

Protein Purification 1: Lysis and Affinity Purification

Day 3: Wednesday 21nd March

Today's talk

- Bacterial protein expression
 - Harvesting cultures
 - Lysing your cells and clearing the lysate
- Mammalian protein expression
 - Harvesting cells/supernatant
 - Lysis and clearing the lysate/supernatant
- Affinity chromatography
 - Immobilised Metal Affinity Chromatography (IMAC)
 - Glutathione S-transferase purification
 - Biotin/Streptavidin and Strep-II/Strep-Tactin
 - Other options
- Engineered proteolysis
 - 3C (PreScission), TEV, Thrombin, SUMO



Harvesting your bacterial culture

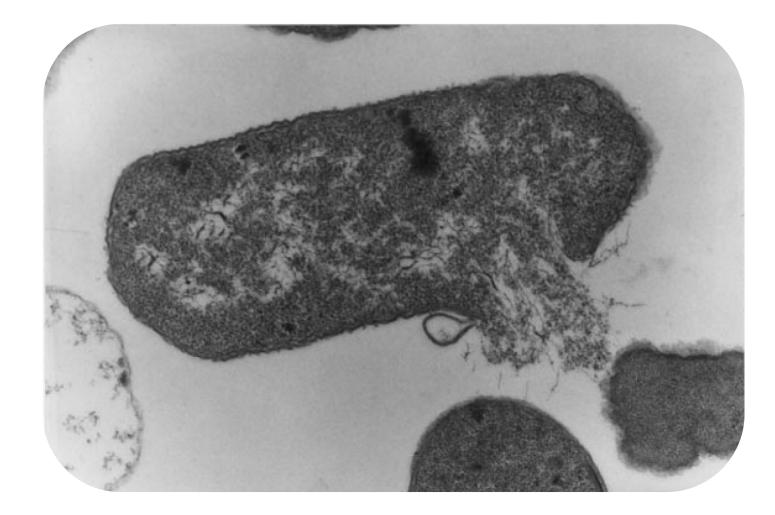
- For cytoplasmic proteins, harvest by centrifugation
 - 5000 × g for 15 min
- Decant off the supernatant
- Scrape pellets into a beaker for immediate processing, or a tube for storage
- Pellets can be stored for months at -20°C for a few months, or at -70°C for years and years and years...





Bacterial cell lysis

- Mechanical
 - Cell disruptor
 - Sonicator
 - Freeze-thaw
- Chemical
 - Detergent-based

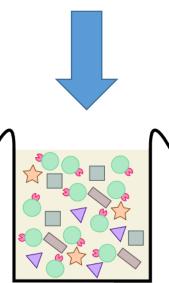




Preparing to mechanically lyse cells

- Cells need to be resuspended in a lysis buffer
 - Important that you control the pH and [salt]
 - Often will include reducing agents and protease inhibitors*
 - *EDTA-free for His-tagged proteins
 - I often include a tiny bit of detergent (0.05% or less)
 - Some proteins like glycerol to be present
 - Lysozyme can also help (especially for non-pLysS cells)
- Lysis buffer should generally be kept cold
 - Bacteria have proteases even expression strains that lack OmpT and Lon
 - Proteases: very active at 37°C, poorly active at 4°C
 - Keep your lysate on ice until the affinity capture and wash to minimise proteolysis of your protein







Preparing to mechanically lyse cells

- Important that cells are **thoroughly** resuspended for efficient lysis
 - No lumps!
- I resuspend cells on a stirring plate in the cold room using a magnetic flea (takes about 15-30 minutes)
 - Can be faster if you pipette up and down
- I lyse up to ~10 g of cell pellet per 50 mL of lysis buffer
 - Main aim is to have a smooth solution with no lumps!





Mechanical lysis: Cell disruptor

- Lyses cells via a combination of pressure, shear force and impact
- Sample is passed through an 100 μm diamond aperture at >20,000 psi
 - Significant shear force to disrupt sample, plus pressure change from high to low pressure
- Jet of liquid hits target and is disrupted by kinetic impact
- Consistent >90% lysis
- Some sample heating
 - Water-cooled jacket to cool samples





Mechanical lysis: French Press or Pressure Homogeniser

- French press:
 - Similar to cell disruptor, large piston generates pressure to force cell suspension through narrow orifice
- Pressure Homogeniser (Emulsiflex)
 - Uses pressurised gas to force sample through a narrow orifice
- Both generate shear force to lyse cells
 - Very efficient cell lysis with some sample heating







Top Image: Glen Mills (https://www.glenmills.com/product-category/cell-disruption-and-culture/cell-disruption/french-press/)

Bottom Image: Avestin (https://www.avestin.com/emulsiflex-c5.htm)

Mechanical lysis: Sonication

- Use ultrasonic waves to disrupt cell membrane
 - Also very efficient at shearing DNA
- Causes localised sample heating
 - Need to sonicate as short bursts, preferably with sample on ice
- Ten cycles of 30s (with 30s rest) at 8 μm amplitude gives very efficient cell lysis
 - You might need to optimise this
- Sonicators can permanently damage your hearing You must wear ear protection!!





