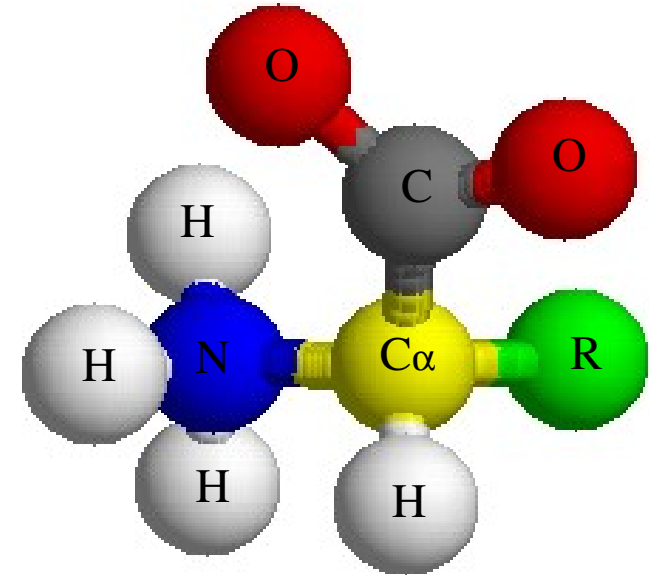


Chapter 4: Amino Acids

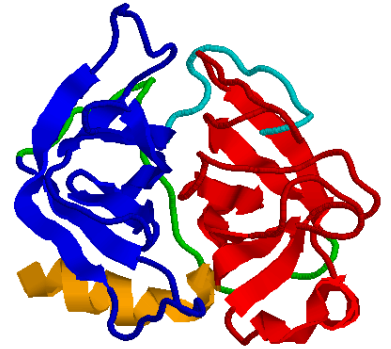


Voet & Voet: Pages 67 - 81

**Any introductory Biochemistry textbook will have
a chapter on amino acids and their properties**

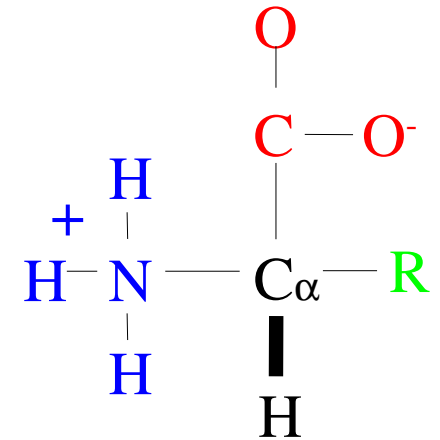
Proteins

- Most abundant macromolecules in cells
- Variable size and physical properties
 - Accounts for diversity of structure and biological function
- Final product of most genes
 - Means of expressing of genetic information
- Linear, heteropolymers of **amino acids**

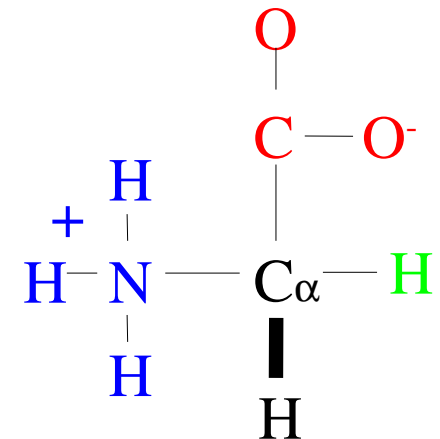


Amino Acids

- **Building blocks of proteins**
 - 20 common amino acids used by all organisms
- **All have an amino group and a carboxylate group covalently attached to a tetrahedral α carbon (C_α)**
- **Only differ at R group (side chain)**
 - Amino acids can be classified based upon the physiochemical properties of the R group
 - eg non polar, polar uncharged, polar charged



Generic amino acid
at neutral pH



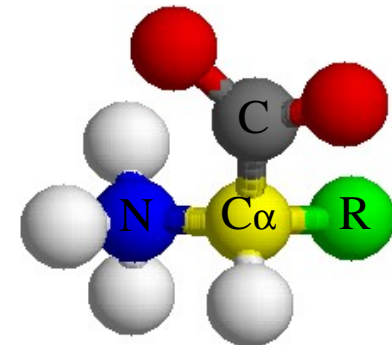
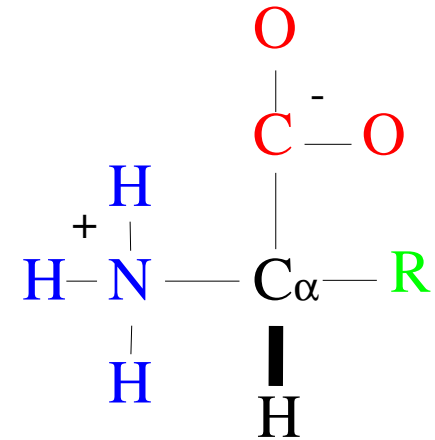
Glycine – a common
amino acid at neutral pH

Amino Acids are zwitterions

In the physiological pH range (5-8), the α -carboxyl and α -amino groups of amino acids are completely ionized

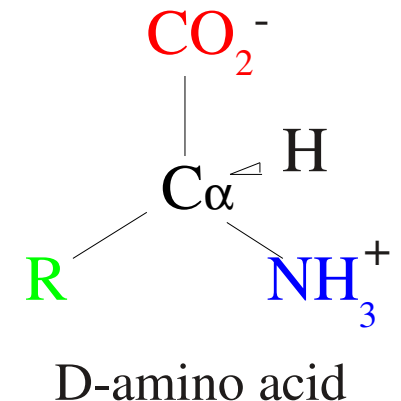
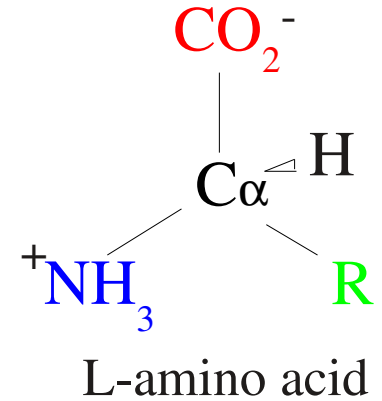
- Compounds with this property are referred to as **zwitterions** (or dipolar ions or ampholytes)
- Zwitterions can act as either an acid or a base

