



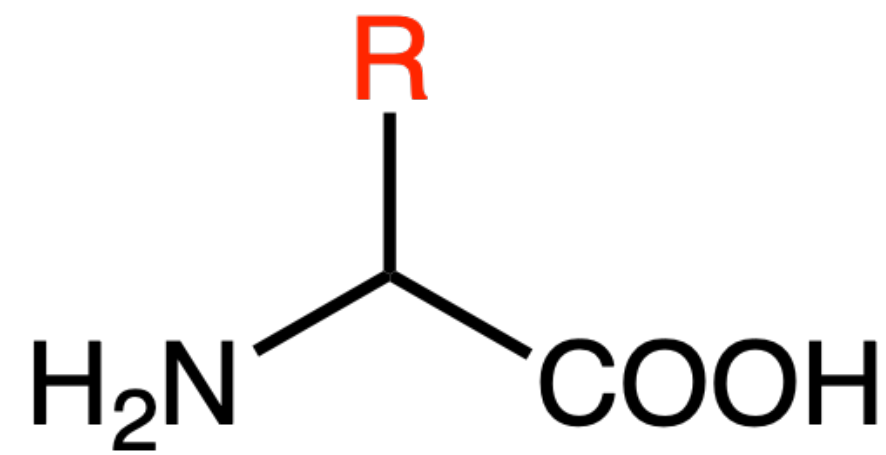
# BIOCHEMISTRY

PROTEINS OR POLYPEPTIDES

CHEMISTRY 165 // SPRING 2020

# The amino acids

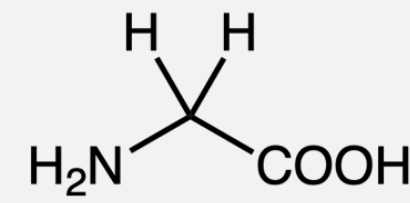
In biochemistry, we are most interested in 20 amino acids (drawn and named to the right) with the general formula  $\text{H}_2\text{NCHRCOOH}$  and the structure:



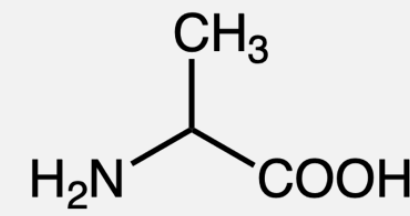
The **R** group is called the sidechain.

[Amino acids combine via condensation reactions to form proteins or polypeptides.](#)

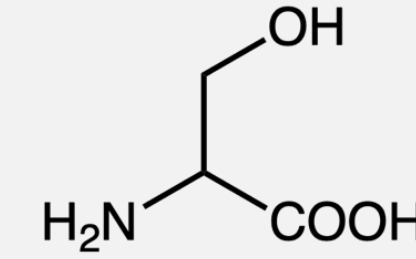
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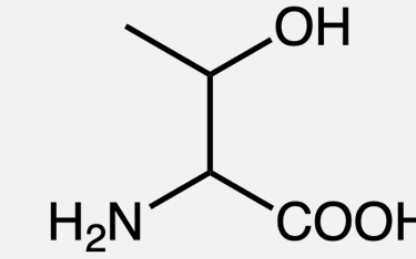
Glycine (Gly, G)



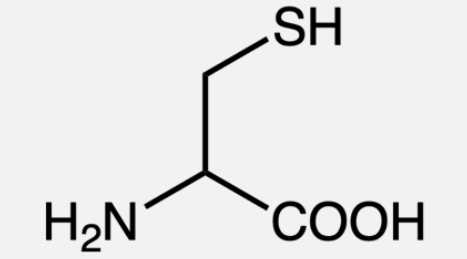
Alanine (Ala, A)



Serine (Ser, S)  
 $pK_a \sim 16$

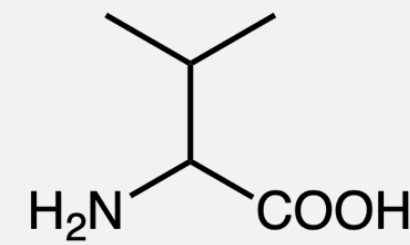


Threonine (Thr, T)  
 $pK_a \sim 16$

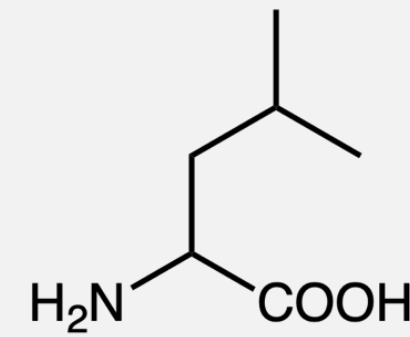


Cysteine (Cys, C)  
 $pK_a = 8.35$

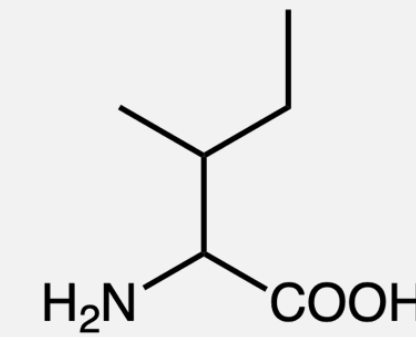
## HYDROPHOBIC



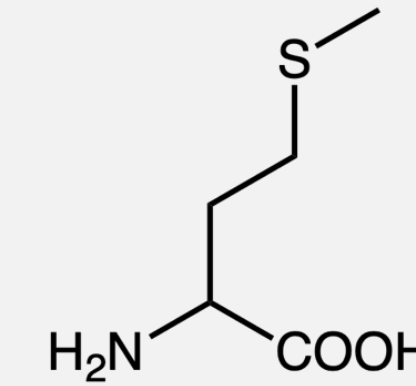
Valine (Val, V)



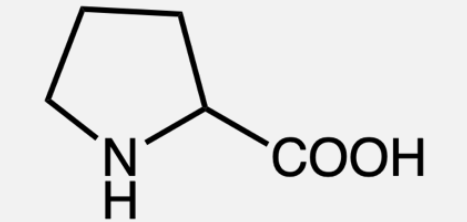
Leucine (Leu, L)



Isoleucine (Ile, I)

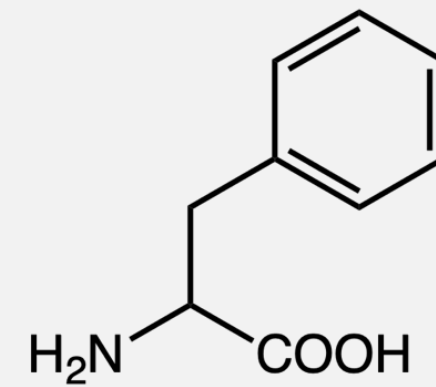


Methionine (Met, M)

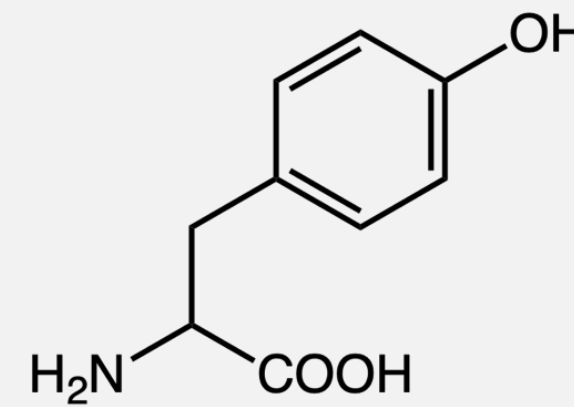


Proline (Pro, P)

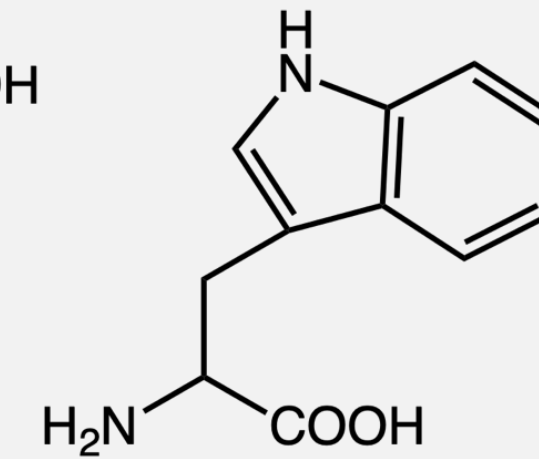
## AROMATIC



Phenylalanine (Phe, F)

