

H-NS Protein purification

pI of H-NS = 5.1 - should require an anion exchange column!

Grow 1L culture to OD650 of 0.600.

Induce with 1 mM IPTG for two hours.

Collect cells and resuspended in buffer A [20 mM Tris-HCl (pH 7.2), 1 mM EDTA, 5 mM β -mercaptoethanol, 10% glycerol], containing 100 mM ~~NaCl~~ ^{NaCl} and 1 mg/ml PMSF.

Sonicate and clear lysate by centrifugation at ~~37000~~ ^{13,000} r.p.m. for ~~25~~ ¹ h at 4°C.

The supernatant was directly loaded onto a ~~PII~~ ^{Heparin} column pre-equilibrated with the same buffer.

A linear ~~NH₄Cl~~ ^{NaCl} gradient up to 1 M was applied and H-NS eluted around ~~800~~ ⁵⁰⁰ mM ~~NH₄Cl~~ ^{NaCl}.

The buffer with NH₄Cl was replaced with buffer A, containing 100 mM NaCl by dialysis-overnight.

Dilute fractions to 50ml with 20mM Tris 1mM EDTA

Subsequently the dialysate was loaded onto a pre-equilibrated Heparin column.

*sample was loaded on QFF column**

H-NS eluted around 550 mM NaCl. H-NS containing fractions were dialysed against a buffer solution containing 20 mM Tris-HCl (pH 7.2), 300 mM KCl and 10% glycerol.

All buffers contained 100 μ g/ml PMSF (Sigma) and 2 mM benzamidine (Sigma) to prevent proteolytic cleavage of the H-NS protein. The purity of the H-NS protein was verified on an SDS-PAGE gel and the protein concentration was determined with the Bio-Rad protein assay (Bio-Rad).

**(Doesn't work well. SFF column better)*

Gel ①

- 1 Marker
- 2 F2
- 3 F33
- 4 F 34
- 5 F 35
- 6 F 36
- 7 F 37
- 8 F 38
- 9 F 39
- 10 F 40
- 11 F 41
- 12 F 42
- 13 F 43
- 14 F 44
- 15 F 45

H-NS ~ 17ml

Gel ②

- 1 Marker
- 2 F20
- 3 F22
- 4 F24
- 5 F25
- 6 F26
- 7 F30
- 8 F31
- 9 F32
- 10 F33
- 11 F34
- 12 F35
- 13 F36
- 14 F37
- 15 F38

H-NS but dirty

=> Try again using SFF column. Works much better