α -carboxyl groups have pK_a s near 2.2 while α -amino groups have a pK_a s near 9.4



Chirality

- Amino acids (except glycine) have a tetrahedral C_{α} bonded to four different chemical groups
 - Consequently, amino acids are optically active or chiral
- Common amino acids are all **L stereoisomers**
 - Amino acid configuration uses the D,L system while synthetic chemistry generally uses the R,S system
- **“CO-R-N”** mnemonic useful for distinguishing L and D stereoisomers
 - Looking down the H – C bond, CO-R-N spelled clockwise indicates the L stereoisomer



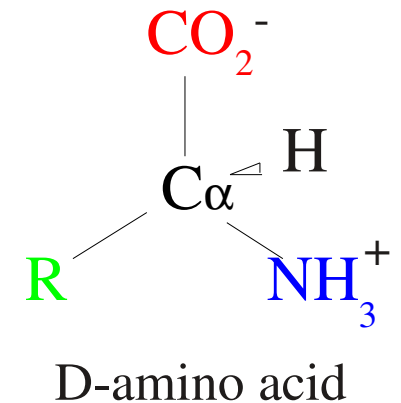
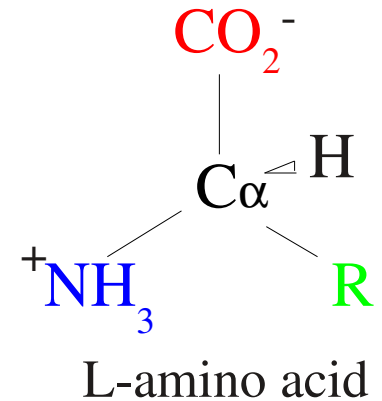
Why the L isomer?

No definitive answer

- D and L isomers have identical energies

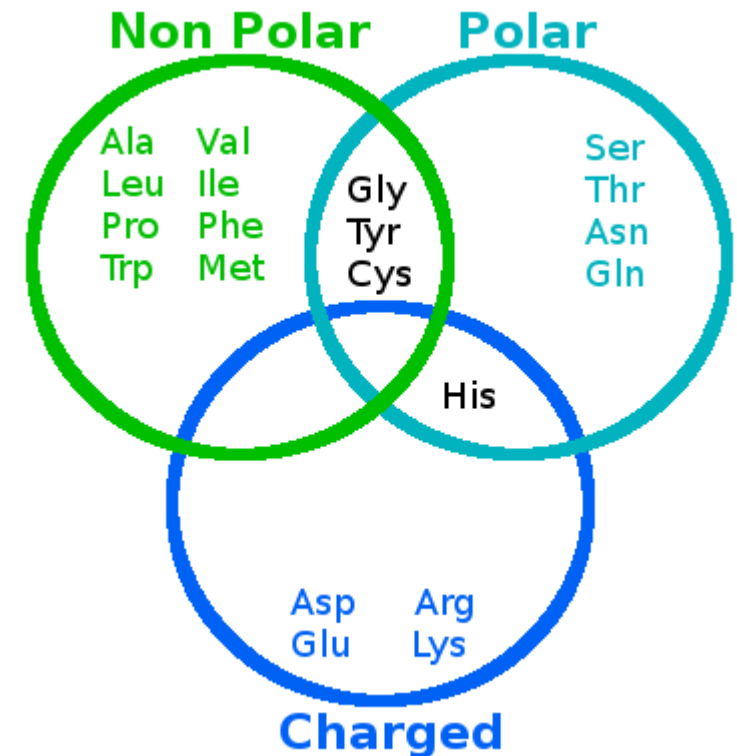
*Repetitive substructure in proteins (helices, sheets, turns) **require** all amino acids have the same configuration.*

- Apparently, living systems evolved from L amino acids based upon an initial random choice.



Classification of Amino Acids

- Amino Acids are classified or grouped according to the physiochemical properties of their R-group
 - Voet & Voet presents one common classification scheme based upon polarity
- **Polarity** is defined as *the magnitude of the dipole induced in the presence of an external electromagnetic field.*

