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## Seifert Lab SOP – Protein Extraction

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We use this protocol and it's working

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## Abstract

This protocol outlines the basics of protein extraction from ear pinna tissue.

## Troubleshooting

## Tissue Collection

- 1 Anesthetize the animals, harvest tissue and dissect the 4mm punch.
- 2 Hold the tissue in the forceps and rinse it quickly in HBSS/PBS.
- 3 Transfer it into the freshly labelled 1.5ml Eppendorf tube and snap freeze in liquid nitrogen.
- 4 Transport to lab and store it at -80°C freeze for further use.

## Protein Extraction

- 5 Note: Before starting, set the centrifuge at 4°C.
- 6 The collected tissue is taken out of -80°C freeze and kept in ice bucket.
- 7 Add 200-400ul of cold protein extraction buffer per sample.
- 7.1 Cold Extraction buffer: In 1ml of RIPA buffer, freshly add 10ul each of protease inhibitor cocktail, Na-orthovanadate and PMSF.
- 8 Use the manual homogenizer to dislodge the tissue (10-15 times). Put it back in ice.
- 8.1 Homogenize the dislodged tissue using sonicator.
- 8.2 You require 1, 15ml falcon of DI water and 1, 15ml falcon of 70% ethanol.
- 8.3 Conditions on sonicator machine: Pulse-Continues, Control cycle-4 and duty cycle-40%.



- 8.4 First wash the sonicator twice with 70% ethanol (10 sec each) followed by twice with DI water (10 sec each).
- 8.5 Now sonicate your protein sample-3 times with 20-25 sec pulse followed by 30 sec in ice.
- 8.6 In between the samples, wash the sonicator with water followed by 70% ethanol for 5 sec each.
- 8.7 After finishing extracting protein from all the samples, clean the sonicator by twice with DI water (10 sec each) followed by twice with 70% ethanol (10 sec each).
- 9 Centrifuge the samples at 13,000 rpm for 15 min at 4°C.
- 10 Collect the supernatant in freshly labelled 1.5ml Eppendorf tubes. From here, either store it at -80°C freeze for further use or go for protein quantification.

## Protein Quantification

- 11 We will take 96-well plate and add different concentrations of BSA standards (2.5ug, 5ug, 10ug, 20ug and 40ug).
- 12 The protein samples will be loaded as 2ul or 5ul/well in triplicates.
- 13 Now add 100ul of BSA reagent in each well.
- 14 Go for plate reading according to their saved protein estimation protocol.
- 15 The estimated protein will be calculated and used accordingly for western blot.