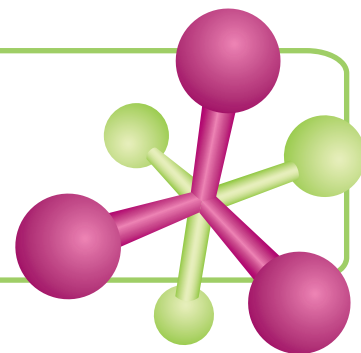


Progenesis SameSpots v4.0



Reproducible 2D gel analysis is possible Be confident in your proteomics data...

Progenesis SameSpots is the 2D gel analysis product of choice for reproducible proteomics results. It enhances your analysis workflow with speed, objectivity and statistical power. You also have features to perform detailed investigations of your data, support spot picking and generate custom reports.

Speed

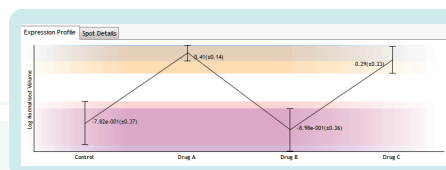
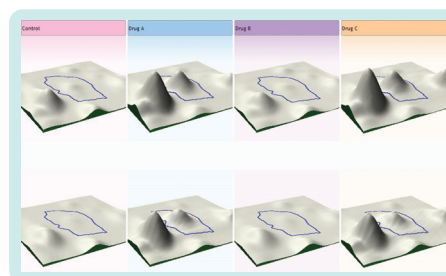
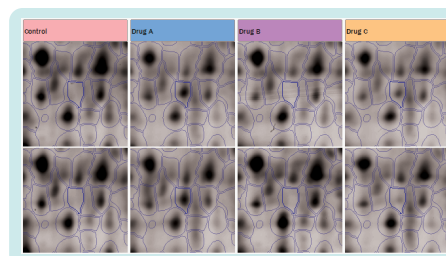
Typical analysis times of 5mins/image* with a fast, streamlined workflow. Faster analysis times means you can run more replicates.

Objectivity

Easy-to-use and proven to give highly reproducible results across-labs. Post analysis editing is eliminated on the majority of experiments.

Statistics

Increased statistical power with 100% matching in your experiment and no missing values, so you can apply valid multivariate statistics.



See what some of our customers have to say:

*Timings based on single-stain experiment with 2 groups, 5 replicates/group, image size of 9MB

John E. Wiktorowicz, Ph.D.,
The University of Texas Medical Branch, USA

"Progenesis SameSpots has increased our throughput - with the demand we experience, this results in tremendous savings in manpower and expense over a year. With the proper tools and attention to detail proteomics can reap enormous benefits whatever your field of biological study."

Dr. Kondo,
National Cancer Research Institute, Tokyo, Japan

"We are currently running more than 1500 2D-DIGE gels per year. Progenesis SameSpots accurately analyses such large gel numbers in a very short period and generates good data for data-mining."

Dr. Hans Voshol,
Novartis Institutes for BioMedical Research, Basel, Switzerland

"We use 2D gels because they provide a reproducible and largely unbiased view of the proteome. SameSpots reinforces that objectivity and reproducibility, since it offers a superb automatic analysis combined with optimal reviewing tools."

Jorgen Ostling,
AstraZeneca R&D, Molndal, Sweden

"..it takes full advantage of the DIGE concept, not only for generating the quantitative data but also by its unique matching technology which generates a 100% match rate, regardless how many samples that are analyzed."

Prof. Steve Pennington, UCD Conway Institute Ireland and VP and President of The British Society of Proteome Research (BSPR)

"By adopting these new (SameSpots) image analysis technologies, our rate of discovery using 2D electrophoresis stands to be significantly enhanced."

Todd M. Umstead,
Penn State University College of Medicine, USA

"I can now complete an analysis in less than one-quarter of the time that it would have taken when using the previous supplier's software, while at the same time obtaining more consistent and reliable results. I have no doubt that SameSpots will very quickly pay for itself in the terms of salary time alone."

