

STANDARD OPERATING PROCEDURE

Title: Trypsin digestion of serum and enrichment of N-glycopeptides using an automated liquid handler

Version #: 1

Author: Hui Zhang Lab

Date: 05/08/2014

Purpose

The purpose of this document is to describe the enzymatic digestion of serum and the subsequent solid phase enrichment of N-glycopeptides using an automated liquid handler to prepare samples that are compatible with mass spectrometry analysis.

Scope

This procedure is used to reduce, alkylate, proteolyze and enrich N-linked glycopeptides from serum.

Responsibilities

It is the responsibility of person(s) performing this procedure to be familiar with laboratory safety procedures. The interpretation of results must be done by a person trained in the procedure and familiar with such interpretation.

Equipment

- Versette liquid handling system (Thermo Fisher Scientific)
- Vacuum centrifuge

Materials

- Disposable automation research pipette tips (Thermo Fisher Scientific)
- 96-well plate
- Affi-gel hydrazide gel (Bio-Rad)
- Frits (2 mm diameter, 1 mm thick, 15 – 45 μ m pores) (POREX)
- 7k MWCO Zeba spin desalting columns (Thermo Fisher Scientific)
- Human Serum (Sigma-Aldrich)

Reagents

- Sodium periodate (Bio-Rad)
- Urea (Thermo Fisher Scientific)
- Tris(2-carboxyethyl)phosphine (Thermo Fisher Scientific)
- Iodoacetamide (Sigma-Aldrich)
- Sequencing-grade trypsin (Promega)
- PNGase F (New England Biolabs)
- Sodium acetate (Sigma-Aldrich)
- Sodium chloride (Sigma-Aldrich)
- Aniline (Sigma-Aldrich)
- Ammonium bicarbonate (NH_4HCO_3) (Sigma-Aldrich)
- Acetonitrile (Fisher Scientific)
- Formic acid (Fisher Scientific)
- Water (Fisher Scientific)
- Trifluoroacetic acid (Thermo Pierce Scientific)

Solutions

- Coupling buffer: 500 mM sodium acetate, 0.3 mM sodium chloride, pH 5
- Oxidation buffer: 15 mM sodium periodate
- 100 mM aniline
- Urea buffer: 8 M urea in 0.4 M NH_4HCO_3
- 10 mM TCEP
- 12 mM iodoacetamide
- 1.5 M sodium chloride
- 80% acetonitrile
- 50% acetonitrile
- 25 mM NH_4HCO_3
- 100 mM NH_4HCO_3

Procedure

1. Prepare hydrazide pipette tips
 - a. Push one frit into a disposable automation research tip (DART)
 - b. Load 200 μL hydrazide resin (50% slurry) into each pipette tip
 - c. Force any residual liquid out of the tip
 - d. Push one 5 mm diameter frit into the tip to secure the hydrazide resin between the two frits
 - e. Wash each tip 5x with 200 μL deionized water

- f. Condition each tip 5x with coupling buffer
 - g. Note: Discard tips with slow flow rates due to high resistance
- 2. Couple glycoproteins to hydrazide resin
 - a. Note: The loading capacity of hydrazide beads is ~ 40 µL serum/200 µL hydrazide beads (50% slurry), and the average glycopeptide yield per 40 µL of serum is 20 µg.
 - b. Dilute serum 1:1 with oxidation buffer
 - c. Oxidize with oxidation buffer containing 100 mM aniline for 1 h at room temperature in the dark
 - d. Buffer exchange with coupling buffer using Zeba spin desalting column
 - e. Slowly aspirate serum sample into hydrazide tips and dispense into 96-well plate for 30 min using Versette
- 3. Denature, reduce, alkylate and proteolyze glycoproteins
 - a. Wash serum coupled to the hydrazide tips via their oxidized glycans with 3 mL of urea buffer
 - b. Reduce disulfide bonds with 10 mM TCEP for 30 min
 - c. Alkylate with 12 mM iodoacetamide for 15 min in the dark at room temperature
 - d. Wash with 3 mL of urea buffer
 - e. Digest with trypsin (1:120) in 100 mM NH_4HCO_3 for 1 h
- 4. Remove residual non-glycopeptides released by trypsin digestion
 - a. Wash hydrazide tips (containing conjugated glycopeptides) with 6 mL of 1.5 M sodium chloride
 - b. Wash hydrazide tips with 6 mL of 80% ACN
 - c. Wash hydrazide tips with 6 mL of deionized water
 - d. Wash hydrazide tips with 6 mL of 25 mM NH_4HCO_3
- 5. Release N-linked glycopeptides
 - a. Release glycopeptides with 1500 U PNGase F in 25 mM NH_4HCO_3 for 1 h at room temperature
 - b. Wash tips 3x with 50% ACN
 - c. Combine eluents and dry in a vacuum centrifuge
- 6. Reconstitute samples
 - a. Re-constitute in 1% TFA
 - b. Measure peptide concentration using NanoDrop
 - c. Dilute to working concentration with 2% ACN/0.2% formic acid

Referenced Documents

Chen J, Shah P, Zhang H. Solid phase extraction of N-linked glycopeptides using hydrazide tip. *Analytical Chemistry*. 2013 (85):10670-10674. PMID: 24079330