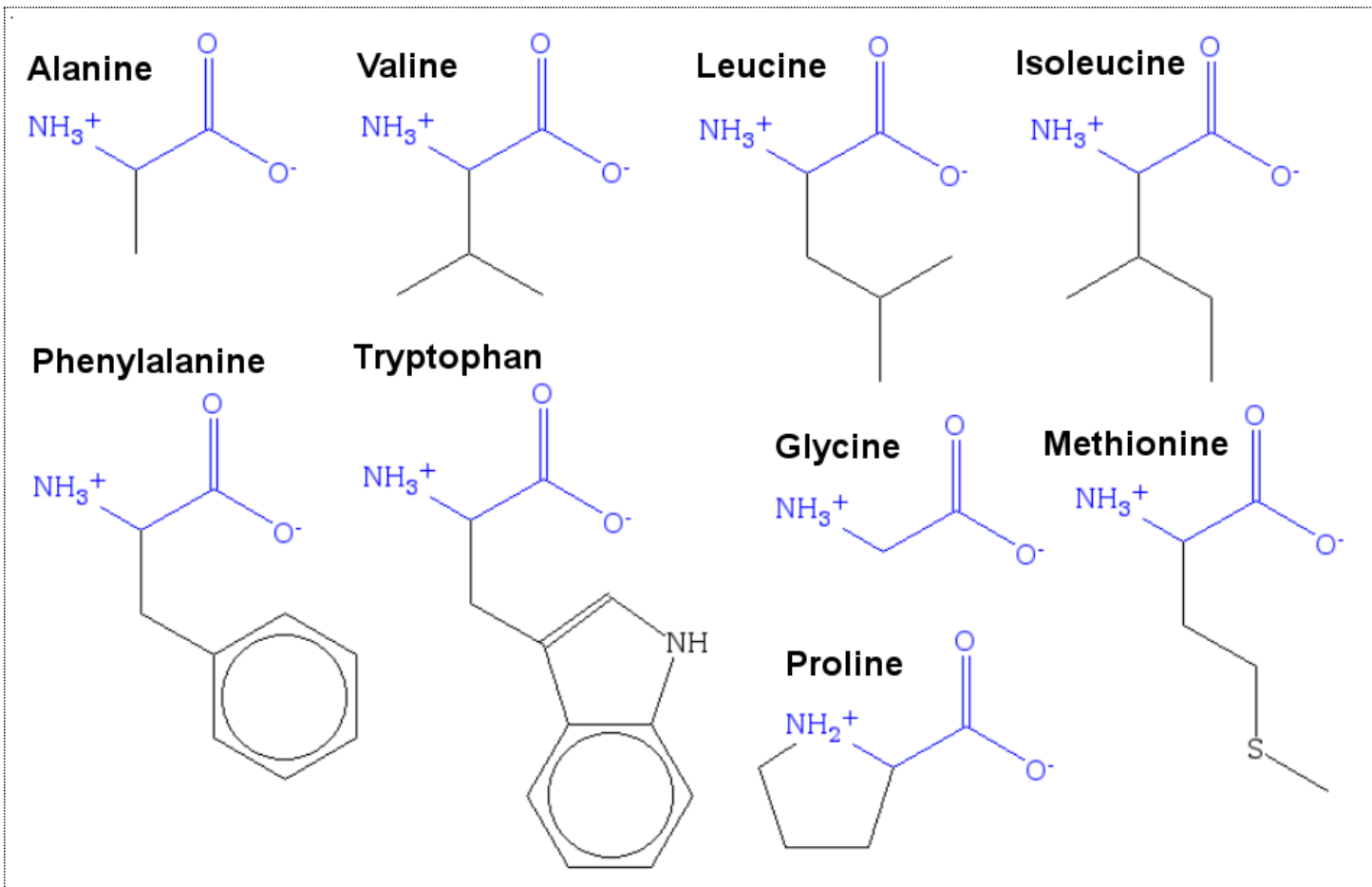


Amino acids grouped by Polarity (at neutral pH).

Classification schemes are an aid to recalling structure and properties of amino acids

Non Polar Amino Acids

R-group is a hydrocarbon (also includes Trp & Met)



Aromatic amino acids
absorb UV light

Proline R-group is
covalently bonded
to α -amine

Glycine can also be
classified as polar
uncharged

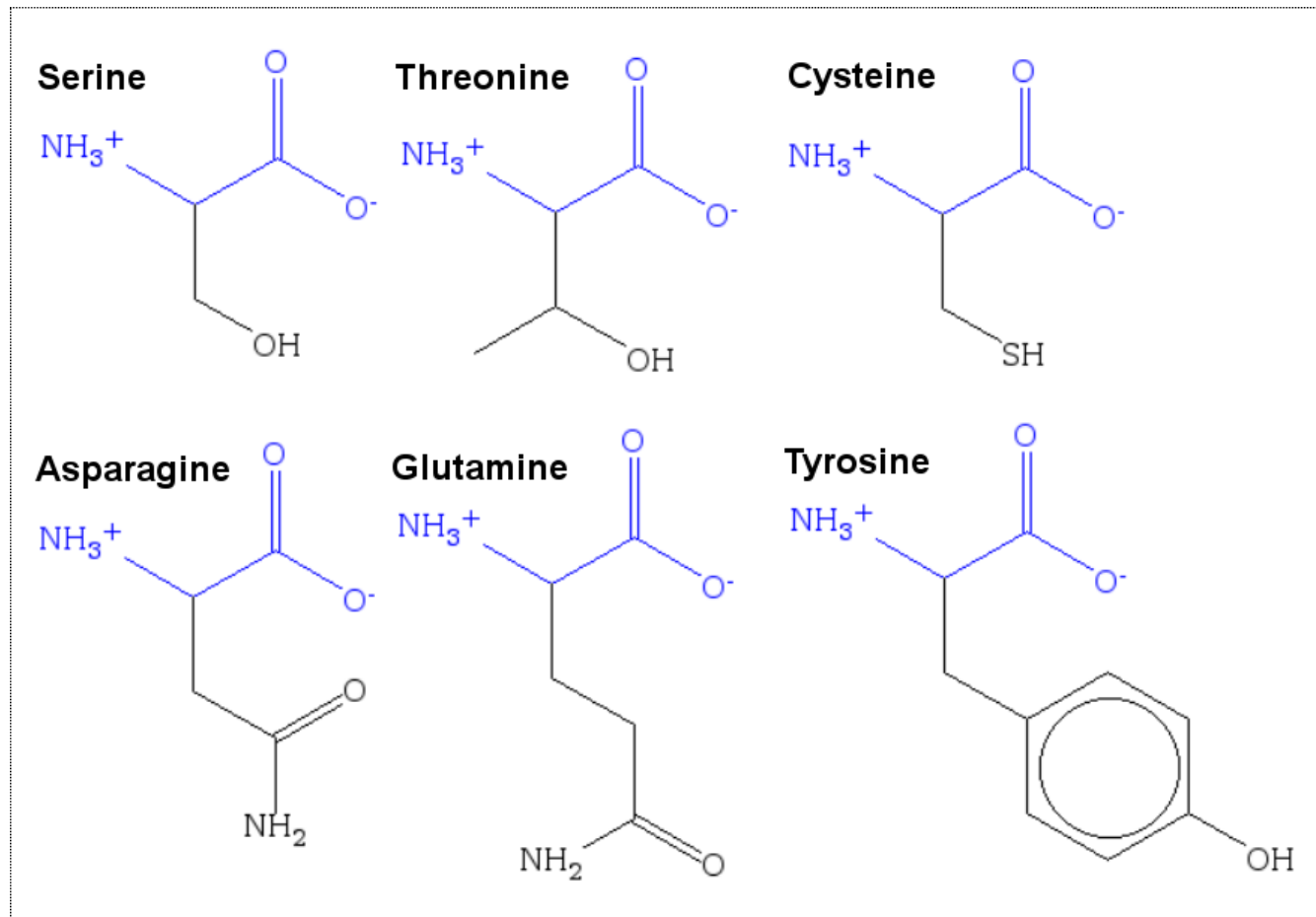
Polar (uncharged) Amino Acids

R-group contains polar (hydroxyl, thiol or carboxyamide) functional group and has neutral charge at pH 7

Ser, **Thr** and **Cys** are often involved in enzymatic reactions

Cys can form disulfide bridges (cystine)

Tyr and **Cys** can also be classified as non polar.

