

Your Comp Analysis So

At ProteinSimple, we're changing the way scientists analyze proteins. Our innovative product portfolio helps researchers reveal new insight into proteins, advancing their understanding of protein function. We enable cutting-edge research to uncover the role of proteins in disease and provide novel approaches to develop and analyze protein-based therapeutics. We empower you to make your next discovery by eliminating common protein analysis workflow challenges.



many challenges—labor-intensive protocols increase time to result and increase user error and data variability. At best, you end up with Western results when what you really need, and deserve, is highly reproducible

analysis problem solver. Jess automates the protein separation and immunodetection process, eliminating many of the tedious, error-prone steps. Just load your samples into the plate, and Jess does the rest. She separates your protein by size and precisely detects, incubates, washes and even the detection steps. Come back to fully analyzed results in just 3 hours. With Jess's chemiluminescent detection you'll get picogram-level sensitivity to analyze the data you get from your sample. Go further with multiplexing—her fluorescent detection gives you the information you need in one shot. Best of all, you get reliable and reproducible results proportional to the amount of protein loaded.

Want to identify whether a protein is present or absent? Jess gives you the qualitative results you need, from a single click to seeing. Even better, she'll quantitate the results for you too. With just a few minutes and Jess's standard curves and precisely quantifying your protein. Jess, she's like a one-stop Western.

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.



HIGH THROUGHPUT

With her 13 analysis cartridges, you'll get the results you need, with minimal hands-on time. Use Jess's high-throughput and multiplexing capabilities for higher throughput.



FAST

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 standard curve view option to quickly analyze your data. Dive deep for full protein size and intensity information. In just a few clicks, you'll be able to generate standard curves for your protein.

Do protein analysis your way, quickly.

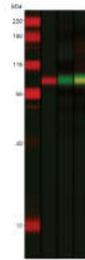
DIFFERENT WAYS

reprobing? Maximize your information you need

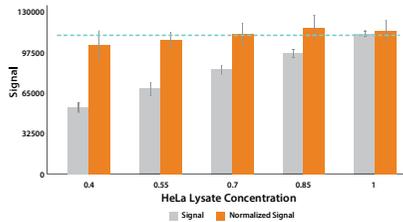
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ce targets or precious samples?
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to see if your samples contain
ust load her in-capillary protein
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proprietary fluorescent protein
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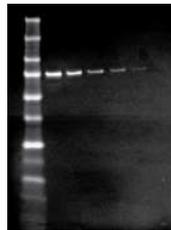
rn's? Snap! Get the picture
em.



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.



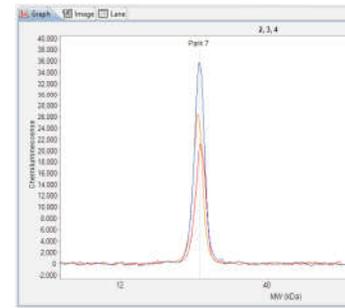
With protein normalization, you can take your protein load comparison and transform your data to effectively normalize your samples, increasing your confidence in your data interpretation.



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 intensity or dive deeper into analysis of protein size and concentration. Dive deeper to compare protein expression between protein isoforms or size changes. Want to analyze expression changes between samples? Jess's protein normalization will give you the confidence you need in your analysis.



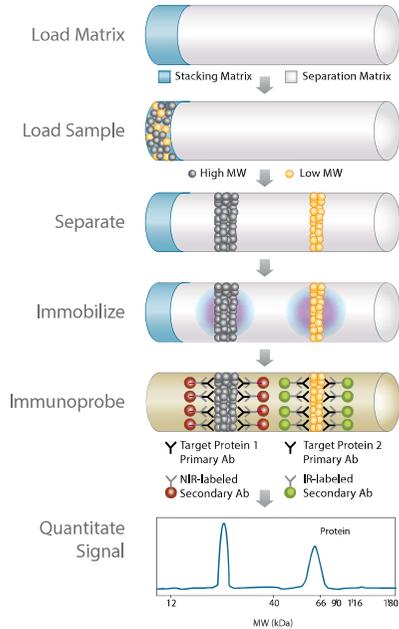
Sample	Primary	Secondary	Cap	Peak	Name	Position	MW (kDa)	Height	Area	% Area	Width
Hela Lysate	Park 7	GAM	2	1	Park 7	311	27	35508.8	316798	50.0	8.4
Hela Lysate	Park 7	GAM	3	1	Park 7	307	27	25799.8	228724	50.0	8.3
Hela Lysate	Park 7	GAM	4	1	Park 7	306	27	20867.5	193841	50.0	8.7

How does it work?

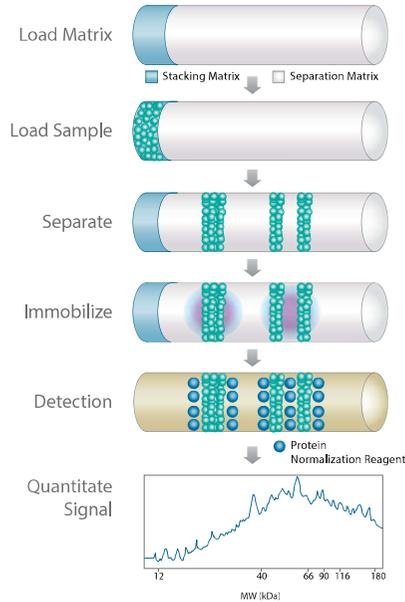
Detection



Fluorescent Detection



Protein Normalization



Specifications

DESCRIPTION	TOTAL PROTEIN SPECIFICATION	CHEMILUMINESCENCE SPECIFICATION	FLUORESCENCE SPECIFICATION
Sample required	0.3-1.2 µg	0.6-1.2 µg	2-4 µg
Volume required	3 µL/well		
Size range	Molecular weight (MW) ladder ranges from 2–440 kDa		
Sizing CV	<10%		
Intra-assay CV	<15%		
Inter-assay CV	<20%		
Resolution (± percent difference in MW)	± 15–20% for MW <20 kDa ± 10% for MW >20 kDa		
Quantitation CV	<20% (total protein, chemiluminescence and fluorescence)		
Dynamic range	2-3 logs	3-4 logs	3-4 logs
Sensitivity	ng	Low pg	High pg
Capillary	5 cm, 100 µm, 400 nL		
Runtime	<3 hours		
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		



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