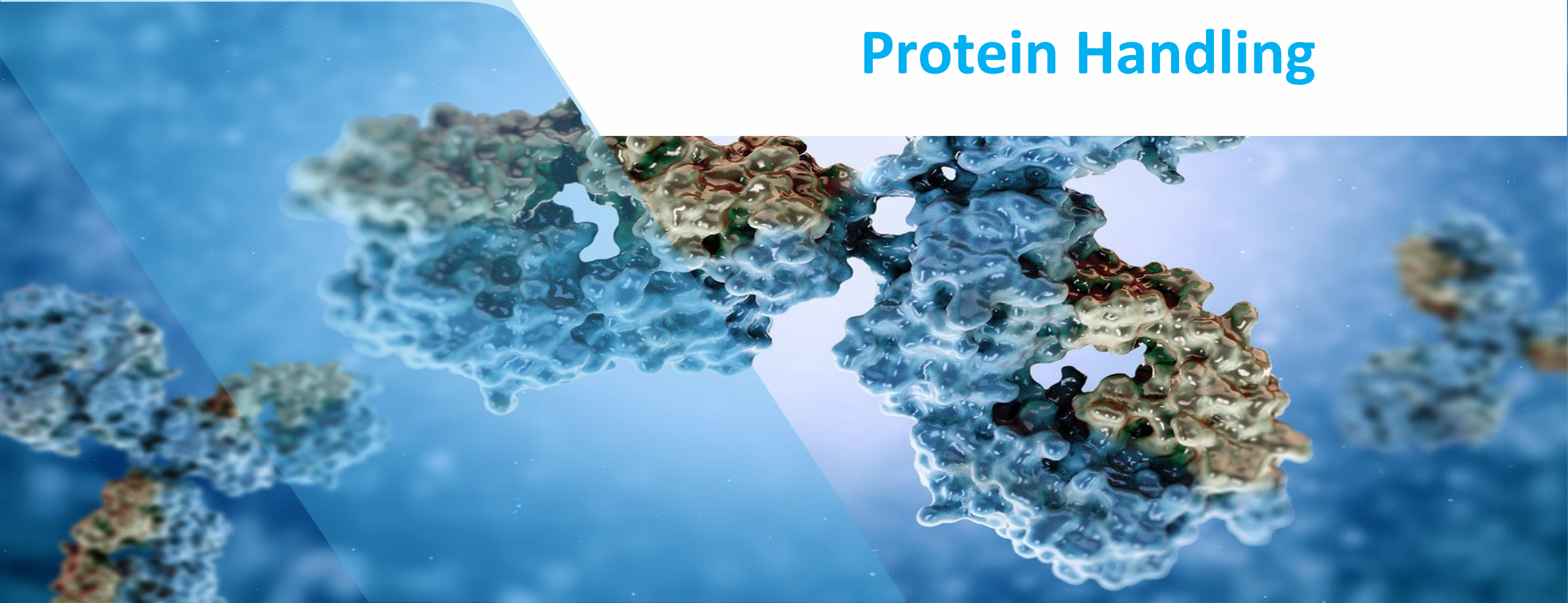


# Protein Handling





# Learning Objectives

1. Describe the chemical structure of proteins

2. Describe the relationship between protein structure, stability and function

3. Define how manufacturing/processing conditions significantly impact biopharmaceutical stability

4. Explain why protein stability is critical to the production of biopharmaceuticals

# Topics



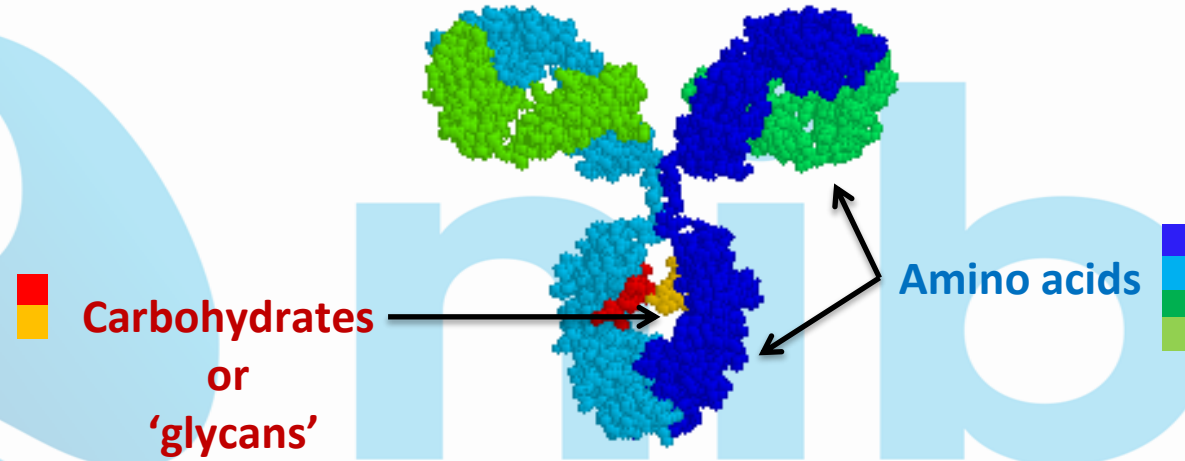
Introduction to Protein Chemistry

Protein Structure

Post Translational Modifications

Protein Stability

# What are Proteins?

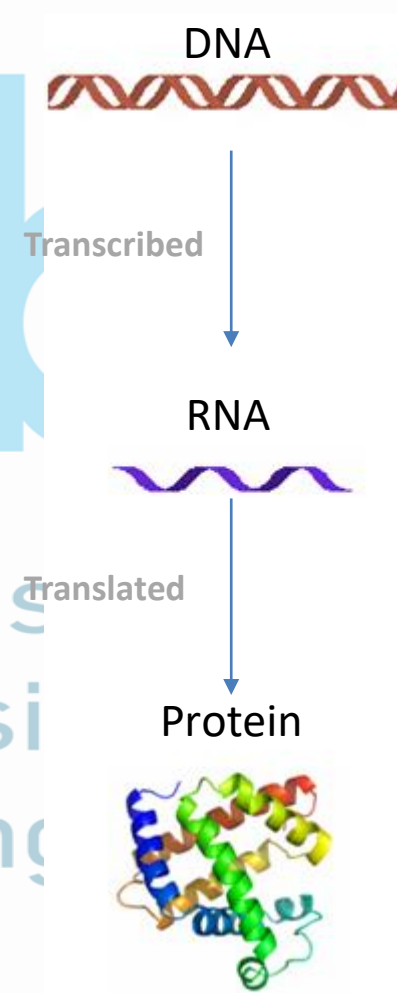


- Proteins are large, 3D, molecules that have many functions inside the body
- Proteins are primarily composed of molecular building blocks called '**amino acids**'
- Many biopharmaceuticals also contain carbohydrates (sugars) called '**glycans**'

# How are Proteins Made in the Body?

- The 'recipe' for a protein is encoded as DNA (master copy of the recipe)
- The DNA is **transcribed** into RNA by the cell (working template)
- RNA can be understood by cellular machinery and **translated** into the language of amino acids
- The cell then assembles a protein molecule from the code

