

# Trace Fluorescent Labeling for Protein Crystallization Screening

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## Hands-on Practical Outline.

- I. Introduction to covalent modification of proteins.
  - A. Several groups are commonly available as targets.
    1. Amines – side chain and N-terminal.
    2. Sulfhydryls – less commonly available.
    3. Carboxyls
  - B. Easiest is Amines – use pH control to select type.
    1. pH ~7.5, N-terminal amines
    2. pH ~>8.5, side chain amines.
    3. I prefer side chain, less chance of systematic effects.
- II. Solutions & Materials needed.
  - A. Reaction buffer – 0.05M NaBorate, pH 8.75 used here. Must be free of primary and secondary amines (tertiary OK)!
  - B. Reactive probe solution – carboxyrhodamine-SE used (5 mg in 1 mL DMF).
  - C. Centrifugal desalting column – 0.5 mL (Pierce). Note – max speed is 1500xG.
  - D. Crystallization buffer solution – whatever your protein is equilibrated in. I'm using 0.05 M NaHepes, 0.1 M NaCl, pH 7.5.
- III. Procedure
  - A. Equilibrate desalting column in reaction buffer (2-3 passes).
  - B. Prepare protein solution – demo uses 0.8 mLs at 15 mg/mL as the stock solution.
  - C. Pass 80 uL of protein solution thru desalting column.
  - D. Add 0.5 uL of reactive probe solution.
  - E. Interlude – dismantle microscope and show parts.
  - F. Re-equilibrate column in crystallization buffer (2-3 passes).
  - G. Pass derivatized protein solution thru desalting column.
  - H. Add protein back to stock solution.
  - I. Bring volume up to 1.0 mLs.
  - J. Use microscope to verify fluorescence.
    1. Solution should be slightly colored at most.
    2. Should see more color in the desalting column – the unbound probe.
    3. From experience – this results in ~0.25% of the protein being labeled.
  - K. Use protein as normal in setting up crystallization plates.

When viewing the plates – the crystals may show some slight color to those with more discerning eyes (not mine). Crystals are obvious as they give a much brighter emission than the solution or amorphous precipitate. For leads, look for precipitate that has 'bright spots', that cannot be obviously determined to be crystals.