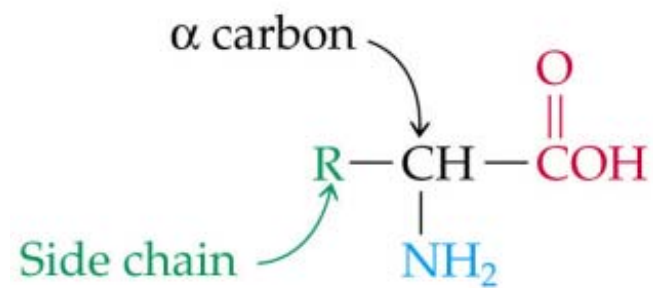
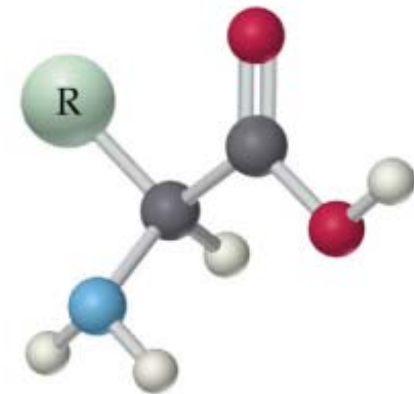


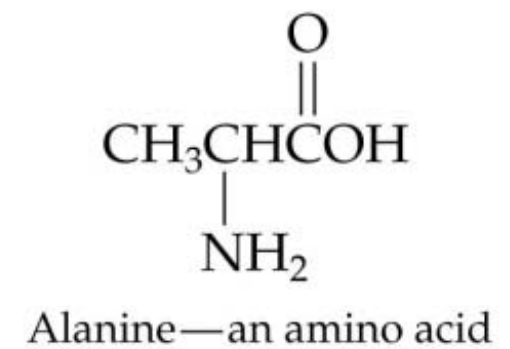
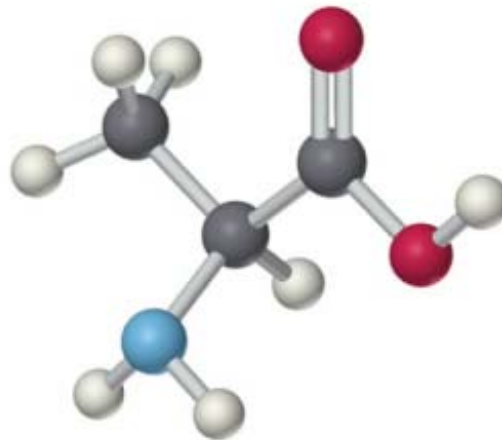
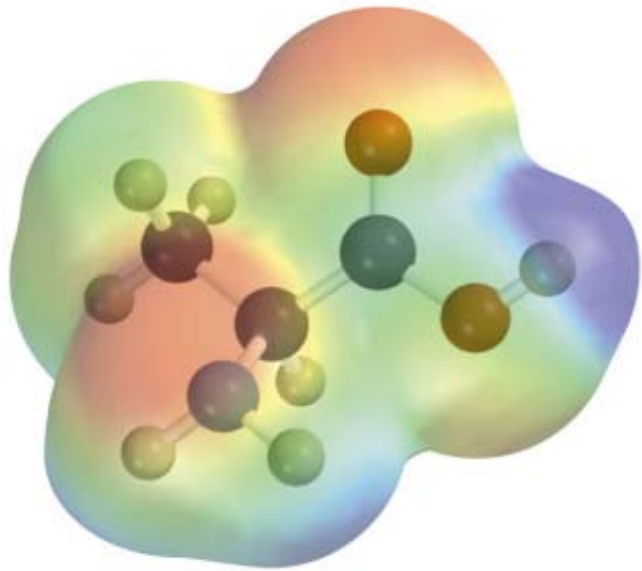
Prinzipielle Struktur von alpha-Aminosäuren