Chemical and other methods of cell lysis

- Freeze-thawing bacterial cells can cause significant cell lysis
- Hen egg white lysozyme can degrade the peptidoglycan of the bacterial cell wall, promoting lysis
- Detergents can be used to lyse bacterial cells
 - 1% TWEEN-20
- Combining all three can give reasonably efficient lysis (especially for small-scale tests)
 - But detergents and multiple freeze/thaws can also disrupt protein folding, lowering yield
- Proprietary cell lysis cocktails exist
 - Expensive for large/frequent protein preps



Clearing cell lysates

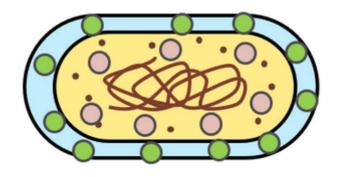
- Remove chromosomal DNA, membranes and insoluble proteins
- Centrifuge at >20,000×g for ≥30 min
 - I routinely use 40,000×g for 30 min
- Pellet should be relatively small (<10% of total volume)
 - Large pellets are caused by inefficient lysis or presence of **inclusion bodies**
- Keep samples cold
- Reserve sample of cleared cell lysate for SDS-PAG

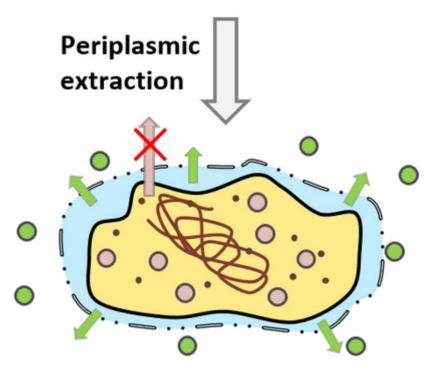




Periplasmic bacterial expression

- Aim to liberate protein in periplasm of bacteria
 - Lyse outer cell wall but not inner cell wall
- Incubate with a hypertonic buffer (500 mM sucrose) to swell cells
- Transfer to hypotonic buffer (125 mM sucrose) to lyse periplasm
- Centrifuge to precipitate cell pellet
- Filter supernatant containing periplasmic proteins







Mammalian cell lysis (cytoplasmic proteins)

- Mammalian cells are much larger and more fragile than bacterial cells
 - Harvest at 200×g for 5 min to prevent premature lysis
- Very easy to lyse
 - Passage through a 23G needle six times
 - Osmotic shock in hypotonic buffer (10 mM Tris pH 7.5)
 - Incubation with detergent (1% Triton X-100)
 - Dounce homogeniser
- Clarify lysate by centrifugation (40,000×g, 30 min)





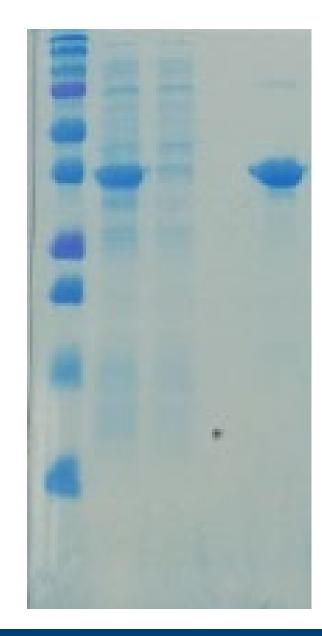
Preparing mammalian cell supernatants

- Spin out cells (200×g for 10 min)
- Clarify lysate (40,000×g for 30 min)
- 0.2 μm filter if storing for a long time
 - Some secreted mammalian proteins like antibodies can be stored for years and years at 4°C...



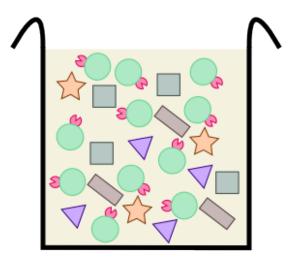


- Selective resin (beads) that can capture your protein
- Separates your protein from the vast majority of other cellular proteins
 - Including proteases
- Concentrates your protein ready for further purification steps, if required (tomorrow's seminar)



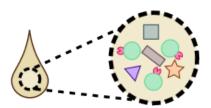


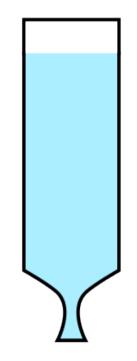
• Start with clarified cell lysate (or supernatant)





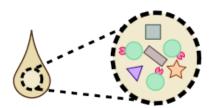
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- Apply to selective resin