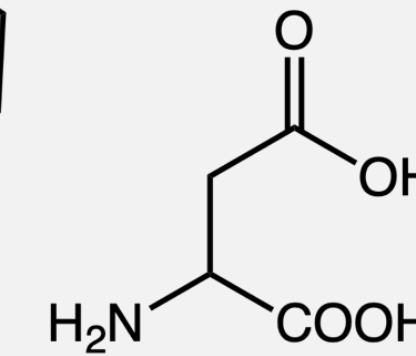


Tyrosine (Tyr, Y)

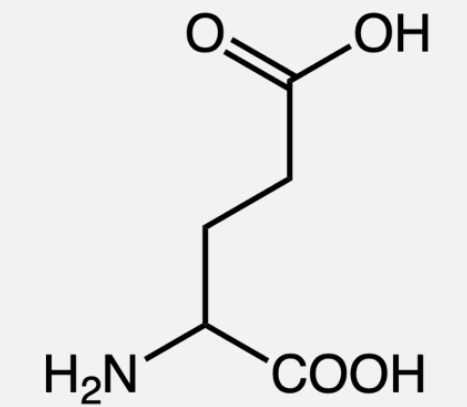


Tryptophan (Trp, W)

## ACIDIC

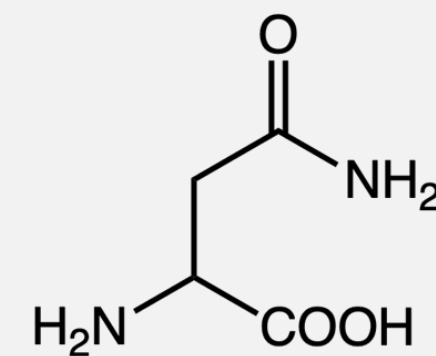


Aspartic Acid (Asp, D)  
 $pK_a = 3.9$

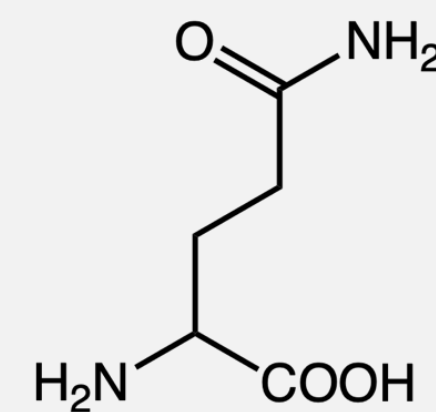


Glutamic Acid (Glu, E)  
 $pK_a = 4.07$

## AMIDE

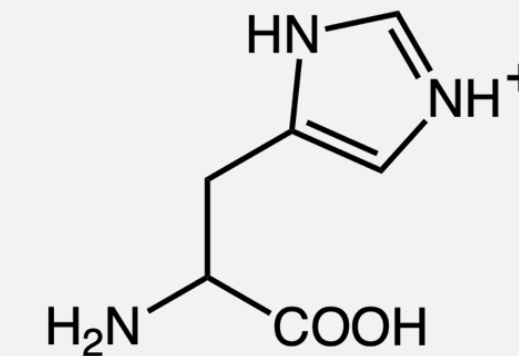


Asparagine (Asn, N)

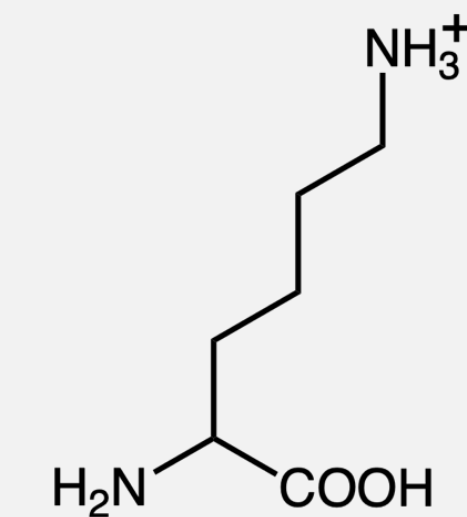


Glutamine (Gln, Q)

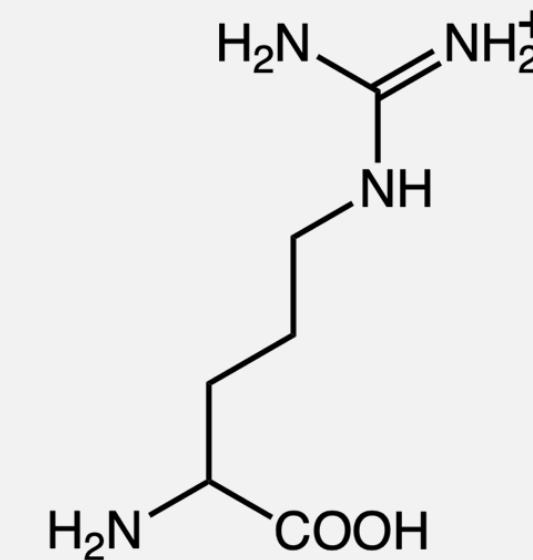
## BASIC



Histidine (His, H)  
 $pK_a = 6.04$



Lysine (Lys, K)  
 $pK_a = 10.79$

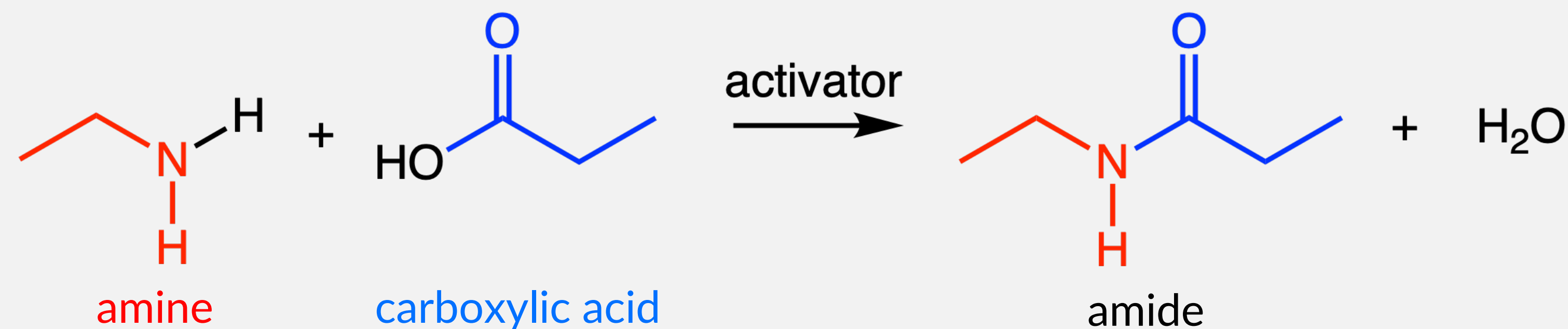
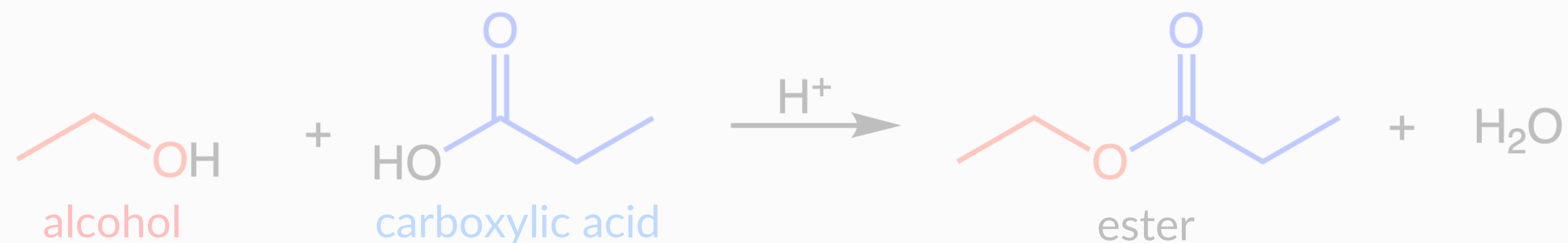
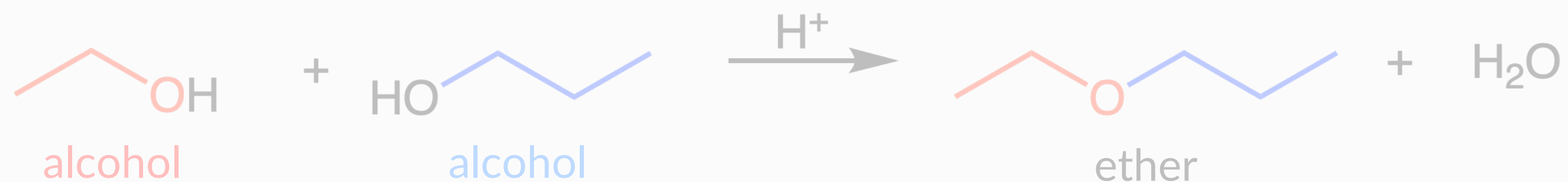


