

THE SIMPLE WAY TO

Your Next Breakthrough

Evolve Your Protein Analysis with Simple Western Systems



03

Move Beyond Traditional Western Blots

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 semi-quantitative results when what you really need, and deserve, is highly reproducible immunoassay quantitation. Discover more and solve your protein analysis problems with Simple Western™ Systems.



Meet Simple Western Systems

- Identify whether a protein is present or absent
- · Quantitate protein expression changes
- · Identify and analyze phosphorylated isoforms
- Characterize post-translational modifications
- Normalize target expression to protein load
- Delve deeper into isoform analysis with charge separation

What Does the Data Look Like?

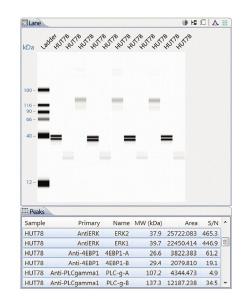
Simple Western size-based assay data is processed automatically in Compass™ for Simple Western™

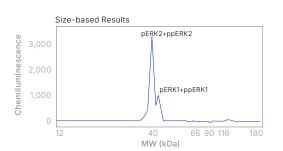
Software for you. Sample data is displayed by lane in a virtual-blot like image similar to traditional results with one big exception—not only do you get more information, you get it as soon as the assay is complete. Quantitative results such as molecular weight, signal intensity (area), % area, and signal-tonoise for each immunodetected protein are presented in the results table automatically.

For those who are more accustomed to capillary electrophoresis, we offer traditional electropherogram view that allows you to visualize your results with ease.

Just need total protein? We've got that covered too!

FIGURE // 01





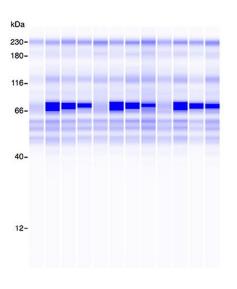
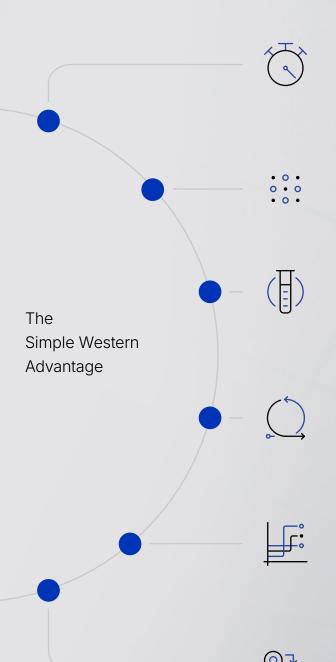


FIGURE 1. Total Protein detection. Decreasing concentrations of DNAK in Hela lysate and negative controls (15, 7.5 & $3.75 \mu g/mL$).

How Can Simple Western Technology Help You?



Fast and Fully Automated

With Simple Western Systems, it's pipette, run and done! Simply load your sample, antibodies and reagents into the plate, insert your plate and cartridge into the Simple Western Instrument instrument and press start. With as little as 3 hours of hands-free run time, you can be analyzing data and making decisions with your data in no time.

Highly Sensitive Multiplex

Get pico-gram level sensitivity in chemiluminescence and fluorescence channels and the advantage of **flexible multiplex assays** including total protein normalization of immunoassay results.

Low Sample volume

Use as little as 3 μ L of sample per well so you can save your precious samples and maximize data from each sample.

Reproducible

By eliminating the variability from manual steps of traditional western blotting protocols including sample loading, gel-to-membrane transfer of proteins, and user and site differences, Simple Western technology is highly reproducible offering <15% intra-assay CVs.

Quantitative

With Simple Western Systems, protein quantitation is a breeze. With 25 or 100 independent sample throughput per run, Simple Western enables inclusion of standard curves, QC standards, and controls to design robust fully quantitative protein assays.

Differentiate

Simple Western charge assays enable the identification of charge isoforms such as mono- or di-phosphorylated states of your target protein. Get another dimension to your data beyond size.



How Simple Western Works

01

Prepare Your Reagents

Reagent prep with Simple Western System isn't complicated. Pipette, mix, spin...Soon you'll be a protocol prodigy.

02

Load the Plate

After preparing the reagents, simply load the reagents and samples into a Simple Western Sample Plate.

