



الجمهورية الجزائرية الديمقراطية الشعبية
PEOPLE'S DEMOCRATIC REPUBLIC OF ALGERIA



وزارة التعليم العالي والبحث العلمي
MINISTRY OF HIGHER EDUCATION AND SCIENTIFIC RESEARCH

جامعة باجي مختار - عنابة
BADJI MOKHTAR UNIVERSITY - ANNABA
كلية العلوم
FACULTY OF SCIENCES
قسم البيوكيمياء
DEPARTMENT OF BIOCHEMISTRY

Structural and Metabolic Biochemistry

TCSNV 2nd

Fondamental unit S3

AMINO ACIDS / PROTEINS

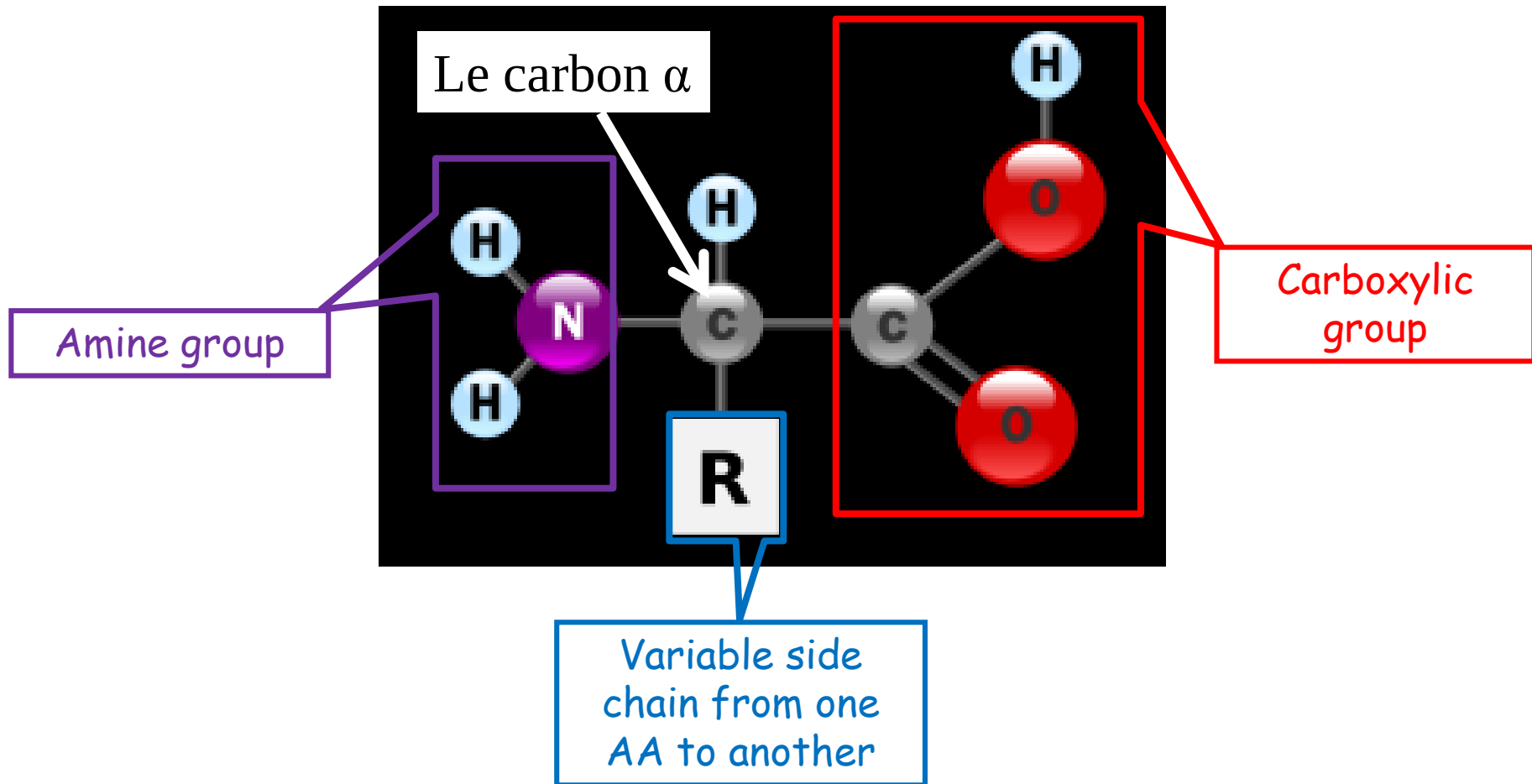
Pr. KADI-BIREM

Protein → proteide → peptide → amino acid

To date, 22 amino acids have been discovered in the proteins of living organisms. Among these, 20 are included in the genetic code of animals. Of these 20 amino acids, 8 are essential: they cannot be synthesized de novo by the organism and must therefore be obtained through the diet.

Isoleucine, Leucine, Lysine, Methionine, Phenylalanine,
Threonine, Tryptophane, Valine

General structure of an amino acid



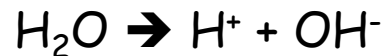
Properties of amino acids

Amphoteric properties

→ Substance that can act as an **acid** and as a **base**

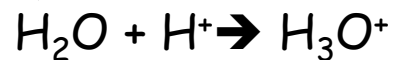
Ex: H₂O

Acid = Donates a proton



In a basic medium
(Low-proton environment)

Base = Accepts a proton

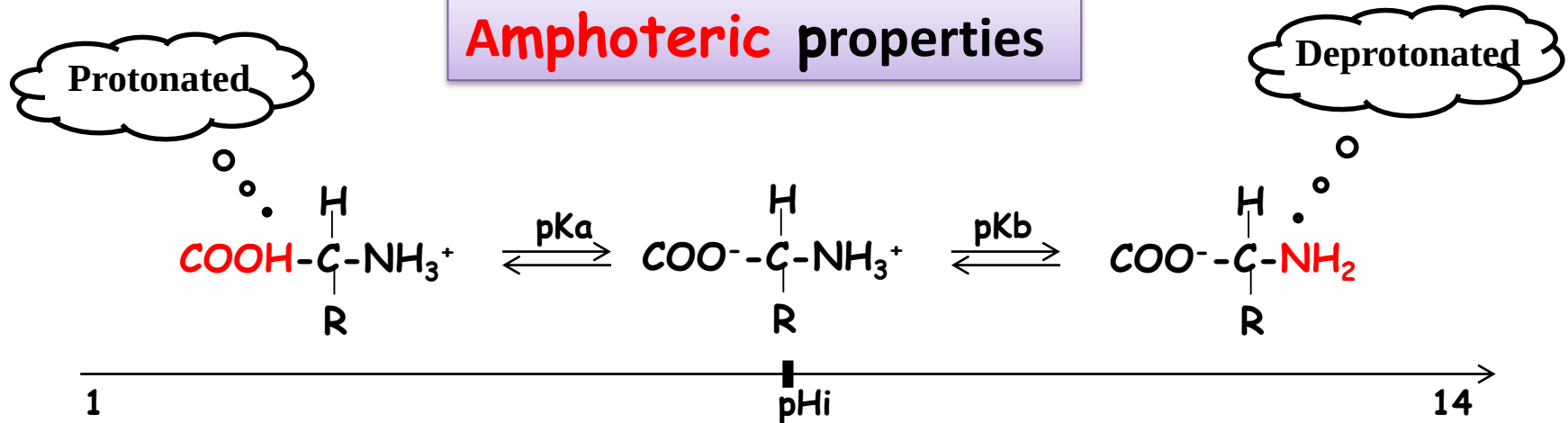


In an acid medium
(Rich-proton environment)

Just like water, amino acids
have amphoteric properties

Properties of amino acids

Amphoteric properties



- **pK → a specific pH**

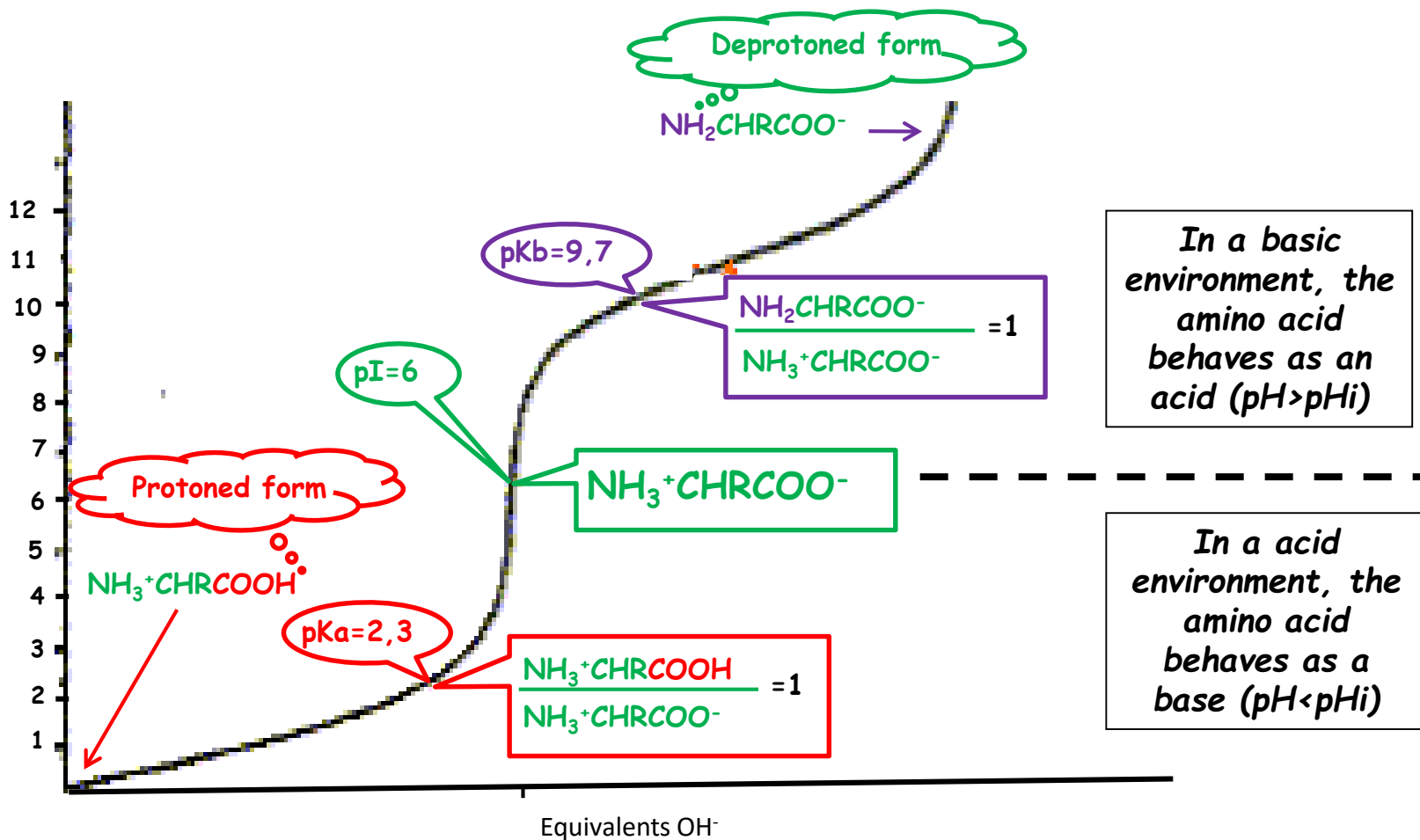
Every amine and carboxylic group have a pK (pK_a / pK_b)