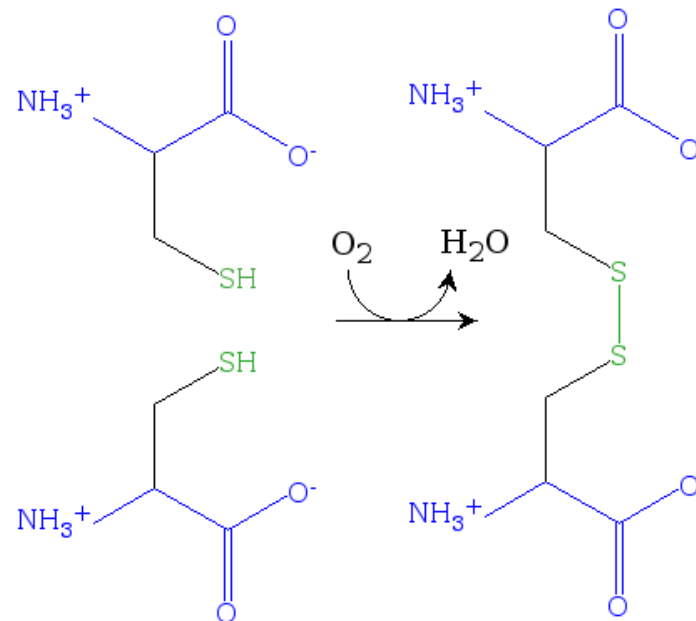


Cysteine

- Polar uncharged amino acid due to its thiol group
- Cysteines spontaneously form a covalent bond (**disulfide bridge**) between their thiol groups in the presence of oxidizing agents

Disulfide bonded cysteine residues are referred to as **cystine**.

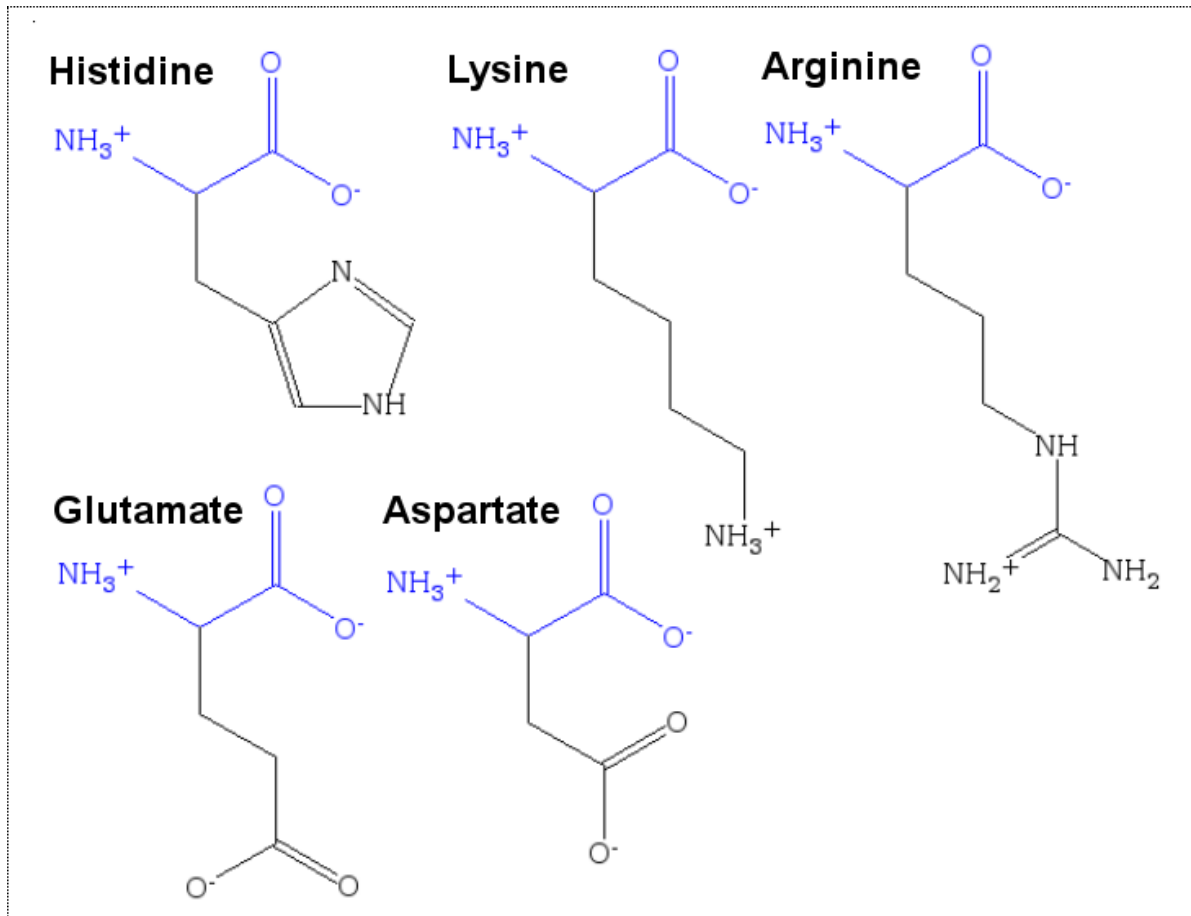
Cysteine is polar
(uncharged) ...



