## S1 PROTECTIONS

## <u>Materials:</u>

1. 10 X Hybridization Buffer

10 X = 4M NaCl 0.4M Pipes 6.8 0.02M EDTA

- 2. S1 Digestion Buffer 66 mM Na0Ac 0.3 M NaCl 4mM ZnS04
- 3. tRNA 10 mg/ml
- 4. Formamide

## **Procedure:**

- 1. Ethanol precipitate RNA sample, end labeled template, and carrier tRNA. Total amount of RNA should be 100 µg.
- 2. Resuspend pellet with  $10 \ \mu l \ H_2 0$ . Vortex.
- 3. Add 80 µl formamide, vortex.
- 4. Add 10 g 10X hybridization buffer. Vortex. Finger Spin.
- 5. Incubate at 68°C for 15 minutes. (65 to 75°C temp blot).
- 6. Transfer to 53°C and incubate for >4 hours. Upon transfer, tubes should not cool below 53°C. (Bring temp. block to water bath overnight.)
- 7. To each tube:

Remove from 53°C; Add 300µl ice cold S1 buffer; Quickly finger mix; Place tube in ice; Next tube, etc.

- 8. Add 0.5-1 μl S1 nuclease to each tube. If you have many samples you can add the S1 nuclease to the S1 buffer and add both of these in step 7.
- 9. Incubate 37°C, 60 minutes.
- 10. To stop digestion phenol extract.
- 11. Add 1ml ethanol (only) to precipitate. Dry pellets well (spin vac if necessary).
- 12. Resuspend pellets in  $10 \ \mu l H_2 0$ .
- 13. Remove an aliquot and add formamide dye. Because you are not loading your entire sample, you have the option to go back and re-digest your samples if you find that digestion was incomplete.
- 14. Incubate dye containing samples at 95°C for 5-10 minutes. Quick chill on ice.
- 15. Load on urea-polyacrylamide gel. Run 25 mA.
- 16. Fix gel with 7% acetic acid, 15 minutes.
- 17. Soak in H<sub>2</sub>O or 1X TBE, 15-20 minutes.

18. Dry and expose.

## **Comments:**

1. Originally, the amount of S1 nuclease required was titrated to digest 100 µg of total RNA. Conveniently, tRNA can be added at Step #1 to achieve this. However, if using large amounts of tRNA bothers you, there are two options:

a. In Step #1, leave out the carriertRNA. Then, before phenol extracting in Step #10, add 20  $\mu$ g tRNA, and

--*or*--

- b. As little as 20  $\mu$ g carrier can be used in Step #1.
- 2. You must be absolutely sure that you are in template excess. There is an easy way, and a hard way, to confirm this.
  - Hard way: Do a series of protections with a constant amount of template and increasing amounts of test RNA. Over a range from 1 µg to 100 µg, your signal should increase proportionally.
  - Easy way: Load a 1:100 dilution of undigested template on gel along with your samples and compare the intensity of this band to those of your protected samples.
- Sure fire way to make template: Short version. (See end labeling section for specific details.)
  Digest 50-75 μg plasmid DNA. Phosphatase, kinase, etc. Resuspend final fragment in 100-200 μl TE. 1 μl is sufficient per S1. (200-400 cps per μl of template is ideal.)
- 4. For endogenous protections, use 5-10 μg of cytoplasmic RNA. For protections of transfected genes, use 10-100 μg of RNA.