pH=pK → 50 % protonated form and 50 % deprotonated form

- **Isoelectric point (pHi)**

$$\text{pH} = (\text{pK}_A + \text{pK}_B) / 2$$

Titration of amino acids



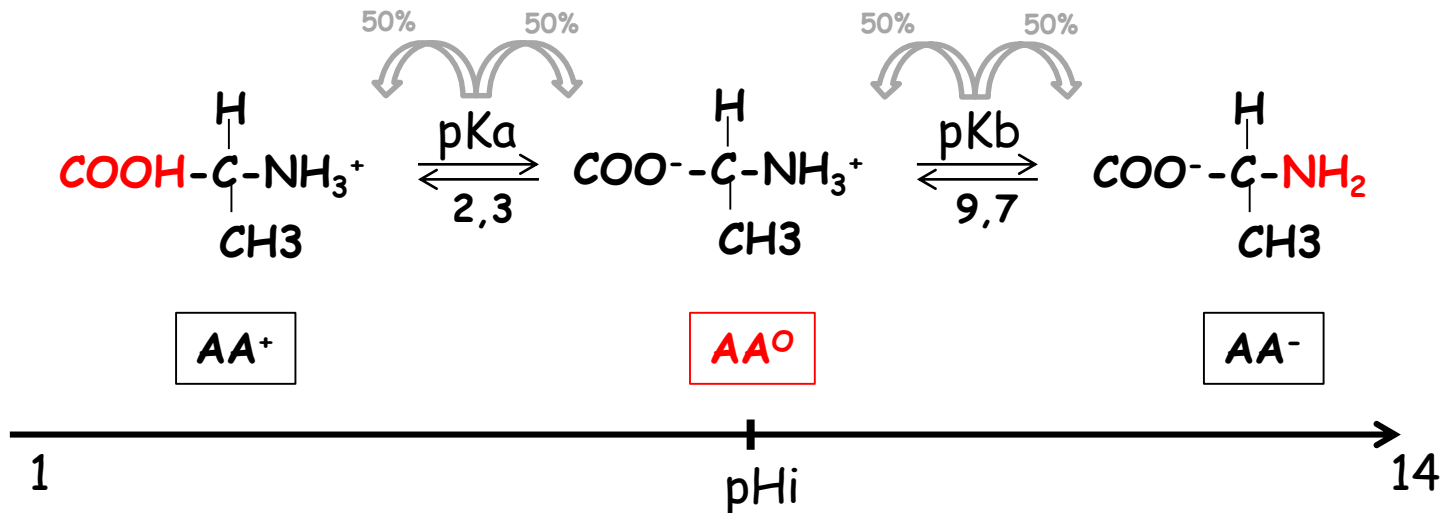
Properties of amino acids

Amphoteric properties

For an amino acid with a **non ionizable** side chain

Ex: Alanine (ALA) (A)

$pK_a=2,3$ $pK_b=9,7$



$$pH_i = pK_a + pK_b / 2$$

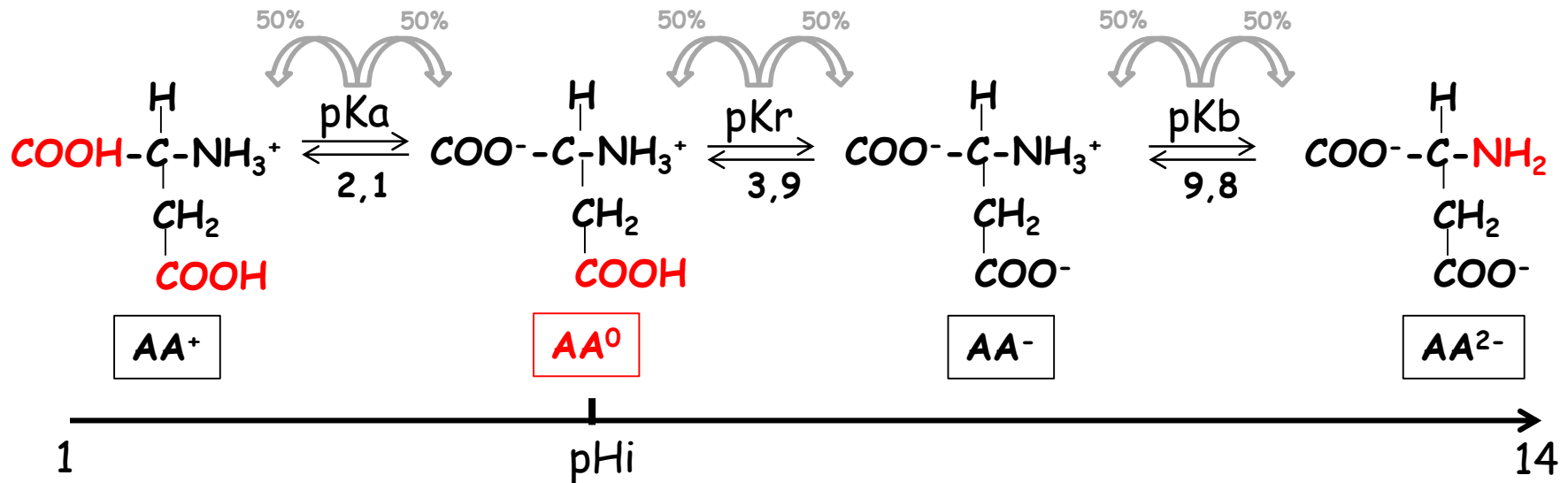
Properties of amino acids

Amphoteric properties

For an amino acid with **ionizable** side chain

Ex1: Acide aspartique (ASP) (D)

$pK_a=2,1$ $pK_b=9,8$ $pK_r=3,9$



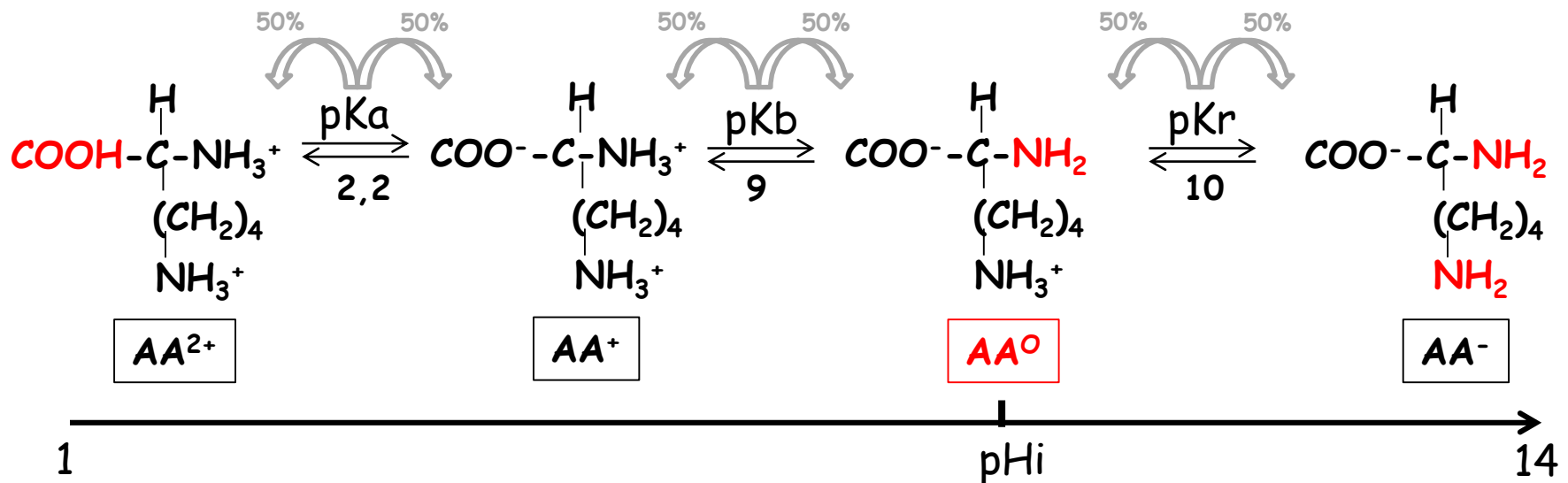
Properties of amino acids

Amphoteric properties

For an amino acid with **ionizable** side chain

Ex2: Lysine (LYS) (K)

$pK_a=2,2$ $pK_b=9$ $pK_r=10$



$$pH_i = pK_b + pK_r / 2$$

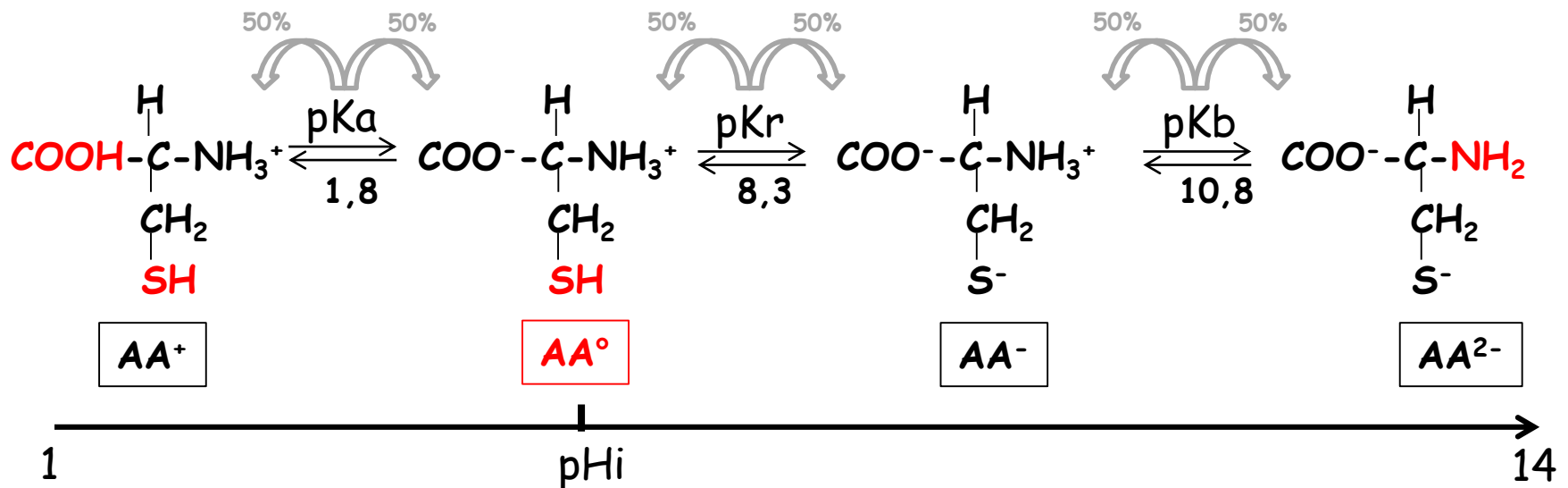
Properties of amino acids

Amphoteric properties

For an amino acid with **ionizable** side chain

Ex3: Cystéine (CYS) (C)

$pK_a=1,8$ $pK_b=10,8$ $pK_r=8,3$



$$\text{pH}_i = \frac{\text{pK}_a + \text{pK}_r}{2}$$

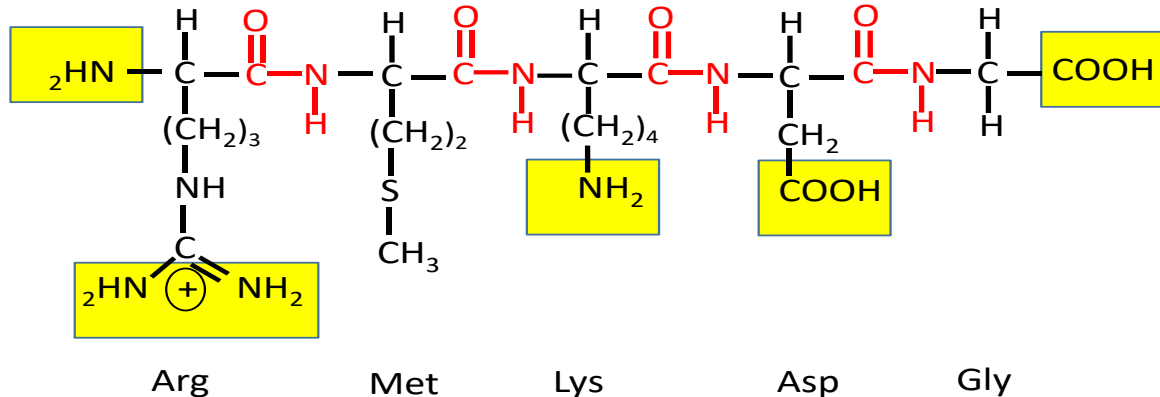
Properties of amino acids

Amphoteric properties

Ionisation of a peptide

Example : at $pH = 10$

To determine the ionization state of a peptide, we must first identify the ionizable groups.