... and Cystine is
(very) non-polar

Polar (charged) Amino Acids

R group contains (carboxylate or amine) functional group that is partially or fully ionized at neutral pH



His has a pK_a near neutral

Amines are neutral or positively charged

Carboxylates are neutral or negatively charged



Names, Abbreviations, and more

Residues are named by replacing the **-ine** suffix of amino acids with **-yl**

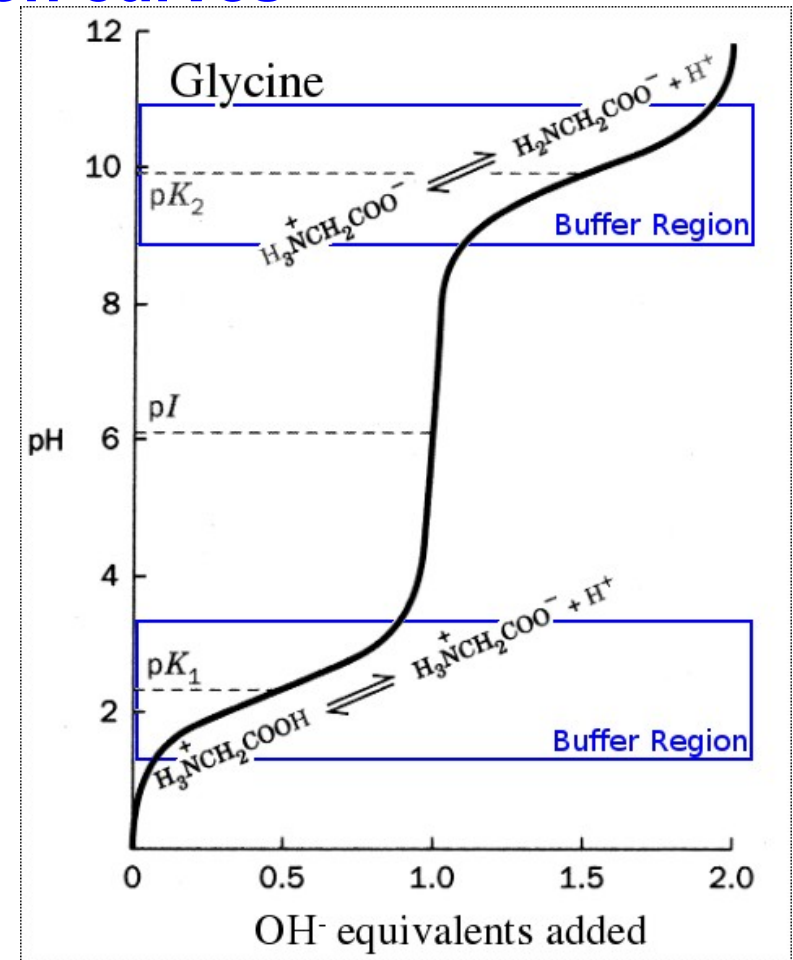
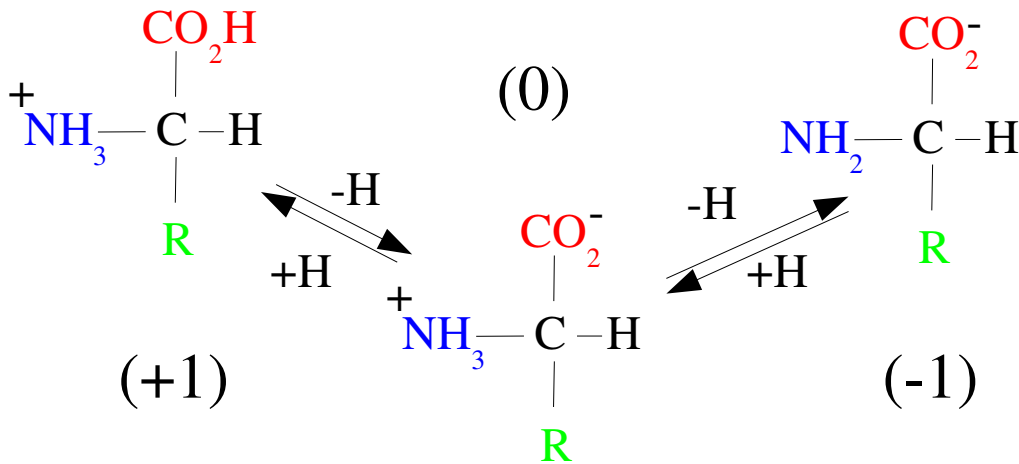
- Need to know amino acid names, 3 letter code, R-group pK_as and residue names

Amino Acid	Residue	3-letter Code	1-letter Code	R-group pK _a	Frequency (%)
Nonpolar					50.1
Glycine	Glycyl	Gly	G		7.1
Alanine	Alanyl	Ala	A		8.3
Proline	Prolyl	Pro	P		4.7
Valine	Valyl	Val	V		6.9
Leucine	Leucyl	Leu	L		9.7
Isoleucine	Isoleucyl	Ile	I		6.0
Tryptophan	Tryptophanyl	Trp	W		1.1
Phenylalanine	Phenylalanyl	Phe	F		3.9
Methionine	Methionyl	Met	M		2.4
Polar uncharged					24.0
Serine	Seryl	Ser	S		6.5
Threonine	Threonyl	Thr	T		5.3
Cysteine	Cystyl	Cys	C	8.4	1.4
Asparagine	Asparagyl	Asn	N		4.0
Glutamine	Glutaminyl	Gln	Q		3.9
Tyrosine	Tyrosyl	Tyr	Y	10.5	2.9
Polar charged					25.9
Histidine	Histidyl	His	H	6.0	2.3
Lysine	Lysyl	Lys	K	10.5	5.9
Arginine	Arginyl	Arg	R	12.5	5.5
Aspartate	Aspartyl	Asp	D	3.9	5.4
Glutamate	Glutamyl	Glu	E	4.1	6.8

Ionization States of Free Amino Acids

Free amino acids have 2 (or more) acid/base groups and produce **complex acid/base titration curves**