<https://www.youtube.com/watch?v=gG7uCskUOrA>





# Topics



Introduction to Protein Chemistry

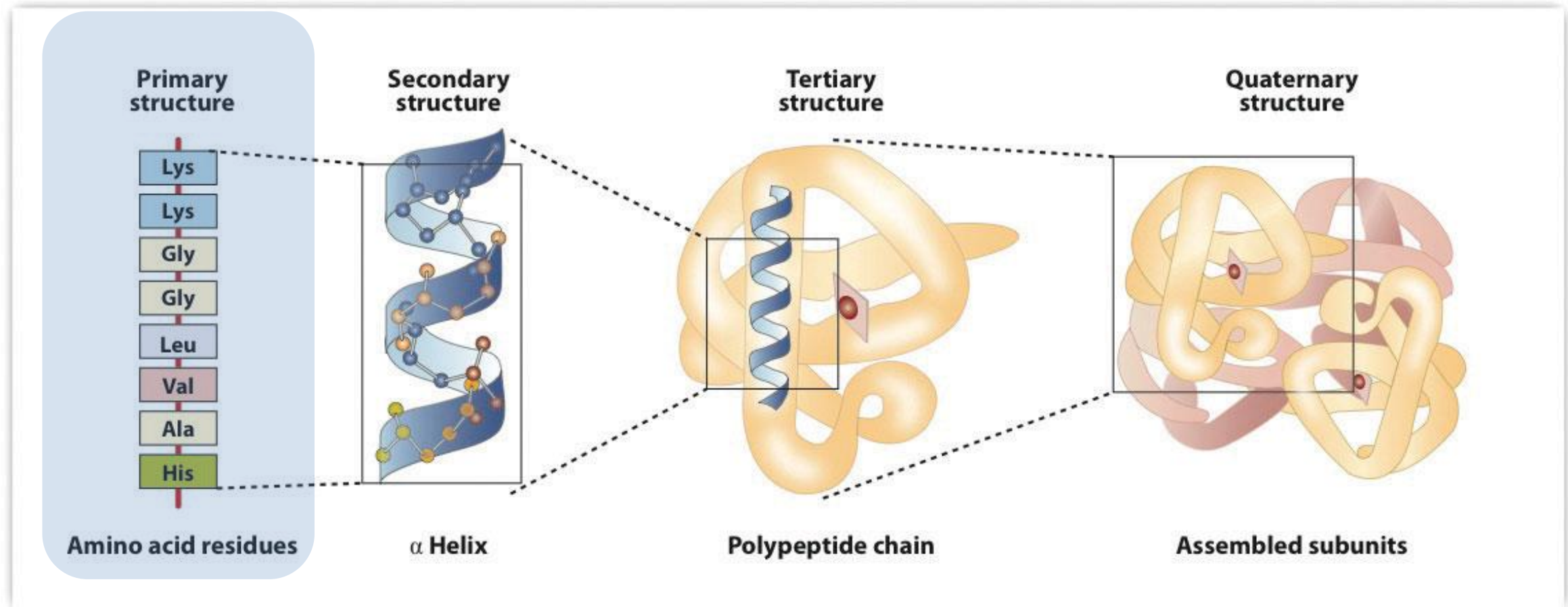
Protein Structure

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Protein Stability

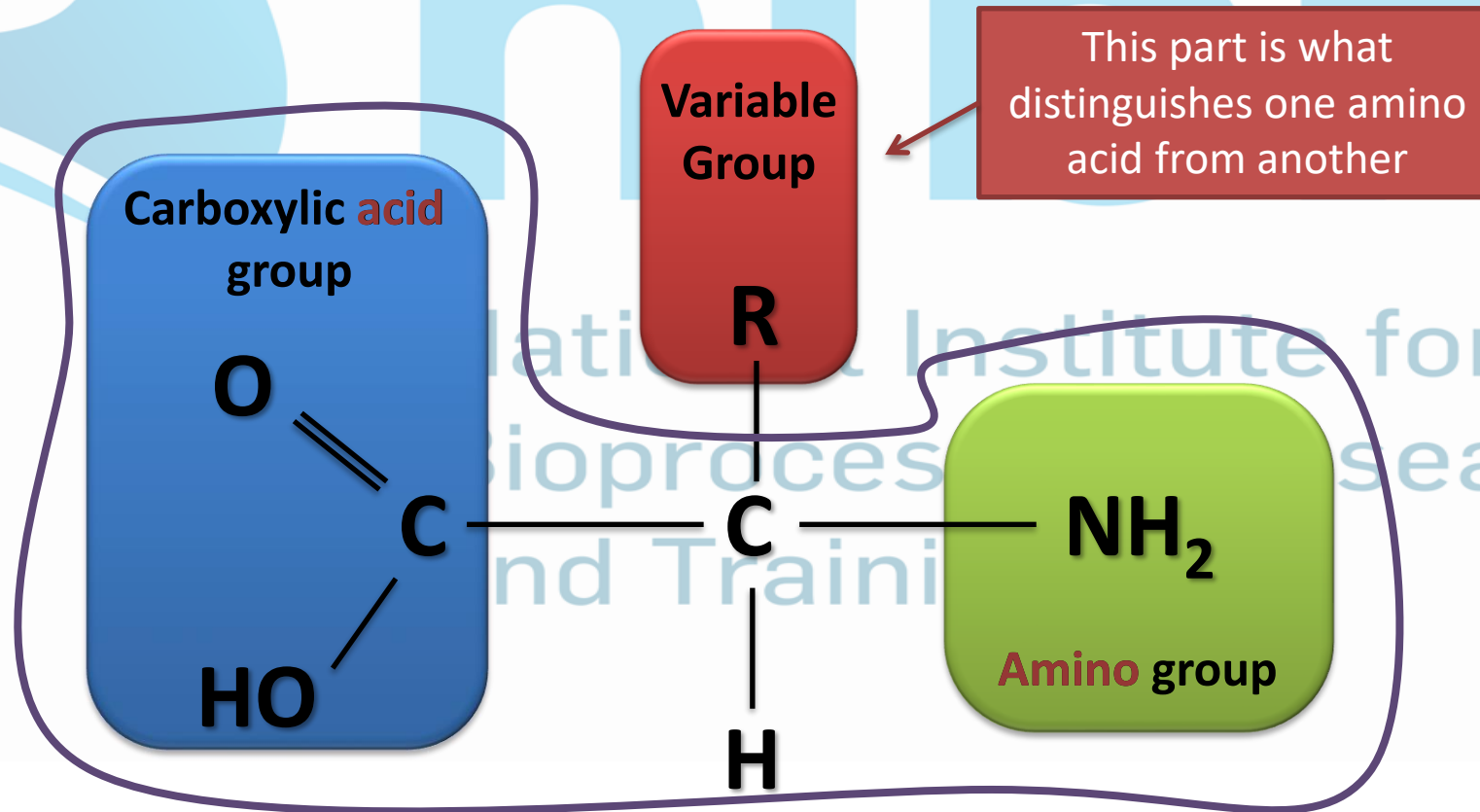
# Protein Structure: Four Levels of Organisation

Proteins have complex 3D structures. This structure is very important – it allows the protein to do its job.



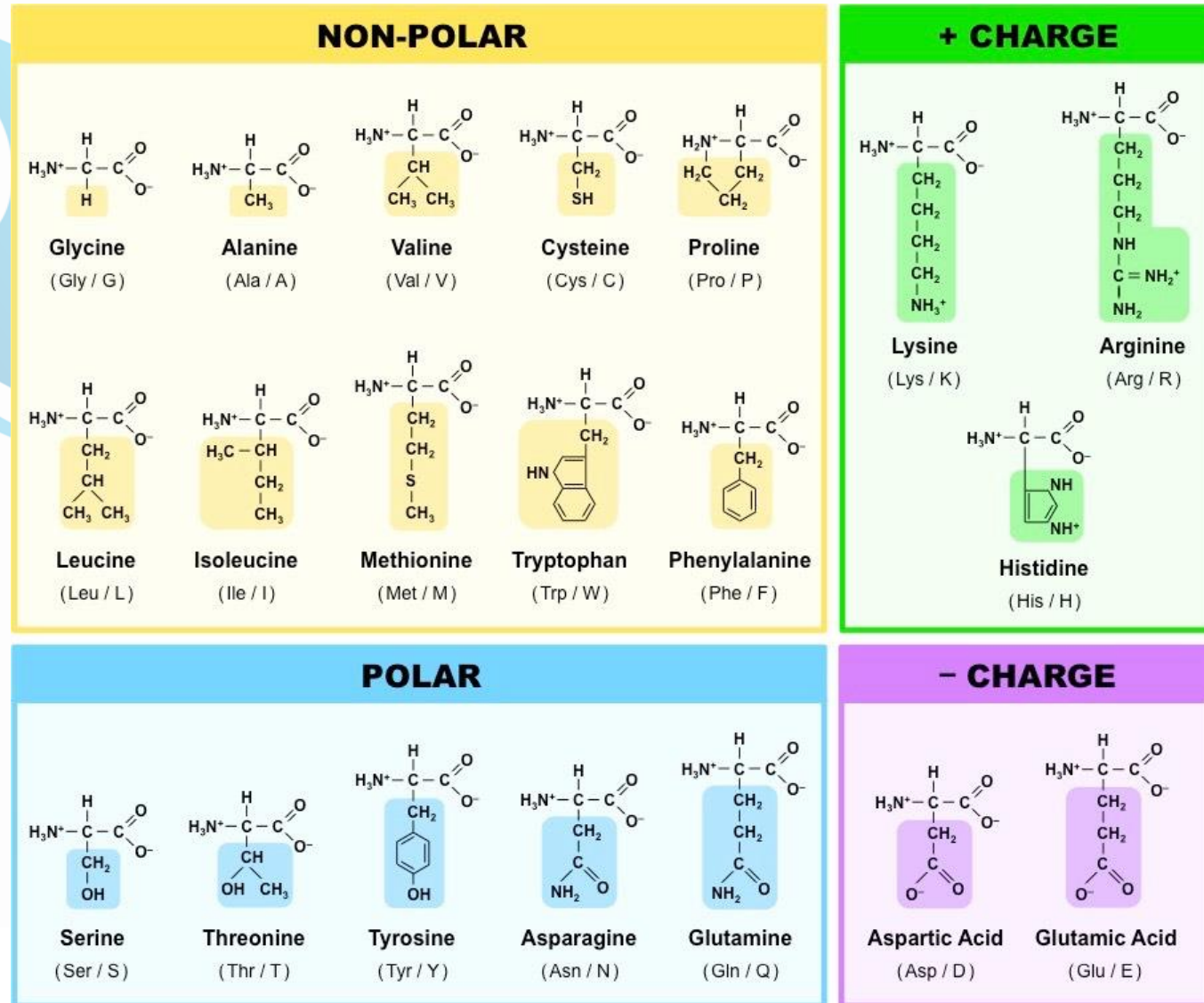
# Primary Structure

- Proteins consist of chains of **amino acids**
- Amino acids are molecules consisting of a carbon with a primary amine, a carboxylic acid, a hydrogen & a **variable side chain group** (R)



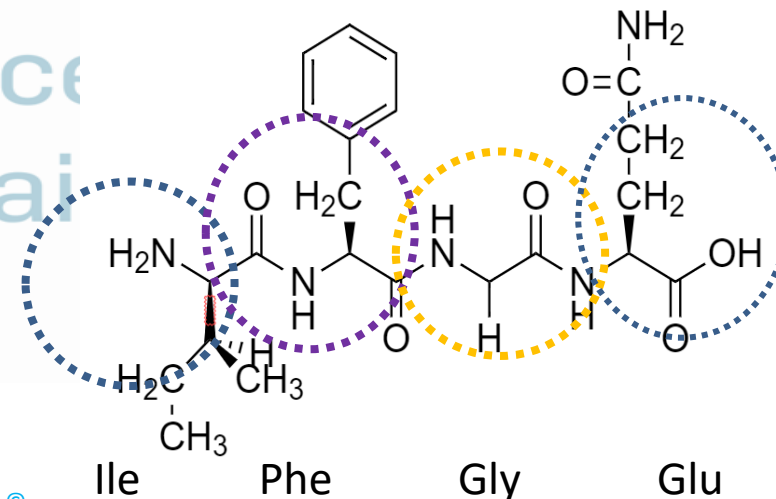
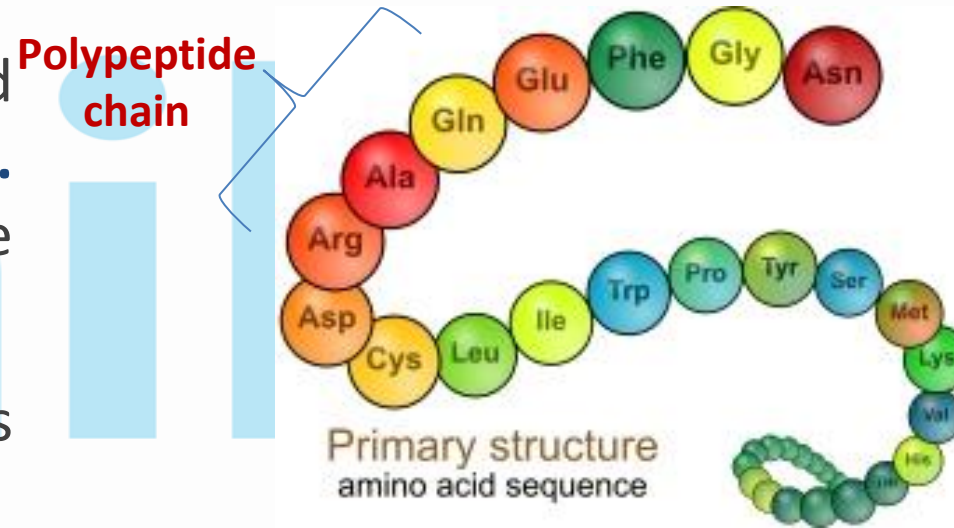