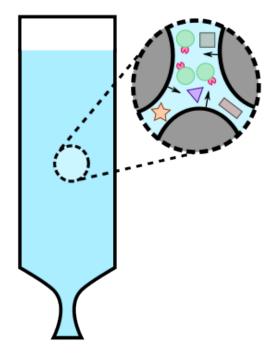






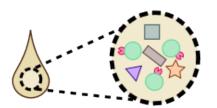
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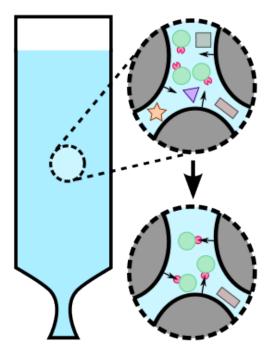






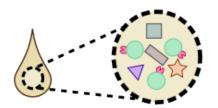
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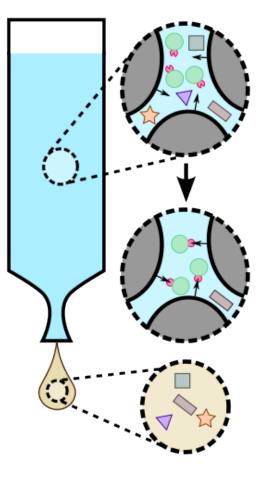






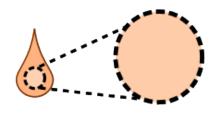
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- Apply to selective resin
- Selective resin will capture tagged protein
- Other proteins will **flow through** column

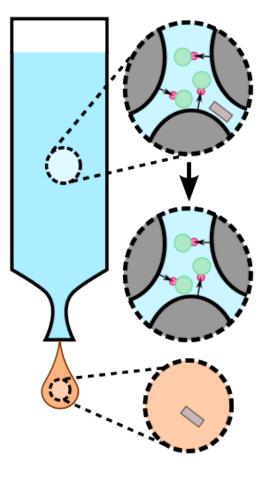






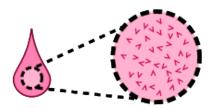
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- Wash column with wash buffer to remove residual unbound protein

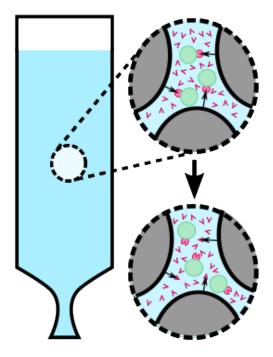






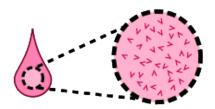
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- Apply elution buffer
 - Contains molecule that competes with tag:column interaction

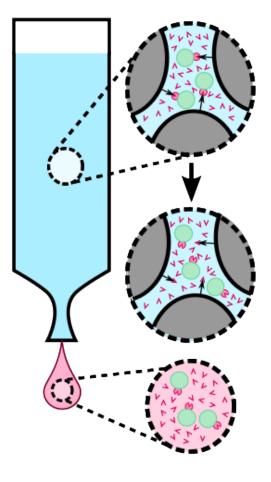






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- Apply elution buffer
 - Contains molecule that competes with tag:column interaction
- Protein is eluted in elution buffer

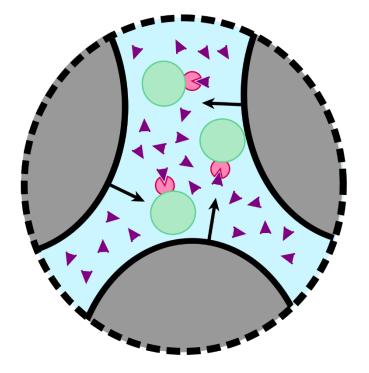






Affinity chromatography elution

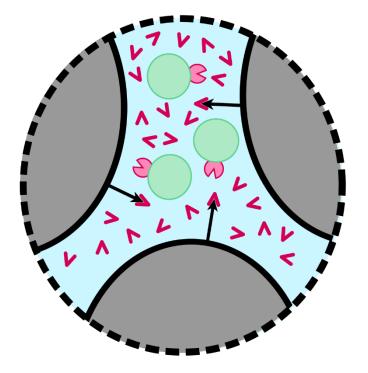
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 - Molecule that mimics the resin: competes with resin for binding to the tag





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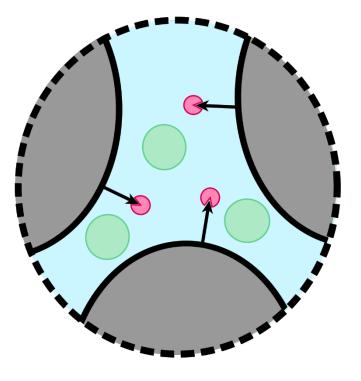
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Affinity chromatography elution

