7475
Europe | Middle East | Africa TEL +44 (0)1235 529449
China info.cn@bio-techne.com TEL +86 (21) 52380373

For research use or manufacturing purposes only. Trademarks and registered trademarks are the property of their respective owners.