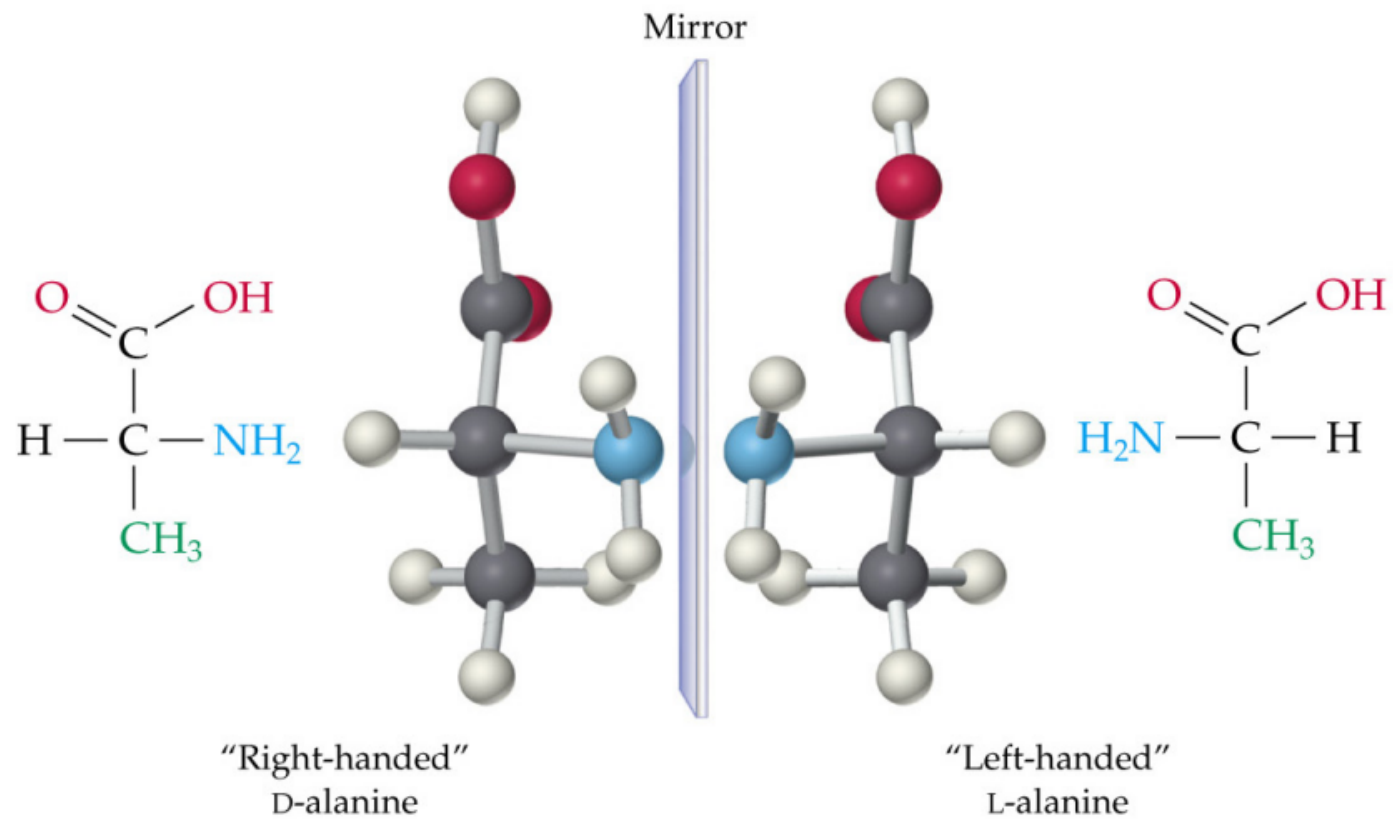
Generalized structure of
an α -amino acid



Die einfachste optisch aktive alpha-Aminosäure: Alanin

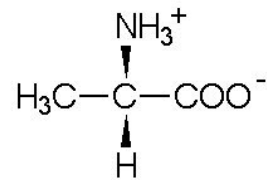


D- und L-Form von Alanin (Fischer-Projektion)

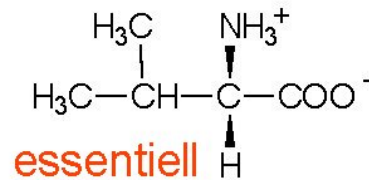


Aminosäuren mit unpolaren Resten = Seitengruppen

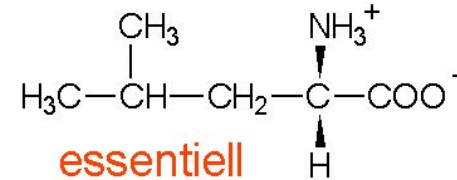
A. mit unpolaren Seitengruppen:



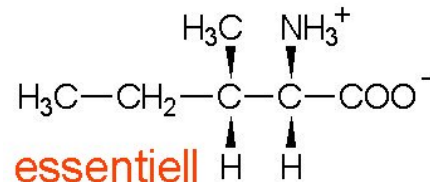
L-Alanin
(Ala, A)



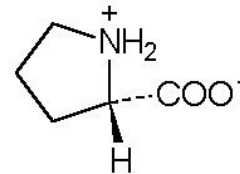
essentiell
L-Valin
(Val, V)



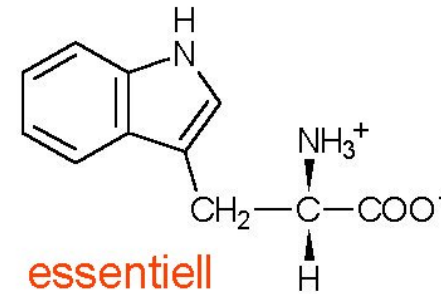
essentiell
L-Leucin
(Leu, L)



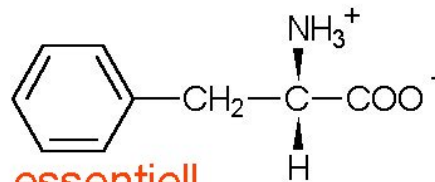
essentiell
L-Isoleucin
(Ile, I)



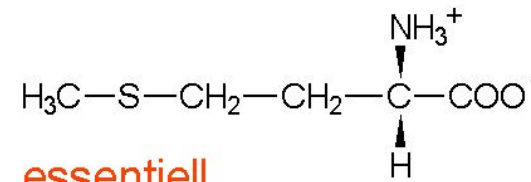
L-Prolin
(Pro, P)



essentiell
L-Tryptophan
(Trp, W)



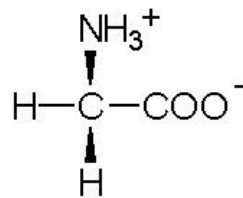
essentiell
L-Phenylalanin
(Phe, F)



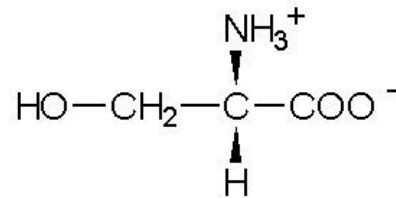
essentiell
L-Methionin
(Met, M)

Aminosäuren mit polaren ungeladenen Resten = Seitengruppen

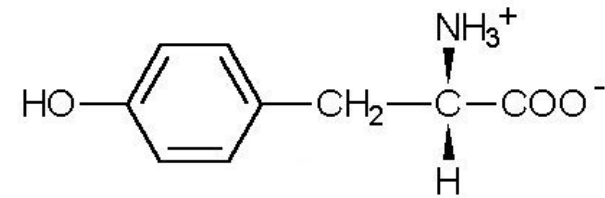
A. mit polaren ungeladenen Seitengruppen:



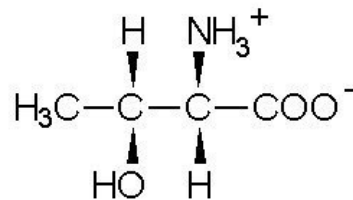
Glycin
(Gly,G)



L-Serin
(Ser,S)

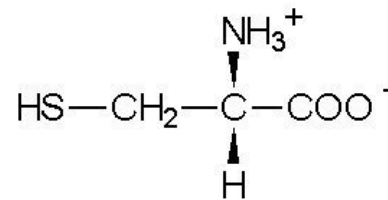


L-Tyrosin
(Tyr,Y)

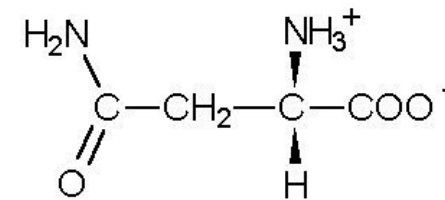


L-Threonin
(Thr,T)