03

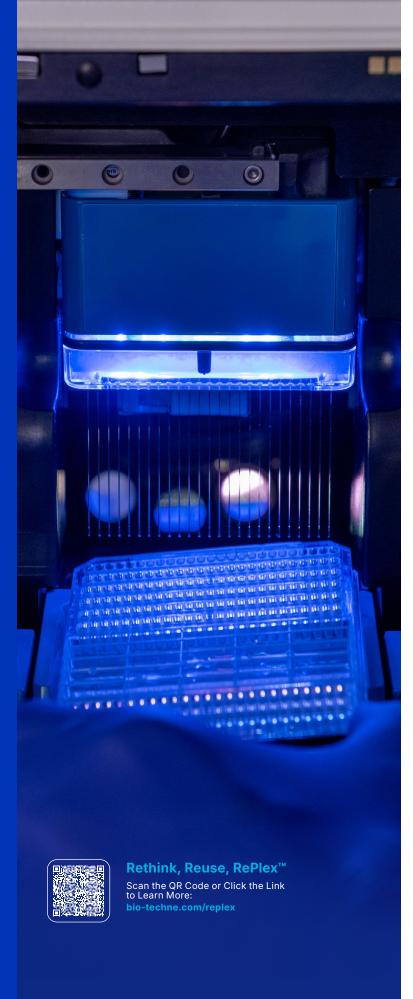
Start the Run

Place the Sample Plate, any additional Pre-filled Reagent Plate(s) needed and the capillary cartridge(s) into the Simple Western Instrument and use Compass Software to start a run. Walk away and take back your lost time.

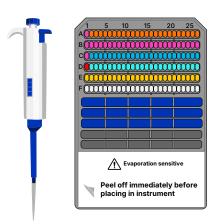
04

Your Simple Western System Does the Rest

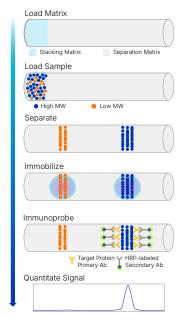
Simple Western System analyze proteins through separation and immunoprobing in a capillary-based system. Each step of the process is precisely controlled, ensuring the highest quality results. Come back to fully analyzed, quantitative results.



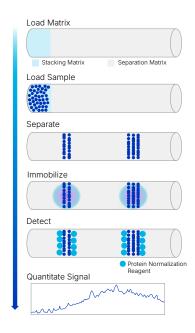
It's pipette, run & done! Simple Western does the rest.



Sized-Based Assay:Chemiluminescent Detection

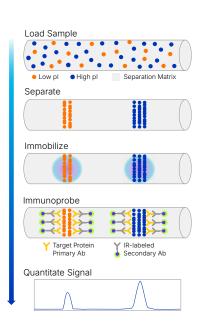


Sized-Based Assay: Fluorescent Protein Normalization



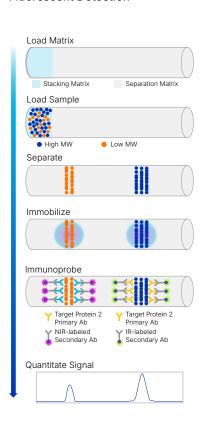
Charge-Based Assay:

Chemiluminescent Detection



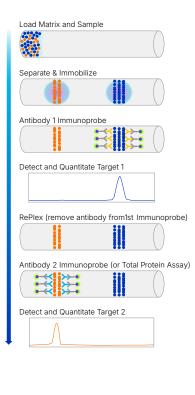
Sized-Based Assay:

Fluorescent Detection



Replex Assay

with Leo, Jess & Abby



06

Stop, Analyze, & Wow!

At the end of your run, you'll have multiple options for data viewing. For size-based runs, use the laneview option to compare band intensity like in traditional Western blotting (FIGURE 2) or view the electropherogram of the protein separation by size and intensity (FIGURE 3). Easily turn these qualitative views into fully quantitative tables or standard curves to dive deeper into quantitative analysis, allowing you to compare protein expression changes and analyze protein isoforms or size changes (TABLE 1).

With NanoPro 1000™ System, distinguish phosphorylation changes in proteins using the cIEF separation. View your results in electropherogram format and take your analysis further by examining the signal intensities across the isoelectric point (pI) range (FIGURE 4).

FIGURE // 02

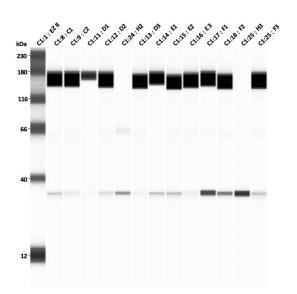


FIGURE 2. The virtual lane view option in Compass Software for Simple Western Technology lets you compare protein separation and band intensity, like in traditional western blotting.

Want to analyze expression changes between samples or compare runs? Choose either a chemiluminescence-based Total Protein Module or fluorescence-based Protein Normalization kit to normalize your immunoassay data so you can have the confidence you need in your analysis.

Use RePlex[™] Assay with Leo[™], Jess[™] and Abby[™]
Systems to run two serial immunoassays within
the same capillary so you can quantify expressed
phosphorylated target and total target levels.

Get all your rich protein characterization data from just one sample!

FIGURE // 03

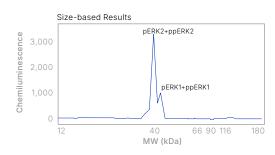


FIGURE 3. Electropherogram graph view of protein separation by size and intensity in Compass for Simple Western Software.