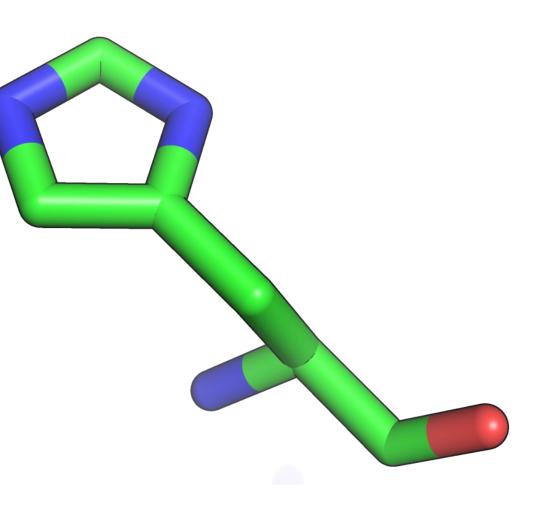
- Elution buffer can be:
 - Molecule that mimics the resin: competes with resin for binding to the tag
 - Molecule that mimics the tag: competes with tag for binding to the resin
- You can also elute protein by using a protease that cleaves between the protein and the tag
 - Perform on column proteolysis using engineered proteases and cleavage sites
 - Protein is eluted
 - You can then then regenerate column using small molecules as above





Immobilised Metal Affinity Chromatography (IMAC)

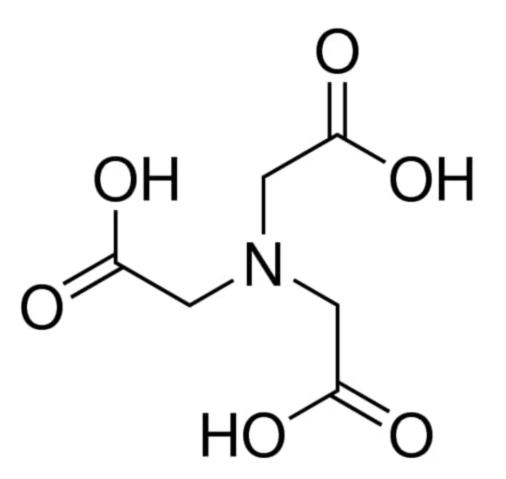
- Exploits the affinity of histidine side chains for metal ions
- Encode a short polyhistidine tag at the N or C terminus of your protein
 - Usually His₆





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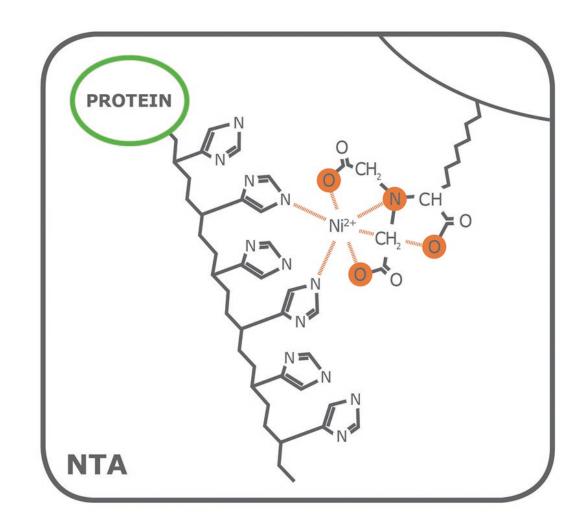
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 - E.g. Nitrilotriacetic acid (NTA)





Immobilised Metal Affinity Chromatography (IMAC)

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- Resin has a metal chelating group
 - E.g. Nitrilotriacetic acid (NTA)
- Most common metal ion is Ni²⁺
 - Co²⁺ is also used





Ni-NTA IMAC

- Protein is eluted using imidazole
 - >200 mM imidazole
 - Competes with His side chain for binding Ni-NTA
- NiNTA columns have (relatively) high non-specific binding
 - Use low concentration (10-20 mM) imidazole in wash buffers
 - Use high salt in wash buffers
- Imidazole is a chaotropic agent
 - Unfolds proteins
 - Remove as soon as possible after elution (dialysis, etc)



Glutathione S-transferase (GST) affinity chromatography

- Glutathione S-transferase (GST) is a small (26 kDa) protein that catalyses conjugation of reduced glutathione (GSH) to other chemicals
 - Detoxification of foreign substances
- GST binds to GSH with high affinity
- GST tags are appended to the N or C termini of proteins
 - Can increase solubility of tagged protein
 - Efficiently captured by GSH conjugated to a resin

