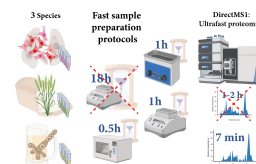


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# DirectMS1 Ultrafast proteomics: Protein extraction & digestion

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From Days to Minutes: Accelerated Protein Digestion for Fragmentation-Free Ultrafast Mass Spectrometry

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**We use this protocol and it's working**

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**Keywords:** ultrafast proteomics, microwave digestion, accelerated protein digestion, fragmentation-free mass spectrometry, proteomics, sample preparation, directms1 ultrafast proteomic, digestion ultrafast proteomic workflow, throughput analysis in clinical proteomic, clinical proteomic, bottleneck of overnight protein digestion, high reproducibility in protein quantitation, overnight protein digestion, protein extraction, protein quantitation, rapid sample preparation, trypsin digestion, protein, mass spectrometry, yeast quantitation standard, reducing preparation time, employing liquid chromatography, preparation time, liquid chromatography

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## Abstract

Ultrafast proteomic workflows, employing liquid chromatography-mass spectrometry, demand equally rapid sample preparation. To overcome the bottleneck of overnight protein digestion, this study optimized reduction, alkylation, and trypsin digestion using ultrasound, microwave irradiation, and elevated temperatures, achieving completion within 30-60 minutes. Validation using various sample types (*S. cerevisiae*, A549 cells, winter wheat) and human-yeast quantitation standards confirmed the reproducibility of label-free quantification. The microwave-assisted method, reducing preparation time 35-fold, showed high reproducibility in protein quantitation, demonstrating the potential for high-throughput analysis in clinical proteomics.



## Guidelines

For troubleshooting help, feel free to message to the corresponding author [iatarasova@gmail.com](mailto:iatarasova@gmail.com).

Avoid salts and non-volatile buffers in your final purification steps, if possible. In most cases, dry your sample down in a speed-vac and re-suspend it in 50mM ammonium bicarbonate pH 8.3 for a tryptic digestion.

The list of reagents can be found in the 'Materials' section



## Materials

	A
	Acetonitrile LC-MS
	Ammonium bicarbonate (NH <sub>4</sub> HCO <sub>3</sub> )
	MilliQ water
	SDS
	Dithiothreito l
	Iodoacetami de (single- use vial 56mg)
	ProteaseMA X(TM) Surfactant Trypsin Enhancer, 1mg
	Pierce BCA Protein Assay Kit
	Methanol
	Chloroform
	Urea
	Sequencing Grade Modified Trypsin
	Copure <sup>®</sup> C18 Endcapped Octadecyl
	Trifluoroace tic acid



## Protocol materials

- ✕ Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #23225**
- ✕ Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #23225**
- ✕ Methanol **Merck MilliporeSigma (Sigma-Aldrich) Catalog #M3641**
- ✕ Chloroform
- ✕ Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #23225**
- ✕ Urea **Merck MilliporeSigma (Sigma-Aldrich) Catalog # 57-13-6**
- ✕ SDS **Bio-Rad Laboratories Catalog #1610301**
- ✕ Dithiothreitol **neoFroxx Catalog # 3483-12-3**
- ✕ Iodoacetamide (single-use vial 56mg) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #A3221**
- ✕ Acetonitrile LC-MS **Biosolve Catalog #01207802**
- ✕ Ammonium bicarbonate (NH<sub>4</sub>HCO<sub>3</sub>) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #AC393212500**
- ✕ ProteaseMAX(TM) Surfactant Trypsin Enhancer, 1mg **Promega Catalog #V2071**
- ✕ Acetonitrile LC-MS **Biosolve Catalog #01207802**
- ✕ Ammonium bicarbonate (NH<sub>4</sub>HCO<sub>3</sub>) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #AC393212500**
- ✕ ProteaseMAX(TM) Surfactant Trypsin Enhancer, 1mg **Promega Catalog #V2071**
- ✕ Sequencing Grade Modified Trypsin **Promega Catalog #V5111**
- ✕ Dithiothreitol **neoFroxx Catalog # 3483-12-3**
- ✕ Iodoacetamide (single-use vial 56mg) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #A3221**
- ✕ Sequencing Grade Modified Trypsin **Promega Catalog #V5111**
- ✕ Dithiothreitol **neoFroxx Catalog # 3483-12-3**
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- ✕ Iodoacetamide (single-use vial 56mg) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #A3221**
- ✕ Dithiothreitol **neoFroxx Catalog # 3483-12-3**
- ✕ Sequencing Grade Modified Trypsin **Promega Catalog #V5111**
- ✕ Copure® C18 Endcapped Octadecyl **Biocomma**
- ✕ Copure® C18 Endcapped Octadecyl **Biocomma**
- ✕ Copure® C18 Endcapped Octadecyl **Biocomma**
- ✕ MilliQ water



