

Oct 27, 2025

In solution digestion for protein standards (Urea)

DOI

<https://dx.doi.org/10.17504/protocols.io.ewov19db7lr2/v1>



Cristina CARDENAL PERALTA¹

¹University of Edinburgh



Cristina CARDENAL PERALTA

University of Edinburgh

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.ewov19db7lr2/v1>

Protocol Citation: Cristina CARDENAL PERALTA 2025. In solution digestion for protein standards (Urea). **protocols.io**
<https://dx.doi.org/10.17504/protocols.io.ewov19db7lr2/v1>

License: This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited



Protocol status: Working

We use this protocol and it's working

Created: June 14, 2024

Last Modified: October 27, 2025

Protocol Integer ID: 101821

Keywords: Protein Standards, trypsin, Lys-C, Mass Spectrometry, solution digestion for protein standard, digestion of protein standard, protein standard, different protein standard, spectrometry instrument, solution digestion, digestion, house quality standards for technical calibration, preparation for injection

Abstract

Digestion of protein standards, such as albumin, is a common practice to keep in-house quality standards for technical calibration of Mass-Spectrometry instruments. Here we describe a protocol to digest three different protein standards and their preparation for injection in a Mass-Spectrometry instrument.

Guidelines

-



Materials

Protein Standards

🧪 1 mg Albumin from bovine serum

🧪 1 mg Creatine kinase (rabbit)

🧪 1 mg Beta-lactoglobulin (bovine)

Buffers

[M] 8 Molarity (M) Urea in ABC

[M] 50 millimolar (mM) Ammonium bicarbonate (ABC))

[M] 0.1 % volume TFA (trifluoro acetic acid)

Reagents

[M] 1 Molarity (M) DTT (Dithiothreitol)

[M] 1 Molarity (M) IAA (Iodoacetamide)

Enzymes

Lys-C proteinase

Trypsin proteinase

Equilibration buffers for SEP-PAK C18

Solution A: 0.1 % TFA in Milli Q water

Solution B: 0.1 % TFA, 80 % ACN (acetonitrile)

C18 Cartridge Vac 1cc (100 mg)

Sep-Pak C-18 (Waters) WAT023590

Safety warnings

⚠ Incubation with IAA needs to be performed in the dark.

Before start

Make the Urea and ABC buffers fresh on the same morning of the experiment.



Steps (Day 1)

5h 20m

- 1 Weight 1 mg of each standard and resuspend in 100 μ L of 8 M Urea in ABC. 10m
- 2 Add DTT to a final concentration of 25 mM, incubate for 30 min at Room temperature in a shaker (1200 rpm) 30m
- 3 Add IAA to a final concentration of 55 mM, incubate for 30 min Room temperature in the **dark**. 30m
- 4 Add 5 μ L of Lys-C (stock at [M] 1 μ g/ μ L), for 3-4 hours at 37 $^{\circ}$ C 4h
(This is adding 1 μ g of Lys-C per 20 μ g of protein).
- 5 Dilute sample 5X times (add 400 μ L) with ABC buffer 5m
This step is performed because trypsin does not work in high urea concentrations. Be careful to use the ABC buffer and NOT the one with Urea and ABC when diluting.
- 6 Add 5 μ L of trypsin (stock at [M] 1 μ g/ μ L , resuspend in 20 μ L of 0.1 % TFA) to each sample. Incubate O/N at 37 $^{\circ}$ C 5m
(This is adding 1 μ g of trypsin per 20 μ g of protein).

Steps (Day 2) Sep-Pak C-18 cartridge loading

20m

- 7 Equilibrate the Sep-Pak C-18 column:
Flush the column with 5 ml of solution B or Acetonitrile 5m
- 8 Add 10 ml of solution A. Add the digested sample into the 10 ml of solution A and let it into the Sep-Pak column slowly. 5m
- 9 Wash the sample with 10 ml of Buffer A 5m
- 10 Elute with 2 $\times$ 600 μ l of solution B (collect your purified sample in a 2 mL tube) 5m



- 11 Dry down purified peptides completely in a speed-vac. Resuspend in Buffer A (up to 20 μ M)