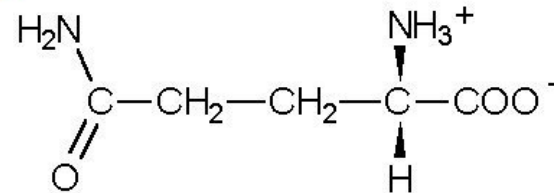
essentiell



L-Cystein
(Cys,C)



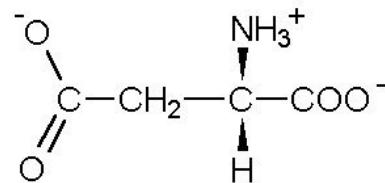
L-Asparagin
(Asn,N)



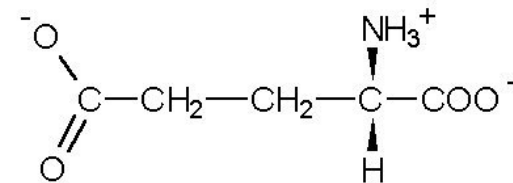
L-Glutamin
(Gln,Q)

Saure Aminosäuren (oben) und basische Aminosäuren (unten)

Saure A. (besitzen neg. geladene Seitengruppen):

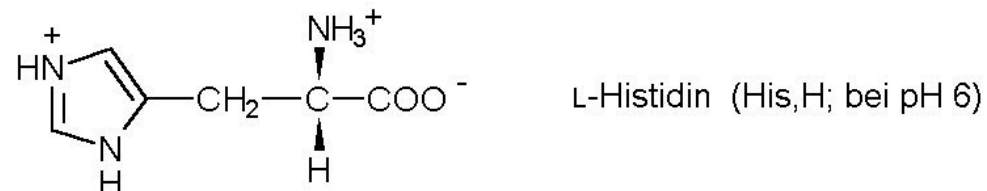
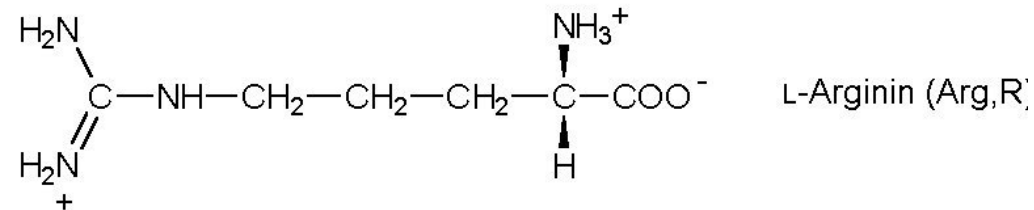
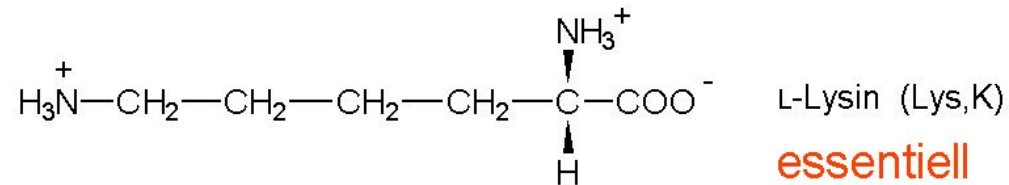


L-Aspartat
(L-Asparaginsäure; Asp, D)

