

**UMR-CNRS 8612 - Institut Galien Paris-Saclay** 

**University Paris Saclay** 

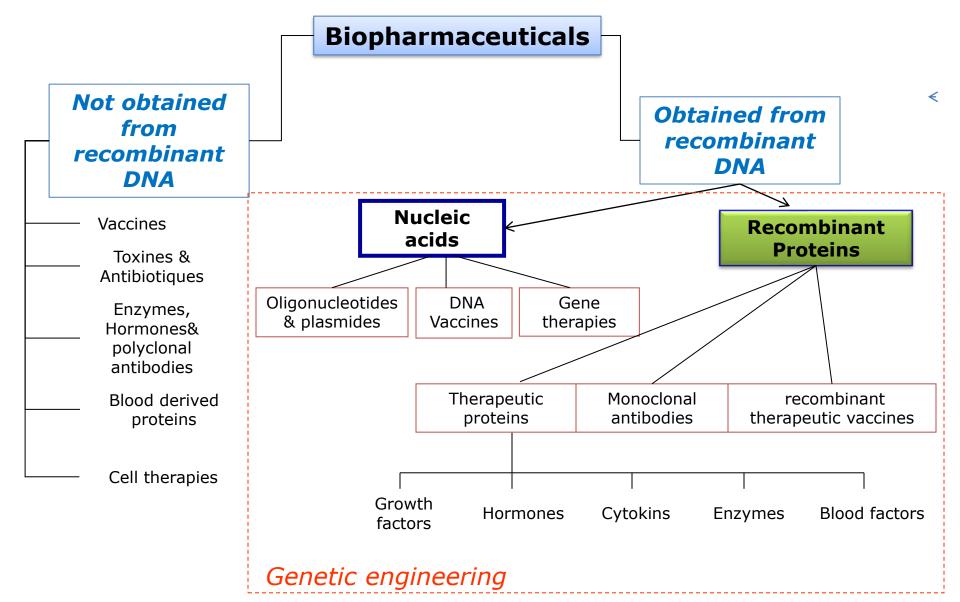
### **Quality control of biopharmaceuticals**

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### Classification of Biopharmaceuticals



### • Different sources of biopharmaceuticals

> **Definition**: Biopharmaceuticals are defined as pharmaceuticals manufactured by biotechnology methods, with the products obviously having **biological sources**, usually **involving live organisms or their active components** 

**1-Extraction** : from tissues, animal or human fluids (blood, urines, cells, tissues, milk...)

**2-Cell culture production** : Prokaryotic cells (bacteria) or eukaryotes (fungi, insects, mammals)

natural secretion : toxins or bacterial enzymes
(Botulinum toxin, streptokinase ...)
after genetic modification : expression of a foreign gene

**3- From transgenic animals or plants** 

4- Live Organisms (larvae; eggs...)

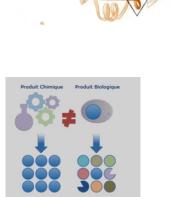




### Specificities of therapeutic proteins

**Therapeutic proteins differ greatly** from chemically synthetized molecules by: Secondary structure • Their size Aspirin, Interferon alfa, MW: 180 D 165AA, MW: 19 625 Da Monoclonal antibody 150kDa βsheets High structural complexity which is mandatory for their activity Feuillet bêta 3 N-linked chains 1 O-linked chain Heterogeneity **Tertiary structure** One glycoprotein = mixture of glycoforms in a given and constant proportion Ь