Arginine (Arg, R)  
 $pK_a = 12.48$

# Recall: Making amides via condensation

In general terms, condensation reactions are:  $A + B \rightarrow A-B + \text{small molecule (such as: H}_2\text{O, NH}_3\text{, HCl, etc.)}$

Reaction: combine two oxygenates into another oxygenate; requires an acid catalyst ( $\text{H}^+$ ) or activator.



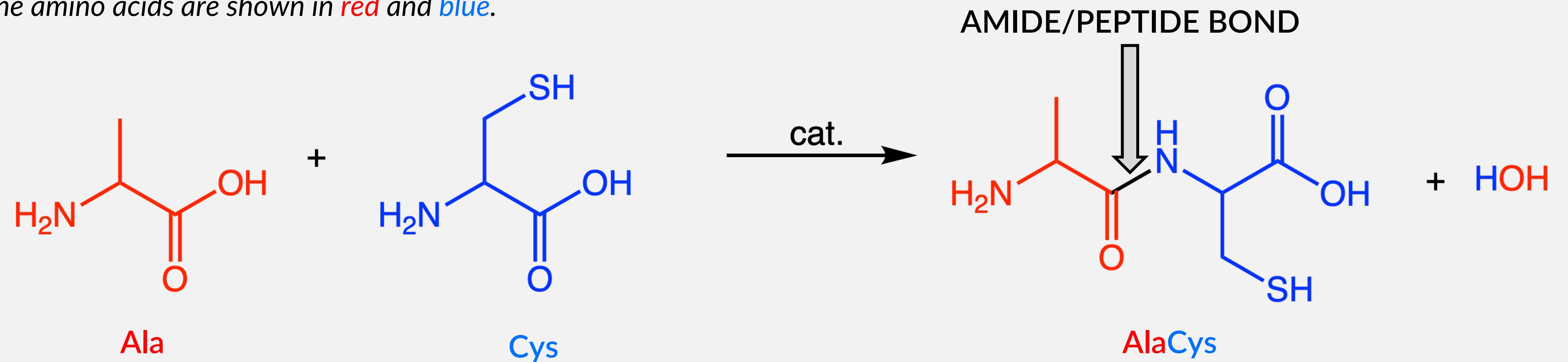


# Making proteins from amino acids

Consider a small protein (a dipeptide) made from condensation of **alanine (Ala)** and **cysteine (Cys)**:

*The amide/peptide bonds are shown with black bonds.*

*The amino acids are shown in **red** and **blue**.*



*These condensation reactions typically require a catalyst, in the form of an enzyme.*

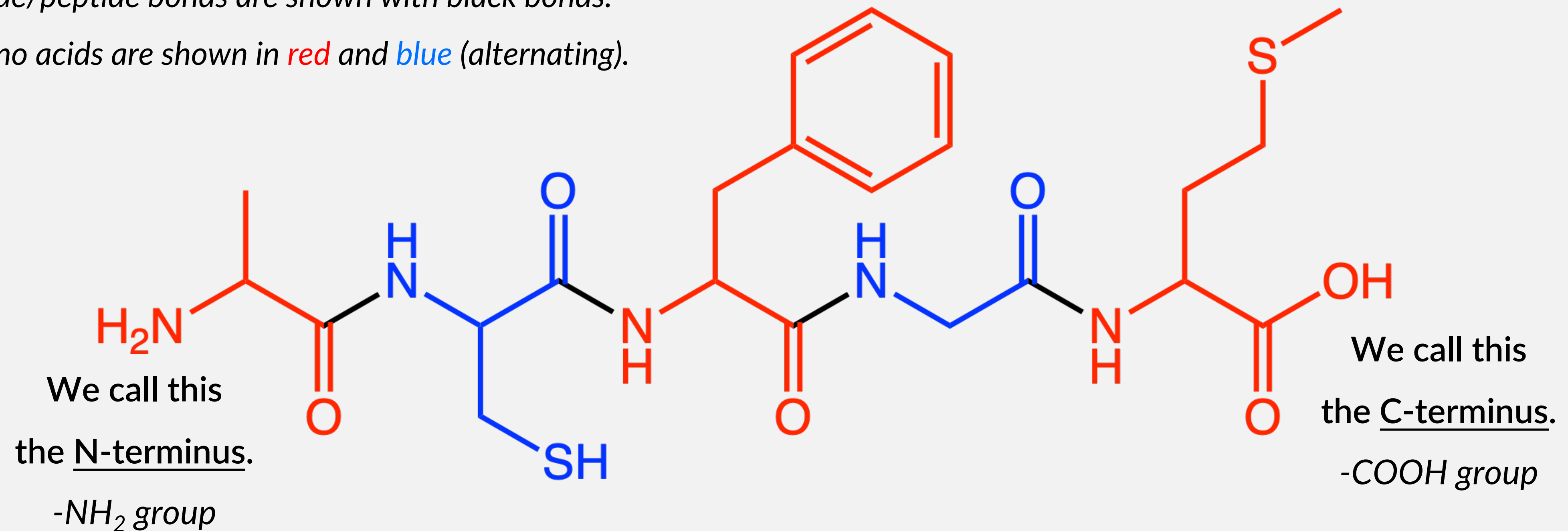
The new amide bond formed between the amino acids is sometimes also called a peptide bond.

# Naming proteins (polypeptides)

Consider now the larger protein (a polypeptide):

The amide/peptide bonds are shown with black bonds.

The amino acids are shown in *red* and *blue* (alternating).

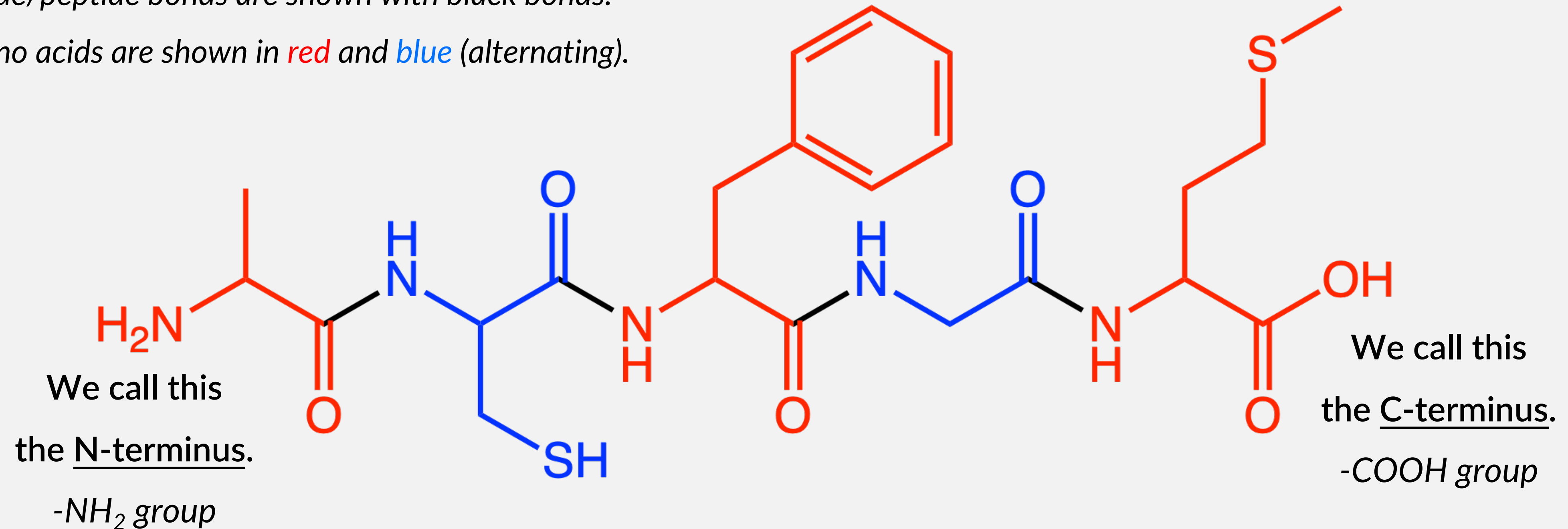


We name proteins from the N-terminus (left) to the C-terminus (right), so: **AlaCysPheGlyMet**

# Properties of proteins: Acid-base + charge

The amide/peptide bonds are shown with black bonds.