⊗ Methanol **Merck MilliporeSigma (Sigma-Aldrich) Catalog #M3641**

⊗ Copure® C18 Endcapped Octadecyl **Biocomma**

⊗ Copure® C18 Endcapped Octadecyl **Biocomma**

⊗ Urea **Merck MilliporeSigma (Sigma-Aldrich) Catalog # 57-13-6**

⊗ Trifluoroacetic acid (TFA)

⊗ Iodoacetamide (single-use vial 56mg) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #A3221**

⊗ Sequencing Grade Modified Trypsin **Promega Catalog #V5111**

⊗ Trifluoroacetic acid (TFA)

⊗ Trifluoroacetic acid (TFA)

⊗ Trifluoroacetic acid (TFA)

⊗ Trifluoroacetic acid (TFA)

## Troubleshooting

## Safety warnings

⚠ To ensure safety, reactions with chloroform, methanol and trifluoroacetic acid must be carried out in a fume hood. The experimenter must be protected with gloves throughout the experiment.


## Ethics statement




The authors declare no competing ethics statement.

## Yeast protein extraction



2h


- 1 5m  


 Sample 1\*10<sup>9</sup> yeast cells.


Add  100 µL Lysis buffer volume to the  Sample 


Lysis buffer:



 10 % volume  Acetonitrile LC-MS **Biosolve Catalog #01207802**

 50 millimolar (mM)


 Ammonium bicarbonate (NH<sub>4</sub>HCO<sub>3</sub>) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #AC393212500**

 0.1 Mass / % volume

 ProteaseMAX(TM) Surfactant Trypsin Enhancer, 1mg **Promega Catalog #V2071**
- 2 1h  

 600 rpm, 25°C, 01:00:00 
- 3 5m  

Sonication for 2 min (10 seconds on/off cycles) at each 30, 60, 80% amplitudes on ice

 0 °C

### Equipment

QSonica SONICATOR SYSTEM Q800R1	NAME
Sonicator	TYPE
QSonica	BRAND
discontinued	SKU



4  15000 rcf, 00:10:00

10m



5 Collected the supernatant

5m

6 Concentration measurement

35m

 Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #23225**

## Equipment

SPECTROstar Omega

NAME

Microplate reader

TYPE

BMG LABTECH

BRAND

None

SKU



<https://www.bmglabtech.com/en/spectrostar-omega/><sup>LINK</sup>

## A549/MCF7 protein extraction






2h 15m

7  A549 or MCF7 cells


5m

Add  100  $\mu$ L Lysis buffer volume to the  Sample

Lysis buffer:


 10 % volume  Acetonitrile LC-MS **Biosolve Catalog #01207802** 50 millimolar (mM) Ammonium bicarbonate (NH<sub>4</sub>HCO<sub>3</sub>) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #AC393212500** 0.1 Mass / % volume ProteaseMAX(TM) Surfactant Trypsin Enhancer, 1mg **Promega Catalog #V2071**



