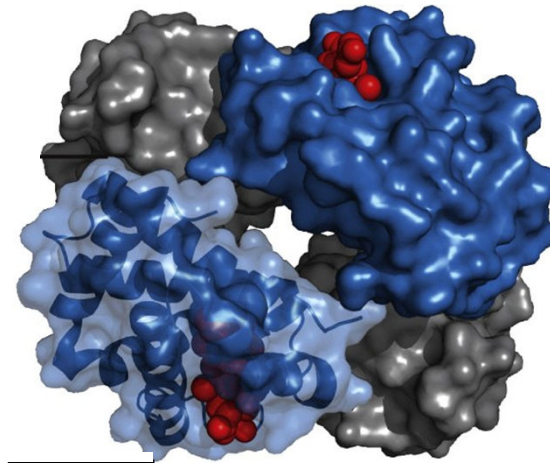
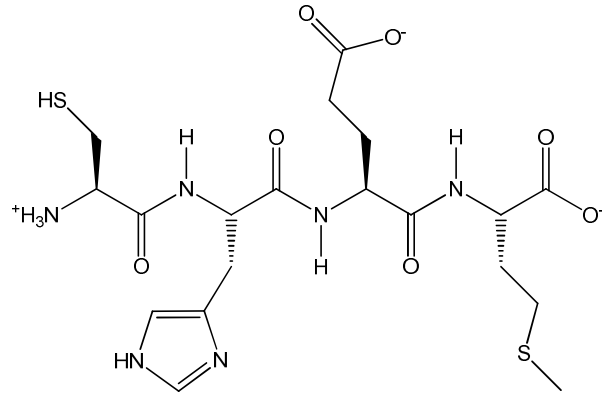
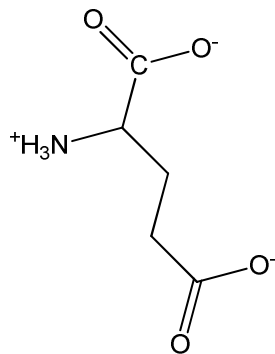


Amino acids have important roles in living organisms



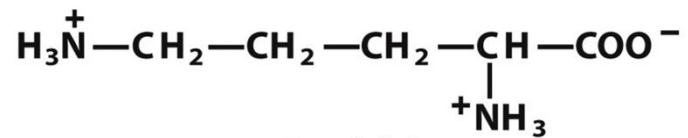
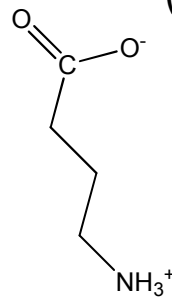
Subunits (building blocks) of peptides and proteins

glutamate

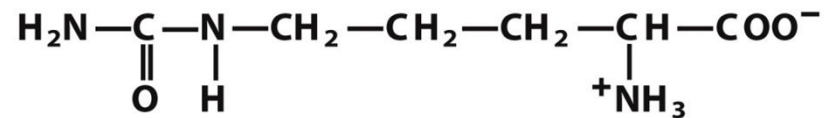


Neurotransmitters

γ -aminobutyric acid
(GABA)



Ornithine



Citrulline

Metabolic intermediates

Proteins are synthesized from 20 'standard' α -amino acids. Their names have 3- and 1-letter abbreviations.

Alanine	Ala	A
Cysteine	Cys	C
Aspartate	Asp	D
Glutamate	Glu	E
Phenylalanine	Phe	F
Glycine	Gly	G
Histidine	His	H
Isoleucine	Ile	I
Lysine	Lys	K
Leucine	Leu	L

Methionine	Met	M
Asparagine	Asn	N
Proline	Pro	P
Glutamine	Gln	Q
Arginine	Arg	R
Serine	Ser	S
Threonine	Thr	T
Valine	Val	V
Tryptophan	Trp	W
Tyrosine	Tyr	Y

TABLE 3-1 Properties and Conventions Associated with the Common Amino Acids Found in Proteins