eg. amino acid without ionizable R group



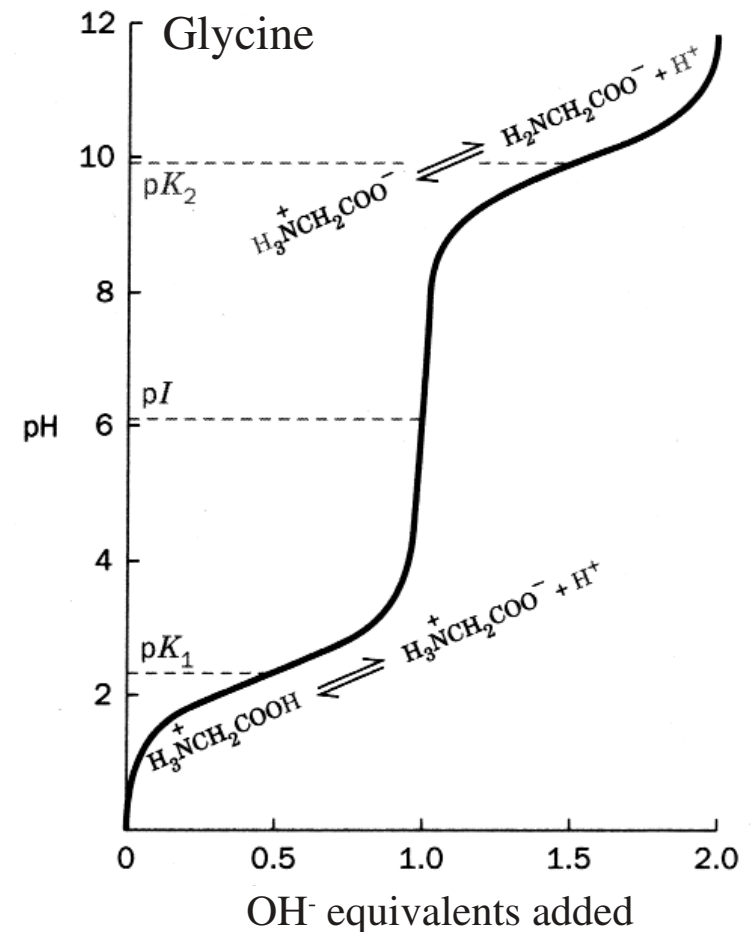


Isoelectric Point

- **Isoelectric point** is the pH at which the total charge of the amino acid (peptide or protein) is zero
 - For amino acids without ionizable R groups, the pI is the average of the pK_as

$$\text{pH} = \text{pK} + \log \frac{[\text{A}^-]}{[\text{HA}]}$$

$$\text{pI} = \frac{1}{2}(\text{pK}_i + \text{pK}_j)$$



More Ionization States (complex case)

A) +2

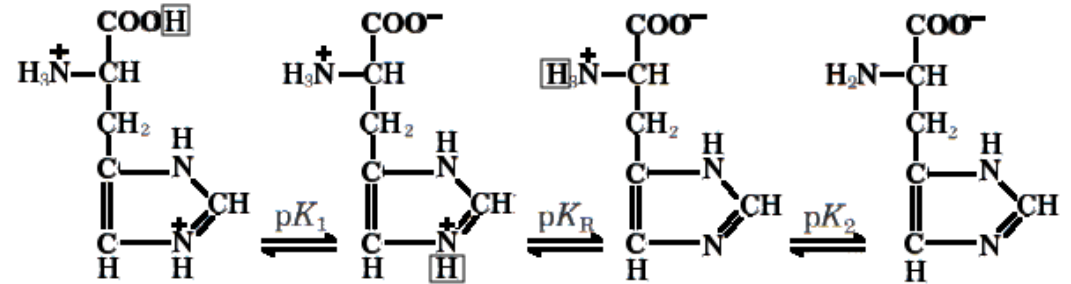
B) +1

C) 0

D) -1

Free amino acid with ionizable R group

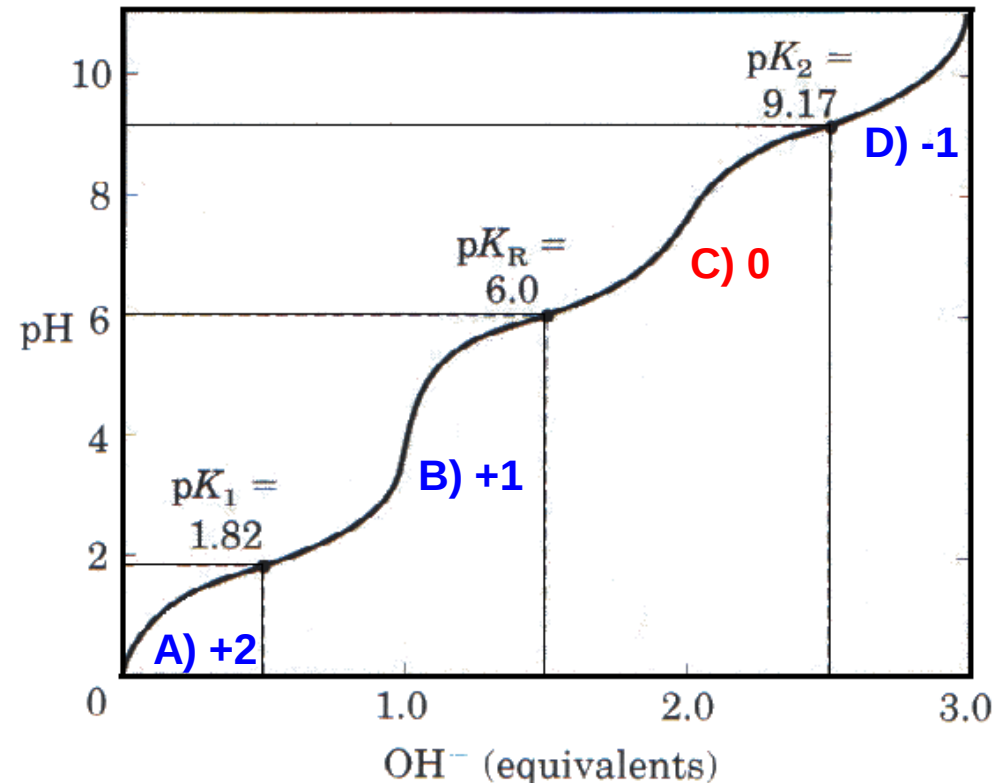
- eg. Histidine titration curve is more complex



So what is the **pl**?

