

Protein Purification with DEAE-Cellulose Anion-exchange Chromatography

Introduction

How to purify C-LytAm7 tagged proteins.

Materials

- 0.05M TRIS Maleic Buffer (pH 7)
 - 1 L
- 1M Maleic Solution
 - 20 mL
- 1.5 M NaCl Buffer
 - 400 mL
- 1.5 M NaCl + 2% Choline Buffer
 - 100 mL
- 1.5 M NaCl Solution
 - 200 mL

Procedure

Preparation for Initial Use

The columns are supplied in a storage buffer of 20% ethanol in water. The counter ion is OH^- . Prior to initial use and after extended storage periods, each column should be conditioned as described below (steps 1-4). Always use HPLC grade reagents and filter and degas buffers. During this operation do not exceed more than 25 % of the maximum recommended flow-rates.

1. Wash with 5 column volumes of dd. Water
 2. Wash with 5 column volumes of 1.5M NaCl
 3. Wash with 5 column volumes of low ionic strength equilibration buffer
 4. Wash with 5 column volumes of high ionic strength limit buffer (1.5 M NaCl buffer)
 5. Wash with 10 column volumes of low ionic strength equilibration buffer.
- The column may now be further equilibrated in the starting buffer at the desired flow rate.

DEAE-Sepharose Column Loading

1. Load 35 mL of your protein solution on the column
 - a. Take 10 mL Fractions and measure protein presence ex. With biorad solution (should be purple colored)
2. Wash with 1.5M NaCl Buffer
 - a. Take 10 mL Fractions and measure protein presence
 - b. Wash step is done when there are no color change (no proteins)
3. Elute your Protein with 1.5M NaCl 2% Choline Buffer
 - a. Take 10 mL Fraction and measure with biorad solution
 - b. 2nd,3rd fraction should be colored if the elution worked