Amino acid	Abbreviation/ symbol	M_r^*	pK_a values			pI	Hydropathy index [†]	Occurrence in proteins (%) [‡]
			pK_1 (—COOH)	pK_2 (—NH ₃ ⁺)	pK_R (R group)			
Nonpolar, aliphatic								
R groups								
Glycine	Gly G	75	2.34	9.60		5.97	−0.4	7.2
Alanine	Ala A	89	2.34	9.69		6.01	1.8	7.8
Proline	Pro P	115	1.99	10.96		6.48	1.6	5.2
Valine	Val V	117	2.32	9.62		5.97	4.2	6.6
Leucine	Leu L	131	2.36	9.60		5.98	3.8	9.1
Isoleucine	Ile I	131	2.36	9.68		6.02	4.5	5.3
Methionine	Met M	149	2.28	9.21		5.74	1.9	2.3
Aromatic								
R groups								
Phenylalanine	Phe F	165	1.83	9.13		5.48	2.8	3.9
Tyrosine	Tyr Y	181	2.20	9.11	10.07	5.66	−1.3	3.2
Tryptophan	Trp W	204	2.38	9.39		5.89	−0.9	1.4
Polar, uncharged								
R groups								
Serine	Ser S	105	2.21	9.15		5.68	−0.8	6.8
Threonine	Thr T	119	2.11	9.62		5.87	−0.7	5.9
Cysteine [§]	Cys C	121	1.96	10.28	8.18	5.07	2.5	1.9
Asparagine	Asn N	132	2.02	8.80		5.41	−3.5	4.3
Glutamine	Gln Q	146	2.17	9.13		5.65	−3.5	4.2
Positively charged								
R groups								
Lysine	Lys K	146	2.18	8.95	10.53	9.74	−3.9	5.9
Histidine	His H	155	1.82	9.17	6.00	7.59	−3.2	2.3
Arginine	Arg R	174	2.17	9.04	12.48	10.76	−4.5	5.1
Negatively charged								
R groups								
Aspartate	Asp D	133	1.88	9.60	3.65	2.77	−3.5	5.3
Glutamate	Glu E	147	2.19	9.67	4.25	3.22	−3.5	6.3

Amino acid pK_a's vary because of attached functional groups (microenvironment influences acidity)