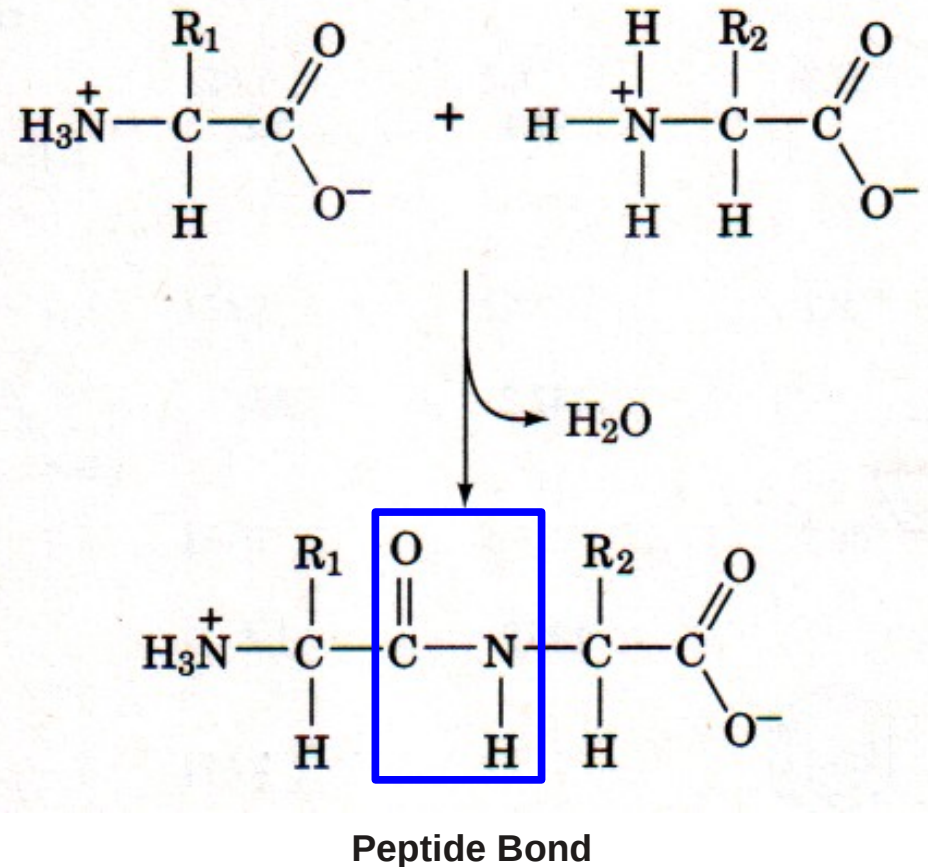
- Average of the pK_a s bounding the molecular species with a net charge of zero

$$(6.0 + 9.17) / 2 = 7.6$$



Peptide Bonds

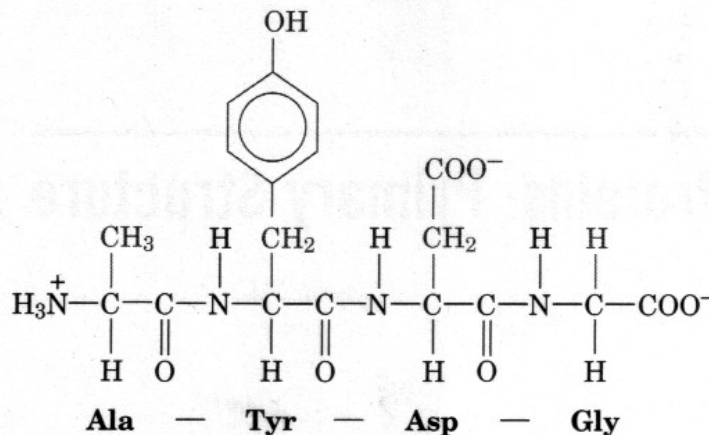
- Amino acids polymerize via **condensation** reactions
 - Carboxylate group of residue 1 forms a covalent bond with the amine group of residue 2
- Linear polymers of amino acids have an **amino (N)** and **carboxyl (C)** terminus
 - Living organism always synthesize proteins from N to C terminus



Amino Acid Polymers

- (1) **residue** – an amino acid (or peptide unit) in an oligopeptide, polypeptide or protein
- (2) **oligopeptide*** – short polymer of residues linked by peptide bonds; up to 10-20 residues.
- (3) **polypeptide*** – longer polymer of residues linked by peptide bonds; larger sizes
- (4) **protein*** – one or more polypeptide chains

*biological polymers are associated with biological function

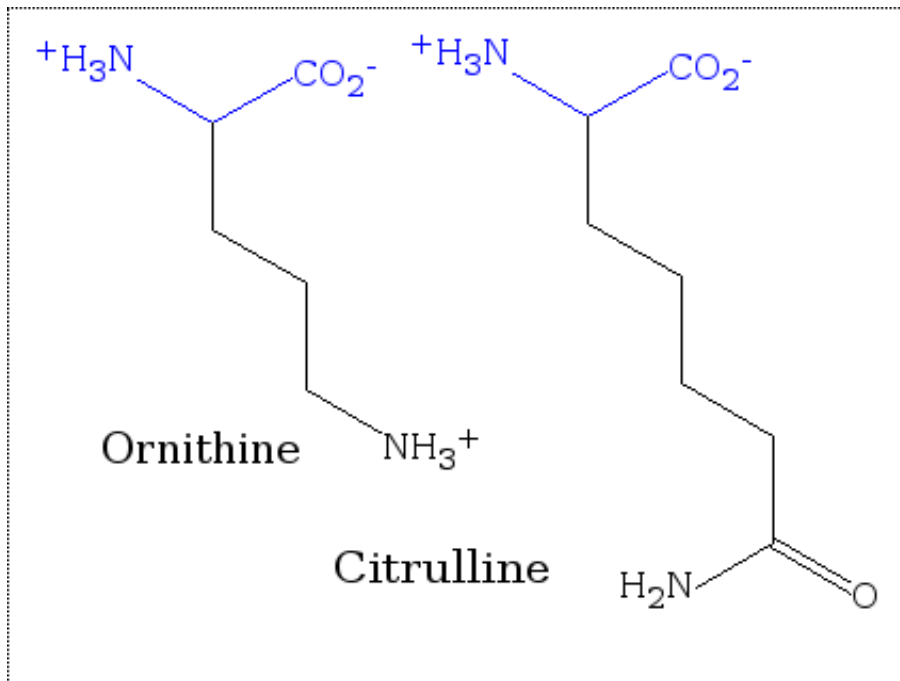


Full name:

Alanyltyrosylaspartylglycine

Non Standard Amino Acids

- > 700 non standard amino acids have been detected in living organisms
 - Many are metabolic intermediates
 - eg. ornithine and citrulline are intermediates in urea biosynthesis