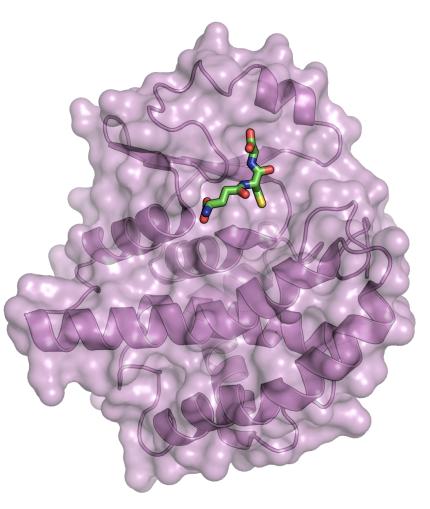
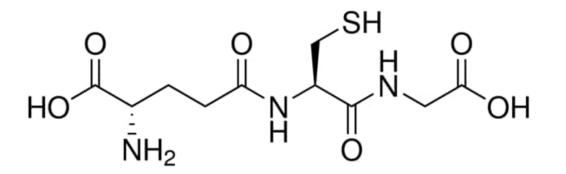




Image: PDB 1UA5 [Kursula et al (2005) Protein Pept Lett 12, 709-12], Stephen Graham (CC BY 4.0)

GST affinity chromatography

- Highly specific binding
 - Low background
- High affinity interaction
 - Allows extensive washing
- Elute with relatively mild conditions
 - 20 mM GSH in normal buffers
- Relatively slow binding kinetics
 - Need longer incubations for binding and elution

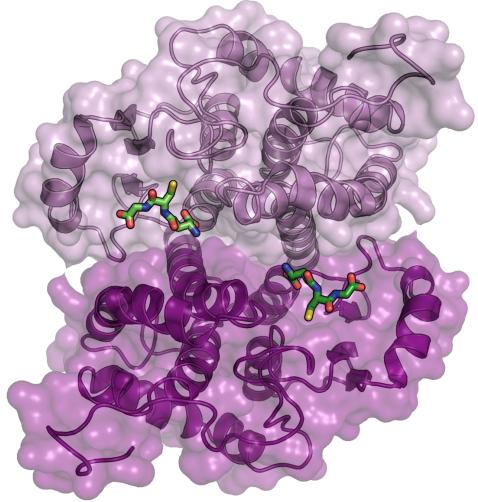


Reduced L-glutathione (GSH) Note the isopeptide bond



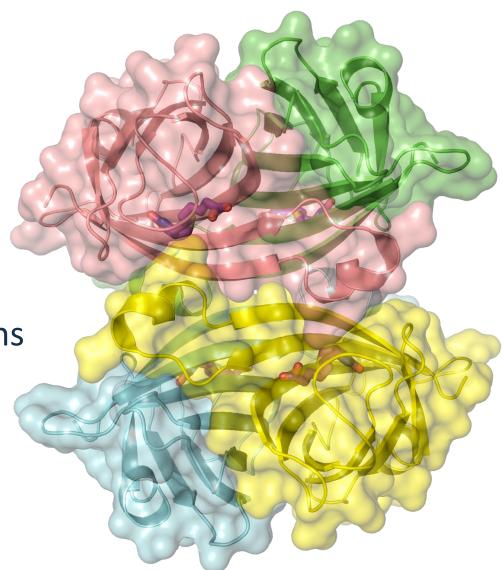
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- Large tag that dimerises
 - Often removed for downstream applications



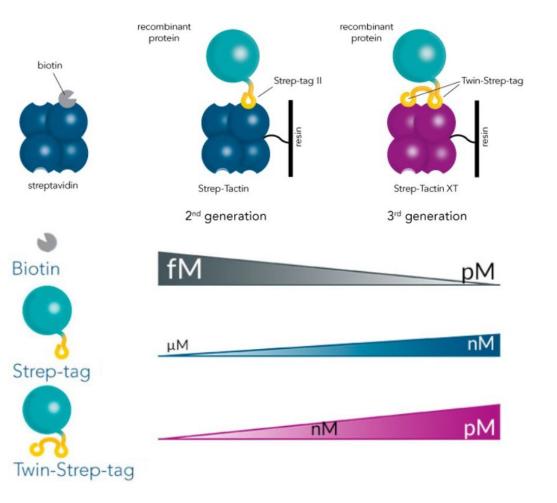
Streptavidin and biotin

- Biotin binding to streptavidin is one of tightest known non-covalent interactions
- Proteins can be covalently biotinylated if conjugated to a biotin acceptor peptide (BAP) sequence
- Can use streptavidin resin to purify proteins
 - Highly specific to biotin
- Some endogenously biotinylated proteins in *E. coli* will co-purify
- Interaction is so strong that elution is very difficult without denaturation