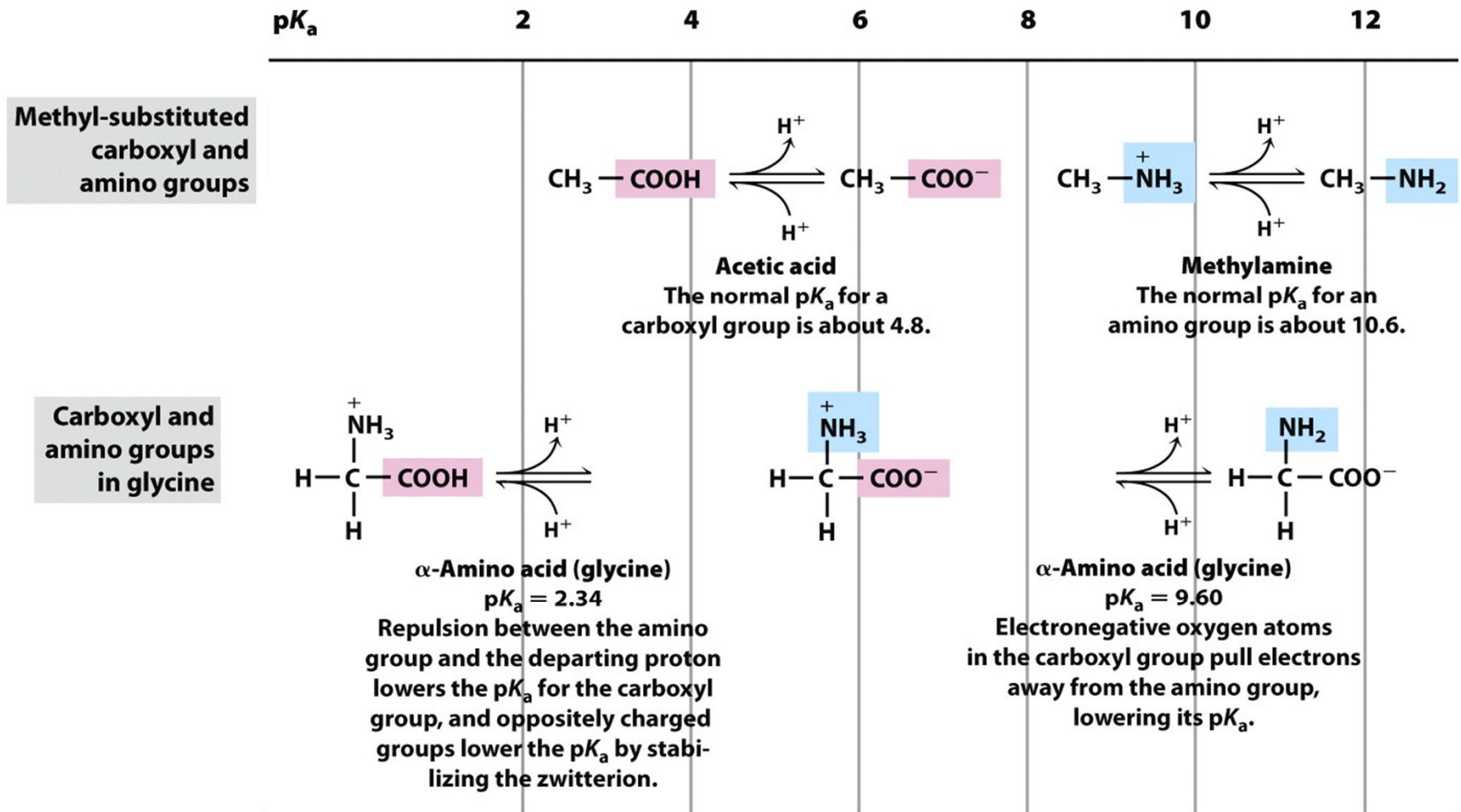
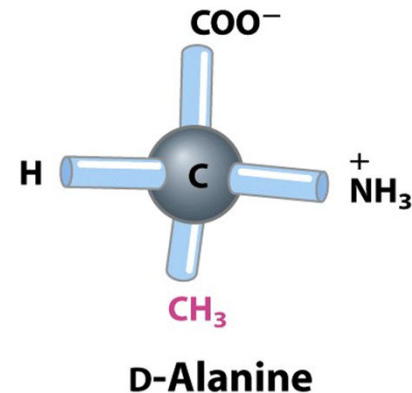
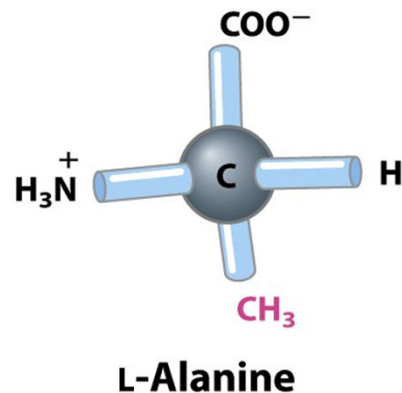
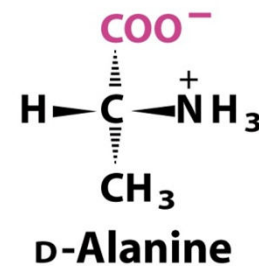
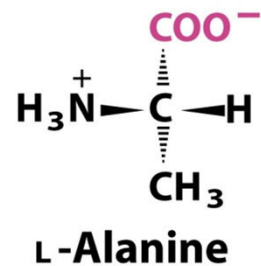
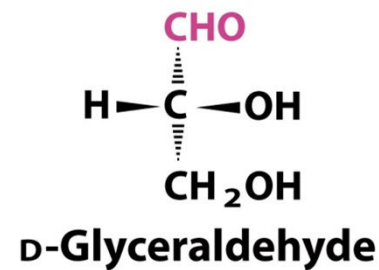
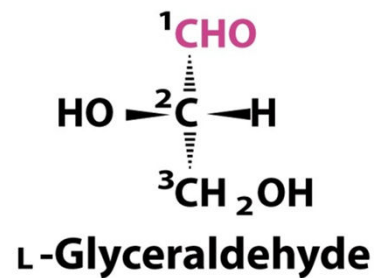