### and by the way to produce them



Hélice alpha,

**α-helix** 

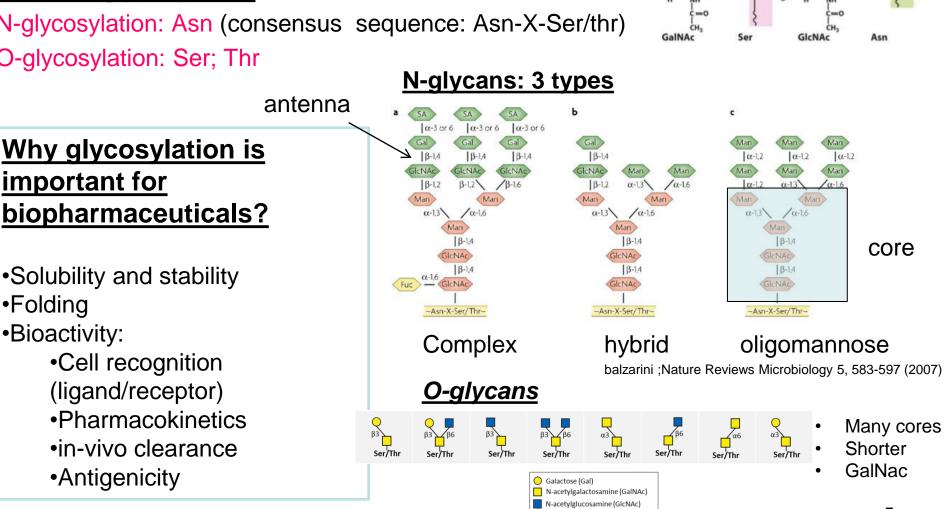
Pas de structi définie



#### Structure/activity: Glycosylation a major posttranslational modification (PMT) (a) O-linked (b) N-linked

#### N and O-glycosylation

N-glycosylation: Asn (consensus sequence: Asn-X-Ser/thr) O-glycosylation: Ser; Thr



### • Quality control of recombinant proteins

- is mandatory to ensure safety and efficiency
- The active product is heterogeneous and structurally complex
- needs several complementary techniques to address :
  - $\circ$  Purity
  - o Identity
  - o Dosage
  - o Stability
  - $\circ$  Activity

### • At which step?

### In-process

In-process tests are performed at critical decision-making steps and at other steps where data serve to confirm consistency of the process during the production of either the drug substance or the drug product.

- Purified protein
- Formulated product
- Compounding product (hospital)

### Quality control - Objectives

- QC testing: To assess and ensure the safety and efficacy of the therapeutic protein
  - To verify lot-to-lot consistency

#### **Identity test**

- To establish that the product batch contains the correct therapeutic protein
- Biological activity, immunogenicity

#### Dosage

- To determine accurately the concentration of the purified protein / in the final product
- To ensure that the correct amount of active pharmaceutical is delivered to the patient

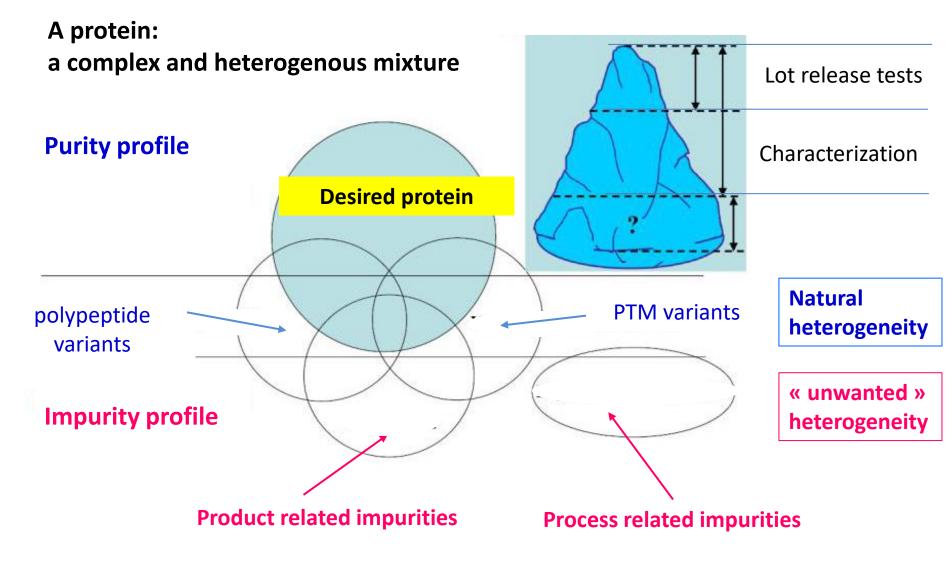
#### **Purity**

- A biopharmaceutical product must be free of contaminating substances
  - Impact the performance of the protein
  - Cause unwanted and serious side effects (immunogenicity)