# Primary Structure



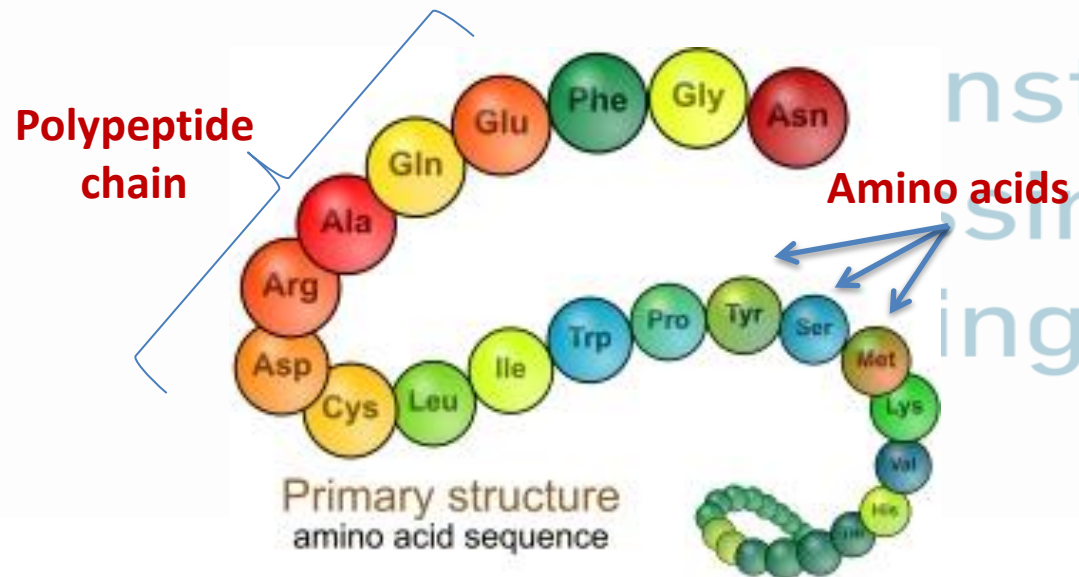
# Primary Structure

- To make a protein, amino acids are linked together in a chain to form a **polypeptide**. This process is carried out by an enzyme complex in the cell
- Each type of protein will have amino acids linked together in a unique order
- The linkage between each amino acid is called a covalent bond called a '**peptide bond**'
- This is an **extremely strong** inter-atomic linkage – not easily damaged during processing



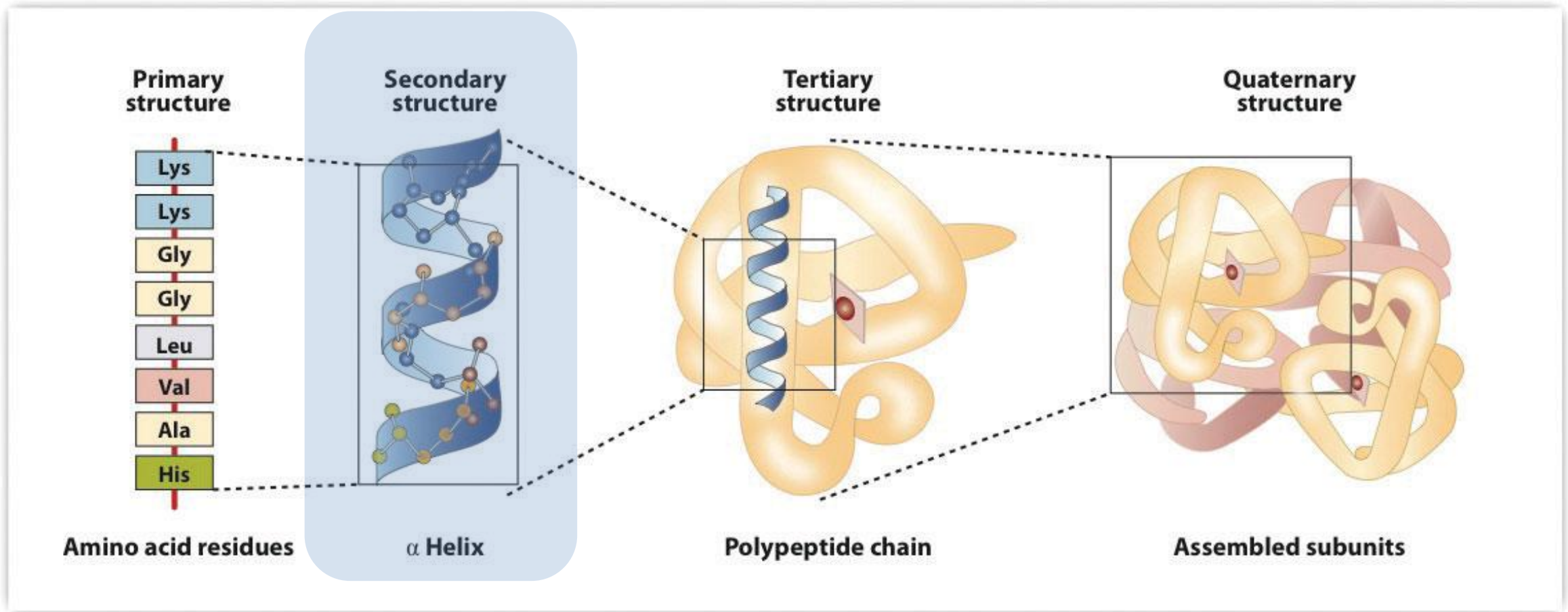
# Primary Structure

- The **sequence** of amino acids in the polypeptide is determined by the **genetic code**
- **Mutations** in the gene will lead to **incorrect amino acid** sequence and ultimately **incorrect protein structure**

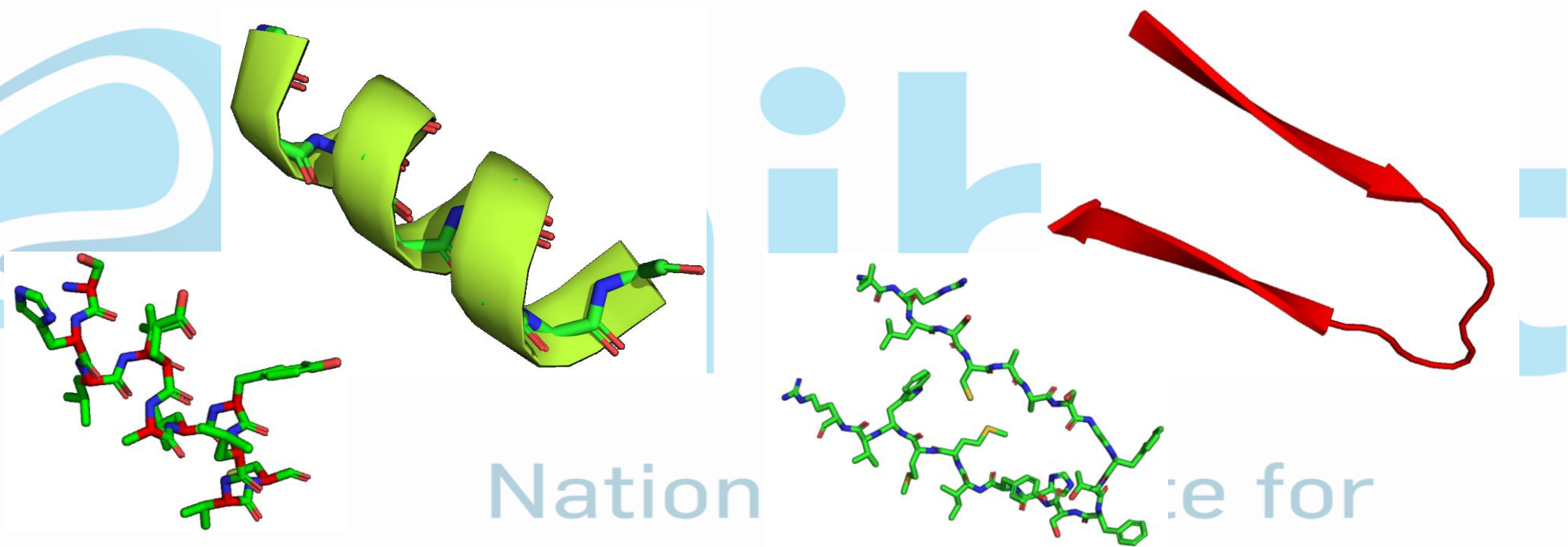


Amino acids can also be chemically changed during processing, formulation, filling storage and sample handling!