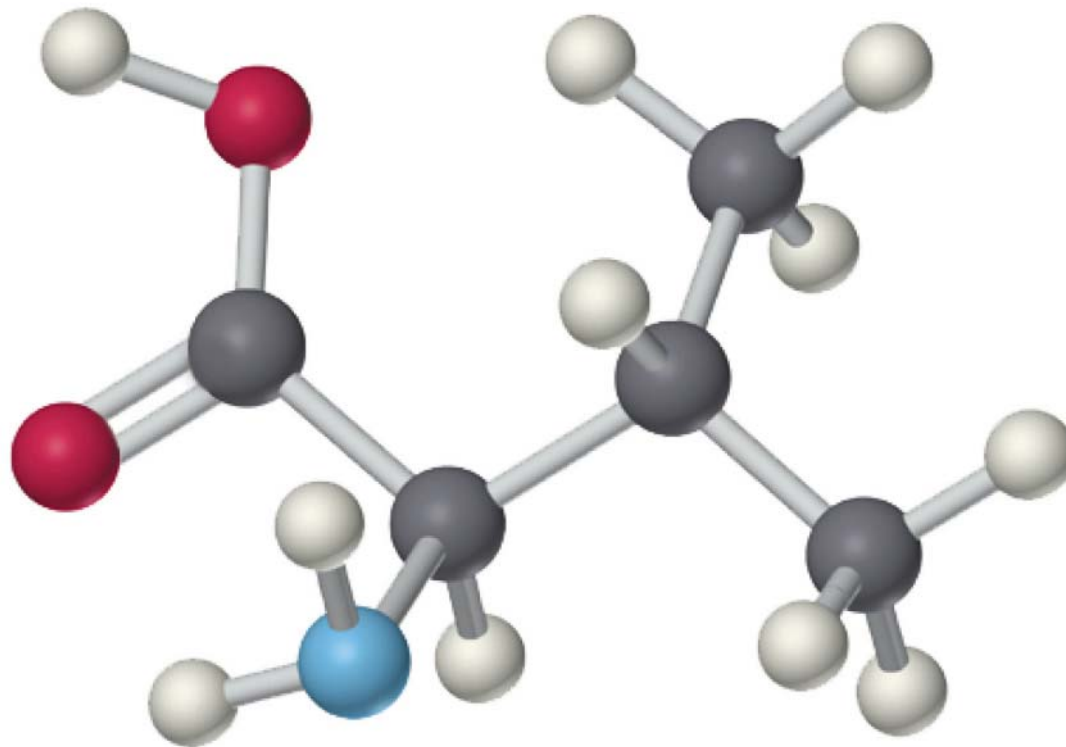


L-Glutamat
(L-Glutaminsäure; Glu, E)

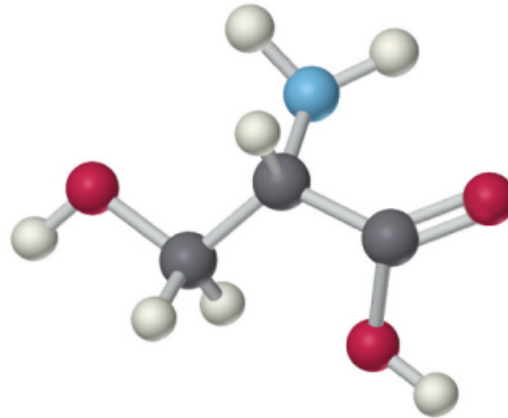
Basische A. (besitzen pos. geladene Seitengruppen):



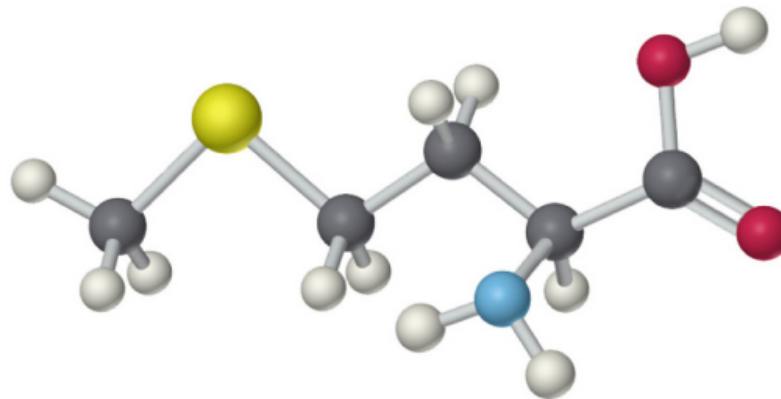
Um welche Aminosäure handelt es sich?



Und um welche Aminosäuren handelt es sich hier?