### **Stability**

- To evaluate the susceptibility of the protein toward different stresses
- To define optimal storage conditions

# • Why the evaluation of the desired product is a real challenge?



Kozlowski S et al, Adv Drug Del Rev 2006

PTM: post-translational modifications

### • What are the impurities?

#### **Process-related impurities:**

Expected to be removed during the downstream process

#### Cell-derived

Host cell proteins, nucleic acids, endotoxins...

#### • Cell-culture medium derived

Inducers, antibiotics, enzymes, serum, bacteria...

#### Downstream derived

Chemicals and biochemicals, inorganic salts, solvents, residual components

### **Product-related impurities:**

#### Related forms and variants

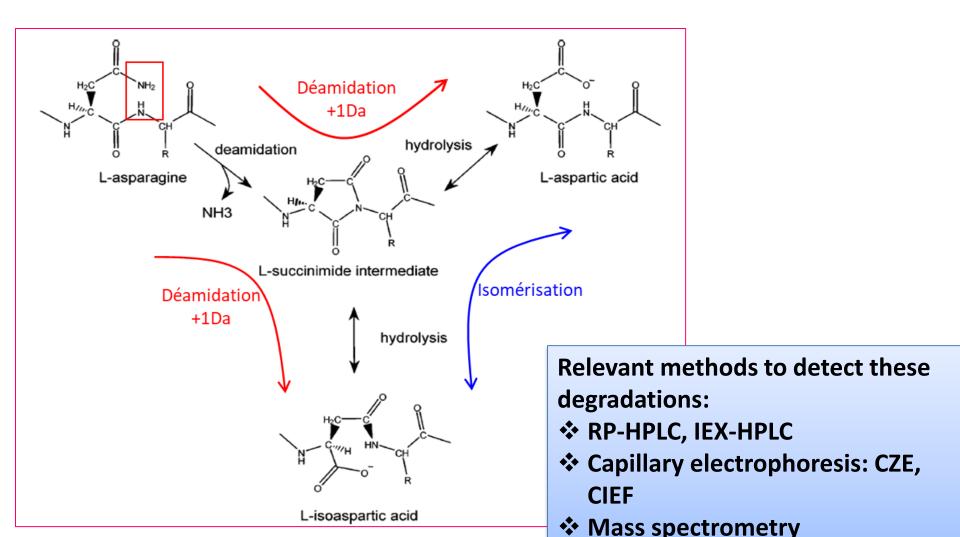
Deamidation, oxidation, isomerisation, racemization, glycated, altered PTMs....

- Truncated forms, fragments;
- Dimers, oligomers, aggregates,
- Misfolded or unfolded forms
- Cys linked variants, scrambling.

#### • Chemical Degradation pathways – charge variants

#### **Deamidation (Gln, Asn), Isomerization (Asp)**

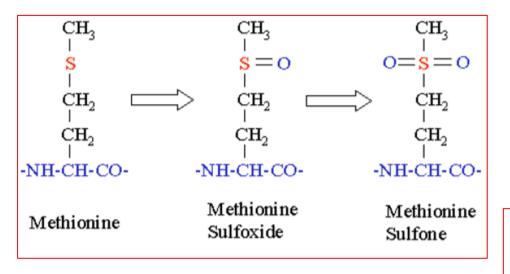
- Very frequent during production, formulation, storage
- Catalyzed by high temperatures and ionic strengths, at neutral and basic pHs