Strep-II tag and Strep-Tactin

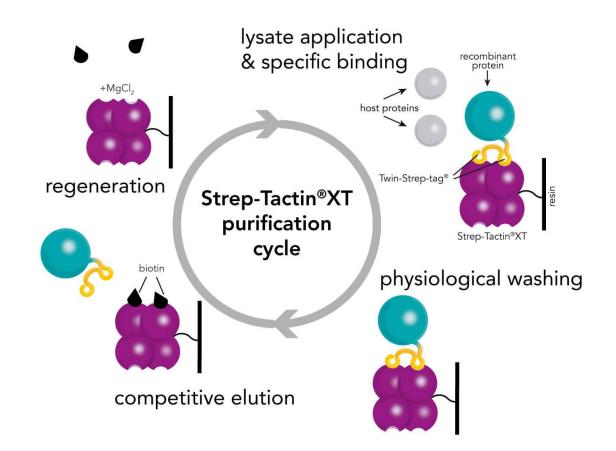
- A short peptide sequence was identified with intrinsic affinity toward streptavidin
 - Strep-II tag: WSHPQFEK
- Streptavidin was engineered to have enhanced affinity for the Strep-II tag and reduced affinity for biotin
 - First generation was Strep-Tactin
 - Latest generation is Strep-Tactin XT
- Can use a twin Strep-II tag for avidity-enhanced binding





Strep-Tactin XT

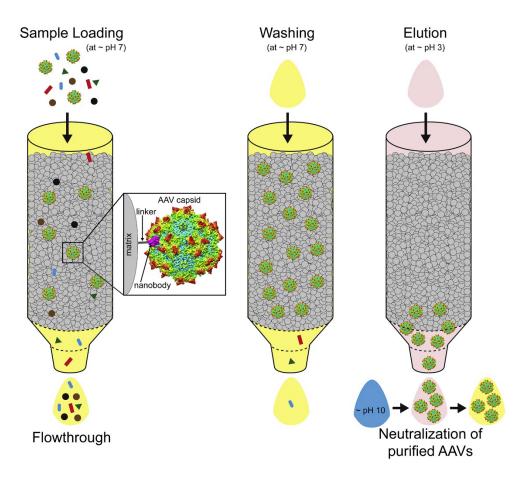
- Combines benefits of a short tag sequence with high specificity and high affinity binding
- Simple protein elution (biotin) and regeneration (MgCl₂)
- High binding capacity (5 mg/mL)
- Resin is quite expensive





Antibody affinity columns

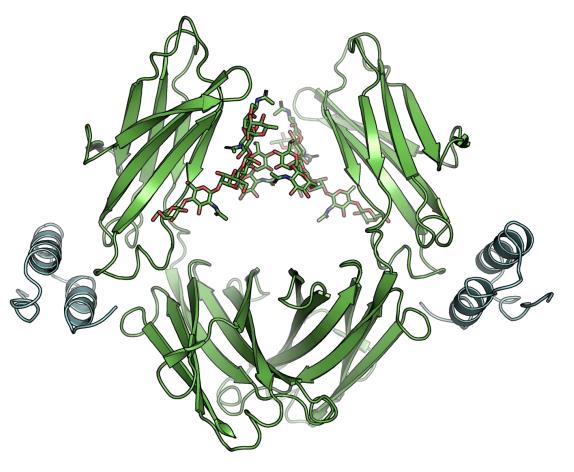
- Antibodies/nanobodies combine high specificity and affinity
- Antibodies or nanobodies covalently coupled to agarose resin can be used as affinity columns
 - Antibodies against your protein of interest
 - Antibodies against an epitope tag appended to your protein (e.g. HA, FLAG)
- Elute using low pH or excess of the engineered epitope (synthetic peptide)
- Quite expensive resin and regeneration





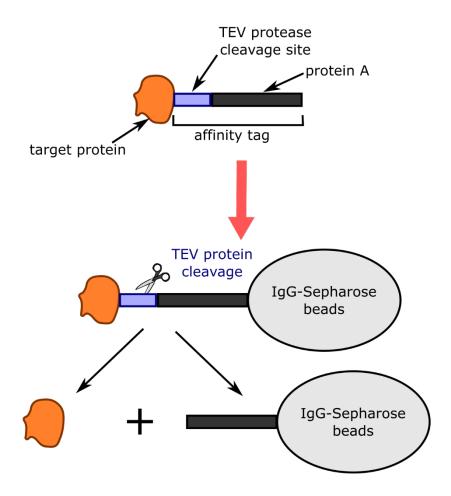
Protein A, Protein G and Protein A/G

- Protein A and Protein G are antibodybinding proteins from *Staphylococcus aureus* and *Streptococcus* bacteria
- Bind the Fc (A) and Fc + Fab (G) domains of mammalian antibodies
 - Different affinities for different isotypes
- Protein A/G is a recombinant fusion of the IgG binding domains of proteins A and G
- Proteins A, G and A/G are excellent for antibody purification



Protein A tags and IgG columns

- Protein A can be used as an affinity tag
 - May enhance solubility of proteins expressed via secretion from mammalian cells
- Can use IgG fused to agarose to capture tagged proteins
- Cleave protein off column using engineered protease
- Retained protein A can be eluted using low pH