The amino acids are shown in *red* and *blue* (alternating).



We name proteins from the N-terminus (left) to the C-terminus (right), so: **AlaCysPheGlyMet**

Because the amino acids that comprise this protein are all pH-neutral, the overall protein is also pH-neutral.

As such, we also expect this protein to be charge-neutral at pH ~ 6 since the constituent amino acids have pI ~ 6.

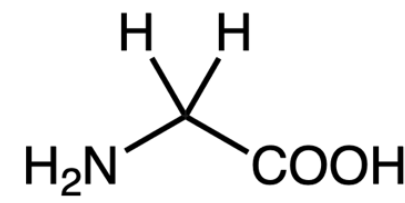
# PRACTICE PROBLEM 1

Estimate the overall charge on the protein at pH = 6:

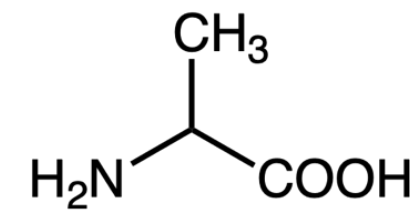
ArgProProGlyPheSerProPheArg.

— answer —

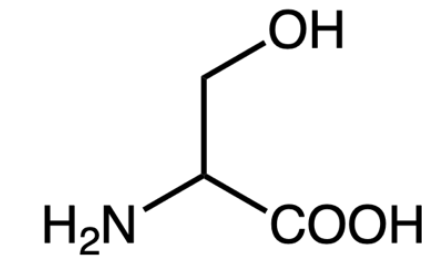
## SMALL



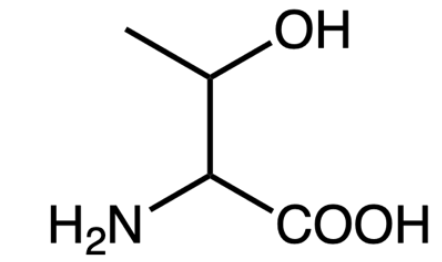
Glycine (Gly, G)



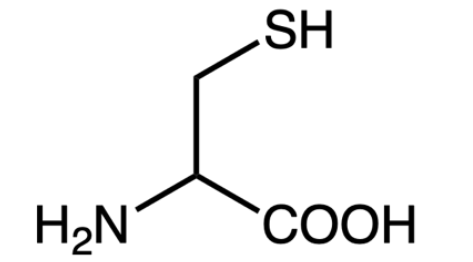
Alanine (Ala, A)



Serine (Ser, S)  
 $pK_a \sim 16$

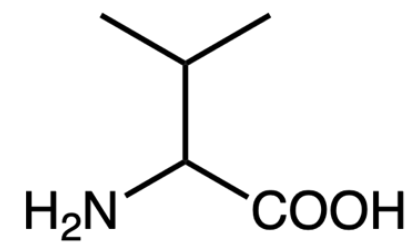


Threonine (Thr, T)  
 $pK_a \sim 16$

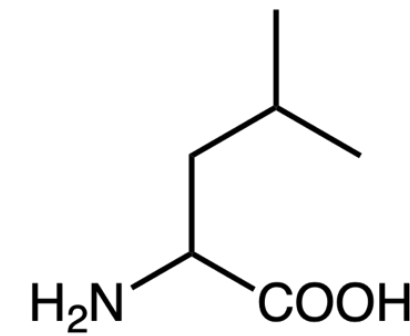


Cysteine (Cys, C)  
 $pK_a = 8.35$

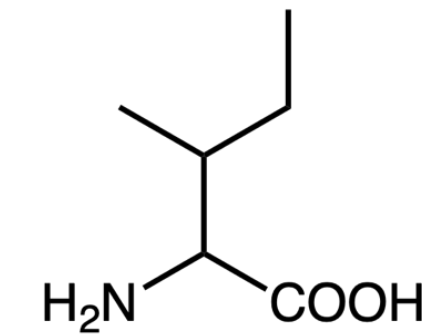
## HYDROPHOBIC



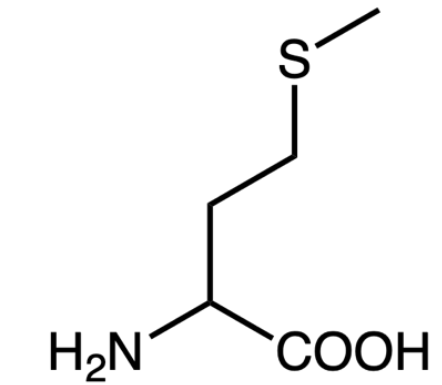
Valine (Val, V)



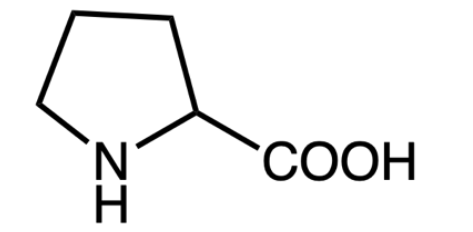
Leucine (Leu, L)



Isoleucine (Ile, I)

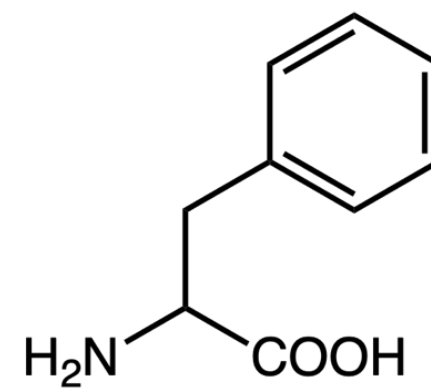


Methionine (Met, M)

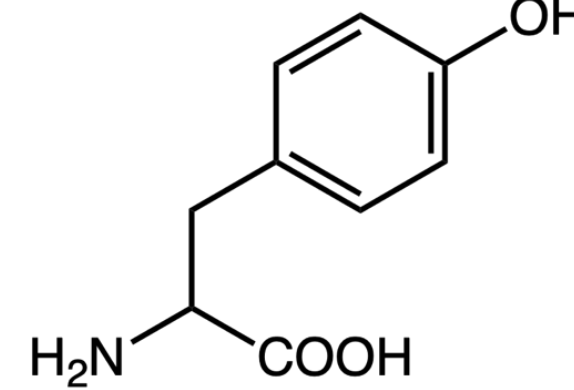


Proline (Pro, P)

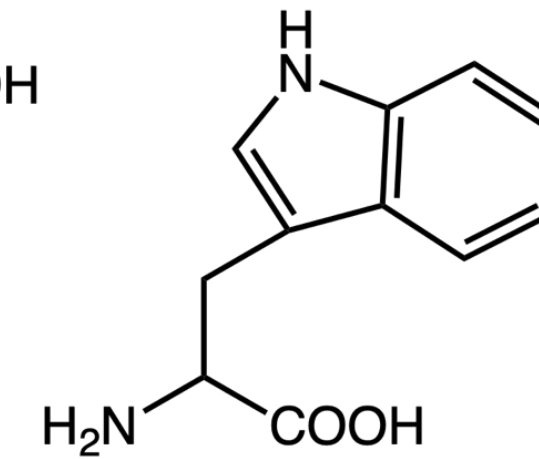
## AROMATIC



Phenylalanine (Phe, F)