This peptide has **5 ionizable** groups.

When studying the ionization of amino acids, the rule is:

- When $pH > pI \rightarrow$ the amino acid is negatively charged (AA^-)
- When $pH < pI \rightarrow$ the amino acid is positively charged (AA^+)
- When $pH = pI \rightarrow$ the amino acid is neutral (AA^0)

In the case of a peptide, we consider the pK_a , pK_b , and pK_r values.

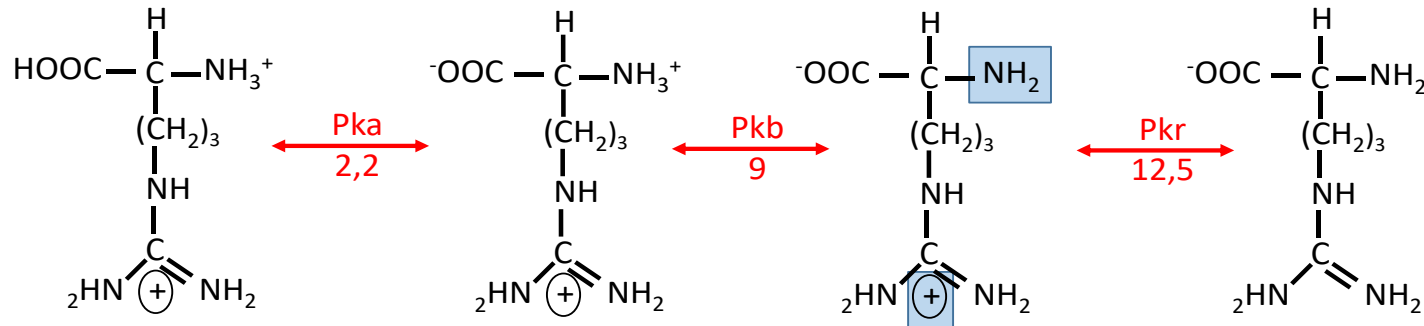
Properties of amino acids

Amphoteric properties

Ionisation of a peptide

We analyze each amino acid individually:

Arginine



- $\text{pH } 10 > \text{pK}_b$, the α -amino group is deprotonated $\rightarrow$ becomes NH_2 .
- $\text{pH } 10 < \text{pK}_r$: the guanidinium side chain is **still protonated** $\rightarrow$ remains positively charged (**Arg⁺**).

Properties of amino acids

Amphoteric properties

Ionisation of a peptide

We analyze each amino acid individually:

Methionine

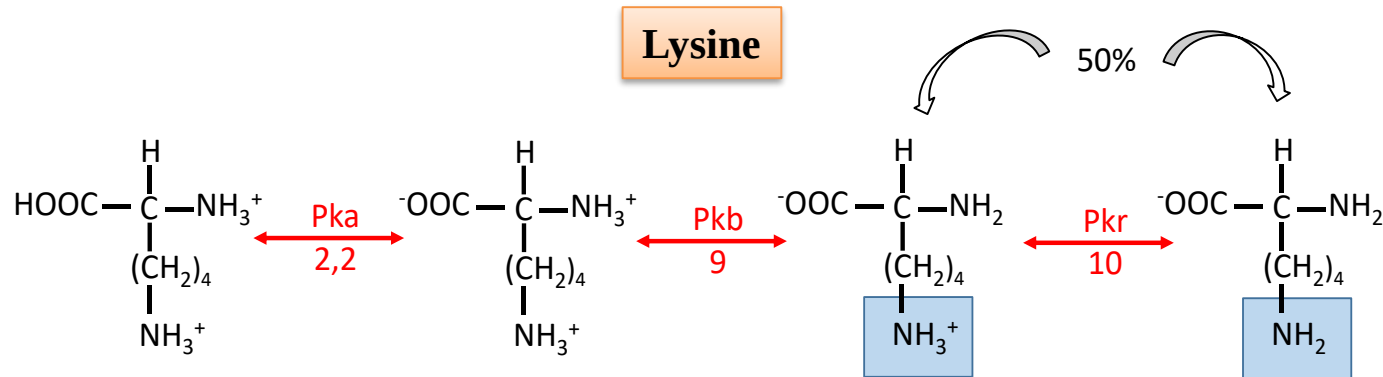
- The α -amino and α -carboxyl groups are involved in the peptide bond.
- The side chain is **not ionizable**.
- This amino acid is **neutral at any pH**.

Properties of amino acids

Amphoteric properties

Ionisation of a peptide

We analyze each amino acid individually:



- The α -amino and α -carboxyl groups are involved in the peptide bond.
- Only the **side chain** ($-\text{NH}_3^+$) is considered.
- **pH = pKr** $\rightarrow$ 50% of side chains are **protonated (+)**, 50% are **deprotonated (neutral)**.

So, we consider **both cases**:

- Neutral: $-\text{NH}_2$
- Positive: $-\text{NH}_3^+$

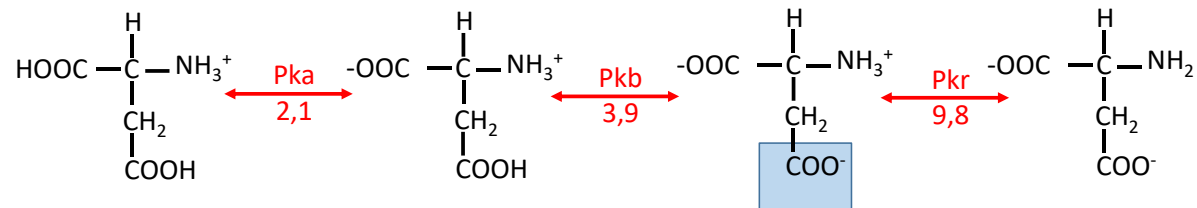
Properties of amino acids

Amphoteric properties

Ionisation of a peptide

We analyze each amino acid individually:

Aspartate



- $\text{pH } 10 > \text{pK}_r$: the carboxyl group is **deprotonated** $\rightarrow \text{COO}^-$

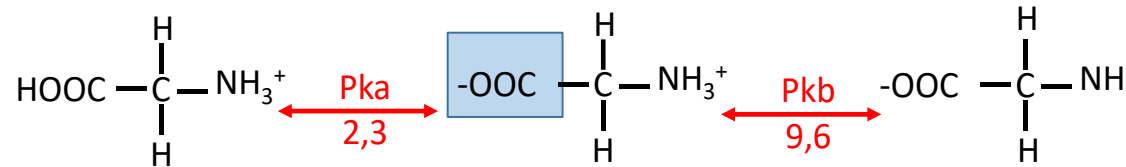
Properties of amino acids

Amphoteric properties

Ionisation of a peptide

We analyze each amino acid individually:

Glycine



- **pH 10 > pKa**, the group is **deprotonated** → **COO⁻**

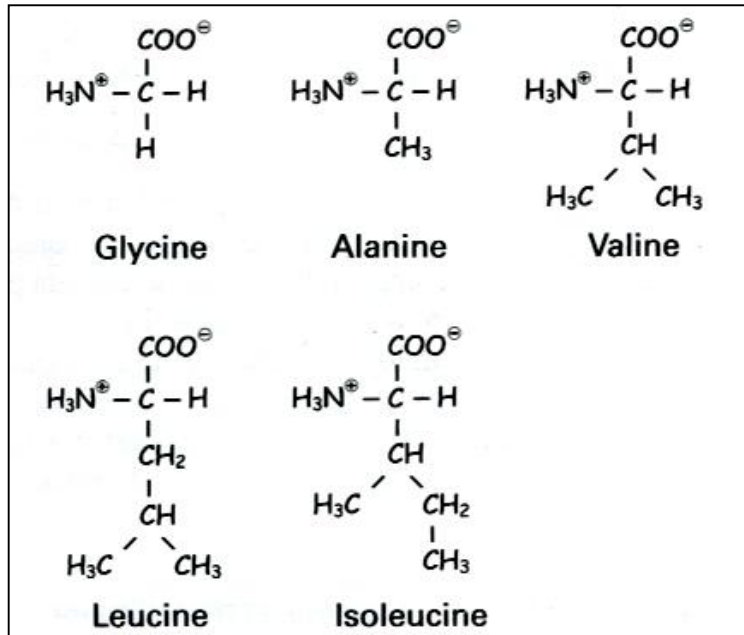
Final result at pH 10:



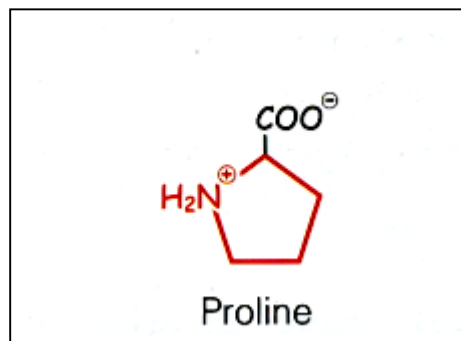
So, the **overall charge** is: **-1 (or 0 if Lys is positively charged)** But assuming the average state, the charge is **approximately -1**.

Nonpolar amino acids (hydrophobes)

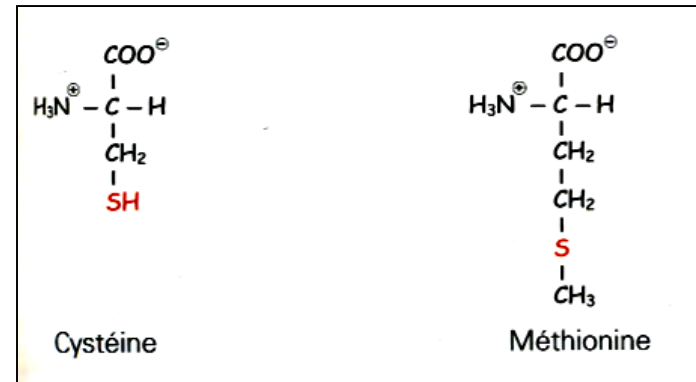
Aliphatic amino acids



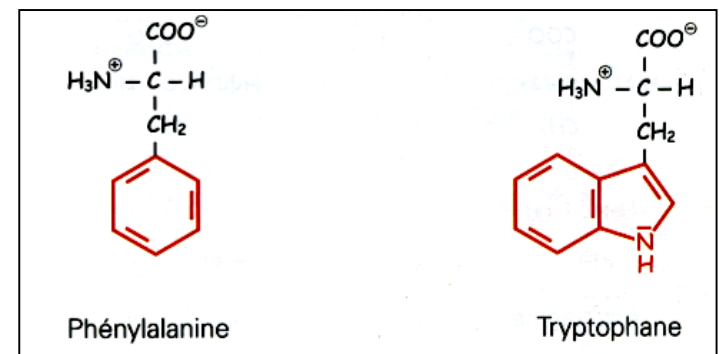
Aliphatic cyclic amino acid



Sulfur-containing amino acids



Aromatic amino acid

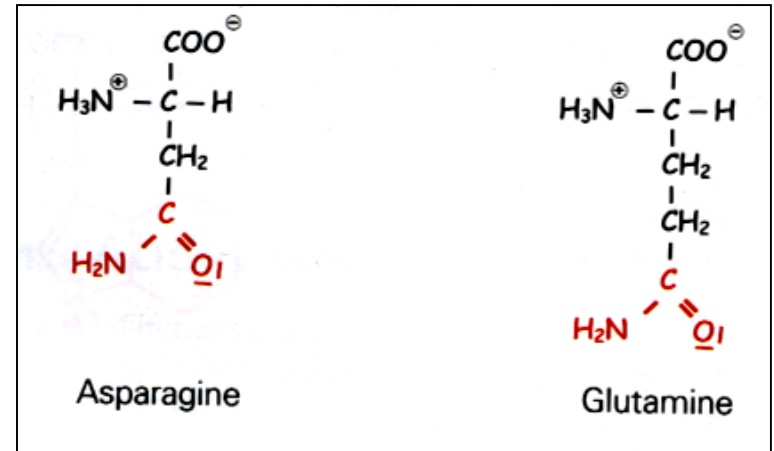
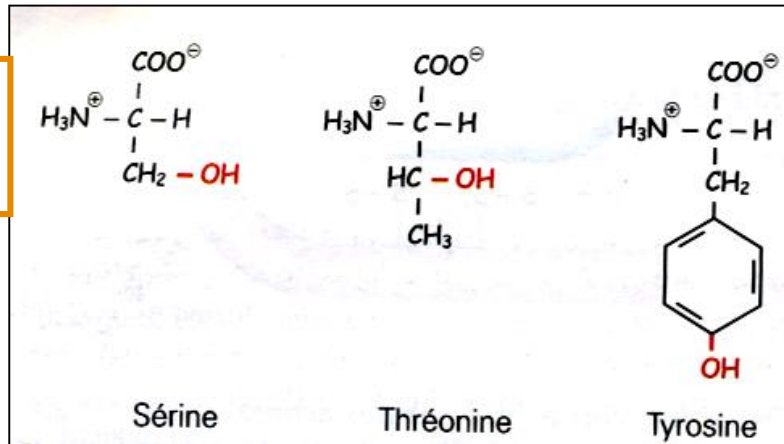