FIGURE // 04

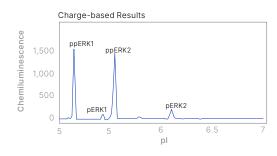


FIGURE 4. Electropherogram graph view of charge-based Simple Western assay data using NanoPro 1000 System. Distinguish protein isoforms like phospho-protein isoforms (shown here) separated by their isoelectric point.



TABLE // 01

Sample	Primary	Cap	Peak	Position	MW (kDa)	Height	Area	Width	S/N	Baseline
C1	anti-G	C1:8	1	430	34	1844.8	17439	8.9	94.1	178.1
C1	anti-G	C1:8	2	601	159	45597.5	537092	11.1	2331.9	255.7
C2	anti-G	C1:9	1	430	34	1119.1	13081	11.0	87.8	186.4
C2	anti-G	C1:9	2	519	63	531.1	7662	13.6	34.5	219.0
C2	anti-G	C1:8	3	601	159	57978.6	653486	10.6	4672.4	281.1

TABLE 1. Compass for Simple Western Software automatically calculates fully quantitative results for each peak detected in your sample and even has a built-in standard curve tool to calculate absolute protein expression levels.

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Simple Western Systems: Pick your Perfect Match



LEO

- Size-based protein analysis
- Up to 100 samples per run
- Fully analyzed results in 3 hours of run time
- Chemiluminescence detection
- RePlex assay
- Chemi-based in-cap total protein normalization



JESS

- Size-based protein analysis
- 13 or 25 samples per run
- Fully analyzed results in 3 hours of run time
- Chemiluminescence & Fluorescence detection
- RePlex assay

- Two approaches for in-cap total protein normalization: fluorescence-based Protein Normalization or chemiluminescence-based Total Protein Assay
- Stellar High Sensitivity NIR/IR detection



ABBY

- Size-based protein analysis
- 13 or 25 samples per run
- Fully analyzed results in 3 hours of run time
- Chemiluminescence detection
- RePlex assay
- Chemi-based in-cap total protein normalization



NANOPRO 1000

- Charge-based analysis
- 96 samples per run
- Fully analyzed results in 11-19 hours of run time
- Chemiluminescence detection

Specifications

Instrument Specifications

System	Leo	Jess	Abby	NanoPro 1000	
Simple Western size assays	•	•	•		
Simple Western charge assays				•	
Max samples per run	100	25	25	96	
Run time	<3 hours	<3 hours	<3 hours	11-19 hours	
Minimal sample volume required	3-5 μL/well	3 μL/well	3 μL/well	5–12 μL/well	
Sample cooling	N/A	N/A	N/A	3° C	
Detection	Chemiluminescence	Chemiluminescence Fluorescence	Chemiluminescence	Chemiluminescence	
RePlex™ (Strip & ReProbe Replacement)	Yes	Yes Yes		No	
Stellar™ (NIR / IR fluorescence detection)	No	Yes	No	No	
Dimensions (H x W x D)	72 x 74 x 74 cm	33 x 33 x 52 cm	33 x 33 x 52 cm	94 x 94 x 61 cm	
Weight	136 kg	23 kg	21 kg	82 kg	
Part Number	004-550	004-650	004-680	004-109	

Immunoassay Size Specification							
Description	Chemiluminescence on Abby & Jess	Stellar Fluorescence on Jess	Classic Fluorescence on Jess	Immunoassay Charge			
Sample required	0.6-1.2 μg	0.6-1.2 μg	2-4 μg	0.6-1.2 μg			
Volume required	3 μL/well	3 μL/well	3 μL/well	5–12 μL/well			
Size range	Molecular weight (MW) ladders range from 2–440 kDa	Molecular weight (MW) ladders range from 2–440 kDa	Molecular weight (MV) ladders range from 2–440 kDa	Widest gradient ranges from pl 3 to pl 10			
Sizing CV	<10%	<10%	<10%	<10%			
Intra-assay CV	<15%	<15%	<15%	<20%			
Inter-assay CV	<20%*	<20%*	<20%*	<20%**			
Resolution (± percent difference in MW)	± 15–20% for MW <20 kDa ± 10%	± 15-20% for MW <20 kDa ± 10%	± 15–20% for MW <20 kDa ± 10%	± 1 pl units			
Quantitation CV	<20%	<20%	<20%	<20%			
Dynamic range	Up to 4 logs	Up to 4 logs	Up to 4 logs	3 logs			
Sensitivity	Low pg	Low pg	Low pg	Low pg			

^{*} Inter-assay CV is with system control

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^{**} Percent peak area



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INCLUDES R&D Systems" Novus Biologicals" Tocris Bioscience" ProteinSimple" ACD" ExosomeDx" Asuragen Lunaphore"

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