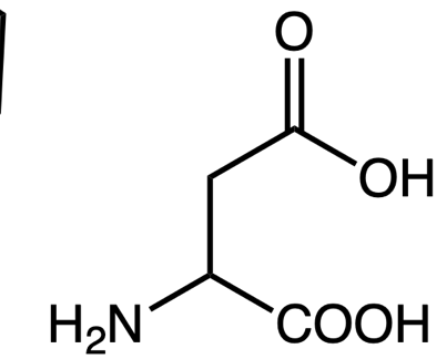


Tyrosine (Tyr, Y)

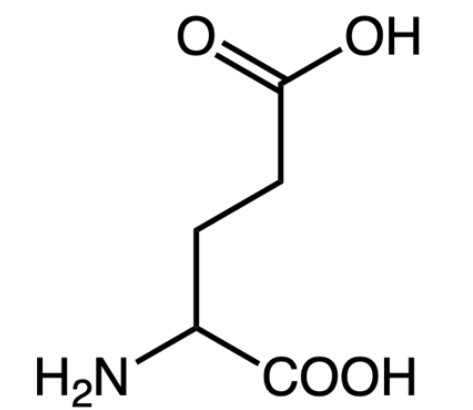


Tryptophan (Trp, W)

## ACIDIC

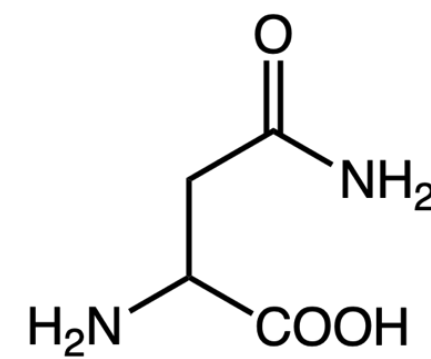


Aspartic Acid (Asp, D)  
 $pK_a = 3.9$

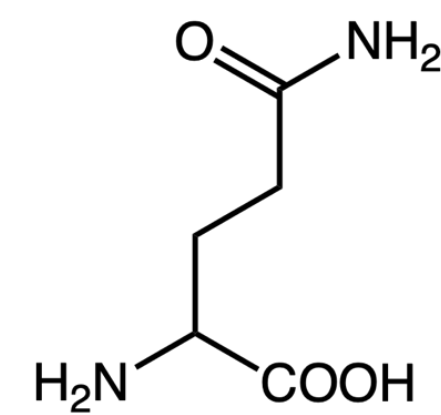


Glutamic Acid (Glu, E)  
 $pK_a = 4.07$

## AMIDE

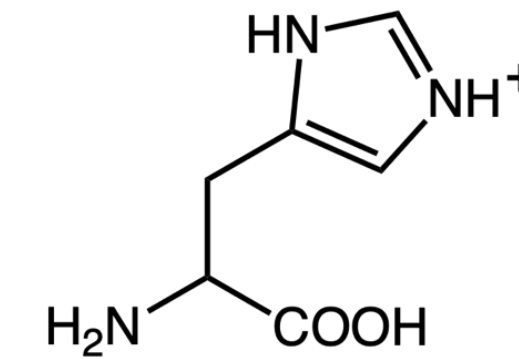


Asparagine (Asn, N)

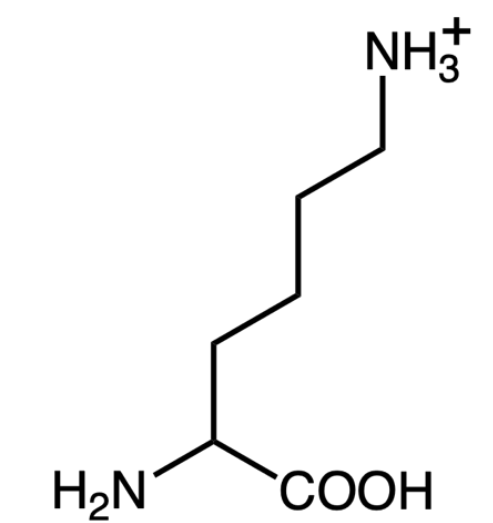


Glutamine (Gln, Q)

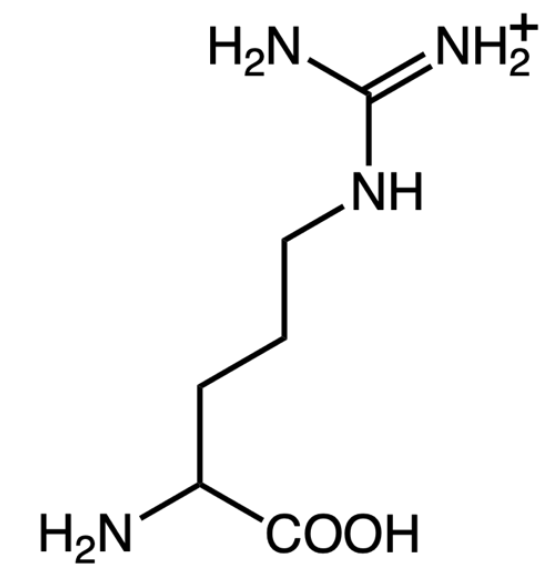
## BASIC



Histidine (His, H)  
 $pK_a = 6.04$



Lysine (Lys, K)  
 $pK_a = 10.79$



Arginine (Arg, R)  
 $pK_a = 12.48$



# PRACTICE PROBLEM 1

Estimate the overall charge on the protein at pH = 6:

ArgProProGlyPheSerProPheArg.

— answer —

Assign charges to each of the amino acids at pH = 6:

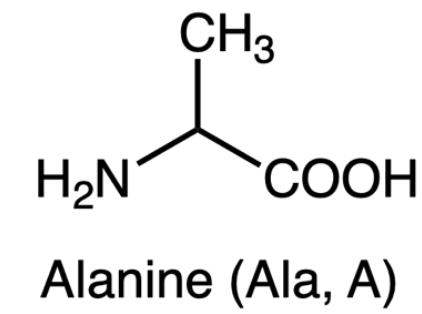
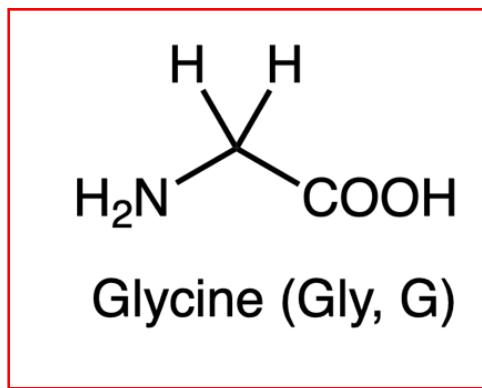
- Arg → pI ~ 11 → charge = +1
- Pro → pI ~ 6 → charge = 0
- Gly → pI ~ 6 → charge = 0
- Phe → pI ~ 6 → charge = 0
- Ser → pI ~ 6 → charge = 0

The overall charge on the protein is the sum of the charges on the individual amino acids. In this case, the two Arg groups contribute a +1 charge each, and all other amino acids contribute 0 charge.