Polar amino acids (hydrophiles)

Hydroxyl amino acids

Amide amino acids

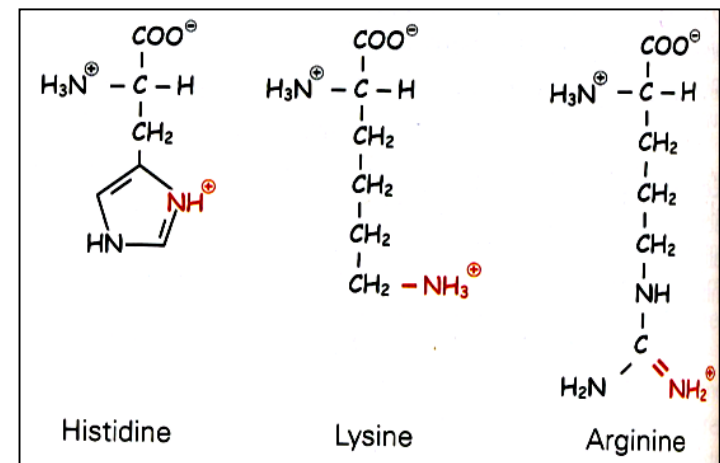
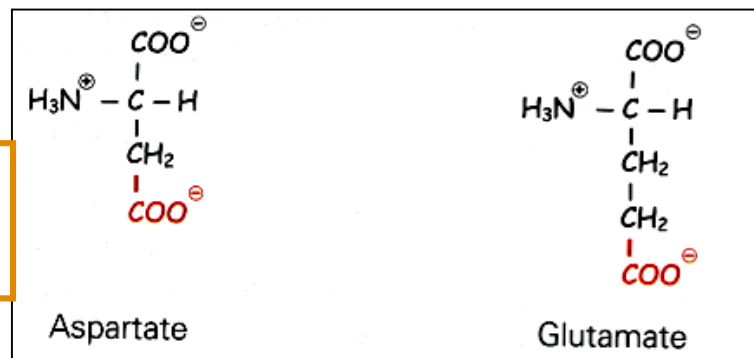
Uncharged
amino
acids



Dicarboxylic amino acids

Basic amino acids

Charged
amino
acids



Fractionation and dosage of amino acids

- ✓ Certain amino acids have specific reactions (coloration or spectrum) that allow for their quantification.
- ✓ Here are techniques for identifying amino acids:

Thin-Layer Electrophoresis (TLE)

Ion exchange column chromatography

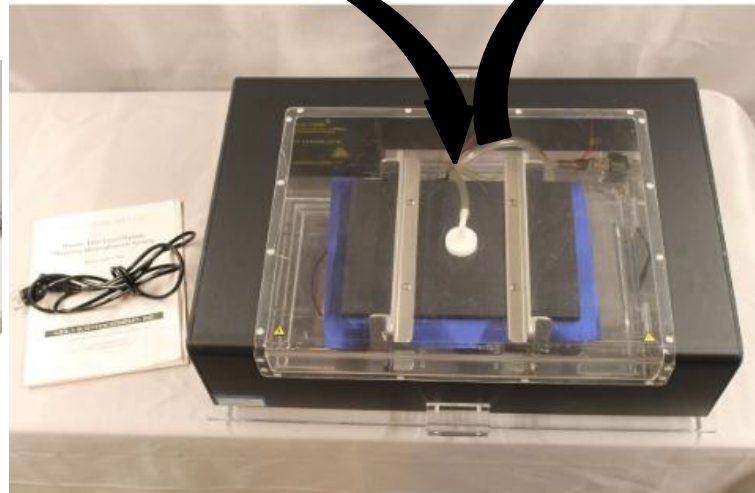
Separation and quantification of amino acids

Thin layer Electrophoresis (TLE)

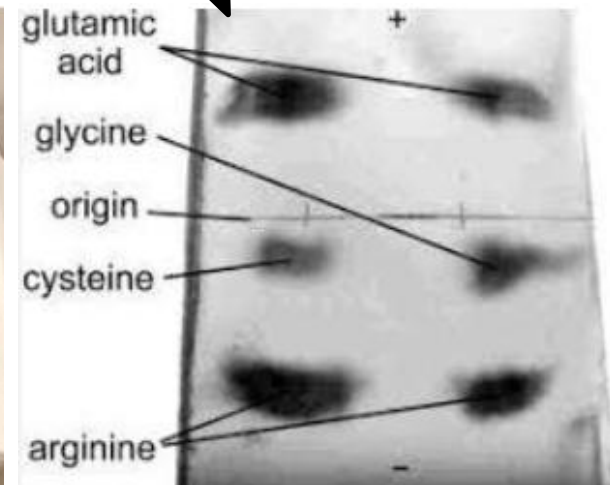
Allows for the separation of amino acids on a **stationary phase (membrane)** using an electric field based on their charges, with **different solvents (mobile phase)** and varying pH levels.



Cellulose layer



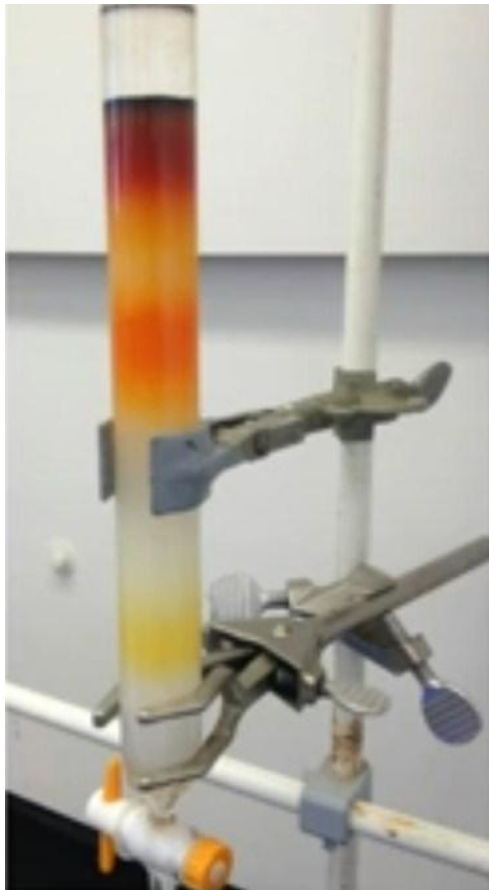
Electrophoresis chamber



Amino acids visualization

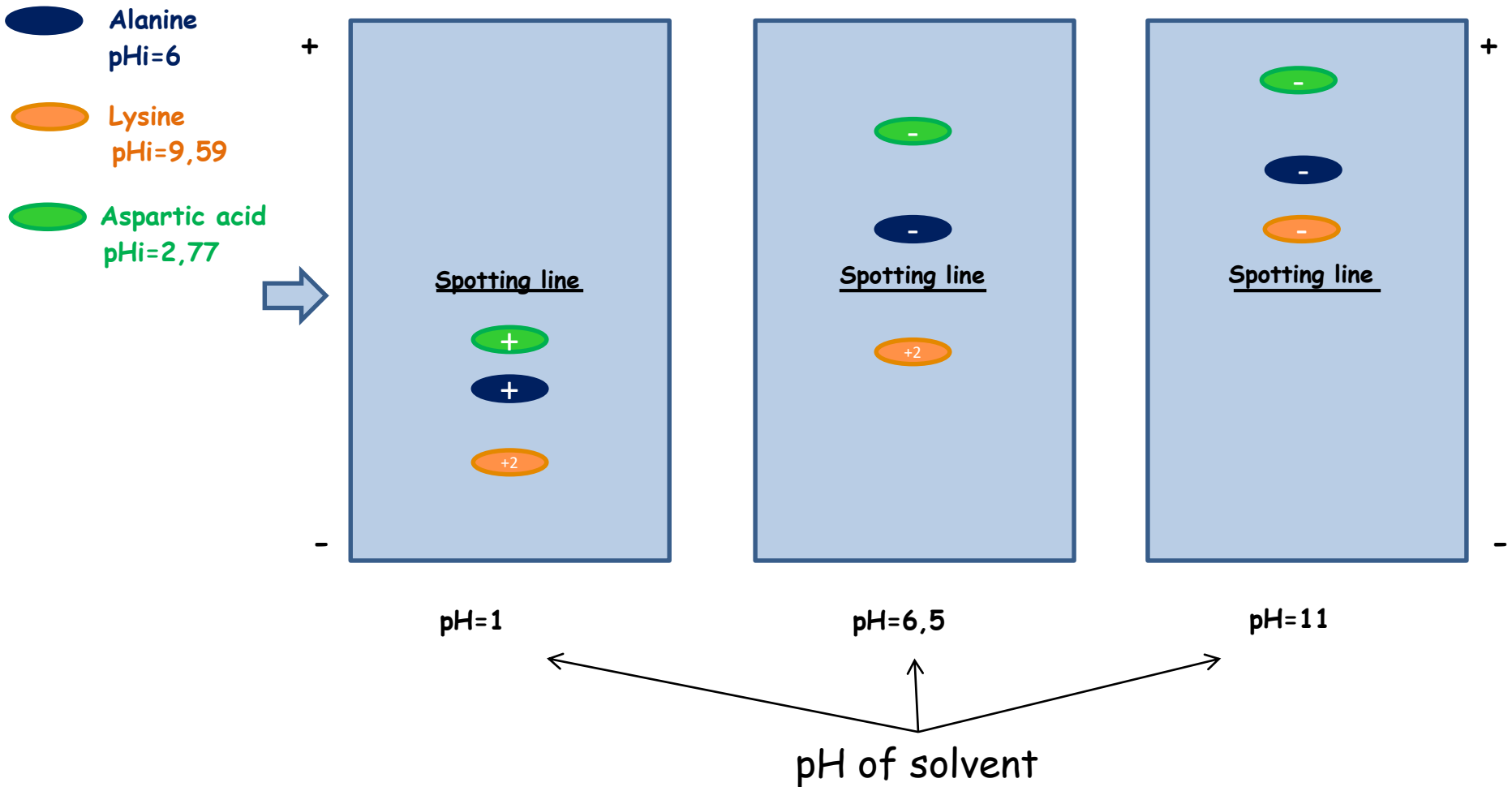
Separation and quantification of amino acids

Ion exchange chromatography (chromatographie sur colonne de résine échangeuse d'ions)

