-SDS-PAGE -Bradford,

- Quantitative +
qualitative assessment
of purity | concentration
- Total protein

Sodium Dodeay l Sulfate -Poly acrylamide Gel Electrophoresis

SDS-PAGE -> Separate proteins according to size ONLY (Shape, charge, 4° has no effect)

SDS Detergent CH3 (CH2) CH2-80-5-0- Not hydrophobic hydrophilic fail Turpose adenature structure of mass ratio of proteins is made constant

charge to mass smay elongated profein big globular protein

Denature protein
N
(big protein)
N SDS
(small
15DS molecule / 2 amino acids.
- same shape
- same charge: mass
- Se parate based on Size

2X sample buffer

1. SDS

2. \(\beta - Mercapto ethano\)
(BME)

reducing agent

- Break disulfide
bonds (ALL)

3. Glycerol

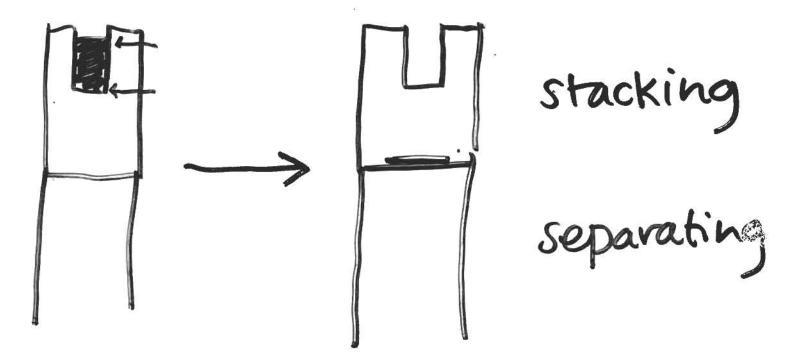
- increases density
of sample so that it
doesn't float away

4. Bromophenol Blue Dye (Blue) 5. Tris (BUFFER)

heat @ 95°C Anatomy of the SDS-PAFE Gel

STACKING 6.8 4-5% acryl. SEPARATING PH 8.8 7.5% (5-18%) (RESOLVING **FUNNING** Tris 6 (yaine SDS

Stacking gel



Allows all profeins in sample to start @ the same starting point.

Separating gel _1 / polyacry lamide than stacking gel. - PH 8.8. > Allows proteins to separate according to Size

Biggest proteins migrate ShowEst Smallest proteins migrate fastest

Staining + Destaining Staining solution - Coomassie Blue (binds nouspecifically to profeins - Acetic Acid -Methanol

AfterStaining Destain - Acctic acció - Methanol

specific activity 4/8 PURER 4/5 MPG/2 SPECIFIC BRADFOR Stecific Total Activity x100/ Activity Total amount of protein

2. Bradford assays

- Measures TOTAL protein in your sample (quantitatively)

-> Allows measurement
of purity(specific
activity)

- Commassie Blue Dye that binds to proteins non-specifically Bradford reagent
- comes red/brown + protein -> BLUE -> Use specto quantitate A595, 0.1-> 0.6

BSA Standard Curve

Asas 0.25 0.1: Thear [BSA] mg/mL range