8  600 rpm, 25°C, 01:00:00

1h



9  550 rpm, 95°C, 00:05:00

5m



10 15 minutes in ultrasonic bath 50 Hz

15m

Equipment

PSB-1335-03

NAME

Ultrasonic bath

TYPE

PCB-GALS

BRAND

1335-03

SKU

11  10000 rcf, 00:10:00

10m



12 Collected the supernatant

5m

13 Concentration measurement

35m

 Pierce BCA Protein Assay Kit **Thermo Fisher Scientific Catalog #23225**



### Equipment

SPECTROstar Omega

NAME

Microplate reader

TYPE

BMG LABTECH

BRAND


None

SKU


<https://www.bmg-labtech.com/en/spectrostar-omega/> <sup>LINK</sup>

## Wheat protein extraction

2h 49m

14  Sample 250 mg of plant tissue

5m

15 Disrupt the  Sample using

3m

### Equipment

CryoMill

NAME

Retsch GmbH, Berlin, Germany

BRAND


20.749.0001

SKU

for 3 min at 30 Hz (3 cycles)

16 Add  1 mL Lysis buffer

5m


(  8 Molarity (M)


 Urea Merck MilliporeSigma (Sigma-Aldrich) Catalog # 57-13-6



in



[M] 2 % volume     SDS Bio-Rad Laboratories Catalog #1610301    )    to  
 250 mg    of homogenate

17    incubate  1000 rpm, 25°C, 01:00:00    1h  


18     20000 rcf, 00:15:00    15m  


19     Pierce BCA Protein Assay Kit Thermo Fisher Scientific Catalog #23225    35m  




Equipment















Microplate reader	NAME
BMG LABTECH	BRAND
no	SKU
<a href="https://www.bmglabtech.com/en/spectrostar-omega/">https://www.bmglabtech.com/en/spectrostar-omega/</a>	LINK

20    Collect supernatant    5m

21    Chloroform - methanol extraction    






21.1    collect  200 µL    supernatant    2m

21.2    add  800 µL    2m  
 Methanol Merck MilliporeSigma (Sigma-Aldrich) Catalog #M3641
















- 21.3 add  200 µL  Chloroform 2m
- 21.4 add  600 µL  MilliQ water 2m
- 21.5 centrifugate  15000 rcf, 00:15:00 15m  

- 21.6 discard supernatant over interphase 2m
- 21.7 add  600 µL 2m  
 Methanol **Merck MilliporeSigma (Sigma-Aldrich) Catalog #M3641**
- 21.8 centrifugate  7000 rcf, 00:07:00 7m  

- 21.9 discard supernatant 2m
- 21.10 dissolve the pellet in 1M 2 Molarity (M)  200 µL 5m  
 Urea **Merck MilliporeSigma (Sigma-Aldrich) Catalog # 57-13-6** 
- 22 Digestion protocols    
Protocol #1 - overnight protein digestion (19h total time)  
Protocols #2-5 - 30-60 minutes protein digestion time

#### STEP CASE

#### Protocol #1 5 steps

- 23 Add  Dithiothreitol **neoFroxx Catalog # 3483-12-3** up to 1M 10 millimolar (mM) 20m  
 500 rpm, 56°C, 00:20:00   



- 24 Add  Iodoacetamide (single-use vial 56mg) **Merck MilliporeSigma (Sigma-Aldrich) Catalog #A3221**    30m  
up to [M] 20 millimolar (mM) Incubate in the dark for  00:30:00
- 25 Add  Sequencing Grade Modified Trypsin **Promega Catalog #V5111**    18h  
[M] 1/75 %(m/m)  
 500 rpm, 37°C  Overnight
- 26 Add  Trifluoroacetic acid (TFA) up to [M] 1 % volume   5m
- Desalting** 20m
- 27 C18 desalting  Copure® C18 Endcapped Octadecyl **Biocomma** 20m

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