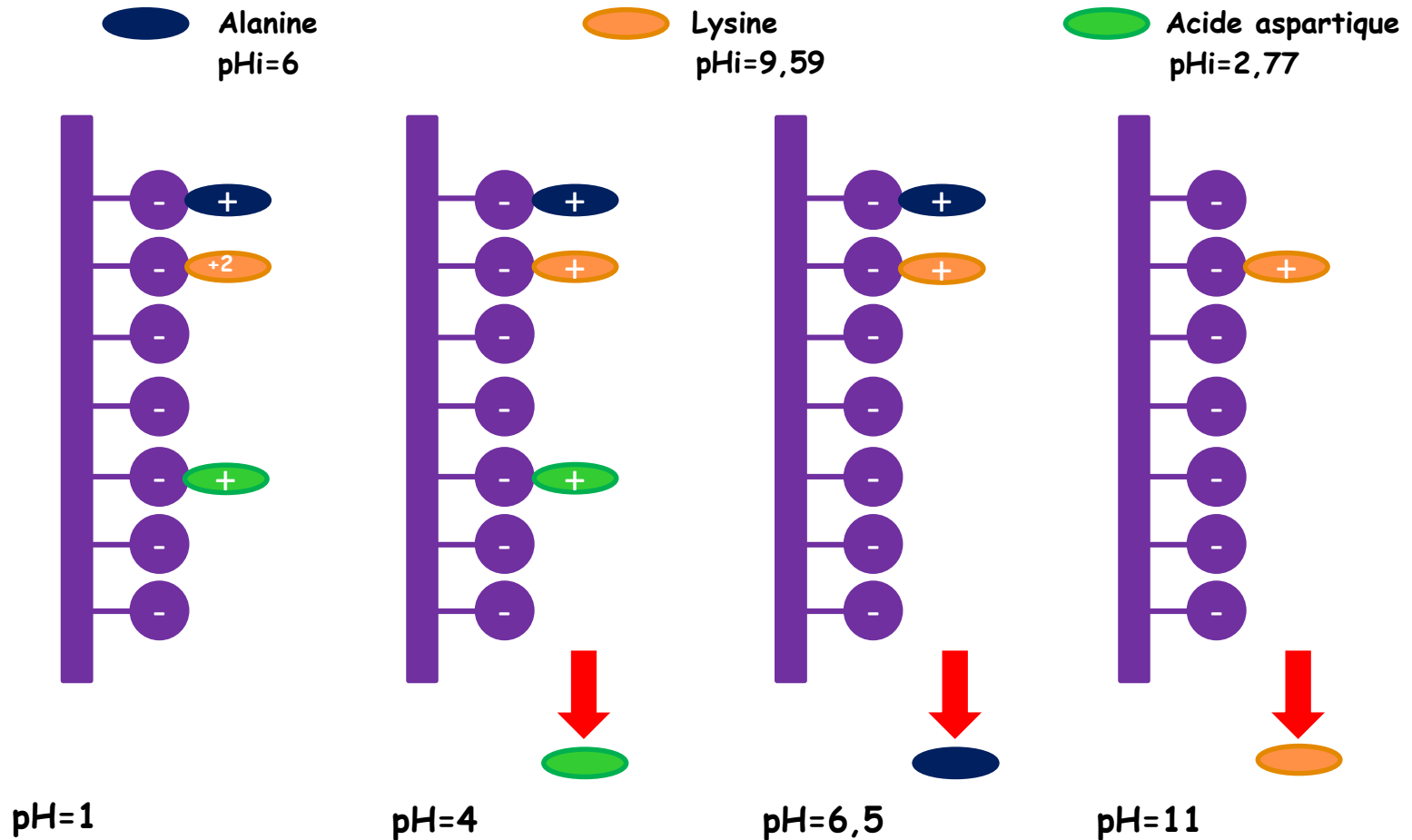


- The column contains a resin with **charged groups $\text{SO}_3\text{H}^- \text{Na}^+$** capable of exchanging ions with those in the solution.
At an acidic pH amino acids behave as bases (charge +). They displace some of the Na^+ ions, replacing them and remaining attached to the column via **electrostatic forces**.
- a **buffer with increasing pH** is passed through the column. When the pH of the eluent exceeds the pI of the amino acid, its net charge changes, causing it to detach and move through the column. This process is called **elution**.

Separation with thin layer chromatography (CCM)



Separation with ion exchange chromatography



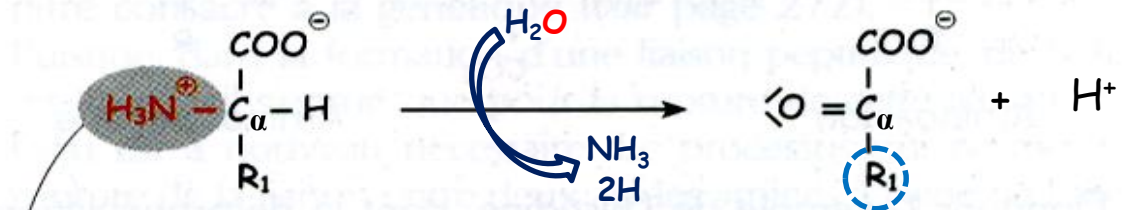
Reactions of amino acids

Desamination and transamination

Crucial for the degradation
of amino acids

Essential for amino acid
biosynthesis

Aminoacide oxydase

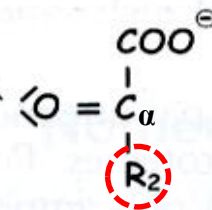


Non essential amino acid

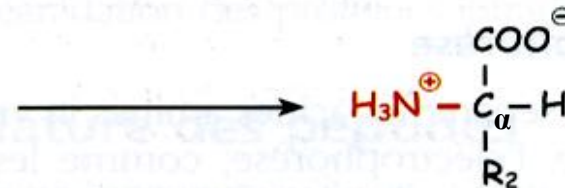
Alpha-keto acid

Transaminase

Amino group (NH_2) is
transferred from one
amino acid to an alpha-
keto acid



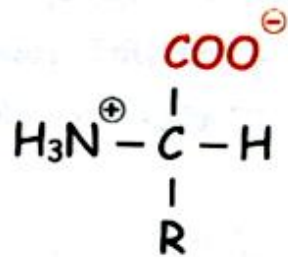
Alpha-keto acid



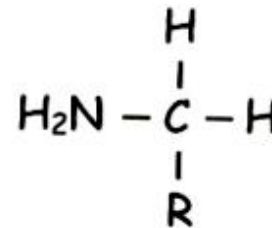
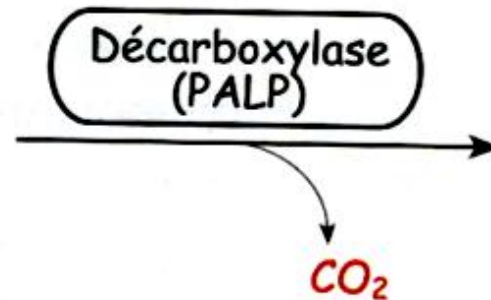
Essential amino acid

Reactions of amino acids

Decarboxylation



Amino acid



Biogenic amine

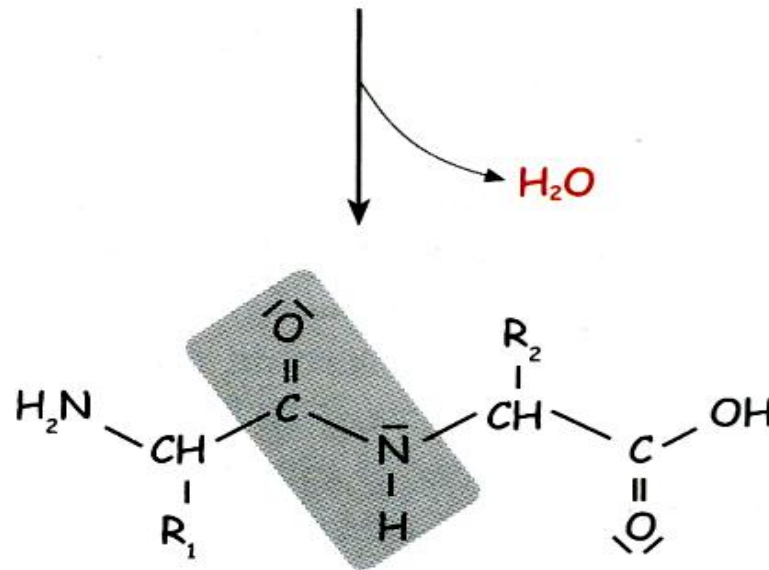
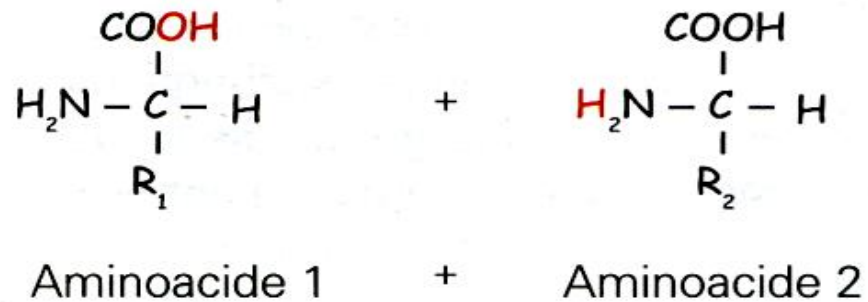
Ex : histidine → histamine acts
as a chemical mediator
(neurotransmitter)

Characteristic constants of amino acids

Name	Code	Letter code	pKa of COOH	pKb of NH3	pKr of Lateral chain	MW
Alanine	ALA	A	2,3	9,7		89,09
Arginine	ARG	R	2,2	9	12,5	174,2
Asparagine	ASN	N	2	8,8		132,12
Acide aspartique	ASP	D	2,1	9,8	3,9	133,1
Cystéine	CYS	C	1,8	10,8	8,3	121,15
Glutamine	GLN	Q	2,2	9,1		146,15
Acide glutamique	GLU	E	2,2	9,7	4,2	147,13
Glycine	GLY	G	2,3	9,6		75,07
Histidine	HIS	H	1,8	9,2	6	155,16
Isoleucine	ILE	I	2,4	9,7		131,17
Leucine	LEU	L	2,4	9,6		131,17
Lysine	LYS	K	2,2	9	10	146,19
Méthionine	MET	M	2,3	9,2		149,21
Phénylalanine	PHE	F	1,8	9,1		165,19
Proline	PRO	P	2	10,6		115,13
Sérine	SER	S	2,2	9,2		105,09
Thréonine	THR	T	2,6	10,4		119,12
Tryptophane	TRP	W	2,4	9,4		204,23
Tyrosine	TYR	Y	2,2	9,1	10,1	181,19
Valine	VAL	V	2,3	9,6		117,15