### Chemical Degradation pathways – size variants

**Oxidation Met, Cys, (Thr, Phe, Lys)** 

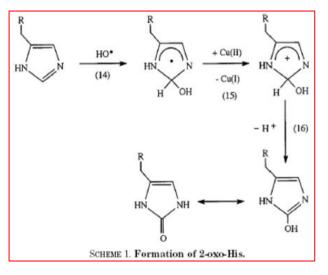
- Very frequent during purification, formulation, storage
- Site specific, metal catalyzed (Met, Cys)
- pH catalyzed (except Met)



**Relevant methods:** 

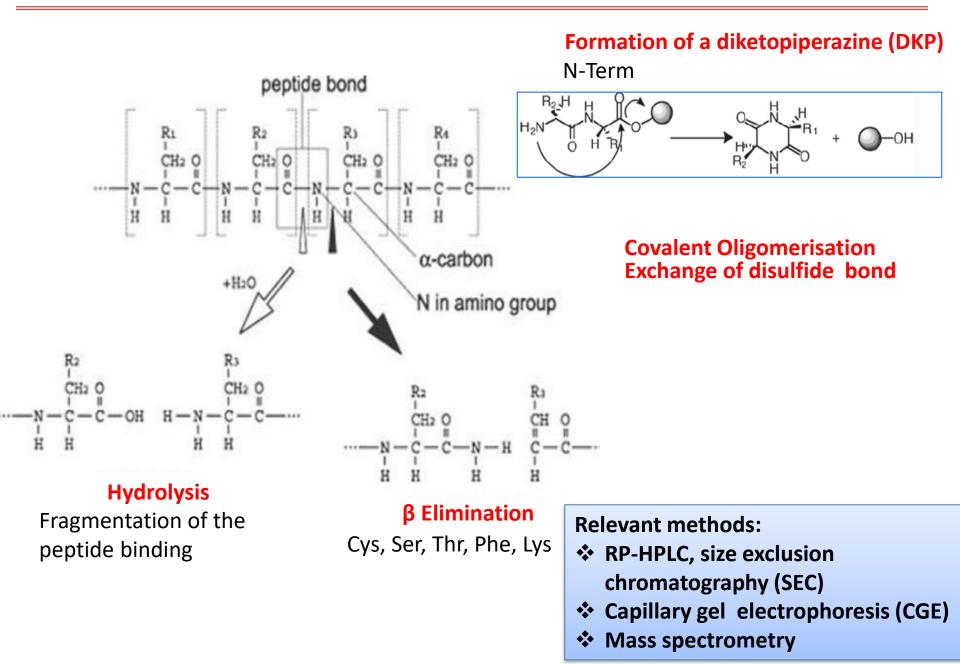
- ✤ RP-HPLC, IEX
- Capillary electrophoresis
- Mass spectrometry

