

21 L- α -Amino Acids proteins polypeptide isoelectric point IEP protolysis pK_a value

At physiologic pH=7, 36 \pm 0.01 carboxylic groups **R-COO⁻** negative charged and amino groups **R-NH₃⁺** positive charged. For example, glutamic acid pK_a reference to physiologic pH value smaller as pK_{aR-COO⁻}=4.25<7.36, pK_{aCOO⁻}=2.19<7.36 and for amine is greater as physiologic pH: 9.67=pK_{a-NH₃⁺}>7.36 .

Table shown constants pK_a of four type parallel protolytic equilibria in each amino acid molecule:

acid	\rightleftharpoons base	+H ⁺ ;	Parallel protolytic equilibria number NpKa average isoelectric point
1. R-COOH	\rightleftharpoons R-COO⁻	+H ⁺ ;	and constant pK _a value IEP= pK _a is calculated as
2. R-NH₃⁺	\rightleftharpoons R-NH₂	+H ⁺ ;	IEP= pK _a = (Σ pK _{aR group} + pK _{a-NH₃⁺} + pK _{a-COOH})/NpKa
3. Tyr-phenol-OH	\rightleftharpoons Tyr-phenol-O⁻	+H ⁺ ;	In <i>Ostwald's dilution law</i> calculate pH of solution
4. Cys-SH	\rightleftharpoons Cys-S⁻	+H ⁺	at concentration C logarithm: pH= $\frac{pK_a - \log C}{2}$ =.....

Amino acid and protein at isoelectric point value pH=IEP sum of total overall **ion** charge is zero 0 — acidic charge (+) ————— zero „0” charge **IEP** ————— in basic medium charge minus (-) ————— pH scale **-COOH** & **-NH₃⁺** positive charge **-COO⁻** & **-NH₃⁺** charge is negative **-COO⁻** & **-NH₂**

<http://aris.gusc.lv/ChemFiles/Albumin/1E7GpI.doc> !E7G.pdb;IgG1.pdb

<http://aris.gusc.lv/ChemFiles/ChromoHem/MyoGlobOxDeoxCoBiliverdin/1MBOaaLin153.doc> ! 1MBO.pdb;

Amino Acid	pKa-COOH	pKa-NH ₃ ⁺	pKa R group
Isoleucine	2.36	9.68	
Valine	2.32	9.62	
Leucine	2.36	9.60	
Phenylalanine	1.83	9.13	
Cysteine	1.96	10.28	8.18
Methionine	2.28	9.21	
Alanine	2.34	9.69	
Proline	1.99	10.96	
Glycine	2.34	9.60	
Threonine	2.11	9.62	
Serine	2.21	9.15	
Tryptophan	2.38	9.39	
Tyrosine	2.20	9.11	10.07
Histidine	1.82	9.17	6.00
Aspartate	1.88	9.60	3.65
Glutamate	2.19	9.67	4.25
Asparagine	2.02	8.80	
Glutamine	2.17	9.13	
Lysine	2.18	8.95	10.53
Arginine	2.17	9.04	12.48

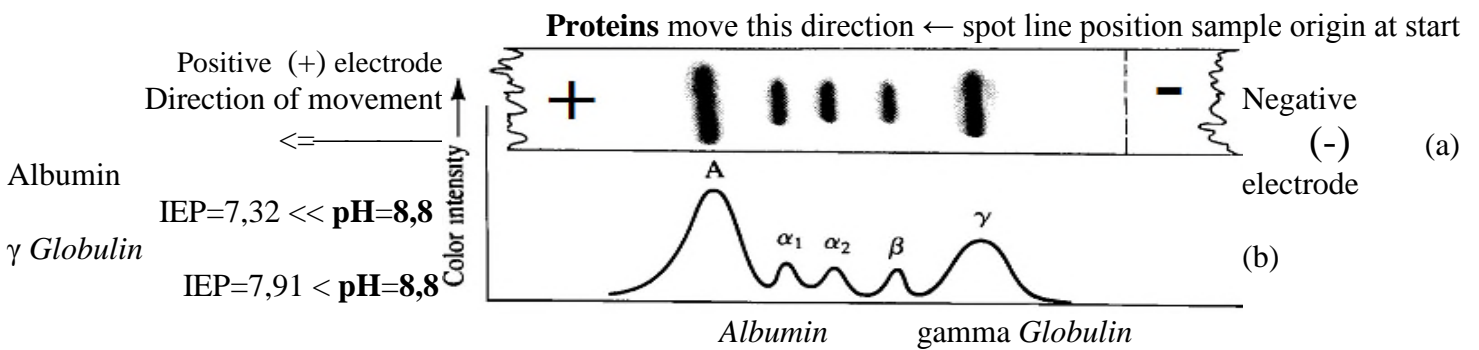
Table5.3 Reginald H. Garrett, Charles M. Grishman, **Biochemistry**, University of Virginia 1995

Myoglobin IEP=7,36 is neutral zero „0” charged molecule, as IEP=7,36 is equal physiologic pH_{blood}=7,36 1MBO.pdb
Albumin molecule E7G.pdb 7,32=IEP 7 fatty acids small (-) charge and 7,40=IEP absent 7 faaty acids (+) positive at physiologic pH=7.36, but gamma *Globulin* IgG1.pdb molecule has positive (+) charge, as at physiologic pH=7.36 is greater IEP=7.91.