Protreins

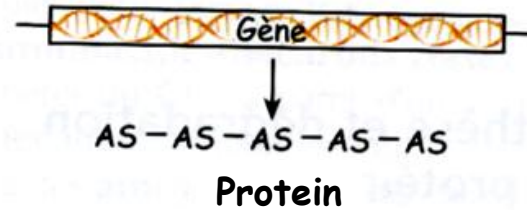
The peptide bond



Peptide bond

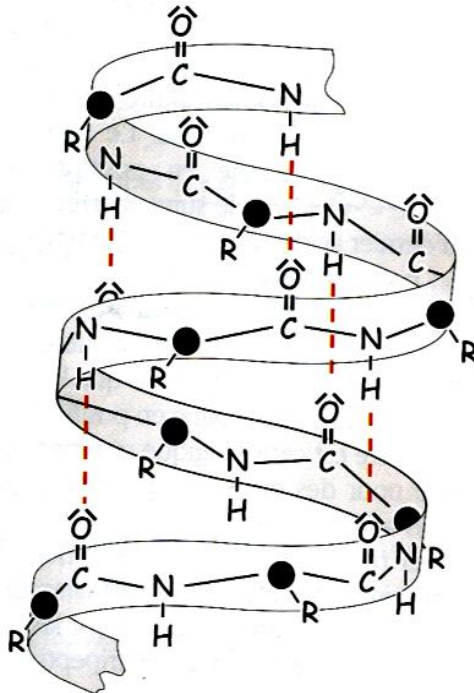
Spatial structure of proteins

Primary structure

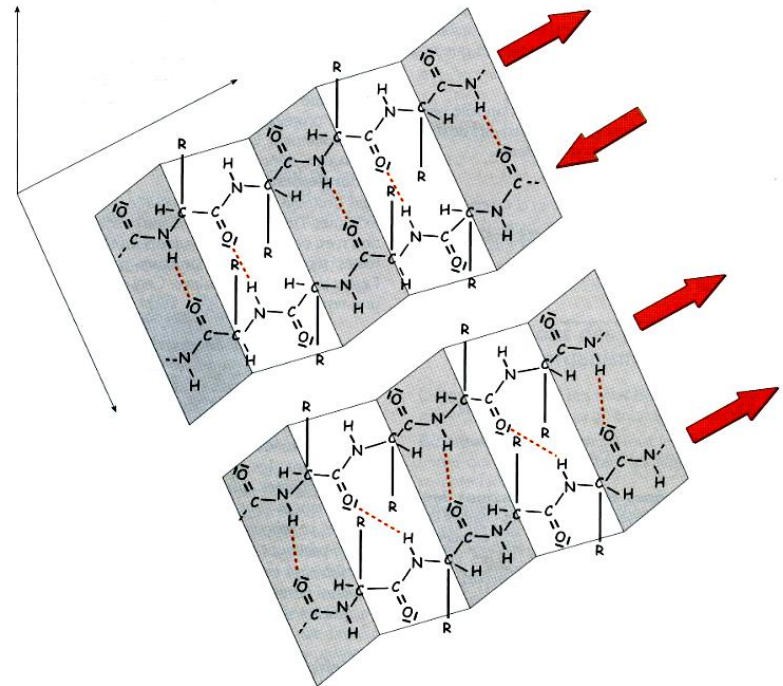


Secondary structure

- α helix

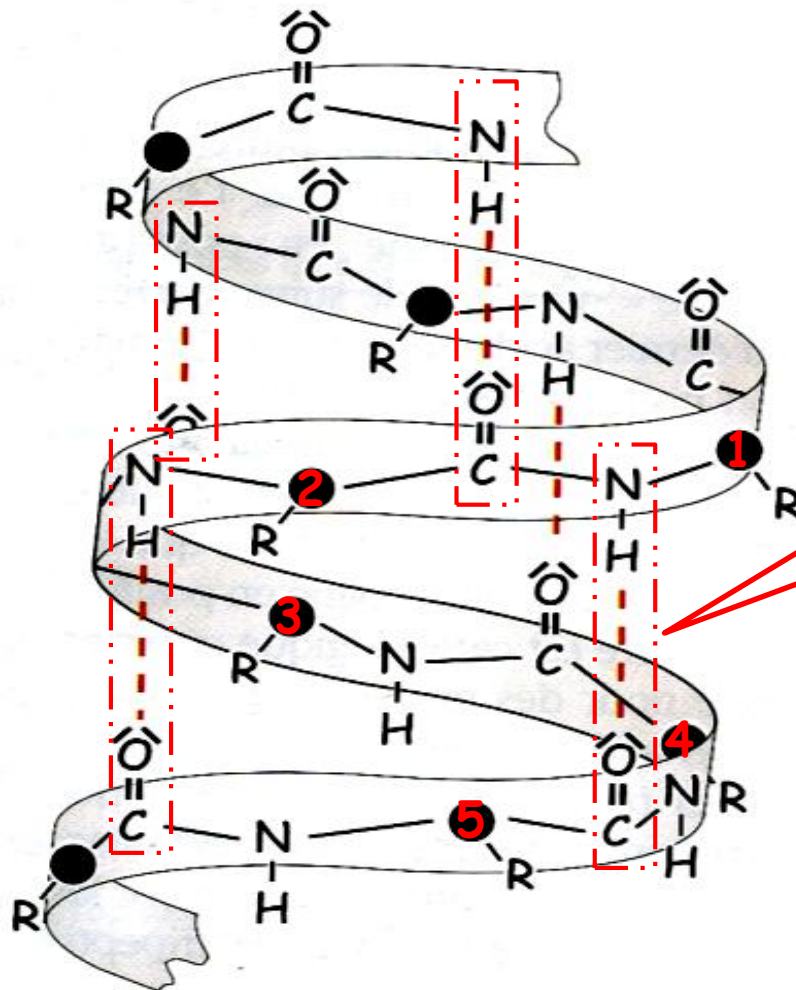


- β sheet



Spatial structure of proteins

Structure of α helix

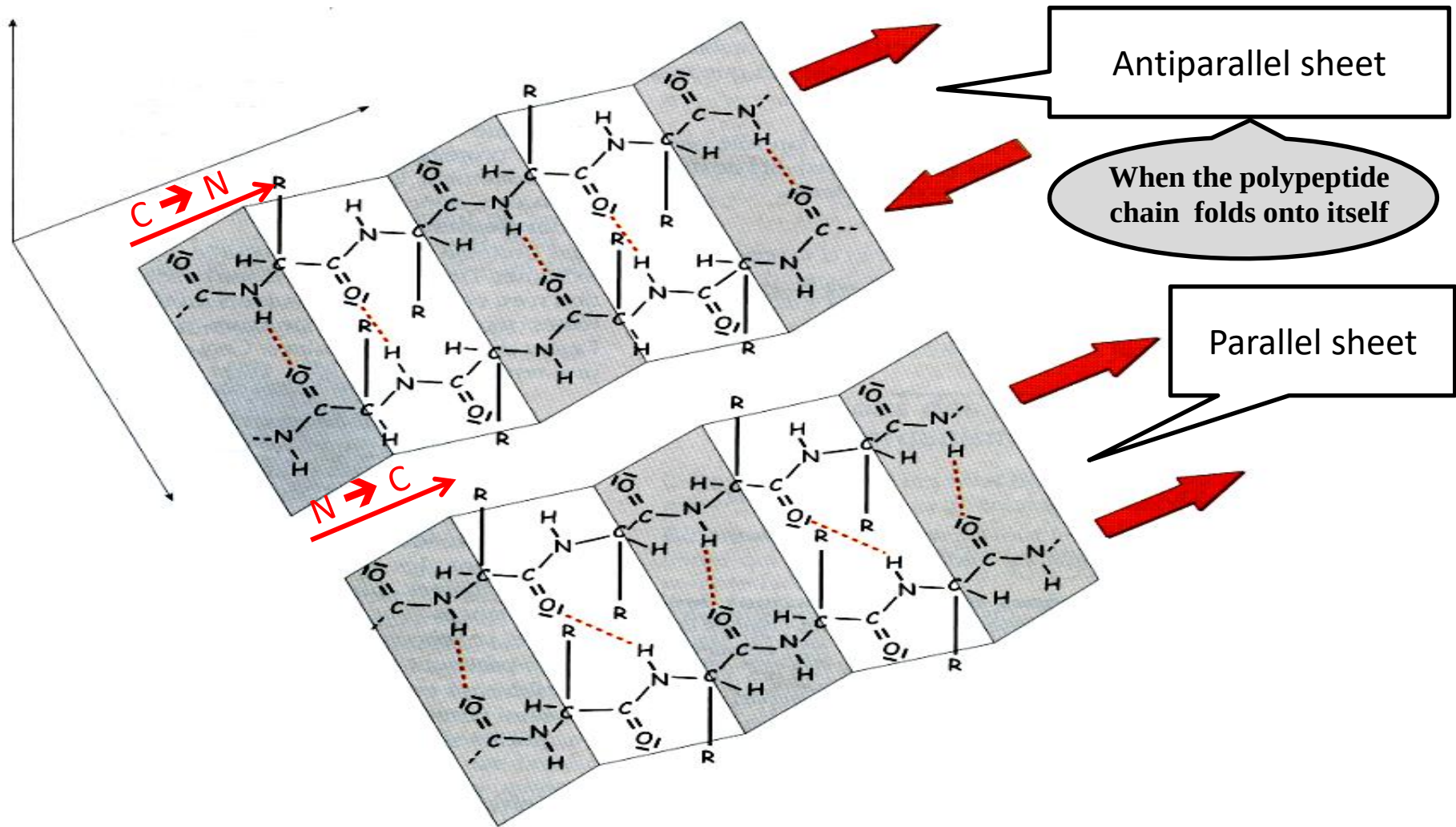


Contain a minimum of 5 residues and a maximum of 40

Hydrogen bonds form between amino acids n and $n+4$

Spatial structure of proteins

Structure of β sheet

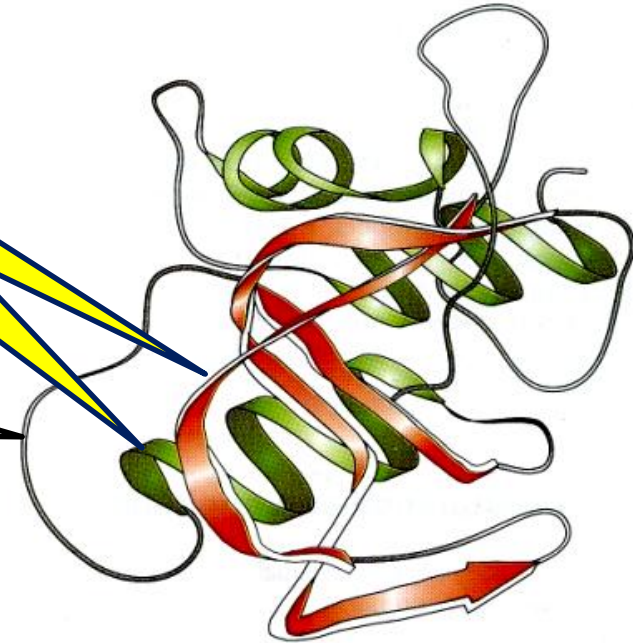


Spatial structure of proteins

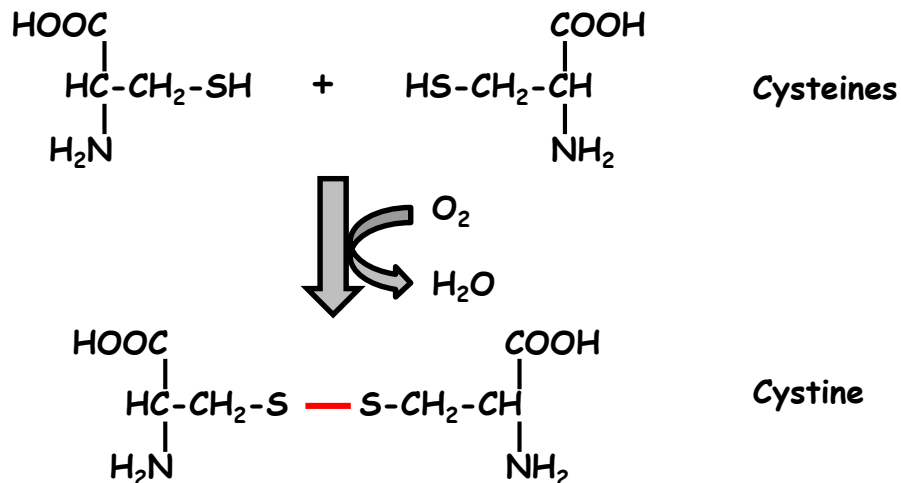
Tertiary structure

A helix + β sheet

Stable structure due to disulfide bridges (covalent bonds) formed between the thiol groups of sulfur-containing amino acids, particularly Cys
 $R-SH + SH-R' \rightarrow R-S-S-R' + 2H^+ + 2e^-$