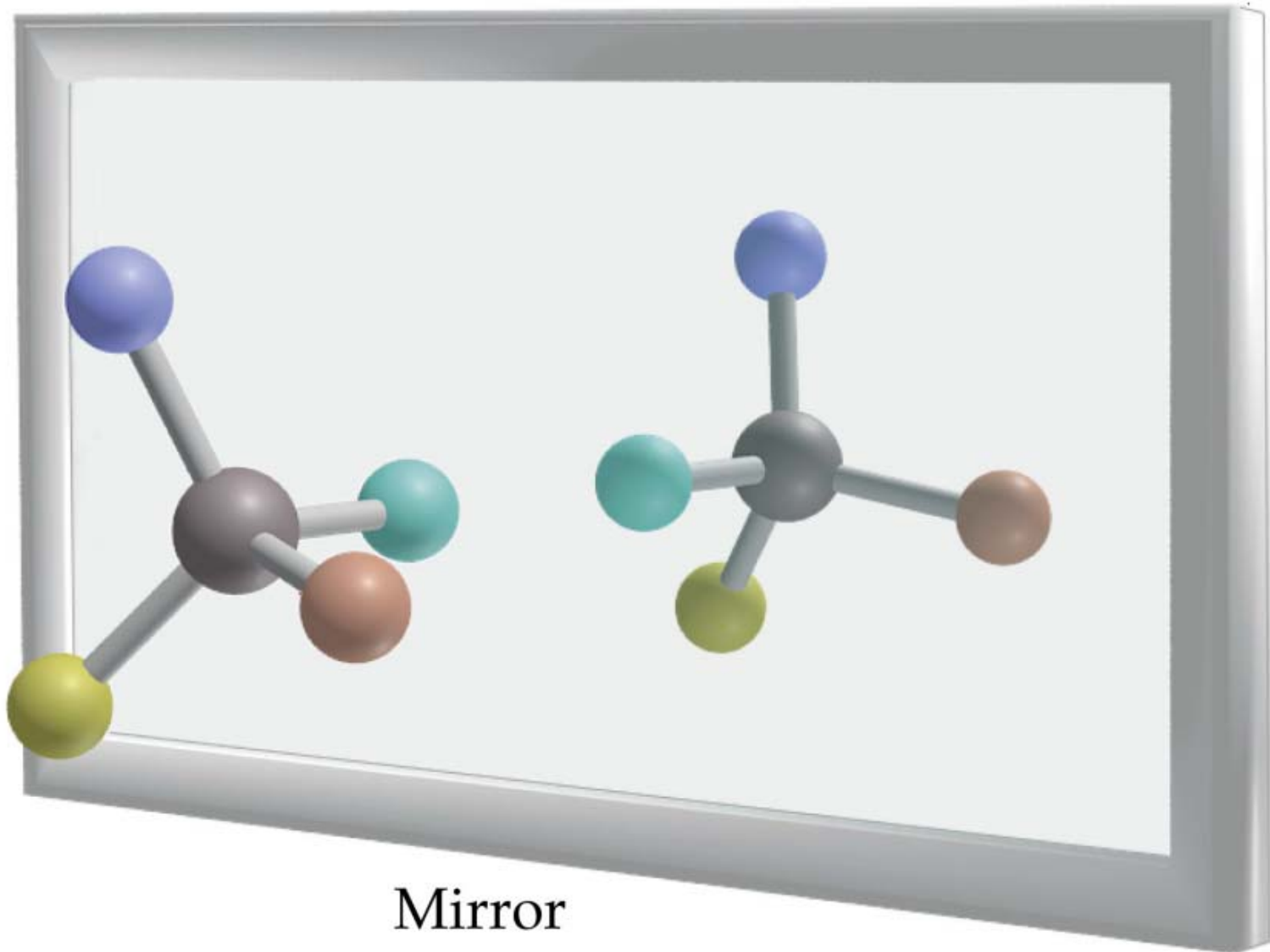


(a)

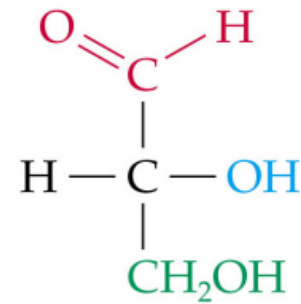
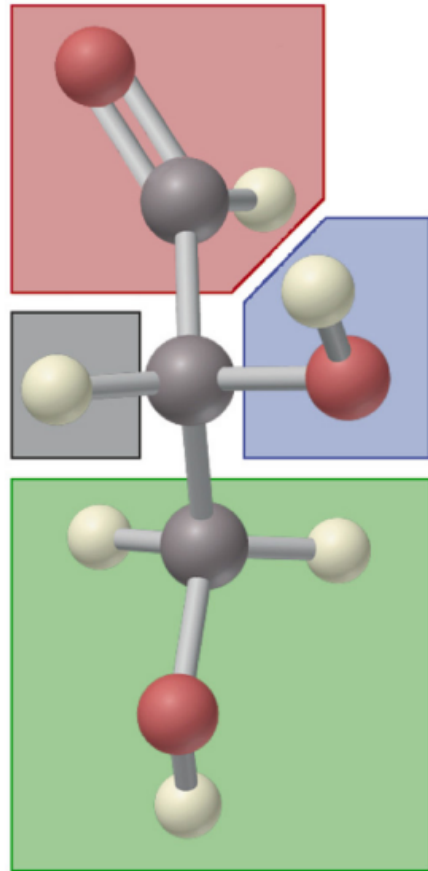