# Protein Structure: Four Levels of Organisation



# Secondary Structure

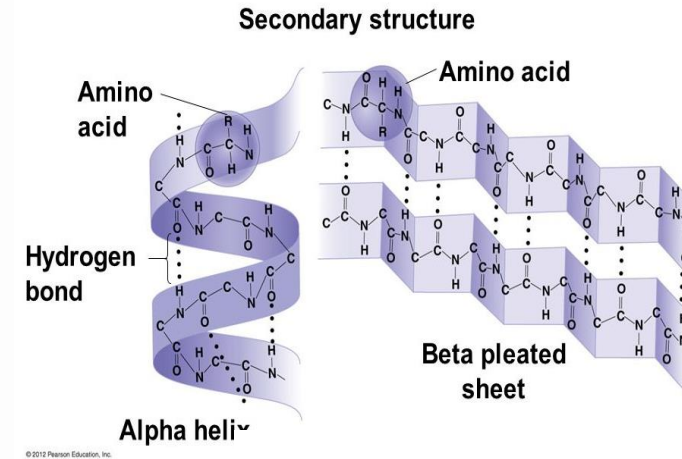


- Amino acids which are **close together** in the sequence (primary structure) start to interact and form secondary structures.
- Secondary structure consists of **repeating patterns**– the peptide winds up into  **$\alpha$ -helicies** or folds into  **$\beta$ -sheets**

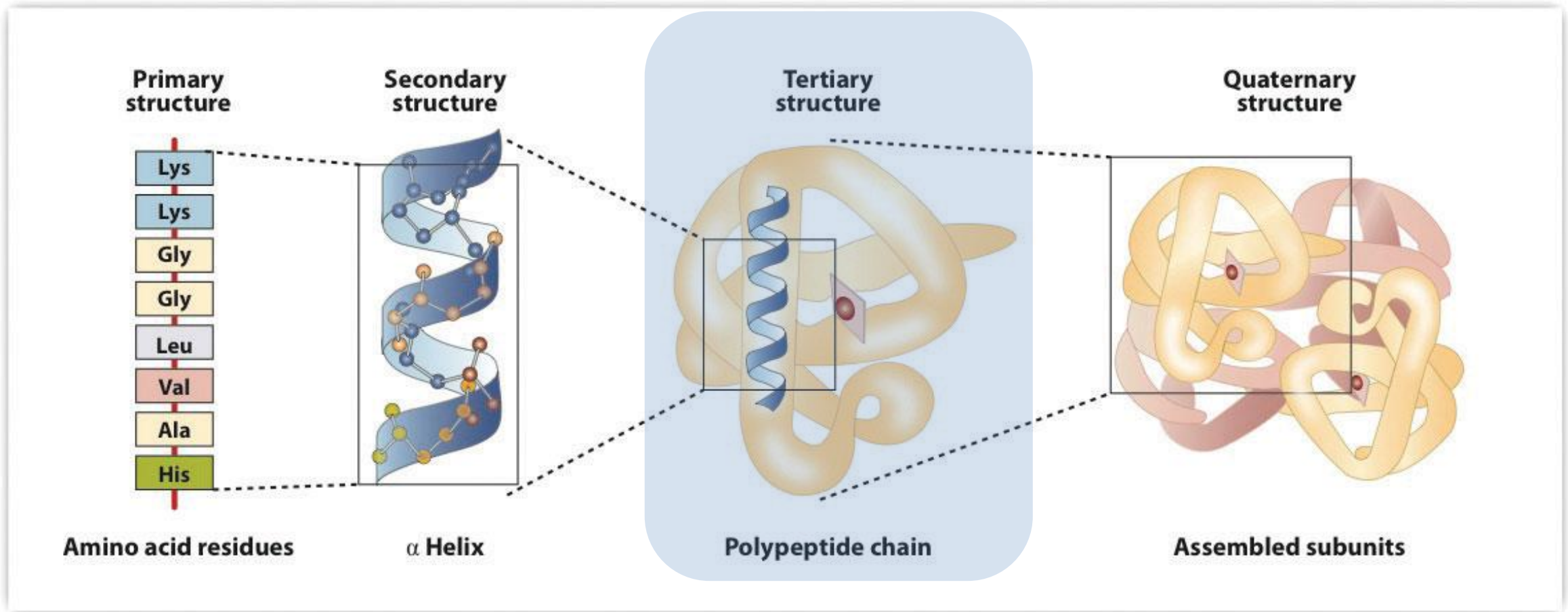


# Secondary Structure

- Secondary structures are held together by hydrogen bonding: **negative** end of one molecule becomes **weakly attached to the hydrogen** of another
- these interactions are called **hydrogen bonds**
- Individual hydrogen bonds are **very weak**, it takes lots of them working together to hold the secondary structure in place
- Hydrogen bonds **can easily be broken** by **incorrect protein handling** during processing

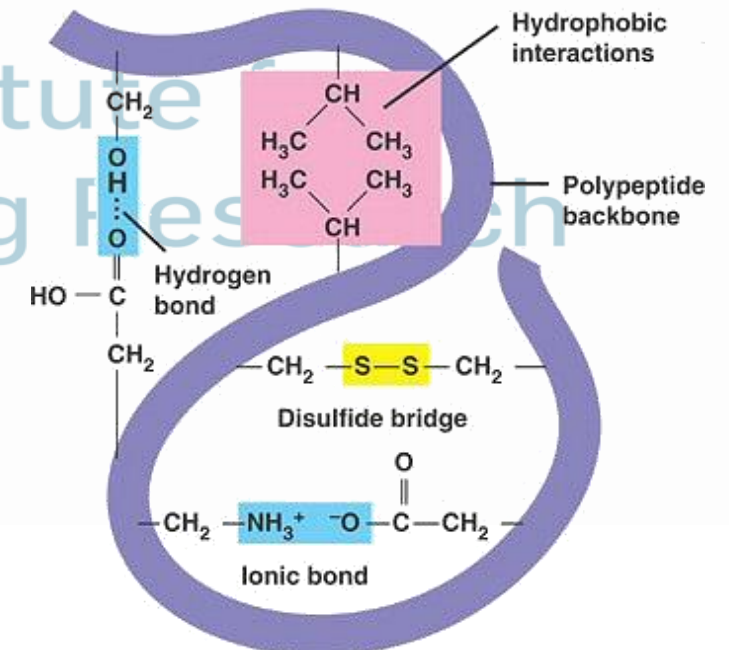
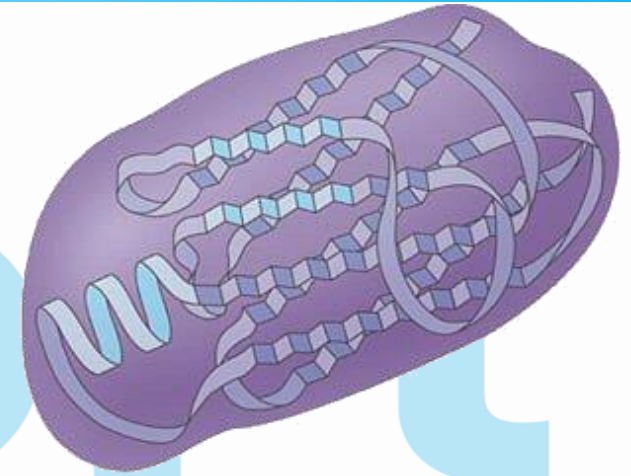


# Protein Structure: Four Levels of Organisation

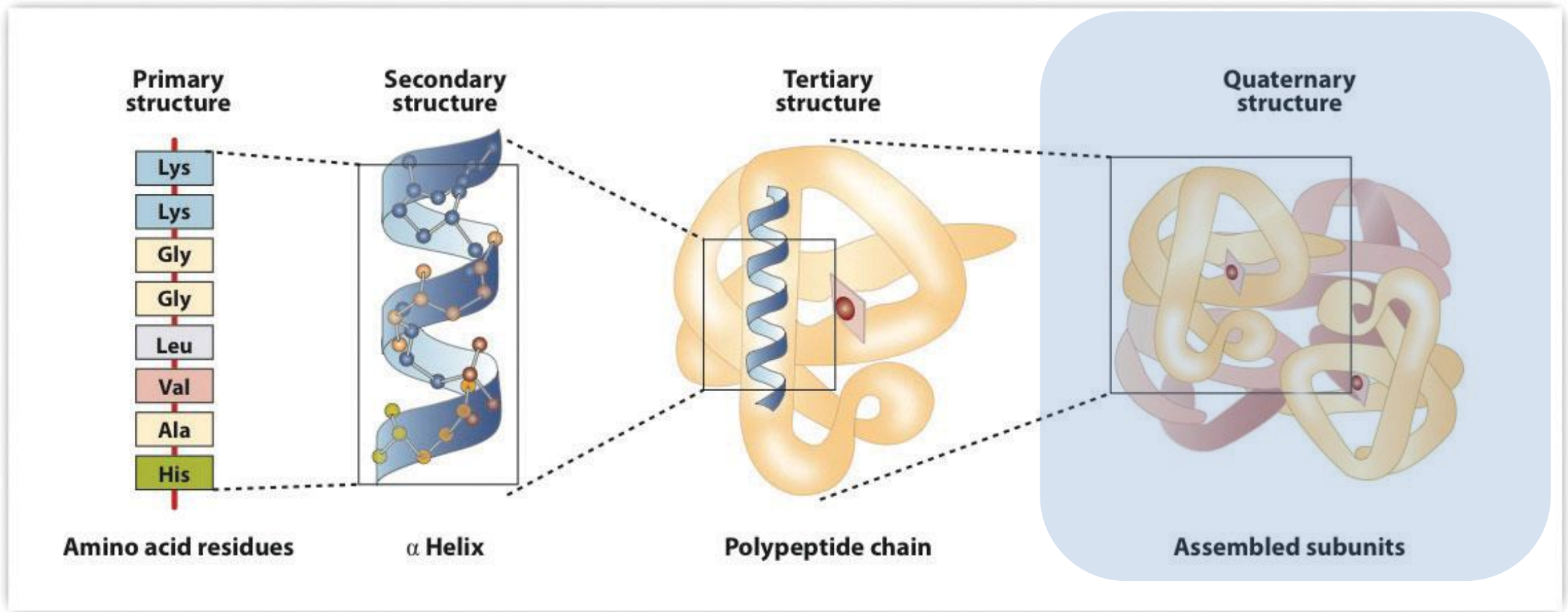


# Tertiary Structure

- How the **substructures** of the polypeptide chain **fold in 3D space**
- The result of interactions:
  - Between amino acid R-groups
  - Between amino acid R-groups and the **fluid environment**
- These interactions include:
  - **Spontaneous formation**
    - Hydrogen bonding
    - Hydrophobic interactions **Relatively weak**
    - Ionic interactions
  - **Enzymatically mediated**
    - Disulfide bridges **Strong**