Tag/Resin	Pros	Cons
His/NiNTA	Small tag High capacity resin (>10 mg/mL) Resin and imidazole are cheap	Higher non-specific binding Lower affinity (can 'over-wash') Elute with chaotropic chemical (imidazole)



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GST/GSH	Highly specific and high affinity High capacity resin (5-10 mg/mL) Resin and GSH are cheap GST helps solubilise proteins	Large protein tag (need to remove for many downstream applications) Slow binding/dissociation kinetics GST helps solubilise proteins



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Strep-II/Strep-Tactin (XT)	Small tag Highly specific (low background) High capacity resin (5 mg/mL) Easy regeneration of resin	Resin is relatively expensive

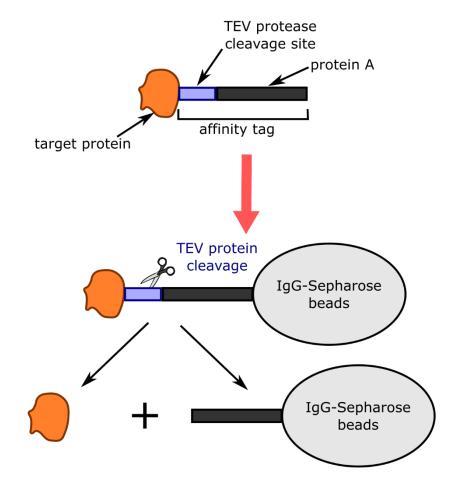


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Biotin/Streptavidin	Super-high affinity	Super-high affinity
Antibody columns	Very specific	Expensive and lower capacity
Protein A and IgG columns	High affinity specific binding Protein A may solubilise proteins	Need to cleave protein off resin Slightly trickier resin storage and regeneration



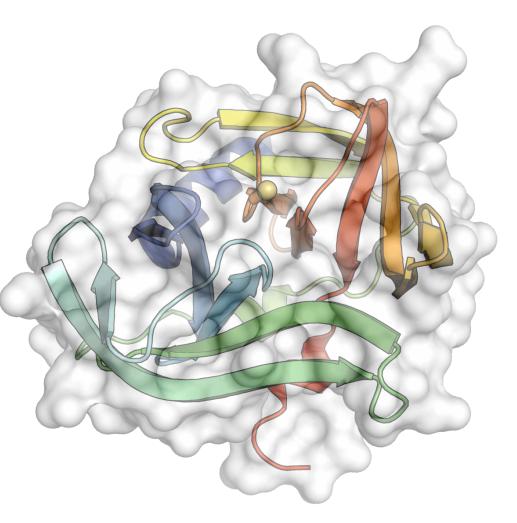
Engineered proteolysis

- Non-specific proteolysis is A Bad Thing
- But proteolytic removal of tags can be helpful:
 - Elute protein from affinity resin without using harsh elution buffers (e.g. low pH)
 - Remove large purification tags that cause oligomerisation or could interfere with function (e.g. GST)
- Can design proteolysis sites into expression constructs and use specific purified proteases to mediate cleavage



3C (PreScission) protease

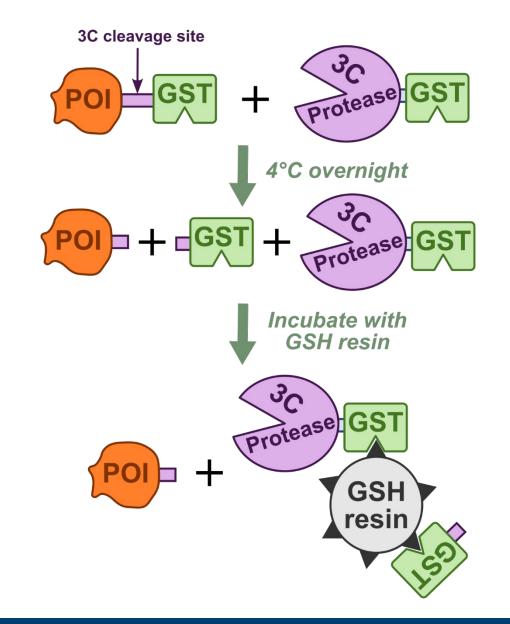
- Cysteine protease
 - Active under reducing conditions, inhibited by divalent cations (e.g. Zn²⁺)
- From human rhinovirus 14
 - Cleaves viral polyprotein
- Recognition sequence: LEVLFQ/GP
- Sold by Cytiva as "PreScission protease"
- Easy to express in *E. coli* and purify