Oxidation of cysteine and the formation of cystine

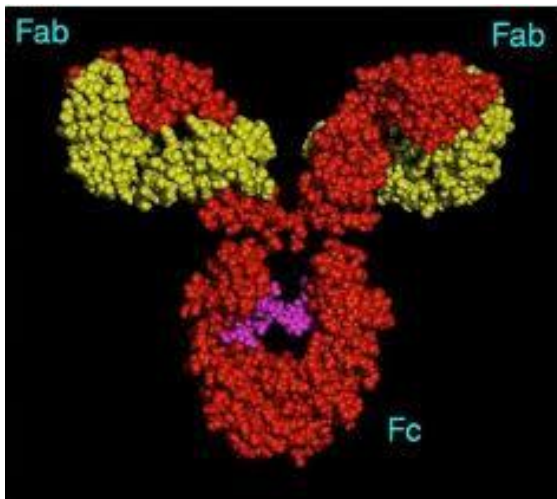


Spatial structure of proteins

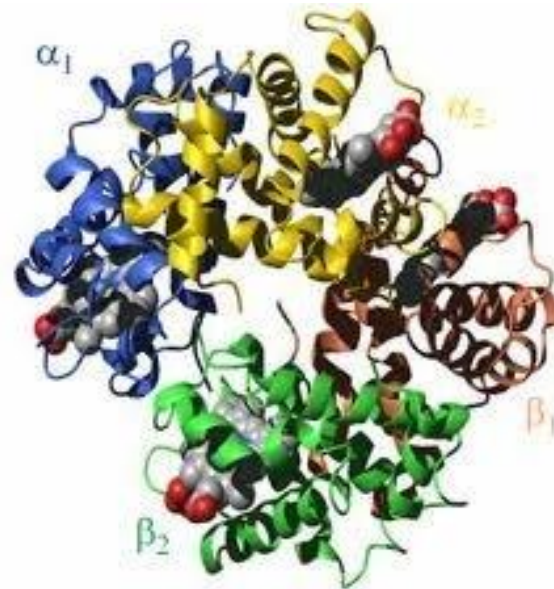
Quaternary structure

Assembly of tertiary structures
→ Protein complex

Antibody



Hemoglobin



Denaturation of protein

The denaturation of protein requires the breaking of:

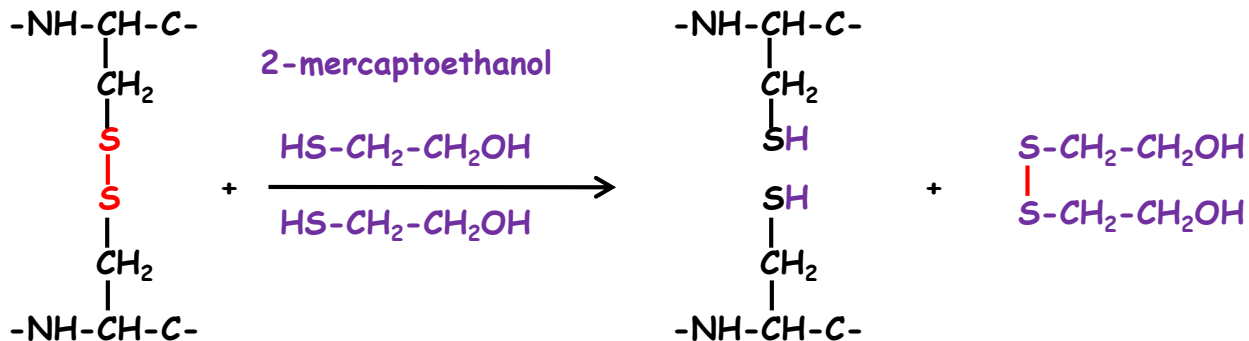
→ **Disulfide bonds**: through reduction (unfolding of the tertiary structure)

→ **Hydrogen bonds**: through heat (unfolding of the secondary structure)



The denatured primary structure migrates in electrophoresis through the pores of a polyacrylamide gel.

Reduction of disulfide bonds



Determination of the amino acid composition of a polypeptide chain

- Fractionated polypeptide chains are separated by **electrophoresis** or **chromatography**.
- For each isolated chain, the amino acid composition is determined.
 - Determination of the N-terminal amino acid
 - Determination of the C-terminal amino acid

Determination of the amino acid composition of a polypeptide chain

Determination of the N-terminal amino acid

Main reagents:

- **FDNB** (1-fluoro 2,4-dinitrobenzene) (1945): **Sanger reagent**
- **PTC** (phenylisothiocyanate): **Edman reagent**
- **Chlorure de dansyl** (acide chloride 1-dimethylaminonaphtalene-5-sulfonique)

Enzymatic method :

- **Aminopeptidase**: acts only on the last peptid bond.

Determination of the C-terminal amino acid

Enzymatique method:

- **Carboxypeptidase**: acts only on the last peptid bond.

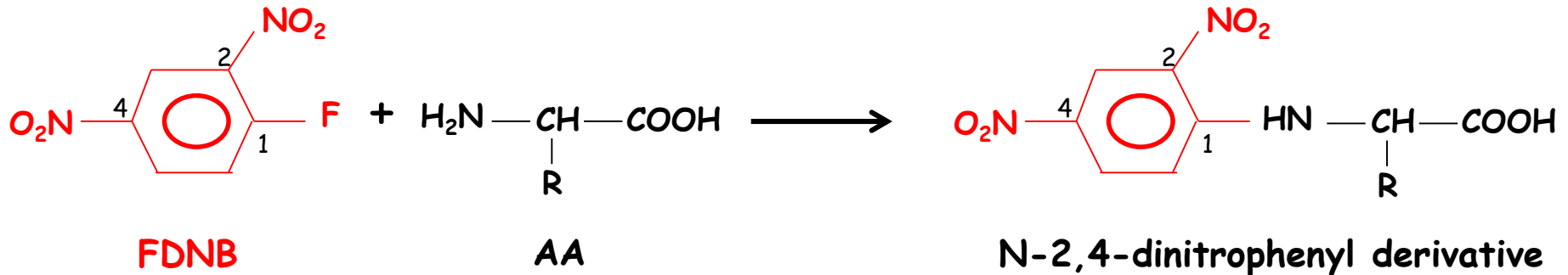
Determination of the amino acid composition of a polypeptide chain

Determination of the N-terminal amino acid

FDNB (1-fluoro 2,4-dinitrobenzene) (1945): Sanger reagent

Non-recurrent method

This method identifies N-terminal amino acid, while the rest of the chain is **degraded** and thus **not reusable**.

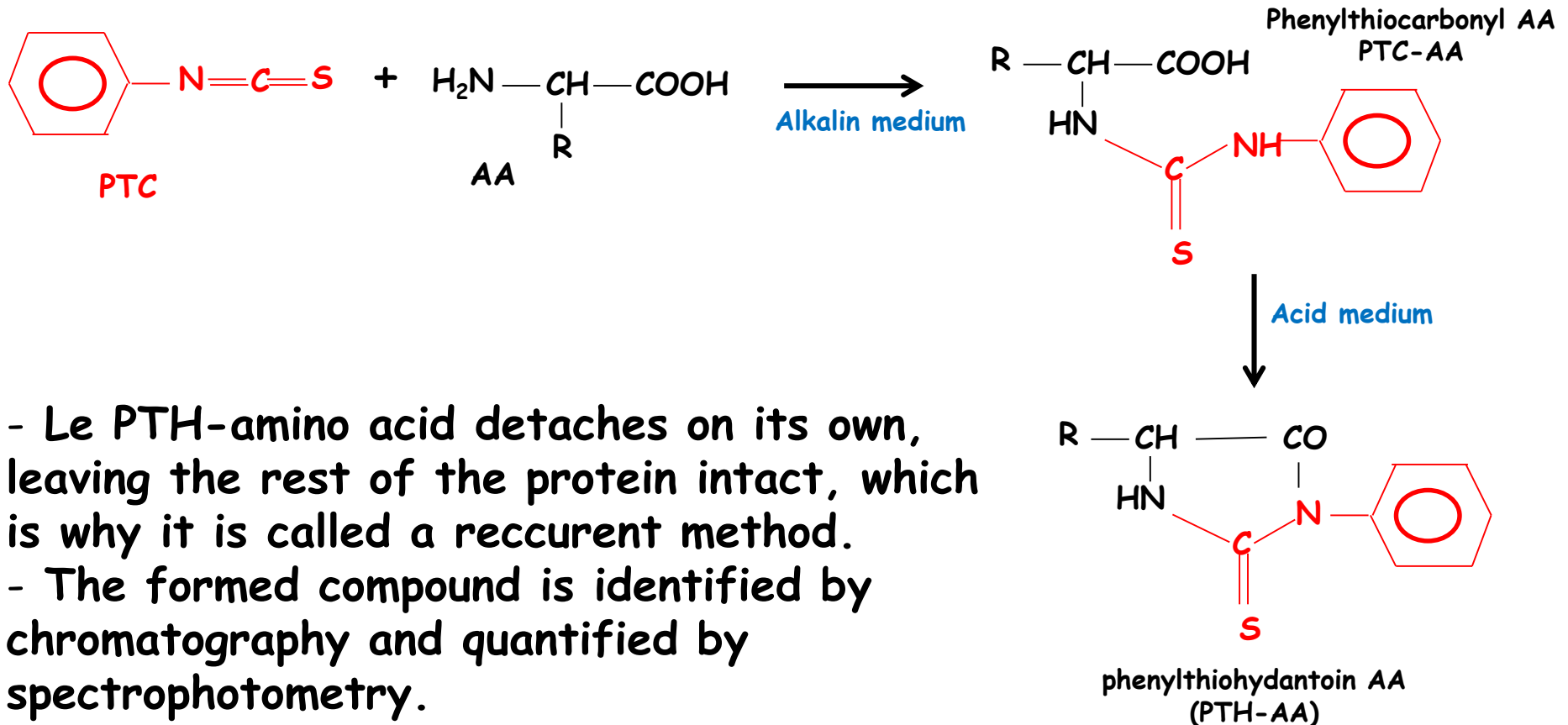


The formed compound is identified by chromatography and quantified by spectrophotometry.