# Protein Structure: Four Levels of Organisation





# Quaternary Structure

- Proteins with **more than one polypeptide chain** have Quaternary Structure
- They can be joined together by
  - Disulfide bridges (mABs)
  - Charge interactions
  - Hydrophobic interactions
- mABs have 4 polypeptide chains in their quaternary structure their heavy chains are held together by a strong bond called a disulfide bond
- Disulfide bonds form between two cysteine amino acids

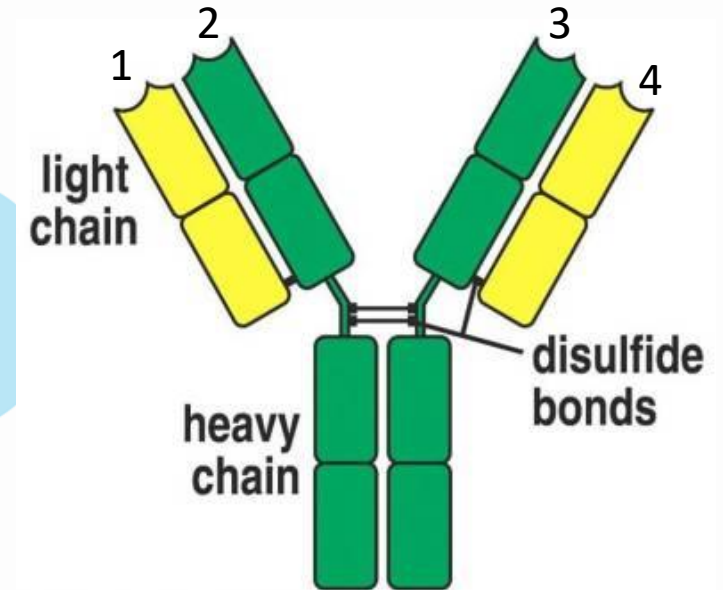
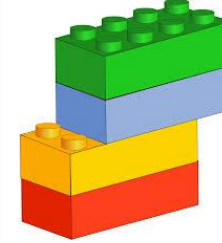
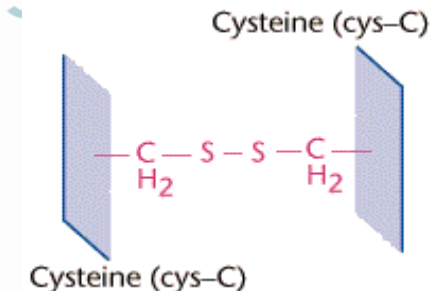


Figure 3-2 Immunobiology, 6/e. (© Garland Science 2005)



<https://www.youtube.com/watch?v=OBJ95upPxuE&t=4s>



# Topics



Introduction to Protein Chemistry

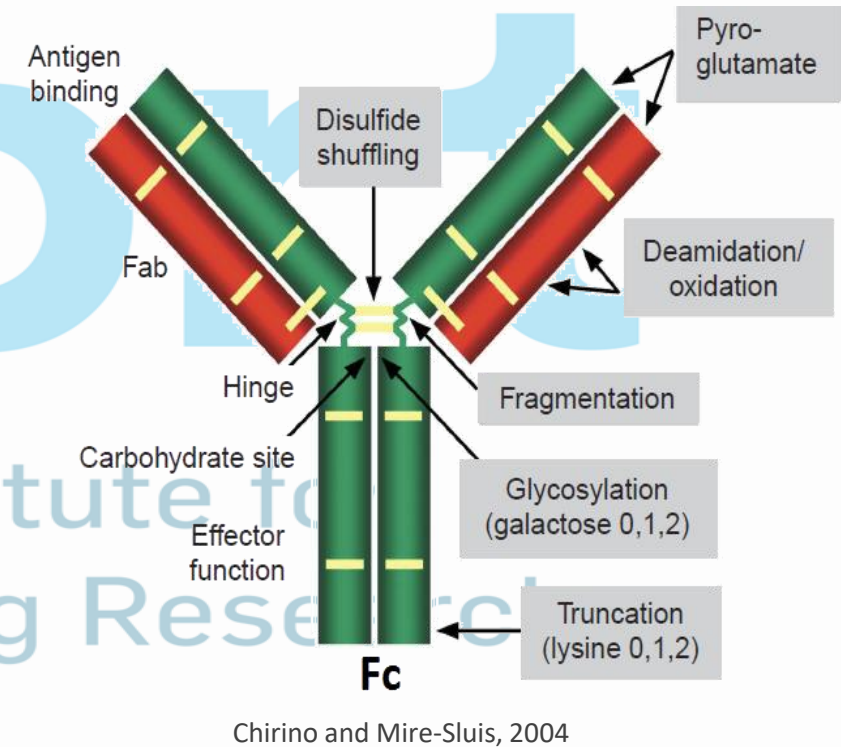
Protein Structure

Post Translational Modifications

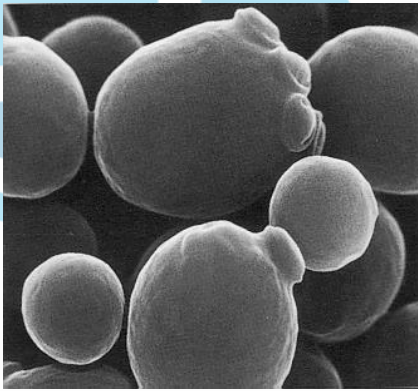
Protein Stability

# Post Translational Modification

- After a cell creates a protein ('translation') **chemical modifications** (PTMs) may occur
  - Performed by enzymes in the cell
- Proteins can undergo 100s of PTMs
- PTMs influence the **structure** and thus the **function** of the protein
- PTMs are species specific
  - Bacteria versus mammalian cells



# Why are post translational modifications important?



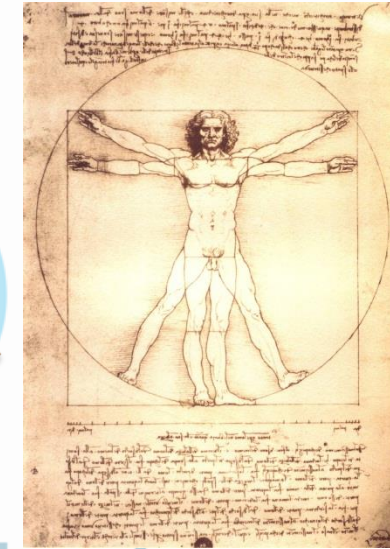
**Yeast**

~6000 genes that  
encode for  
proteins



**Drosophila**

~20,000 genes that  
encode for  
proteins



**Human**

~25,000 genes that  
encode for  
proteins



**PTMs allow for the diversification of the protein pool as organisms  
become increasingly complex**

[www.biochem.if.ua](http://www.biochem.if.ua)

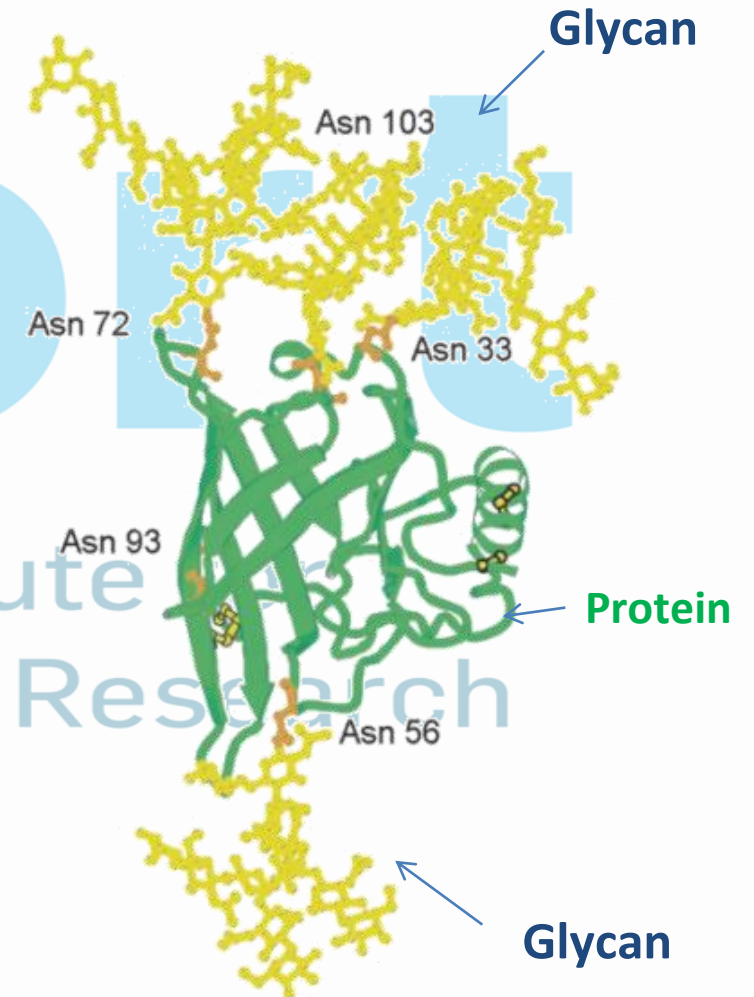
[www.iBioseminars.org](http://www.iBioseminars.org)

[www.mycor.nancy.inra.fr](http://www.mycor.nancy.inra.fr)

