## Desalting of tryptic peptide mixtures using C18 Zip-tips

This step is needed only for the direct analysis of tryptic peptide mixtures by MALDI mass spectrometry. If the sample will be prepared for MS analysis using a nano-HPLC or submitted to LC-ESI mass spectrometry, this step can be omitted.

Materials Trifluoroacetic acid (Fluka) Zip-tips, C18 (Millipore)

1. Dissolve dried peptide mixtures (obtained from steps 14 through 19) in 10  $\mu$ l of 5 % acetonitrile containing 0.1% trifluoroacetic acid.

For not completely dried peptide mixtures, add ultrapure water to make a volume of 8.5  $\mu$ l, and add 0.5  $\mu$ l acetonitrile and 1  $\mu$ l of 1% trifluoroacetic acid.

Check pH of resulting peptide mixture by using a pH paper; confirm and eventually adjust pH of samples to 2-4 using 1 % trifluoroacetic acid.

- Prepare Zip-tip C18 (Millipore). Activate Zip-tip by rinsing 3 times with 10 µl acetonitrile. Equilibrate Zip-tip 3 times with 10 µl of 5 % acetonitrile containing 0.1 % trifluoroacetic acid.
- 3. Bind the peptides by aspirating and dispensing peptide solution; use up to 10 cycles.
- 4. Wash with 10  $\mu$ l of 5 % acetonitrile containing 0.1% trifluoroacetic acid.
- 5. Elute peptides with 10  $\mu$ l of 50 % acetonitrile containing 0.1 % trifluoroacetic acid. Carefully, aspirate and dispense eluant through Zip-tip without introducing air; repeat up to10 times.

Alternatively, elute the digested peptide with 10  $\mu$ l of 20% acetonitrile containing 0.1% trifluoroacetic acid and with 10  $\mu$ l of 60% acetonitrile containing 0.1% trifluoroacetic acid. Combine these two eluants.

- 6. Dry eluant using a Speed-Vac centrifuge at 35 °C.
- 7. If necessary, store at -20 °C (< 24 hr) or -80 °C (for longer periods).
- 8. Prepare sample for MALDI mass spectrometric analysis:

Resolubilize the desalted peptide lyophilate with 1  $\mu$ L matrix solution (10  $\mu$ g/ $\mu$ L a-cyano-4-hydroxy-cinnamic acid in 0.1% trifluoroacetic acid:acetonitrile, 1:1, v/v).

Load a half microliter of sample mixture on a MALDI target plate (with 144 spots for the AB 4700 instrument) and let dry at room temperature. The sample is now ready for MALDI mass spectrometry.

Oregon State University

Maier Lab – Biomolecular Mass Spec

Filename of document: DESALTING OF TRYPTIC PEPTIDE MIXTURES USING C18 ZIP TIPS