His oxidation catalyzed by metal

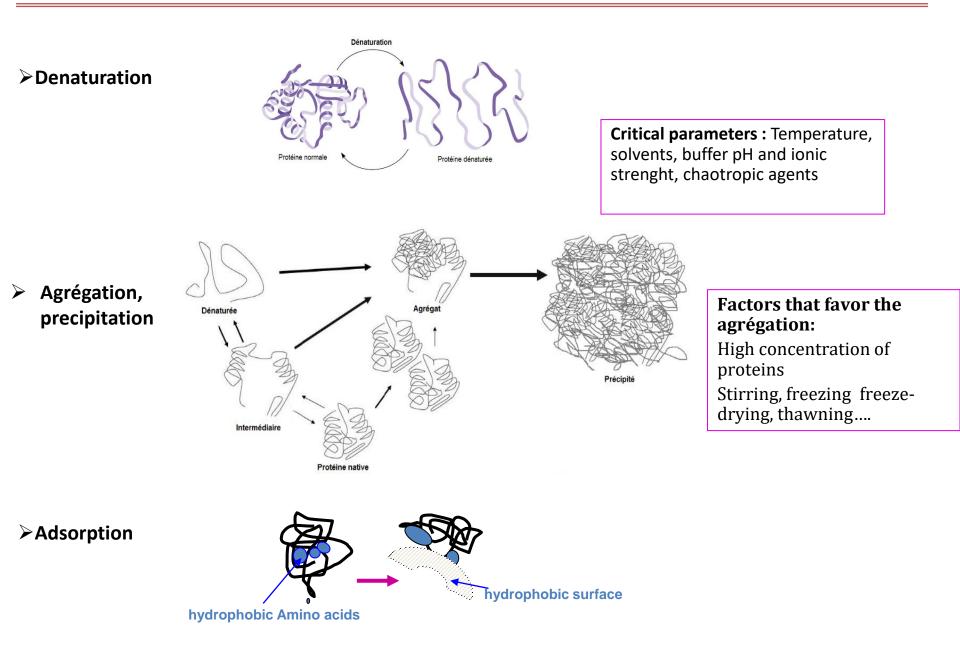


Aliphatic amino acid oxidation $H \rightarrow H_{3} \rightarrow H_{3$ 

#### Chemical degradation pathways : fragments / oligomers

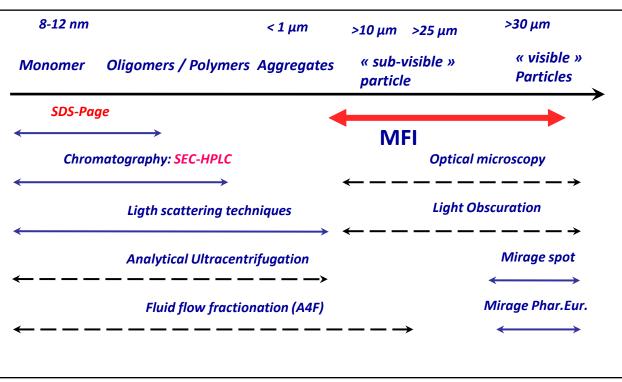


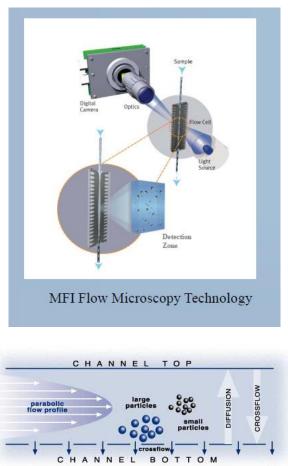
#### • Physical degradation pathways : agregates, oligomers



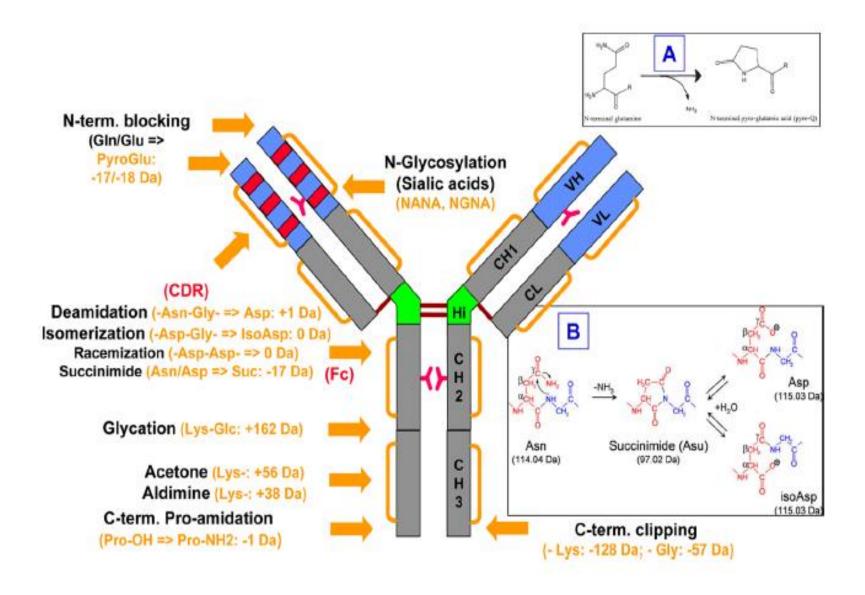
## Aggregates and particules detection

- Aggregates and particles are impurities among others , they can generate undesirable side effects (immunogenicity).
- Analytical techniques to be used depend on their size