(b)

Bild und Spiegelbild



Glycerinaldehyd: Basis für die Zuordnung D- bzw. L- Form bei einer chiralen Verbindung (mit asymmetrischem C-Atom)

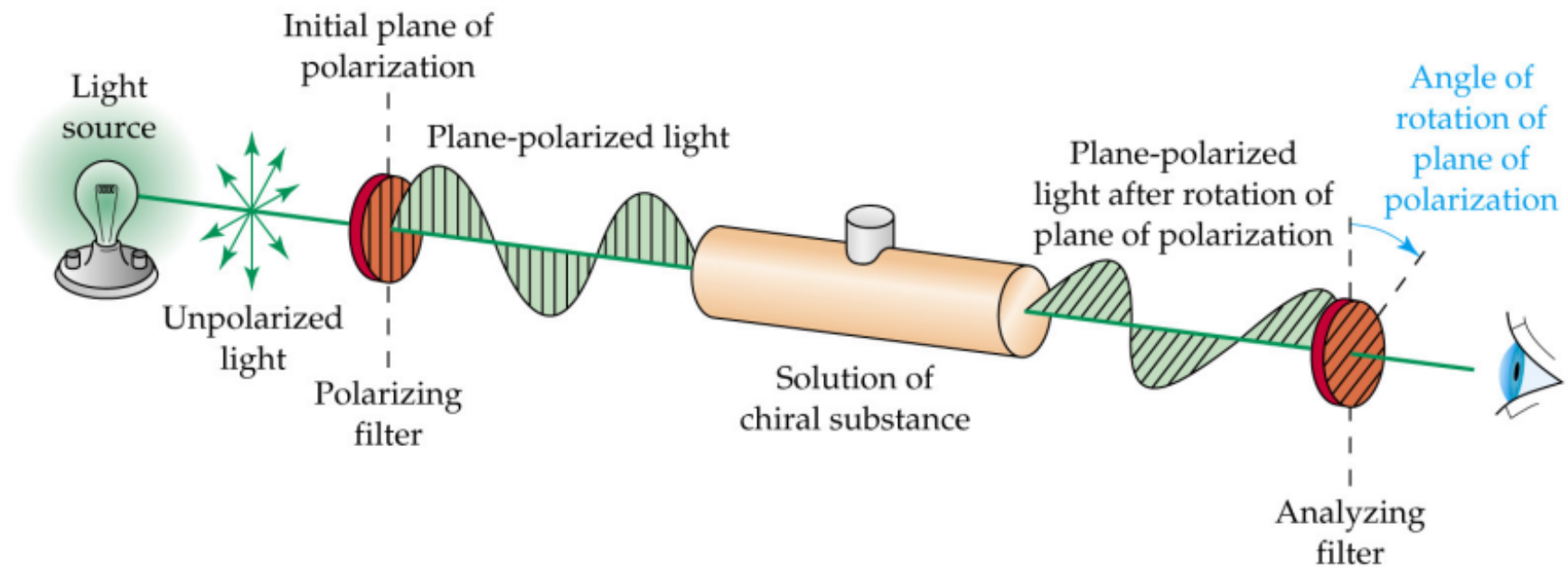