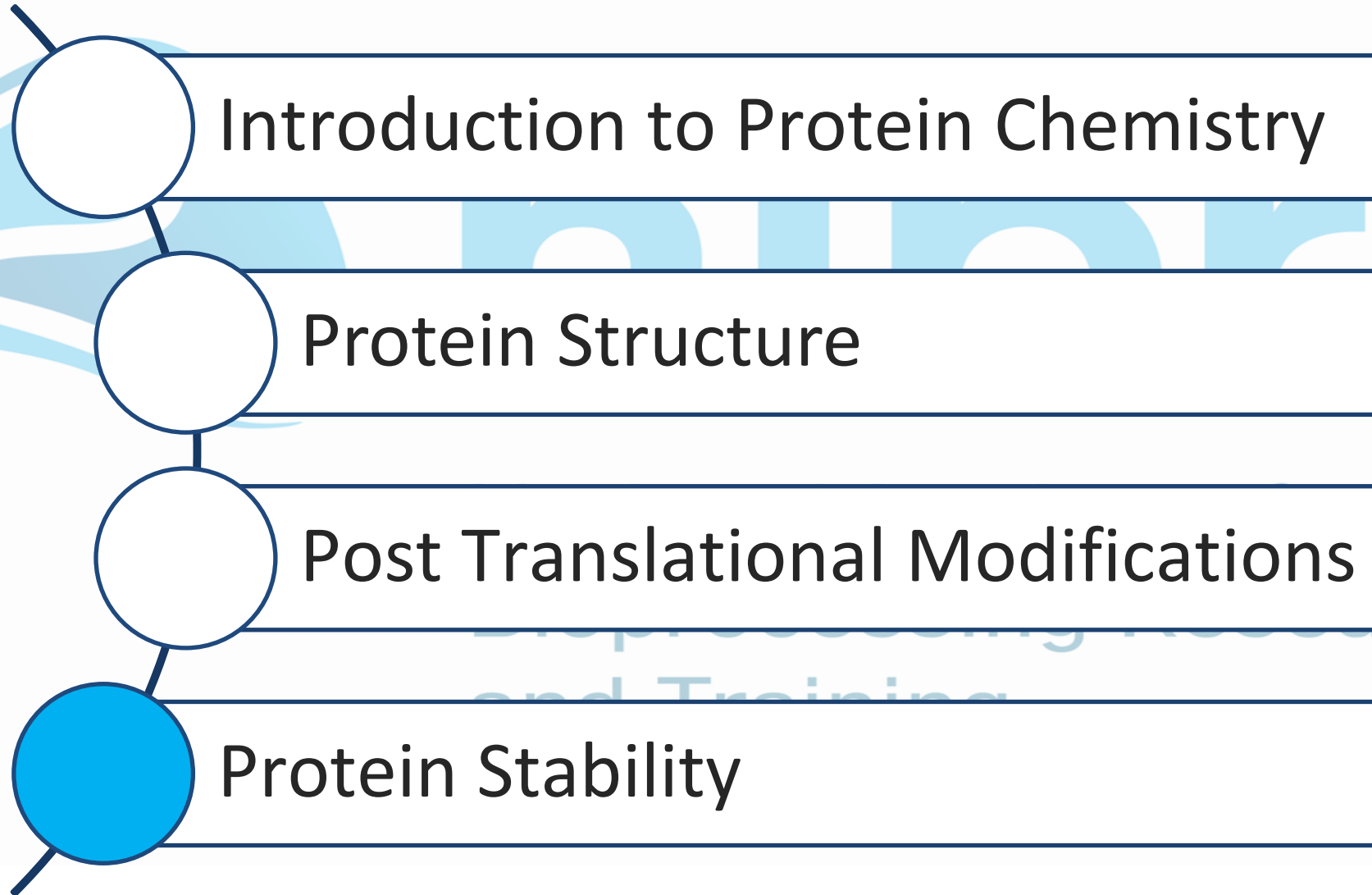
[www.drawingsofleonardo.org](http://www.drawingsofleonardo.org)

# PTM Example: Glycosylation


- The addition of chains of carbohydrates ('glycans') to proteins
  - Performed by 'glycozymes'
- **66% of all biopharmaceuticals are glycosylated**
- Affects many aspects of protein **structure** and **function**
  - Folding, charge, solubility, stability, immune tolerance in patients
- Also affected by bioprocessing!



# Topics





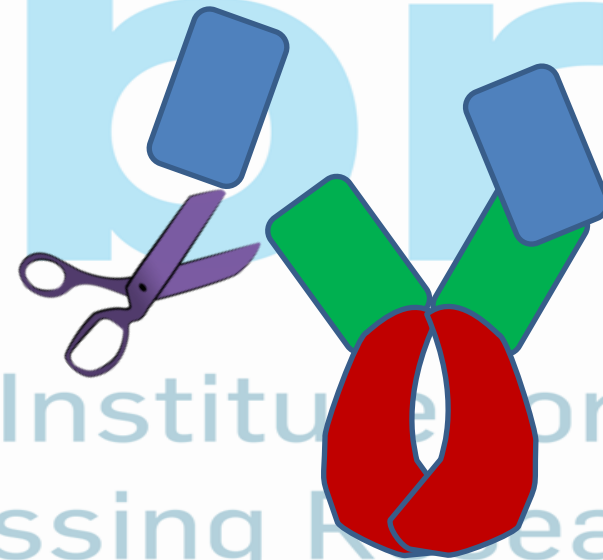


# The structural features of biopharmaceuticals are not fixed!

- Many **physicochemical properties** are subject to change during all steps of manufacturing
- Rigorous **control** of the process and sensitive **testing** of drug **substance**, drug **product** and **raw materials** aim to minimise this
- Batches are still a **heterogeneous mix** of structurally related **isoforms**, but these must be defined and consistent from batch-to-batch

# What happens if a protein becomes unstable?

1. Clipping
2. Denaturation
3. Aggregation
4. Precipitation
5. Chemical alterations



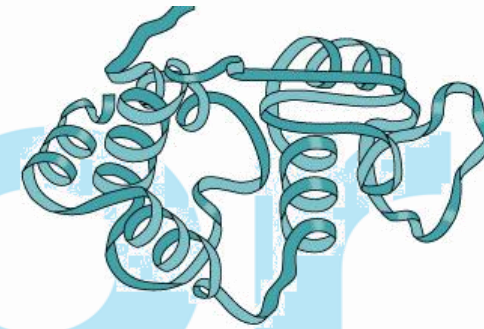
# Clipping (Fragmentation/Truncation)

- Breaking apart of **subunits of a protein**
- Breaking **peptide bonds** between amino acids (hydrolysis) and degrading polypeptide chains
- Caused by:
  - **Contaminating proteases** (more on this later)
  - **Photooxidation**: reaction with light energy
  - Residual **caustic cleaning agents** (sodium hydroxide) in equipment or low pH

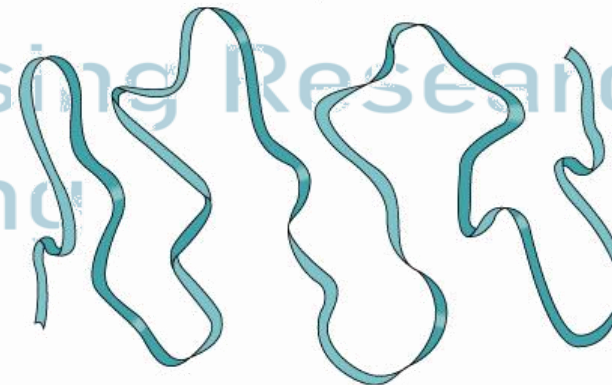


# What happens if a protein becomes unstable?

1. Clipping
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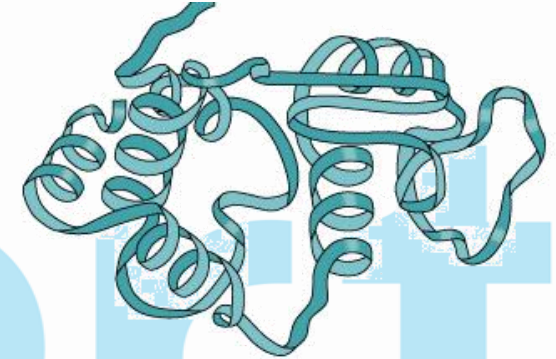
Active (functional) protein



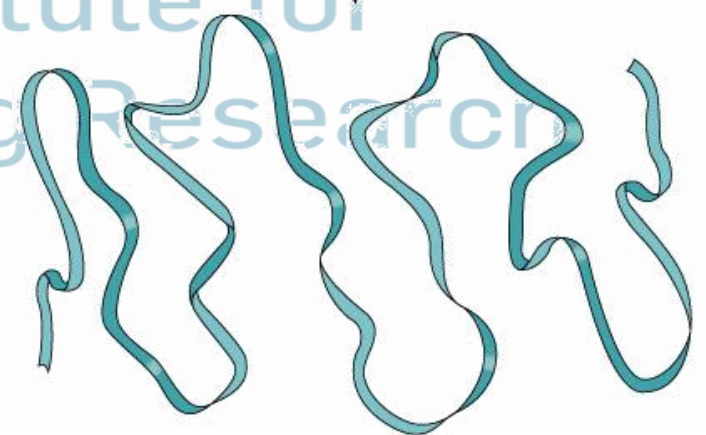
Denatured protein

# Denaturation

- Major change from the original native state without altering the primary structure
- Denatured proteins **lose their function** e.g. no therapeutic action in the patient
- Can be reversible or irreversible



