

**Burridge Laboratory Stress Fiber Isolation Protocol**  
(Adapted from Katoh et al., 2000)

**STOCK Solutions**

4X PBS

250mM TEA (triethanolamine)

20% NP-40

20% Triton X 100

Protease Inhibitors (working conc.)

1. 20 ug/ml aprotinin
2. 1 ug/ml leupeptin
3. 1 ug/ml pepstatin

**WORKING Solutions (prepare fresh)**

PBS + protease inhibitors (PBS)

Low ionic strength extraction buffer (LISE buffer) : 2.5mM TEA + protease inhibitors  
pH 8.2 (dH<sub>2</sub>O)

Extraction buffer 1 : 0.05% NP-40 + protease inhibitors pH 7.2 (PBS)

Extraction buffer 2 : 0.5% Triton X-100 + protease inhibitors pH 7.2 (PBS)

**METHODS**

Wash cells (Ref52 fibroblasts 15-10cm dishes) with PBS containing 1mM MgCl  
(MgCl is used at this step to keep the cells attached to the dish during the extraction in  
LISE buffer.)

Extract with LISE buffer for 20-40 minutes with 5 buffer changes; (Removes soluble  
components-at this point cells swell)

Extract with Extraction buffer 1 for 5 min.

(Removes dorsal surface of cells and nuclei, basal portion of cells remaining as well as  
some nuclei)

Extract remaining components with Extraction buffer 2 for 5 minutes.

Immediately wash with PBS to remove triton (stress fibers are unstable in triton)

Scrape off stress fibers from culture dish and suspend in small volume (<5ml) PBS

(At this point stress fibers still contain cell cortex and focal adhesion components)

Extrude 2X through Z-shaped needle; Separates free individual stress fibers from cortex

Centrifuge at 1000g for 5' to remove large debris

Maybe higher speed to remove focal adhesion complex

Collect stress fibers by centrifugation at 100,000g for 1 hour.

Lipids were removed by Chloroform/Methanol extraction

Pellet was suspended in Ammonium Bicarbonate (AMBIC) and submitted for Mass Spec.

Control stress fiber isolation to identify contaminating proteins in prep.

Ref52 fibroblasts were treated for 30 minutes with 10uM Y-27632 (Calbiochem)

Stress fibers were isolated as above.