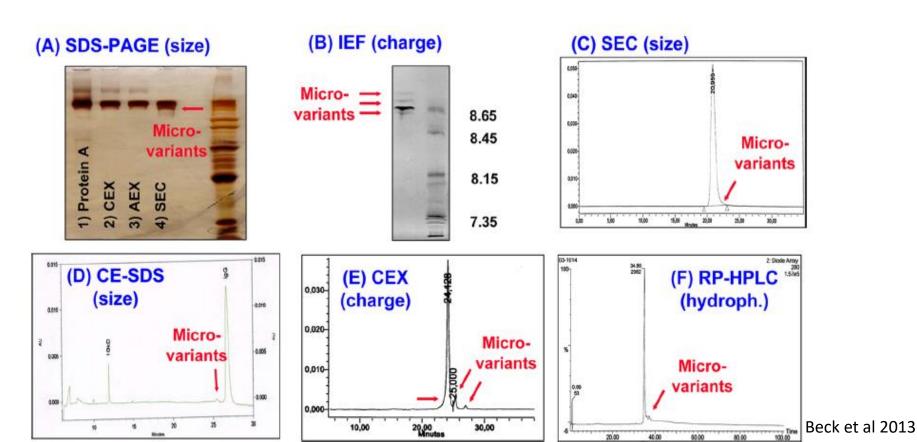


### How many microvariants are we looking for? Exemple of a monoclonal antibody

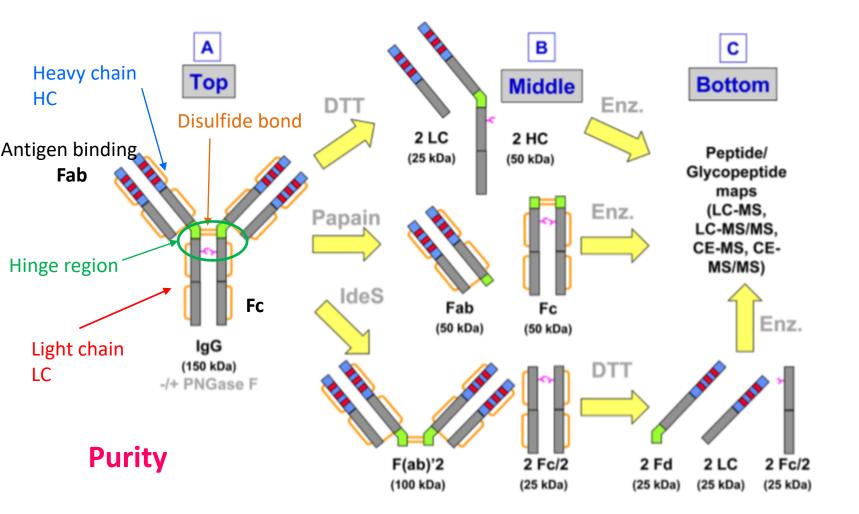


### **Complementary techniques are required for a QC**

- Electrophoresis-based techniques (gel electroph., CE, CE-MS...)
- Liquid Chromatography techniques (all modes/MS)
- **Spectrophotometrics techniques**: Circular dichroism, Fluorescence and UV absorbance, IRTF, DSC, Light diffusion...



### • 3 Approaches for mAbs analysis



#### Top down, Middle up, Bottom up

DTT: dithiothreitol

Beck A et al, Anal Chem2012

#### Middle up approach to detect Isomerisation and succinimide by HIC-chromatohraphy

#### 140 Fab peaks (1\*, 1-6) 120 Fc peaks 100 mAU (214 nm) 80 2 60 ASU 40 isoAsp32 undigested Asu74 material SOASP Asu32 20 nsc, 1 0 15 25 35 10 20 30 40 45 50 Time (min.)