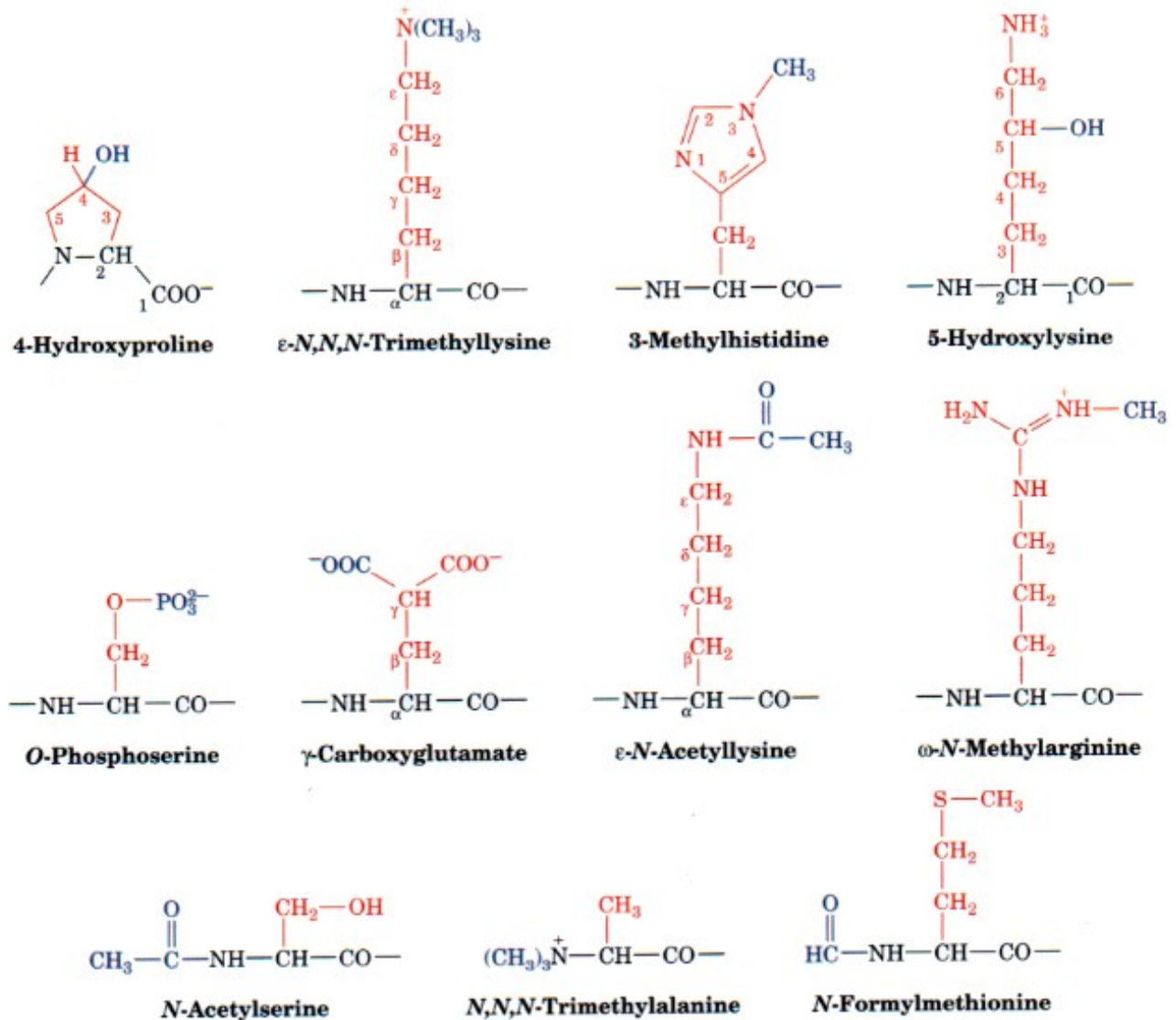


You are not responsible for drawing the structures of the non standard amino acid structures (Slides 18-20)

Non Standard Amino Acids

Non standard amino acids in proteins arise from **post translation modifications**

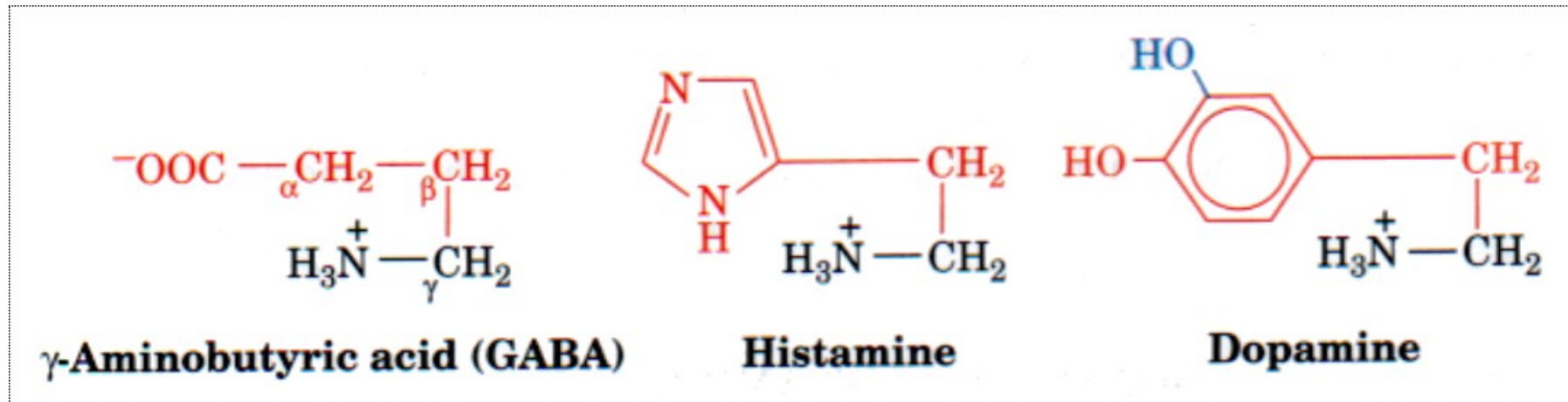
Modification are catalyzed by specific enzymes and target specific residues



Amino Acid Derivatives

Chemical derivatives of amino acids also have important biological functions

eg. Catecholamines (below) lack the α -carboxylate of amino acids



GABA & Dopamine are neurotransmitters. Histamine mediates parts of the immune response.