3C (PreScission) protease

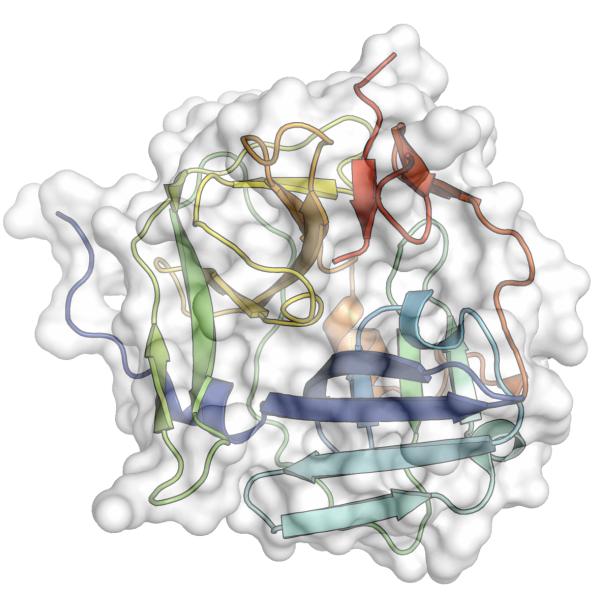
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 - We use GST fusion
 - Allows capture of protease and liberated GST after cleavage





TEV protease

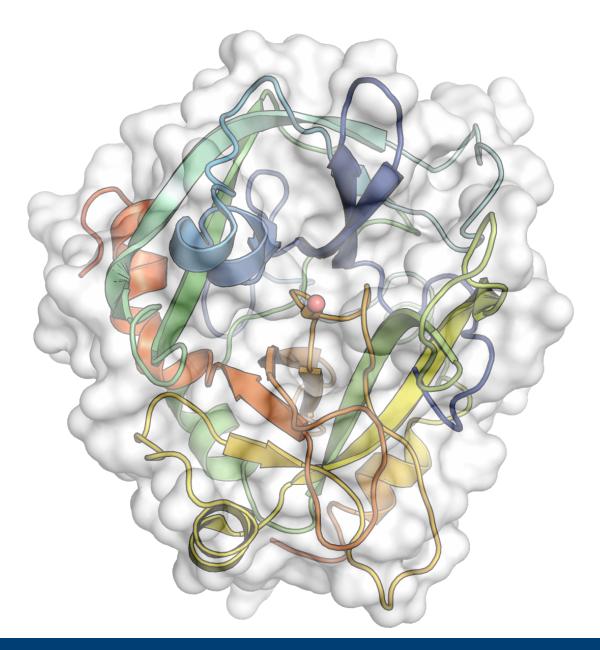
- Cysteine protease
- From tobacco etch virus
 - Cleaves polyprotein
- Recognition sequence: ExxYxQS/G
 - x is any amino acid
- Wild-type TEV:
 - Undergoes autoproteolysis that reduces activity
 - Is poorly expressed in *E. coli* and poorly soluble
- Engineered mutants of TEV exist to overcome these issues
 - And are tagged for easy removal





Thrombin

- Serine protease
 - Inhibited by PMSF or AEBSF
- Acts as a blood coagulation factor
 - Cleaves fibrinogen to fibrin
 - Expressed as a zymogen (pro-enzyme) and activated via cleavage by factor Xa
- Recognition sequence: LVPR/GS
- Abundant in blood of mammals
 - Bovine thrombin can be purchased relatively cheaply
 - Purity varies between vendors...





SUMO protease

- SUMO is a small ubiquitin-like protein
 - Covalently attached to eukaryotic proteins
 - Alters their subcellular localisation and activity
- SUMOylation can be rapidly reversed by deSUMOylases *in vivo* and *in vitro*
 - Recognise folded SUMO protein and cleave C-terminal sequence xGG/x
- N-terminal SUMO tags can enhance protein solubility
- Removed without a 'scar' using SUMO protease
 - Highly active SUMO protease from thermophilic bacteria *Chaetomium thermophilum*

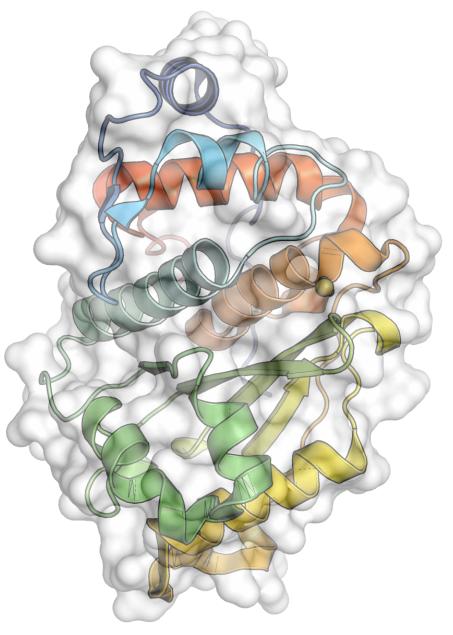


Image: PDB 6DG4 [Lau et al. (2018) *J Biol Chem* 293, 13224-33], Stephen Graham (CC BY 4.0)

Today's talk

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- Mammalian protein expression
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Tomorrow:

Beyond affinity capture: additional techniques to further purify your protein