#### HIC: Hydrophobic interaction chromatohraphy

**Fig. I** HIC elution profile of a papain digested MAbI sample. Peak I\* corresponds to Asu74; nsc = non-specific clip due to papain cleavage, coelutes with peak I\*; Peak 2 is Asp32 with a free thiol group; Peak 3 corresponds to isoAsp32; and Peak 5 corresponds to Asu32.

Pharm Res (2012) 29:187-197

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### Impurities and methods to control the purity

#### 1-Proteines from the Host cell : non spécific methods required

- -SDS-PAGE/Coomassie blue staining
- «host cell proteins» HCP tests

#### 2- Degraded proteins : resolutive techniques

- RP-CLHP, IEX (deamidated, oxidized forms....), HIC (isomerisation)

-isoelectric focusing (IEF), Capillary zone electrophoresis (glycoforms, deamidated, oxidized forms....)

- Size exclusion chromatogaphy (fragments, agrégates)
- -capillary gel electrophoresis (fragments, agrégates, non glycosylated sites...)
- hydrophobic interaction chromatography (HIC) : misfolding , unfolding:
- techniques specific for aggregates and particles detection

#### 3-DNA

-Hybridization method using radiolabelled probes

#### 4- Pyrogenic substances, endotoxins :

- LAL test: in vitro test based on activation of the complement cascade by endotoxin in Limulus Polyphemus

- New coming "RP-HPLC-based technique"

#### 5- Virus

### **Approaches for Identity control**

#### • Primary sequence

- amino acid composition
- automated sequencing (EDMAN)
- peptide mapping

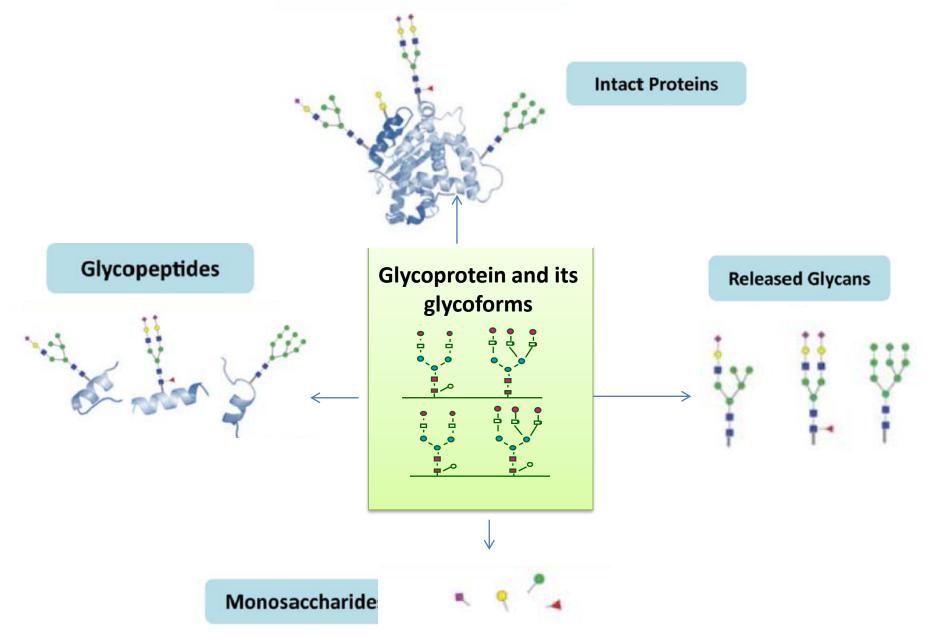
#### Physicochemical parametrers estimation

- Molecular mass: SDS-PAGE, size exclusion chromatography, capillary Gel Electrophoresis mass spectrometry (MALDI HPLC-ESI-MS using ion trap or FTICR mass spectrometry)
- Isoelectric point : gel isoélectrofocusing (IEF) or capillary isoelectrofocusing (CIEF)
- Hydrophobicity : **RP-HPLC**
- Combination of several parameters : PAGE , Capillary zone electrophoresis