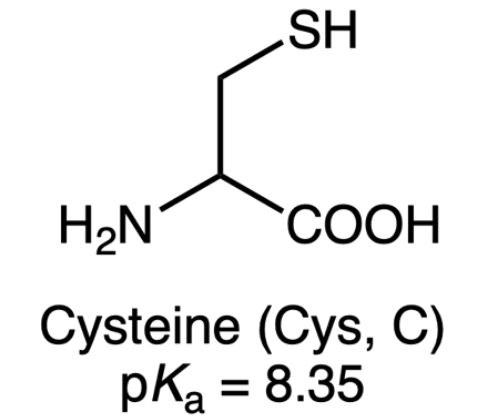
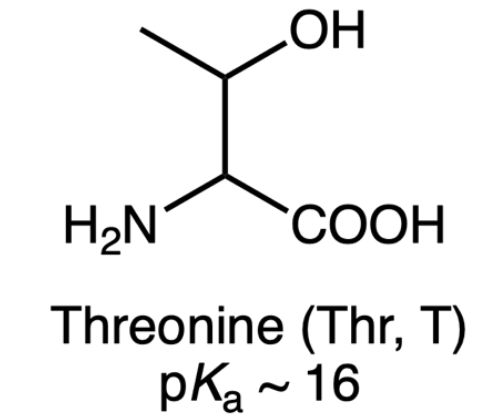
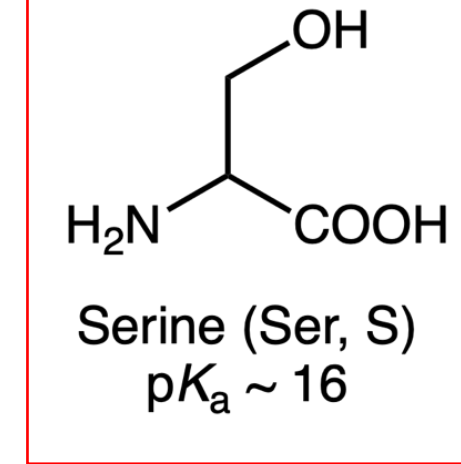
**Overall charge = +2**

The protein is also basic because of the basic Arg groups.

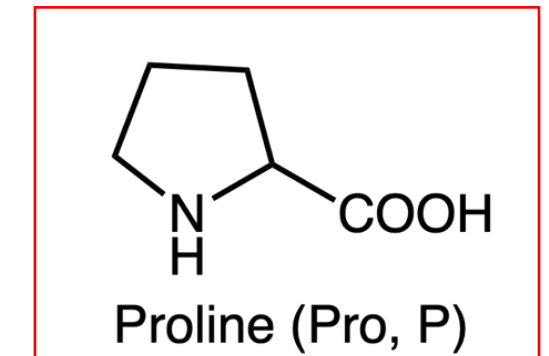
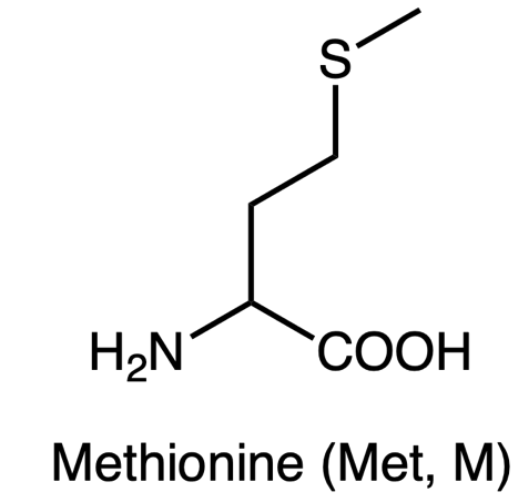
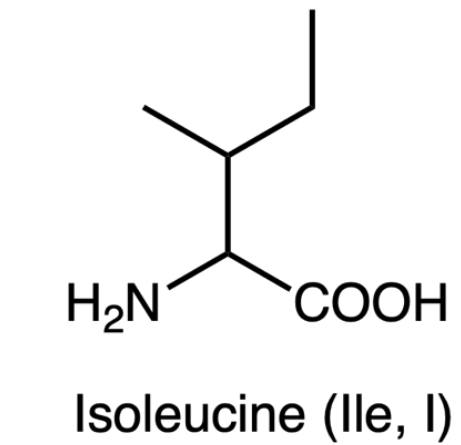
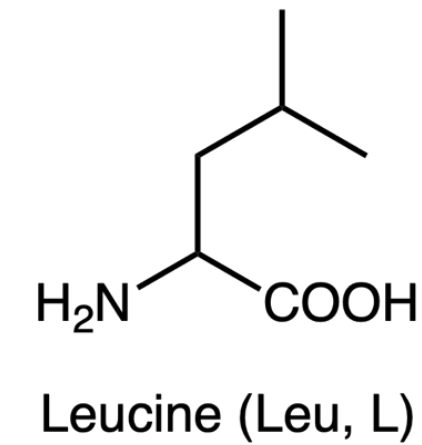
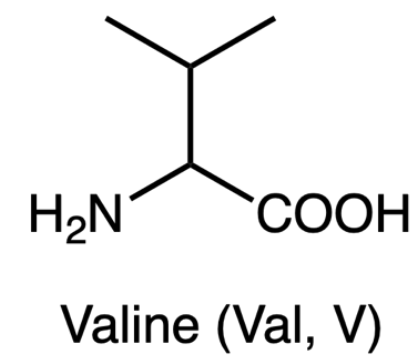
## SMALL



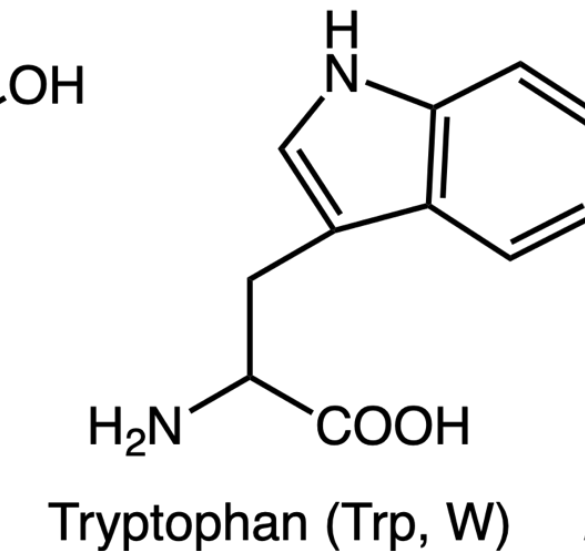
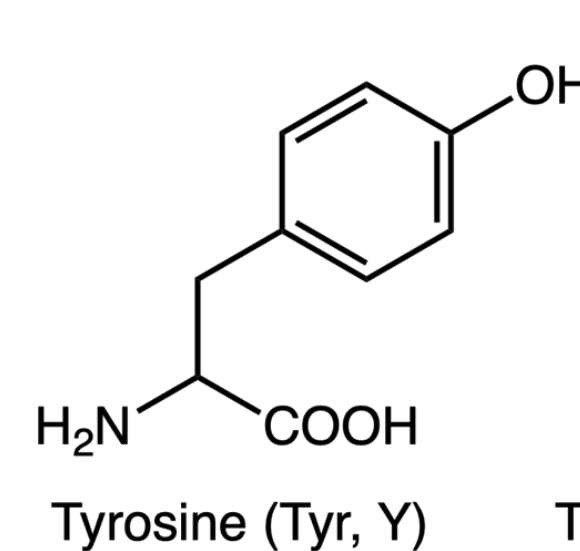
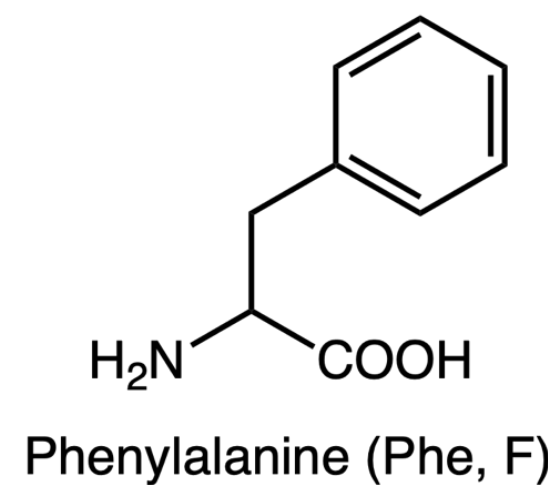
## NUCLEOPHILIC