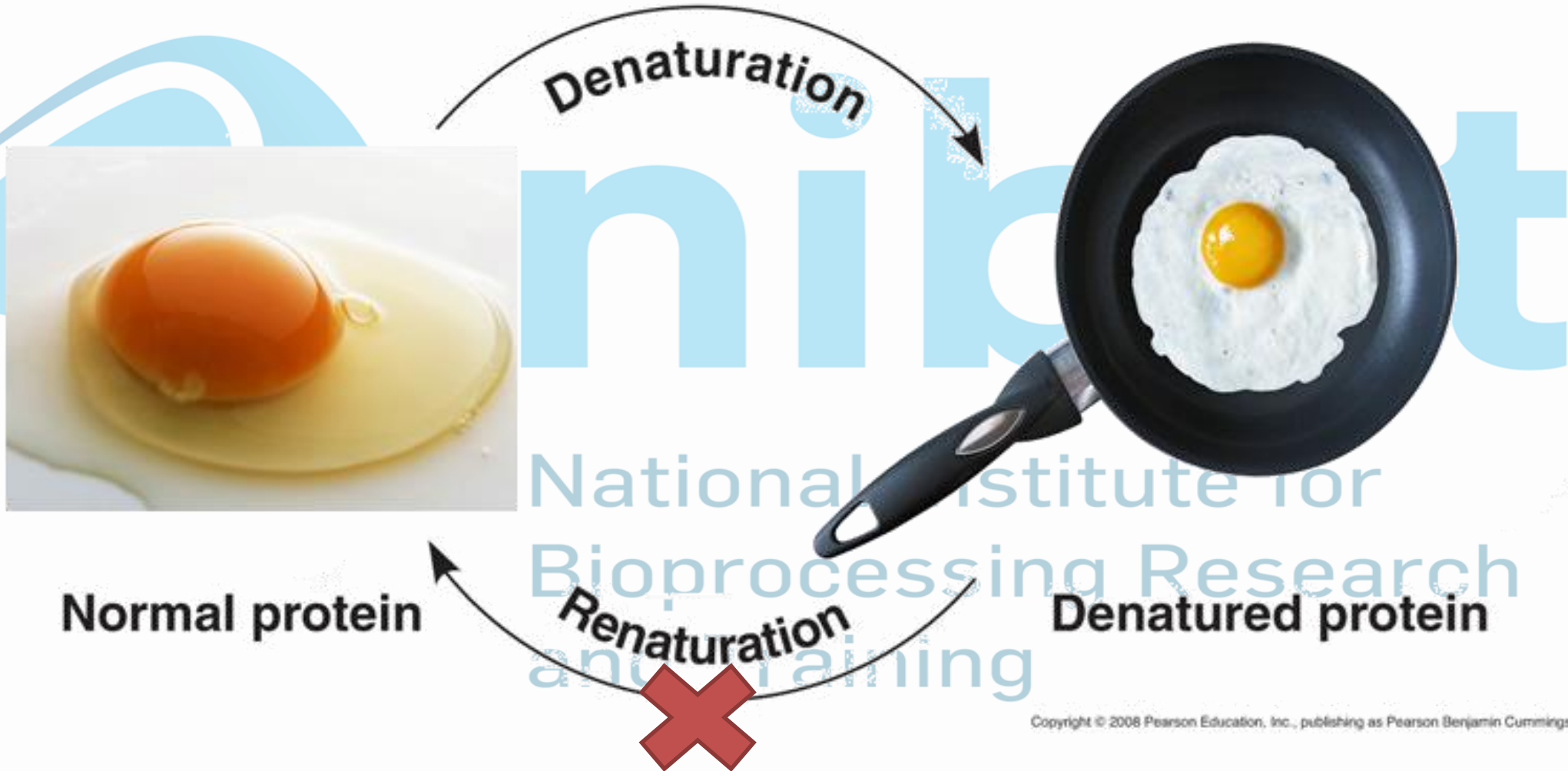
Active (functional) protein



Denatured protein



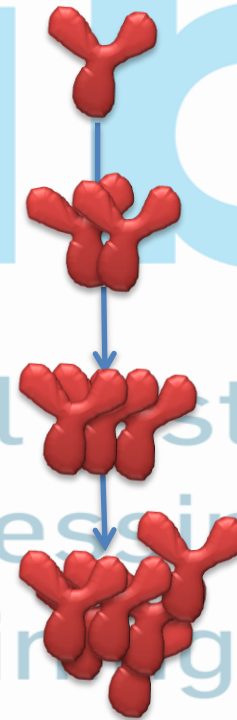
# Denaturation



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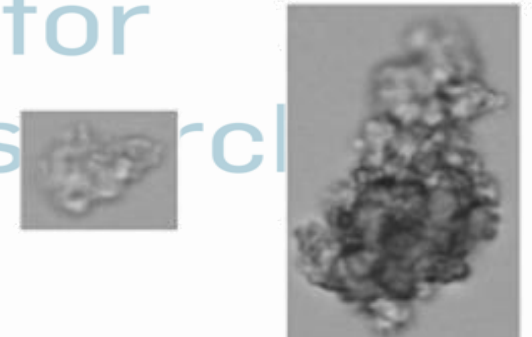
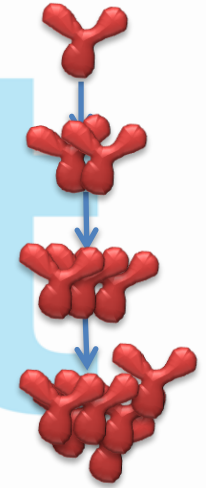
# What happens if a protein becomes unstable?

1. Clipping
2. Denaturation
3. Aggregation
4. Precipitation
5. Chemical alterations



# Aggregation

- Self-association of (usually mis-folded) proteins
- The monomeric (non-aggregated) form of the protein is typically the biologically active form
  - Dimers – 2 protein monomers aggregated
  - Trimers – 3 protein monomers aggregated
  - Multimers – multiple proteins aggregated
- Aggregation can be:
  - Reversible or non-reversible
  - Covalent or non-covalent
  - Visible or subvisible

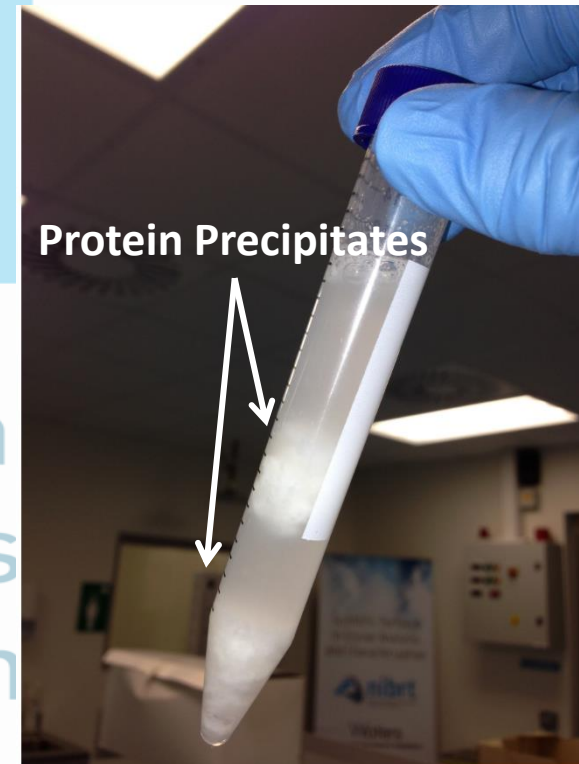


IgG Protein Aggregates by FPIA Imaging

Wim Jiskoot (2010) at FIP Pharmaceutical Sciences World Congress

# What happens if a protein becomes unstable?

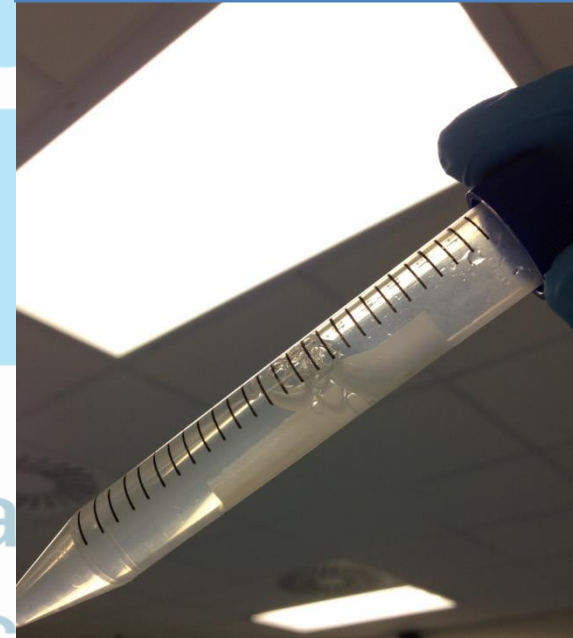
1. Clipping
2. Denaturation
3. Aggregation
4. Precipitation
5. Chemical alterations



# Protein Precipitation

- Protein becomes insoluble
- Result of:
  - Denaturation (heat, pH effects)
  - Excessively high protein concentration
  - Excessively high salt concentration in the buffer
  - Exposure to organic solvents

Protein Solution



Protein is fully dissolved.  
Solution is clear.

+ 2ml HCl



Protein denatured by pH change and precipitates.  
Solution is cloudy.