#### • Secondary and tertiary structure

- Circular dichroïsm
- Infra-Red -based techniques
- Post translationnal modifications : Glycosylation mostly
  - Monosaccharide composition, sialic acid...
  - Glycan mapping
  - Glycopeptide mapping
  - Glycoform analysis: heterogeneity profile
- Biological and immunological activity

### **Glycosylation analysis can be performed at different levels**



### Dosage and activity

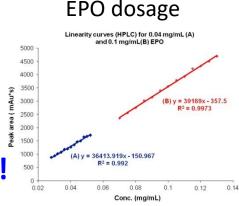
Colorimetric methods

- Méthodes du biuret et dérivées : **Biuret** : CuSO<sub>4</sub> (OH<sup>-</sup>) à 540 nm  $\rightarrow$  0,1 g/l **Lowry** : + Folin-Ciocalteu (Tyr) à 650-750 nm  $\rightarrow$  0,05 g/l **BCA** : + acide bicinchoninique (inverse) à 560 nm  $\rightarrow$  0,01 g/l - Méthodes par fixation de colorants : **Bradford** au bleu de Coomassie G250 (H<sup>+</sup>) à 600-650 nm  $\rightarrow$  0,01 g/l

- Spectrophotometry (UV at 280nm)
- Immunochemical methods : ELISA..
- HPLC, CE techniques can be quantitative too!

#### • Biological activity:

- Administer a known quantity of the product to a biological system (cells, animal...) and measure a quantitative response that will be transcribed into a unit of activity.
- In Vivo, Ex vivo, in vitro (cell cultures), biochemical or enzymatic assays, binding tests (SPR.....)



Rane et al., 2012

# • Analytical techniques for QC of common variants of biopharmaceuticals

	Modifications	Analytical techniques
Charge variants	Deamidation Oxidation Glycosylation	CEX, IEF, CIEF, CZE ; HILIC-HPLC
Size variants	Fragmentation, Aggregation	SEC-HPLC, SDS-PAGE, CGE
Conformers	Misfolding / Denaturation	HIC (not in a QC context)

# Homework for the 5<sup>th</sup> of November

• Prepare **15 min** oral presentation PPT

(groups of 3/4 students; each group assisted by one teacher T1 or T2)

Group 1: Acidic and basic variants analysis of Monoclonal Antibodies by IEX-HPLC- (T1) Group 2: Purity control of therapeutic proteins by Size Exclusion Chromatography (SEC-HPLC) (T1) <u>Group 3</u>: SDS-PAGE for purity and identity controls of therapeutic proteins (T1) Group 8: detection of chemical-Degradation variants of therapeutic proteins by RP-HPLC

Group 5: Peptide mapping of therapeutic proteins by RP-HPLC as an identity Control (T2) <u>Group 6</u>: N- Glycan mapping of therapeutic glycopoteins by HILIC-HPLC (T2) <u>Group 7</u>: Capillary gel electrophoresis for identity and purity control of Therapeutic proteins (T2) Groupe 4: Control of Secondary structure of therapeutic proteins- (T2)

## Oral presentation

- 3-5 slides/ 15 Min/ PPT presentation: use the template for each topic on ecampus
  - Principle/Mechanism of the technique/approach used for this specific application
  - Type of information provided for quality control purposes only , Limits and advantages of the technique
  - A few exemples illustrating this technique/method for quality Control of therapeutic proteins
  - 30 min of group works (now)
     BE CAREFUL : 15 min for presentation is a maximum!
  - Documents available on ecampus plateform; USE PREFERENTIALLY the selected ARTICLES for your presentation (figures or info...)
    - TUO8- biotechnology < supporting files < one folder per topic
  - PPT presentation , to be sent by email to your teacher (T1 or T2) before the 5th of November 2023 12:00am
  - Presentation the 6th of November 9-12am