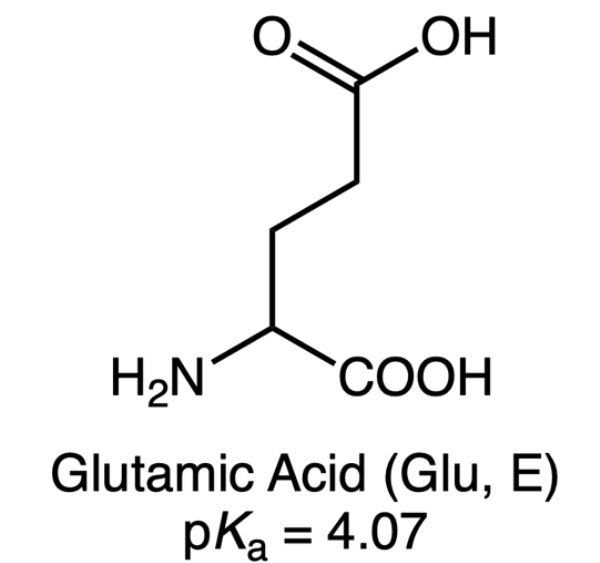
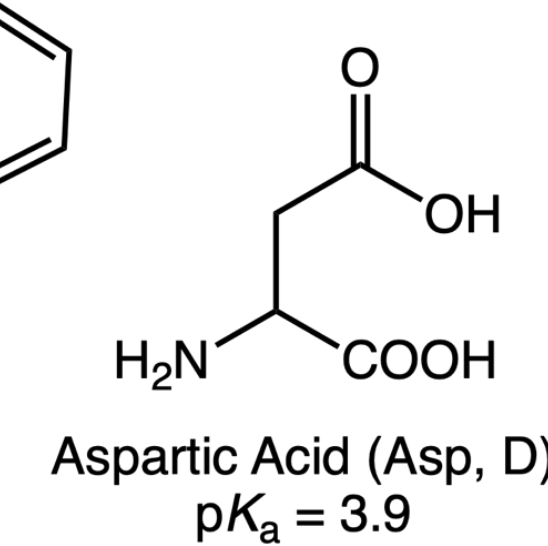
## HYDROPHOBIC



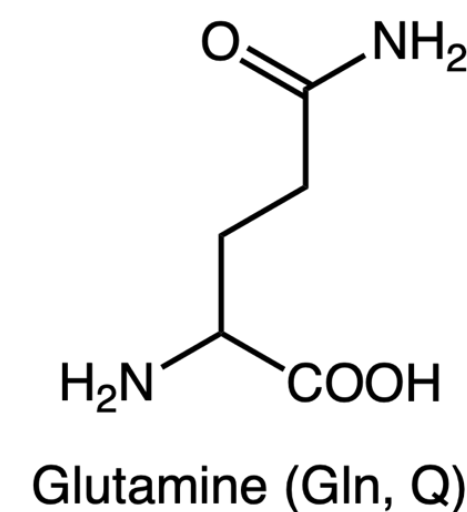
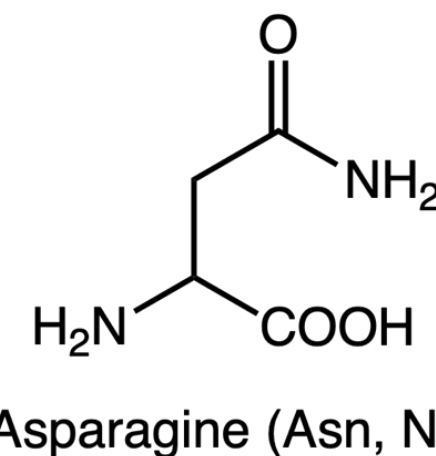
## AROMATIC



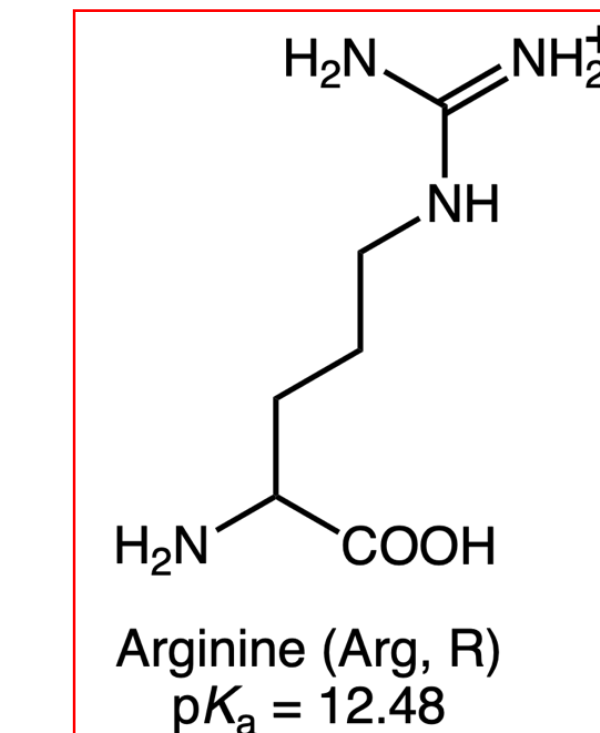
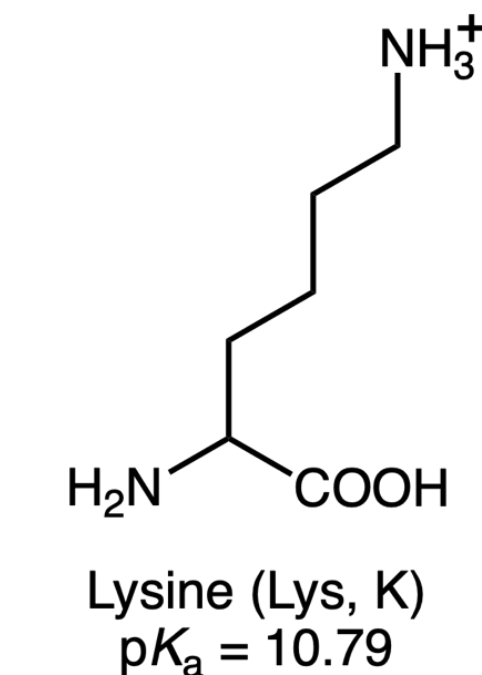
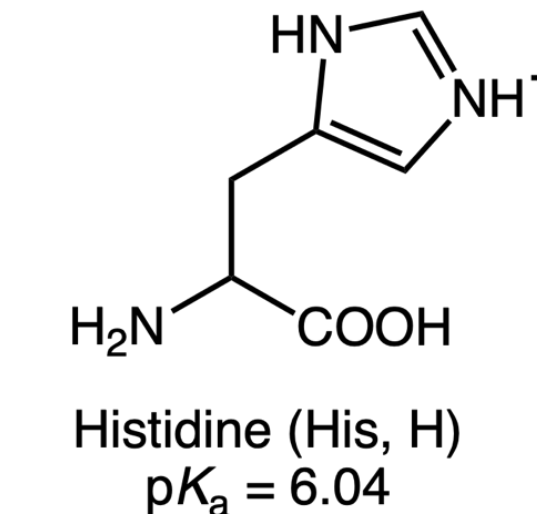
## ACIDIC



## AMIDE



## BASIC



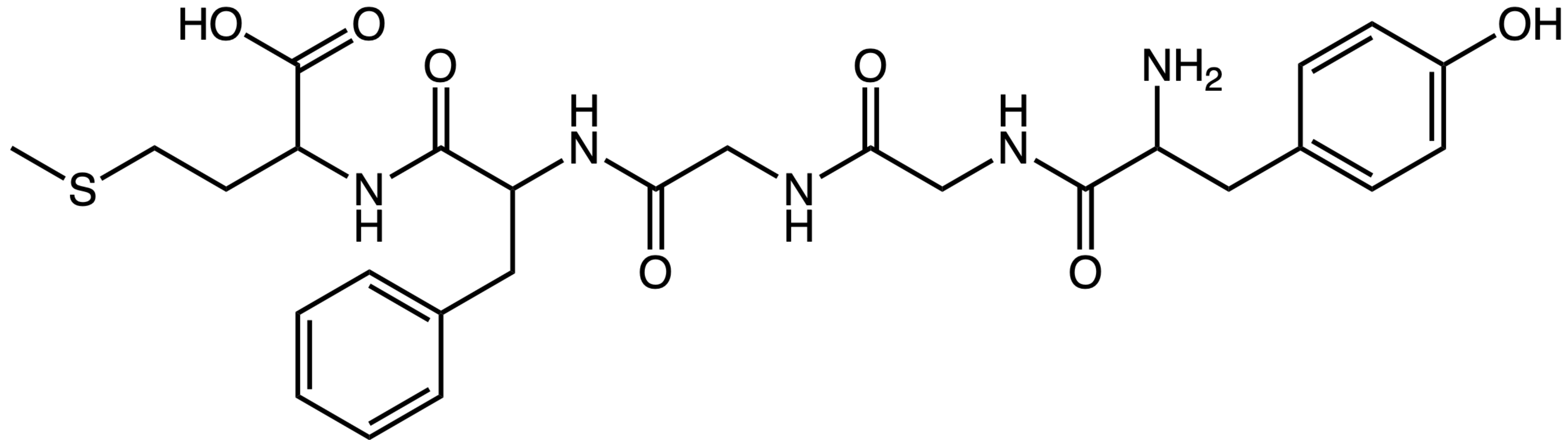


## PRACTICE PROBLEM 2

Identify the amino acids that comprise the following protein.