Iso electric point IEP=pK_a as well protolytic constant pK_a calculates one of side residues R constants sum Σ pK_{aR side residue} plus pK_{aNterminusNH₃⁺} and plus pK_{aCterminusCOO⁻} sum dividing with number NpKa of acidic groups in molecule IEP=pK_a=(Σ pK_{aR side residue}+pK_{aNterminus}+pK_{aCterminus})/NpKa

Figure Separation of serum proteins by electrophoresis.

- (a) A sample is applied as a narrow line at the origin. After **electrophoresis** at pH **8.8**, the paper is dried and stained.
- (b) A plot of color intensity of each spot. γ *Globulin* moves slower as *Albumin* .

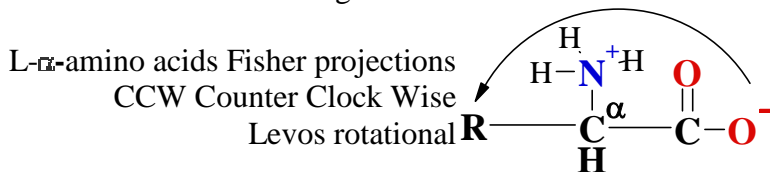
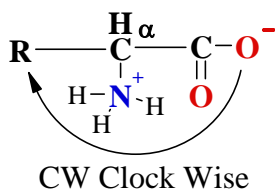


Seleno cysteine, the 21st L- α -Amino Acid

Seleno cysteine is an L- α -amino acid found in a handful of proteins, including certain **peroxidases** and **reductases** where it participates in the catalysis of electron transfer reactions. As its name implies, a selenium **Se** atom replaces the sulfur **S** of its structural analog, cysteine. The pK₃ of seleno cysteine 5.2 is 3 units lower than that of cysteine 8.18. Since seleno cysteine is inserted into polypeptides during translation, it is commonly referred to as the "21st amino acid." However, like the other 20 genetically encoded amino acids, seleno cysteine is specified by a simple three-letter codon **UGA** (see class 16 week Nucleo proteins tRNA 62 codons).

Draw fisher projections for 3D molecules Harper's Biochemistry table-3 on 15-16 page wrong D- to L- α -amino acids to fill table right side: <http://aris.gusc.lv/06Daugavpils/Research/Amineac20L.doc>, as L- α -amino acids! Santa Barbara University 3D molecules: <http://aris.gusc.lv/ChemFiles/MCDB108A/tw-amn/aasframes.htm>.

Harper's Biochemistry Illustrated Table 3-1 shows D-amino acids Fisher projections, which are wrong for human organism proteins. Your task is in **Table** right side to show L- α -amino acids



L- α -amino acids Fisher projections CCW Counter Clock

Table The 20 common L- α -amino acids found in protein.

Physiologic pH=7.36 .

Protein-derived Amino Acids with aliphatic side chains left side	Name	Symbol	Show Fisher projection Structural Formula
1	Glycine	Gly [G]	
2	Alanine	Ala [A]	
3	Valine	Val [V]	
4	Leucine	Leu [L]	
5	Isoleucine	Ile [I]	
With side chains containing hydroxyl (—OH) groups			
left side			
6	Serine	Ser [S]	
7	Threonine	Thr [T]	
18	Tyrosine	Tyr [Y]	Shown below ↓.
With side chains containing Sulfur atoms (—S— ; —SH)			
left side			
8	Cysteine	Cys [C]	
9	Methionine	Met [M]	

Table The 20 common L- α -amino acids found in protein. .

Physiologic pH=7.36 .

	Name	Symbol	Show Fisher projection Structural Formula
With side chains containing Acidic ($-\text{COO}^-$) groups or their Amides ($-\text{CO}-\text{NH}_2$)			
left side	Aspartate		
10	Aspartic acid salt	Asp [D]	
11	Asparagine	Asn [N]	
12	Glutamate	Glu [E]	
13	Glutamine	Gln [Q]	
With side chains containing Basic ($-\text{NH}_n^{(+)}$) Groups			
left side			
14	Arginin	Arg [R]	
15	Lysine	Lys [K]	
16	Histidine	His [H]	
Containing Aromatic Rings			
left side			
16	Histidine	His [H]	Shown above ↑
17	Phenylalanine	Phe [F]	
18	Tyrosine	Tyr [Y]	
19	Tryptophan	Trp [W]	
Imino Acid			
20	Proline	Pro [P]	