Determination of the amino acid composition of a polypeptide chain

Determination of the N-terminal amino acid

PTC (phenylisothiocyanate): Edman reagent
Recurrent method

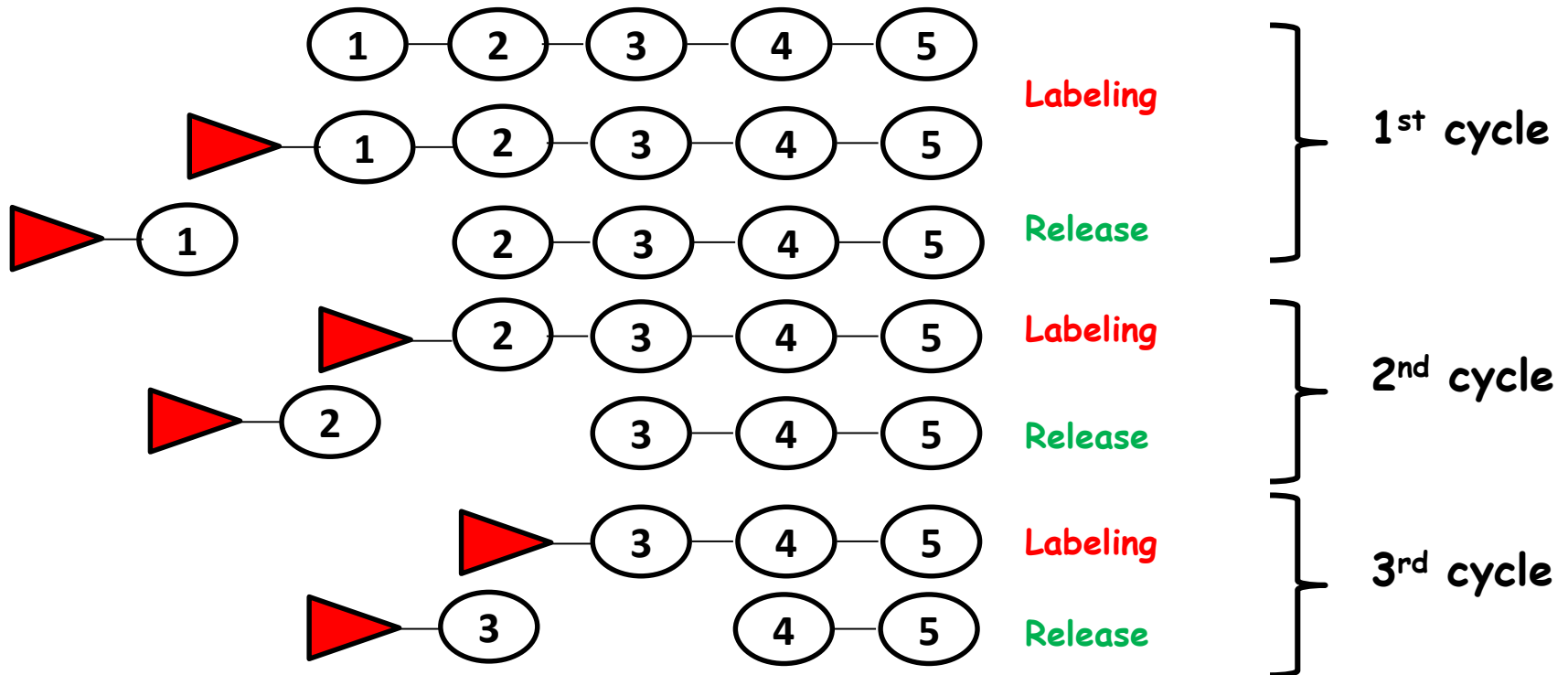


Determination of the amino acid composition of a polypeptide chain

Determination of the N-terminal amino acid

PTC (phenylisothiocyanate): Edman reagent
Recurrent method

Mechanism



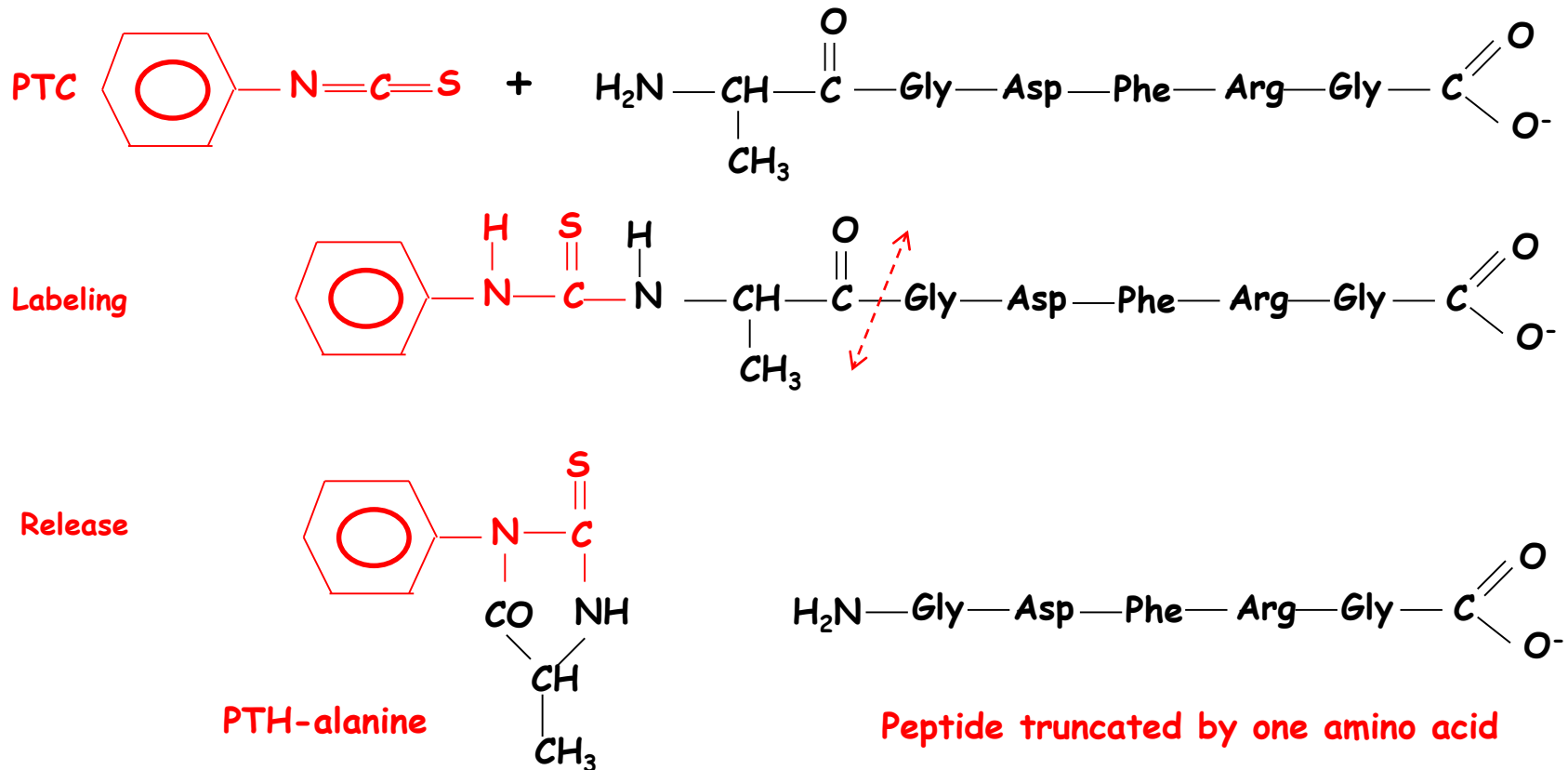
Determination of the amino acid composition of a polypeptide chain

Determination of the N-terminal amino acid

PTC (phenylisothiocyanate): Edman reagent

Recurrent method

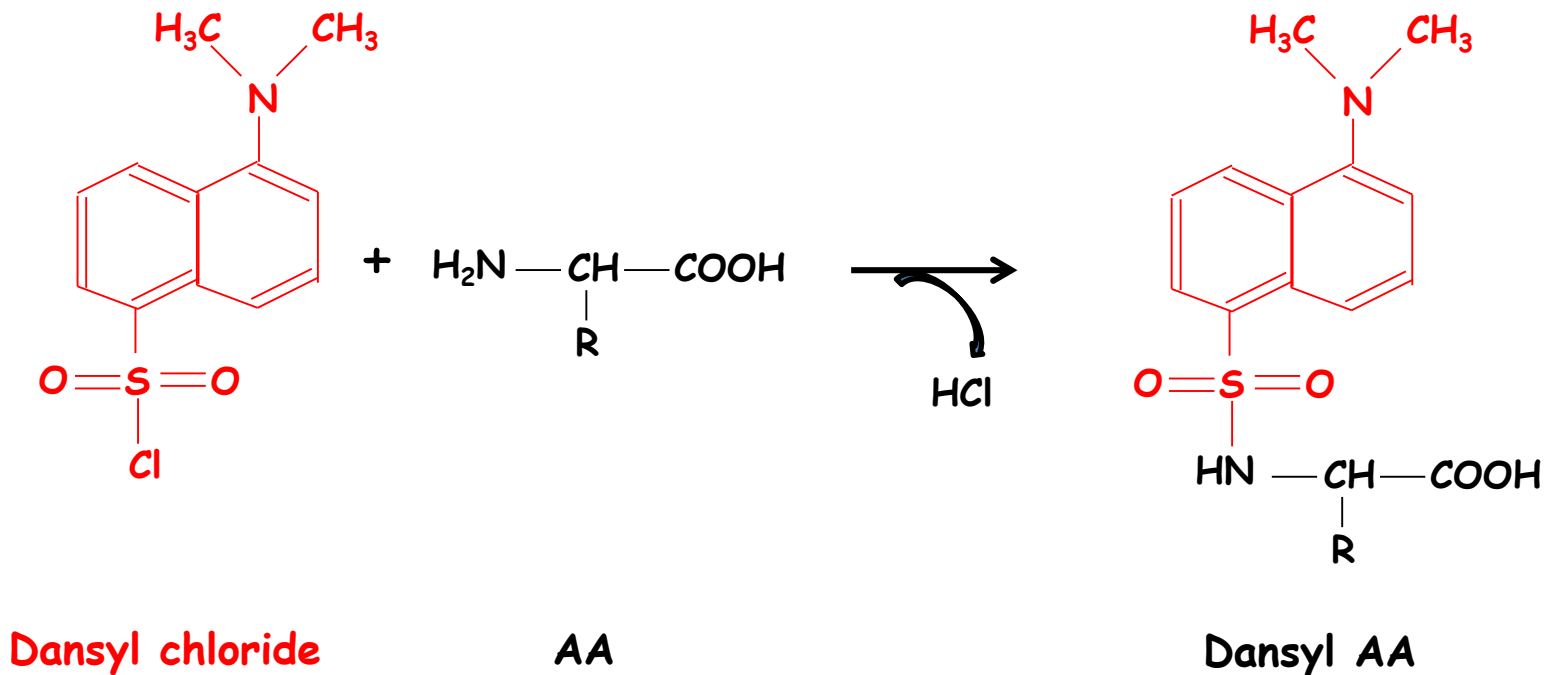
Mechanism



Determination of the amino acid composition of a polypeptide chain

Determination of the N-terminal amino acid

Dansyl chloride (acide 1-diméthylaminonaphtalène-5-sulfonique chloride)



The Dansyl-amino acid is highly fluorescent. It is separated by Thin layer chromatography (TLC) and detected by fluorescence.

Determination of the amino acid composition of a polypeptide chain

Partial hydrolysis

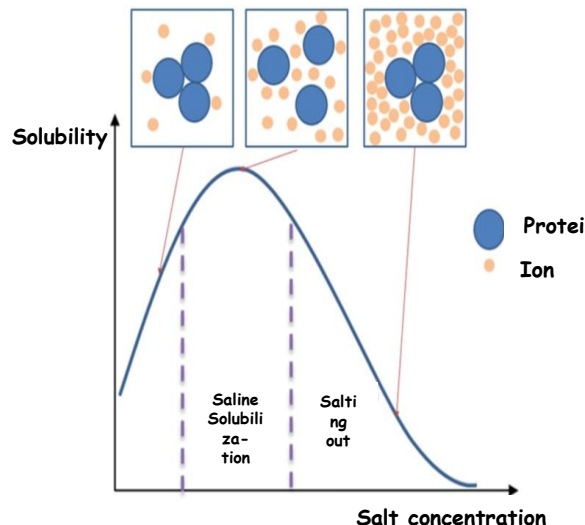
- Obtaining peptides either by chemical agents or by enzymes.
- Enzymatic proteolysis using **endopeptidases**:
 - **The trypsin**: cuts the CO(O) of the **lysine** and **arginine**.
 - **The chymotrypsin**: cuts the CO(O) of **aromatic amino acids** (**phenylalanine**, **tryptophane**) and **leucine** (which are nonpolar amino acids).

Properties of proteins

Solubility

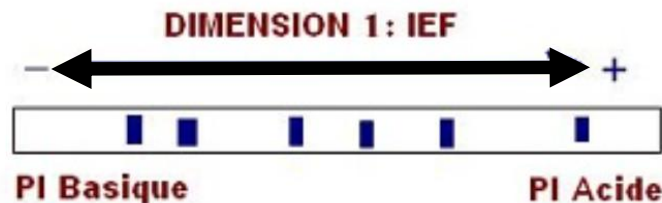
Depends on ionic strength and pH.

- At low ionic strength, monovalent ions promote solubilization by stabilizing the charged groups of proteins.
- At higher strength, ions tend to precipitate proteins. There is competition between the ions and proteins for water molecules, a phenomenon known as « salting out ».



Electrolytic properties

- In a basic medium, proteins behave as anions and migrate toward the anode in an electric field.
- In an acid medium, proteins behave as cations and migrate toward the cathode (-).
- At pI, the net charge is zero, and proteins do not migrate.



Antigenic properties

An antigen is a substance capable of provoking the synthesis of specific antibodies when injected into a living organism.

- The part of the protein involved in the formation of the complex with the antibody is called the **antigenic determinant**.
- Used in protein assays, such as **affinity chromatography**.