Is this protein acidic, basic, or neutral overall?

— *answer* —



## PRACTICE PROBLEM 2

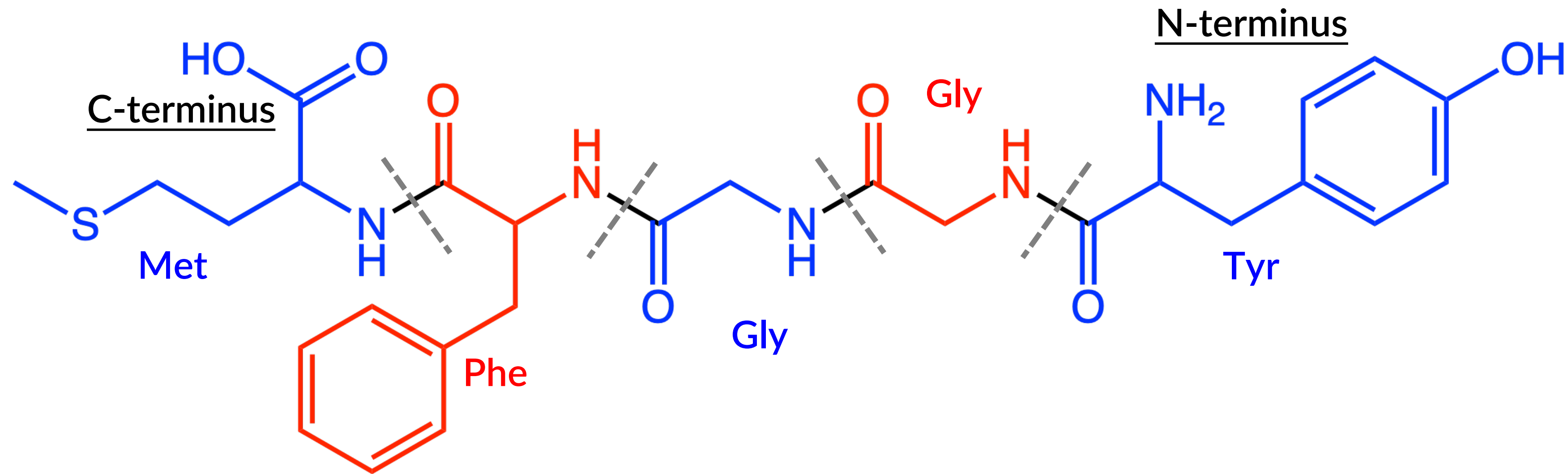
Identify the amino acids that comprise the following protein.

Is this protein acidic, basic, or neutral overall?

— *answer* —

First, find the N-terminus, which will have a free  $-NH_2$  on one of the ends of the protein. In this case, it is on the right side.

Second, cut the protein at the amide/peptide bond: shown as black bonds with dashed gray lines bisecting bond.



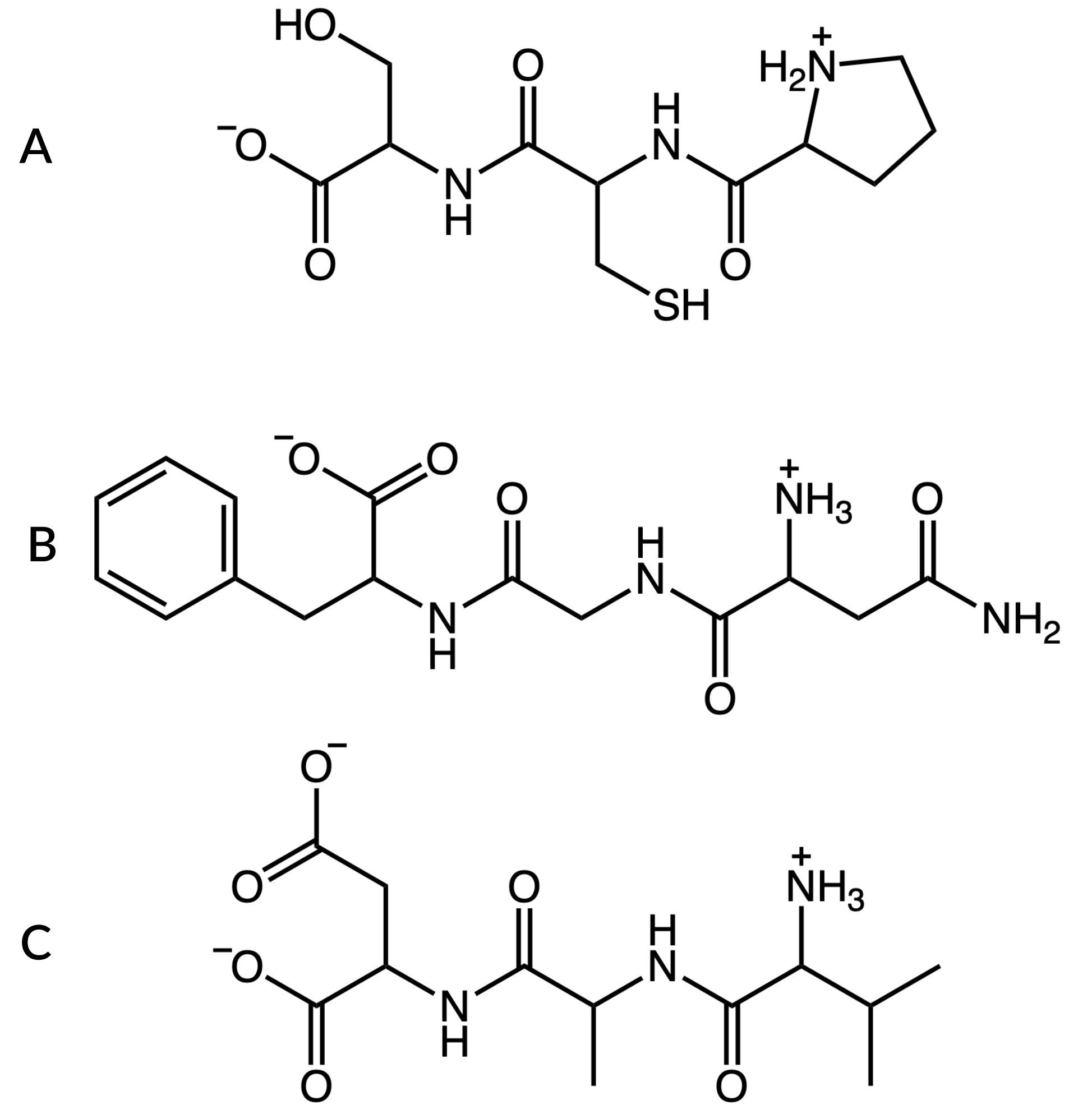
Then identify/name each of the amino acids: **Tyr****Gly****Gly****Phe****Met**.

Since all of the amino acids are pH-neutral, the overall protein is also pH-neutral.

# PRACTICE PROBLEM 3

Which of the following tripeptides is/are acidic?

— answer —



## PRACTICE PROBLEM 3

Which of the following tripeptides is/are acidic?

— *answer* —

For each tripeptide, first identify the amino acids and name the tripeptide starting from the N-terminus (the side a free  $-\text{NH}_2$  group).

- A) ProCysSer
- B) AsnGlyPhe
- C) ValAlaAsp

Because tripeptides A and B are only composed of pH-neutral amino acids (Pro, Cys, Ser, Asn, Ala, Phe) both are pH-neutral tripeptides.

**Tripeptide C is acidic because of the acidic amino acid Asp (aspartic acid).**