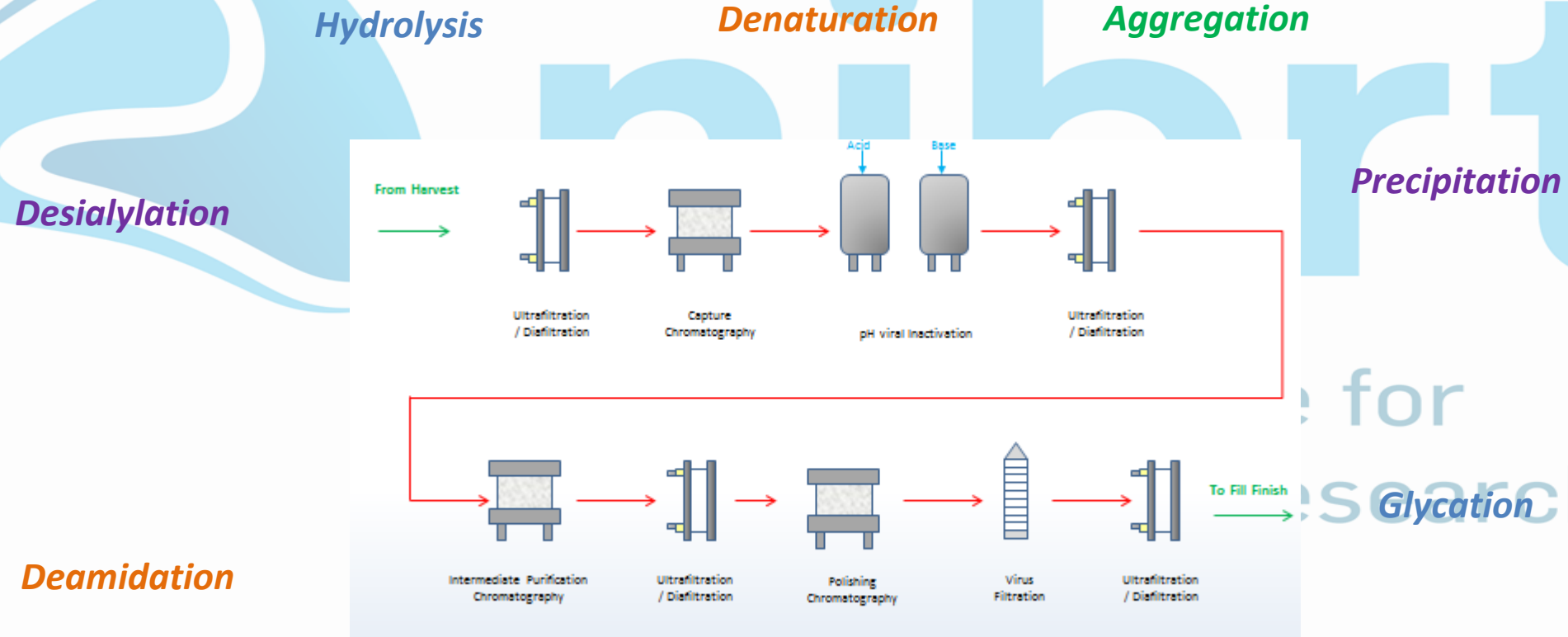
# What happens if a protein becomes unstable?

1. Clipping
2. Denaturation
3. Aggregation
4. Precipitation
5. Chemical alterations

nibrt

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Bioprocessing Research  
and Training

# Downstream processing can chemically alter proteins significantly



Walsh, Gary. *Pharmaceutical biotechnology: concepts and applications*. John Wiley & Sons, 2007.



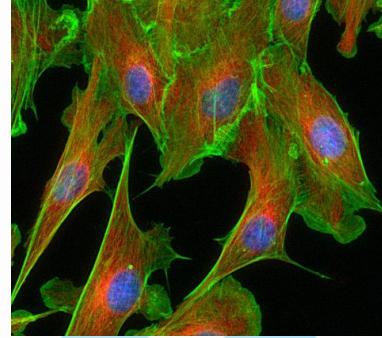
# What happens if a protein becomes unstable?

- Lose therapeutic efficacy
- Cause serious adverse effects in patients
  - Aggregation increases chance of immunogenic reactions
- Reduce production yield

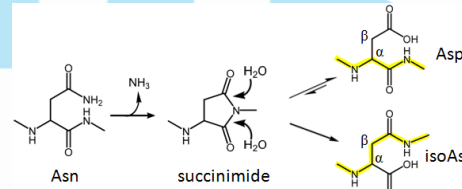
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# Sources of protein instability

1. Biological origin



2. Chemical origin



3. Physical origin



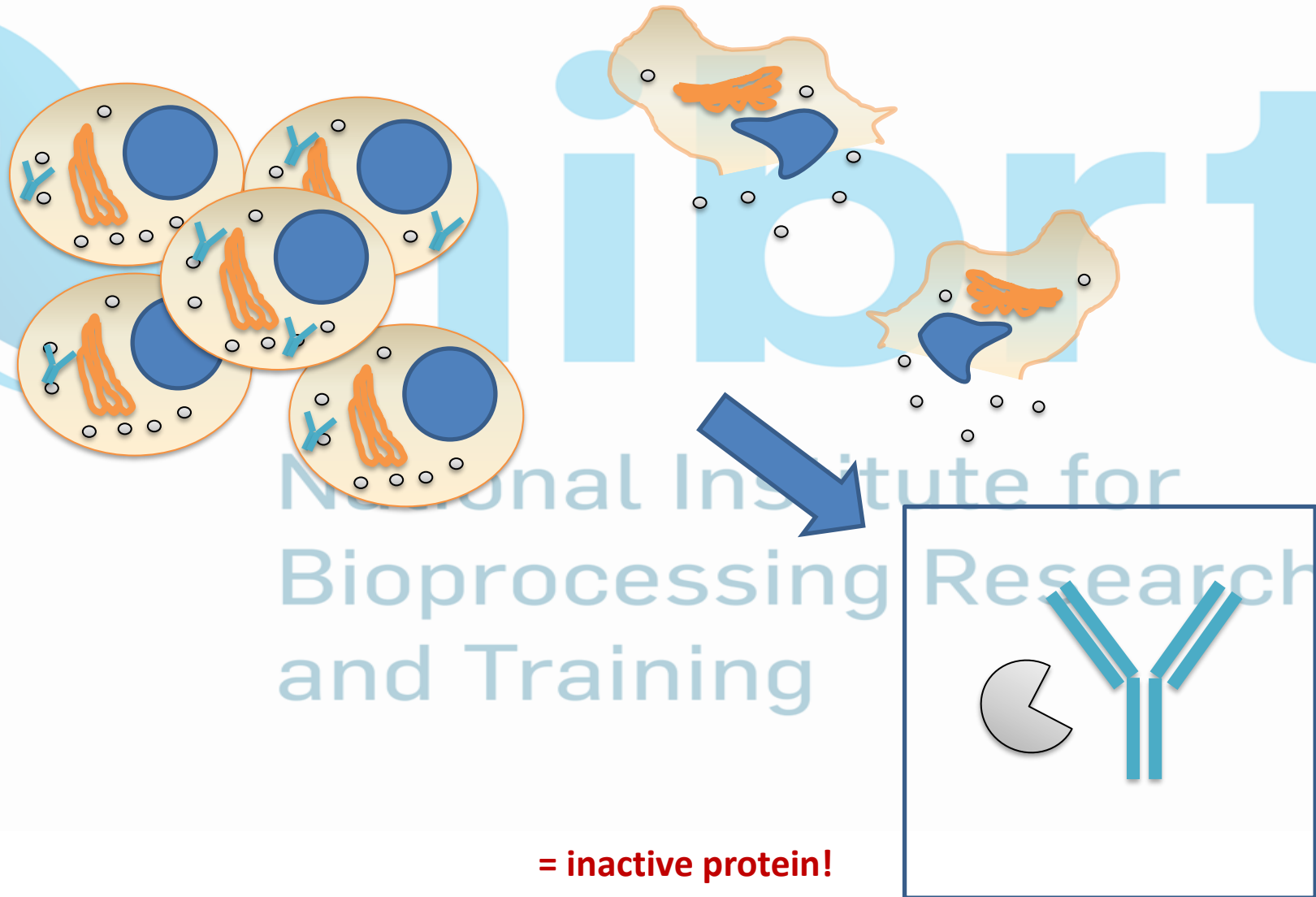


# Biological Origin

- **Mutations** can lead to amino acid substitutions which lead to **structural changes**
- **Media composition** and culture conditions can affect PTMs, particularly **glycosylation**
- **Cell death** in the bioreactor can lead to degradation of the biologic due to release of **proteolytic enzymes**
- **Perfusion processing** can help to quickly separate fragile product from proteolytic enzymes in the bioreactor

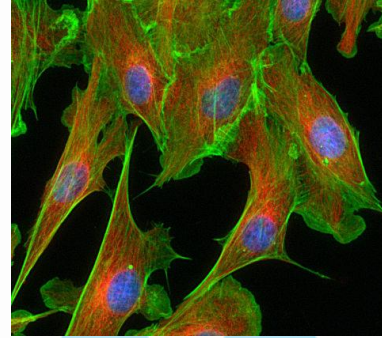


# Proteolytic Enzymes Attack the Biopharmaceuticals in the Bioreactor

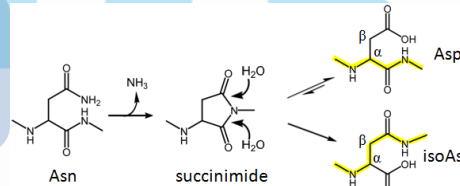


# Sources of protein instability

1. Biological origin



2. Chemical origin



3. Physical origin





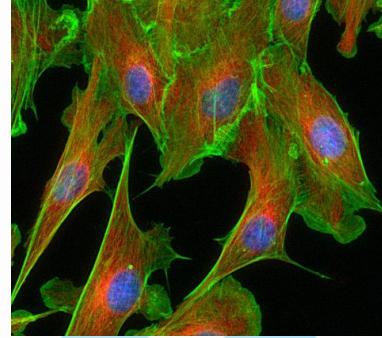
# Chemical Origin

- Certain environmental factors during bioprocessing can change the chemical structure of the protein
  - **pH:** Proteins are very sensitive to pH changes, which can affect how amino acids interact and thus folding
  - **Ionic strength:** salt disrupts protein-protein interactions

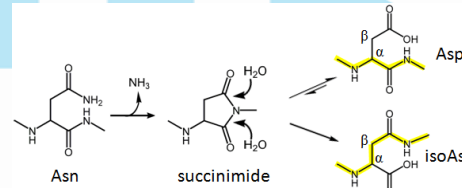
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and Training

# Sources of protein instability

1. Biological origin



2. Chemical origin



3. Physical origin





# Physical Origin

- **Mechanical shear**
  - Vigorous shaking (like whipping egg whites)
- **Temperature**
  - Proteins function at specific temperature
  - Freeze/thaw cycles
- **Protein concentration**
  - Too low: adsorption to container surface
  - Too high: precipitation
- **Light/ionising radiation**
  - Can trigger chemical chain reaction





# Correct Protein Handling

- Prevent shear during transfers
- When purifying protein, maintain physiological conditions as much as possible
- Work at low temperatures if possible (4°C )
- Limit processing timelines
- Avoid protein lingering at/near isoelectric point

Beware of introducing artefacts when handling QC samples!

# Topics



Introduction to Protein Chemistry

Protein Structure

Post Translational Modifications

Protein Stability



# Thank You

