

## In Gel Digestion of Proteins for MALDI-MS

### Buffers and Solutions for In Gel Digests:

#### *500 mM Ammonium Bicarbonate [(NH<sub>4</sub>)HCO<sub>3</sub>]*

- \* (NH<sub>4</sub>)HCO<sub>3</sub>-1.6 g
- \* Millipore water-to 40 mL final volume
- \* Filter through high protein binding filter [Nitrocellulose or PTFE; 0.45 μm pore size]

#### *25 mM (NH<sub>4</sub>)HCO<sub>3</sub>*

- \* 500 mM (NH<sub>4</sub>)HCO<sub>3</sub>-2 mL
- \* Millipore water-to 40 mL final volume
- \* Check pH (should be between 8.0-8.5)

#### *25 mM (NH<sub>4</sub>)HCO<sub>3</sub>/50% Acetonitrile [ACN]*

- \* 500 mM (NH<sub>4</sub>)HCO<sub>3</sub>-2 mL
- \* ACN (HPLC grade)-20 mL
- \* Millipore water-18 mL
- \* Make sure pH stays at 8.0-8.5

#### *500 mM Dithiothreitol [DTT] stock*

- \* DTT-77.1 mg
- \* 25 mM (NH<sub>4</sub>)HCO<sub>3</sub>-to 1 mL final volume
- \* Store at -20 °C in 50 μL aliquots

#### *550 mM Iodoacetamide [IAA] stock*

- \* IAA-101.7 mg
- \* 25 mM (NH<sub>4</sub>)HCO<sub>3</sub>-to 1 mL final volume
- \* Store at -20 °C in 50 μL aliquots

#### *30 mM Potassium Ferricyanide (III) [K<sub>3</sub>Fe(CN)<sub>6</sub>]*

- \* K<sub>3</sub>Fe(CN)<sub>6</sub>-494 mg
- \* Millipore water-to 50 mL final volume
- \* Wrap in aluminum foil and store at 4 °C up to three months.

#### *100 mM Sodium Thiosulfate [Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub>]*

- \* Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub>-237 mg
- \* Millipore water-to 50 mL final volume
- \* Store at 4 °C up to three months

#### *Trypsin stock*

- \* Trypsin [Promega Modified Sequence Grade]-20 μg
- \* 25mM (NH<sub>4</sub>)HCO<sub>3</sub>-1 mL (resuspend gently at room temperature)
- \* Store at -20 °C in 100 μL aliquots

#### *10% N-octyl Glucoside*

- \* N-octyl Glucoside [SIGMA]-250 mg
- \* Millipore water-to 2.5 mL final volume
- \* Store at room temperature

### Gel fragment preparation

\* Excise protein bands. Cut each into 1 mm pieces (not so small that they clog the pipet tip). Place into a low-binding Eppendorf tube (siliconized FISHER brand). Also cut out a gel piece from a protein-free region as control.

\* Wash gel pieces with >10 volumes of Millipore water [~200 μL] for 10 minutes with intermittent vortexing, to wash out acetic acid.

## Destaining

\* For Coomassie Blue / Colloidal staining, destain **two times** (or until the color disappears) for 15 minutes with intermittent vortexing (low setting) with 250  $\mu\text{L}$  of 50% Methanol in 100 mM  $(\text{NH}_4)\text{HCO}_3$ . Remove the solutions and discard. Dehydrate **twice** for 2-5 minutes with intermittent vortexing (low setting) with 200  $\mu\text{L}$  neat Acetonitrile [ACN]. Remove the solutions and discard. The gel slices shrink and become white.

\* For Non-Destructive Silver staining, destain **twice** for 15 minutes with intermittent vortexing (low setting) with 200ul of freshly prepared 1:1 solution of 100mM Sodium Thiosulfate  $[\text{Na}_2\text{S}_2\text{O}_3]$  and 30 mM Potassium Ferricyanide  $[\text{K}_3\text{Fe}(\text{CN})_6]$ . Discard the supernatants. Stop the reaction and wash out silver ions **twice** for 5 minutes with intermittent vortexing (low setting) with 250  $\mu\text{L}$  of Millipore water. Equilibrate slices for 10 minutes with intermittent vortexing in 200  $\mu\text{L}$  25mM  $(\text{NH}_4)\text{HCO}_3$  then dehydrate twice for 10 minutes vortexing with 200  $\mu\text{L}$  25mM  $(\text{NH}_4)\text{HCO}_3/50\%$  ACN. Discard supernatants. The gel slices shrink and become white.

\* For Sypro Ruby staining, no destaining necessary, dehydrate **twice** for 10 minutes with intermittent vortexing (low setting) with 200  $\mu\text{L}$  25 mM  $(\text{NH}_4)\text{HCO}_3/50\%$  ACN. Remove the solutions and discard. The gel slices shrink and become white.

\* Dry gel particles for 10 minutes in a vacuum centrifuge.

## Reduction and Alkylation

\* Rehydrate gel slices in 40-50  $\mu\text{L}$  of freshly prepared 25 mM DTT [in 25 mM  $(\text{NH}_4)\text{HCO}_3$ ]. Reduce the proteins for 30 minutes at 56 °C.

\* Cool the samples to room temperature, pipet off any residual liquid and add 40-50  $\mu\text{L}$  (same volume as previous step) of freshly prepared 55 mM Iodoacetamide [in 25 mM  $(\text{NH}_4)\text{HCO}_3$ ]. Alkylate the proteins for 30 minutes at room temperature in the dark.

\* Wash gel slices with 250  $\mu\text{L}$  of Millipore water for 15 minutes, and discard supernatant. Resuspend gel in 200  $\mu\text{L}$  of 25 mM  $(\text{NH}_4)\text{HCO}_3$  for 10 minutes with intermittent vortexing and discard supernatant. Dehydrate gel pieces **twice** for 2-5 minutes each time with 200  $\mu\text{L}$  of neat ACN. Discard the supernatants.

\* Dry gel particles 10 minutes in a vacuum centrifuge.

## Trypsin Digestion

\* Rehydrate gel slices for 15 minutes at 4°C in 20-30  $\mu\text{L}$  [20 ng/ $\mu\text{L}$ ] Trypsin (Promega Sequence Grade Modified) in 25 mM  $(\text{NH}_4)\text{HCO}_3$  [make sure pH = 8.0-8.5].

\* Overlay the rehydrated gel particles with a minimum amount of 25 mM  $(\text{NH}_4)\text{HCO}_3$  to keep them immersed throughout the digestion. Incubate 16-24 hours at 37 °C.

## Peptide Recovery

- \* Transfer the digest solution to a new low binding microcentrifuge tube.
- \* Extract digested peptides with one volume (40-60  $\mu\text{L}$ ) Millipore water/0.1% TFA by vortexing 20 minutes (high setting) at room temperature. Transfer solution to the low binding microcentrifuge tube.
- \* Perform an additional extraction with 60-70  $\mu\text{L}$  of 5% TFA/70% ACN by vortexing 20 minutes (high setting).
- \* Pool extracted peptides and dry completely in a vacuum centrifuge (approx. 1 hour).

## References

- \* Jimenez, C.R. Current Protocols in Protein Science. 1998; 16.4.1-16.4.5
- \* Gharahdaghi, F. *et al.* Electrophoresis.1999; 20: 601-605.
- \* Hirouki, K. *et al.* Rapid Commun. Mass Spectrom.