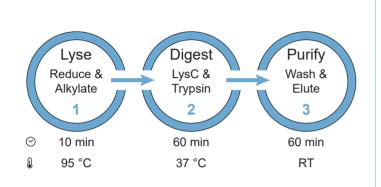


Technologies for your proteomic applications

Standardized sample preparation for MS-based proteomics

iST technology



PreOmics is as a spin-off from the Matthias Mann lab at the provide innovative technologies for mass spectrometry-

Our inStageTip (iST) Kit enables a robust and

reproducible sample preparation with a significant time

advantage (>40 hrs saved) compared to common

protocols. The iST technology is suitable for

integration within semi-automated high-throughput

PreOmics

< 2,5 hours total time

< 1 hour hands-on-time

extracellula

sample preparation workflows.

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4 hours at 37°C

~ 44 hours total time

preparation methods.

-18% -15%

> 4,5 hours hands-on-time

12 hours at -20°C

Common protocol

Products: Sample Cleanup

Phoenix Peptide Cleanup Kit

removes detergents, polymers, salts, lipids and more



Phoenix 96× P.O.00023 | EUR 1599

- exceptional peptide recovery
- simple-to-use, sensitive and fast
- ready-to-go aliquots for best quality

Products: Accessories for iST

for high-throughput sample processing



96well adapter P.O.00002 | EUR 150

- for MTP plate centrifuges
- two adapters



Heating shaker adapter P.O.00004 | EUR 1080

- for MTP plate shakers
- compatible with automated liquid handling platforms
- optimal heat transfer to iST cartridge

All items are available directly from PreOmics, please contact us at order@preomics.com.

We offer worldwide shipping from our warehouses in Germany and the USA.

Am Klopferspitz 19, 82152 Martinsried, Germany Phone: +49 (0) 89 2314 1630 info@preomics.com

Products: Sample Preparation

iST Kits

compatible with label-free and metabolic labeling



iST8× P.O.00001 | EUR 299

- all-in-one for any starting material
- simple-to-use, sensitive and fast
- ready-to-go aliquots for best quality



iST 96× P.O.00027 | EUR 2590

- all advantages of the iST 8×
- high-throughput applications
- compatible with automated liquid handling platforms

compatible with chemical labeling (iTRAQ, TMT)



iST-NHS 12× P.O.00026 | EUR 449

- all-in-one for any starting material
- simple-to-use, sensitive and fast
- ready-to-go aliquots for best quality



iST-NHS 96× P.O.00030 | EUR 2590

- all advantages of the iST-NHS 12×
- high-throughput applications
- compatible with automated liquid handling platforms
- AP-MS Sample preparation of GFP-fusion proteins



iST GFP-Trap 8× gtak-iST-8 | EUR 450

- GFP-Trap coupled to agarose beads
- manage co-IP & MS sample prep
- ready-to-go aliquots for best quality
- available via our partner ChromoTek GmbH at www.chromotek.com

How does the iST Kit work? Benefits of iST Kit

robust & sensitive

- high reproducibility: R² >0.9
- working range: 1-100 μg protein input
- increased peptide & protein identifications

rapid results

- 1 hour hands-on time
- single-step lysis, reduction and alkylation
- all-in-one solution, all reagents provided

cleanest peptide samples

- dual cleanup step removes hydrophilic &
- results in ready-to-measure peptides
- compatible with standard lab equipment

applications

- proteome discovery
- validation of biosimilars
- immunoprecipitations
- absolute & relative quantification

sample types

- cell lines
- fresh & FFPE tissues
- body fluids (serum, plasma, CSF, urine)
- antibodies & host cell proteins
- bacteria, yeast & plant tissues

Customer Statements



"We tested PreOmics` iST Kit and were amazed by the results. The number of identified peptides and proteins was 20-30% higher compared to what is achieved using our standard lab procedure. Using the iST Kit enables you to start processing your samples in the morning and have them ready for LC-MS in the afternoon."

Dr. Manuel Tzouros, Project Leader, Discovery Proteomics, Roche Innovation Center, Basel, Switzerland



"Using this new sample preparation technology, we increased the efficiency of tryptic peptide cleavage and obtained better identification rates in our samples. In addition, the short hands-on-time for the entire workflow, as well as the comparatively low amounts of starting material, now enable us to process more and be more timeefficient."

Dr. Andreas Tebbe, Head of Mass Spectrometry, Evotec Munich



"Our researchers happily rely on the PreOmics iST kit knowing that they will have stabilized the sources of technical variations, permitting them to focus on the biology of our precious samples. The PreOmics team has worked closely with our researchers to consult with us on our unique biological questions and concerns"

Dr. Sasha Singh, Director Proteomics Research, Brigham and Women's Hospital, Harvard Medical School, Boston, USA

high protein identifications



10,000

>15 cell lines



>10 tissues











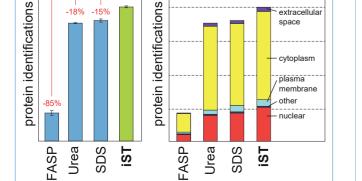




4.500 1,500 >22 conditions

>3 strains

>CSF/urine



Customer validation indicates the highest level of

protein identifications and no functional bias when

compared to other common proteomic sample