Figure 3-11

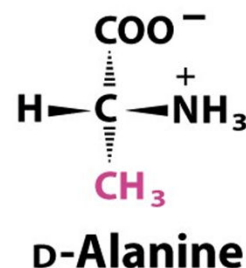
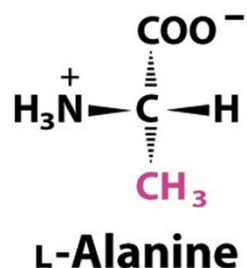
Lehninger Principles of Biochemistry, Fifth Edition

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Most amino acids are chiral, and are designated 'D' or 'L' based on Emil Fischer's nomenclature

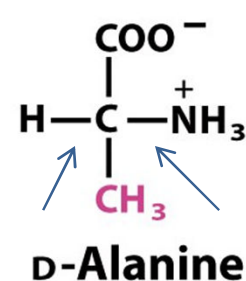
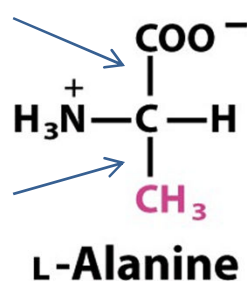


The 'Fischer projection' is a simplified way to depict stereochemistry

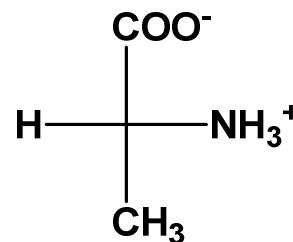
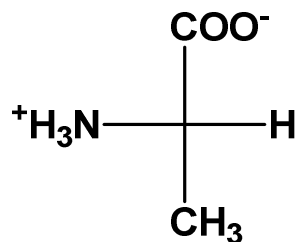


Fisher projection

Vertical bonds point away from viewer (dashed wedges)



Horizontal bonds point toward viewer (filled wedges)

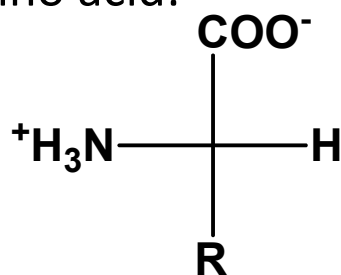


Chiral carbons may be left out entirely

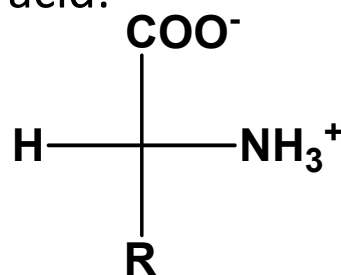
How to identify the stereochemistry of an amino acid (D or L?)

Fisher projection

L-amino acid:



D-amino acid:

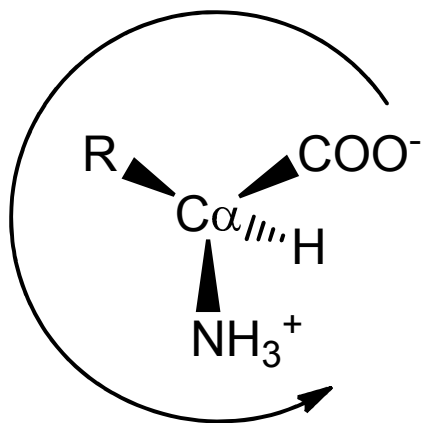


If projection is aligned with:

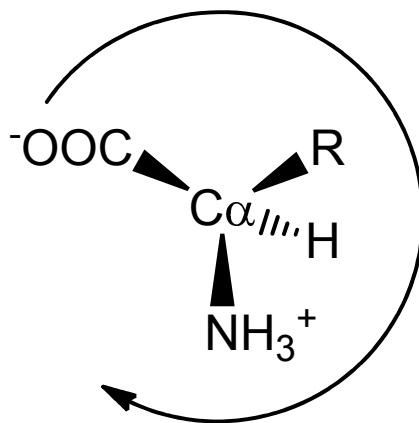
- carbon chain vertical
- carboxylate on top
- amino group on left = L
- amino group on right = D

The CORN rule

L-amino acid:



D-amino acid:



1. Orient the structure to look down the $\text{C}_\alpha\text{-H}$ bond
2. Follow the other C_α substituents to spell CORN (carboxylate, R-group, amino): counterclockwise is L, clockwise is D

Nonstandard amino acids are found in modified proteins and as free metabolites