Groups bonded to C2

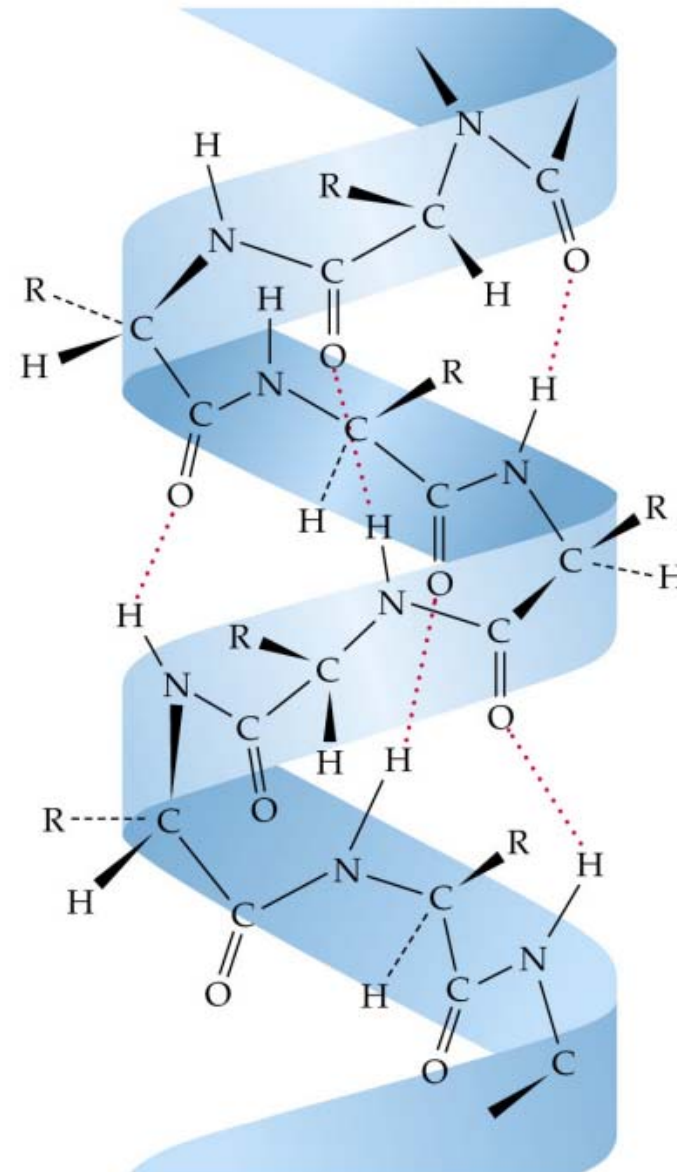
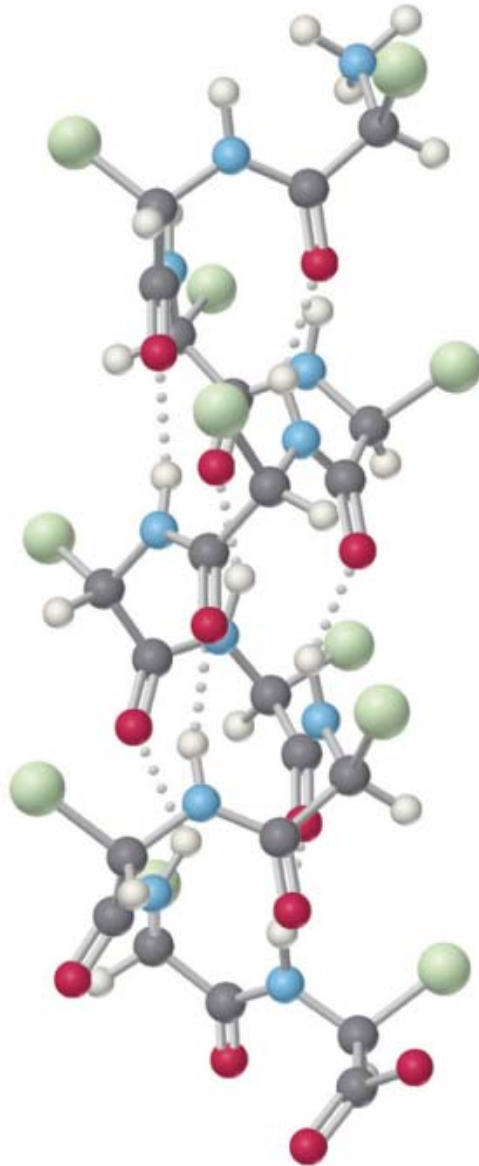
- 1.—CHO
- 2.—H
- 3.—OH
- 4.—CH₂OH

D-Glyceraldehyde

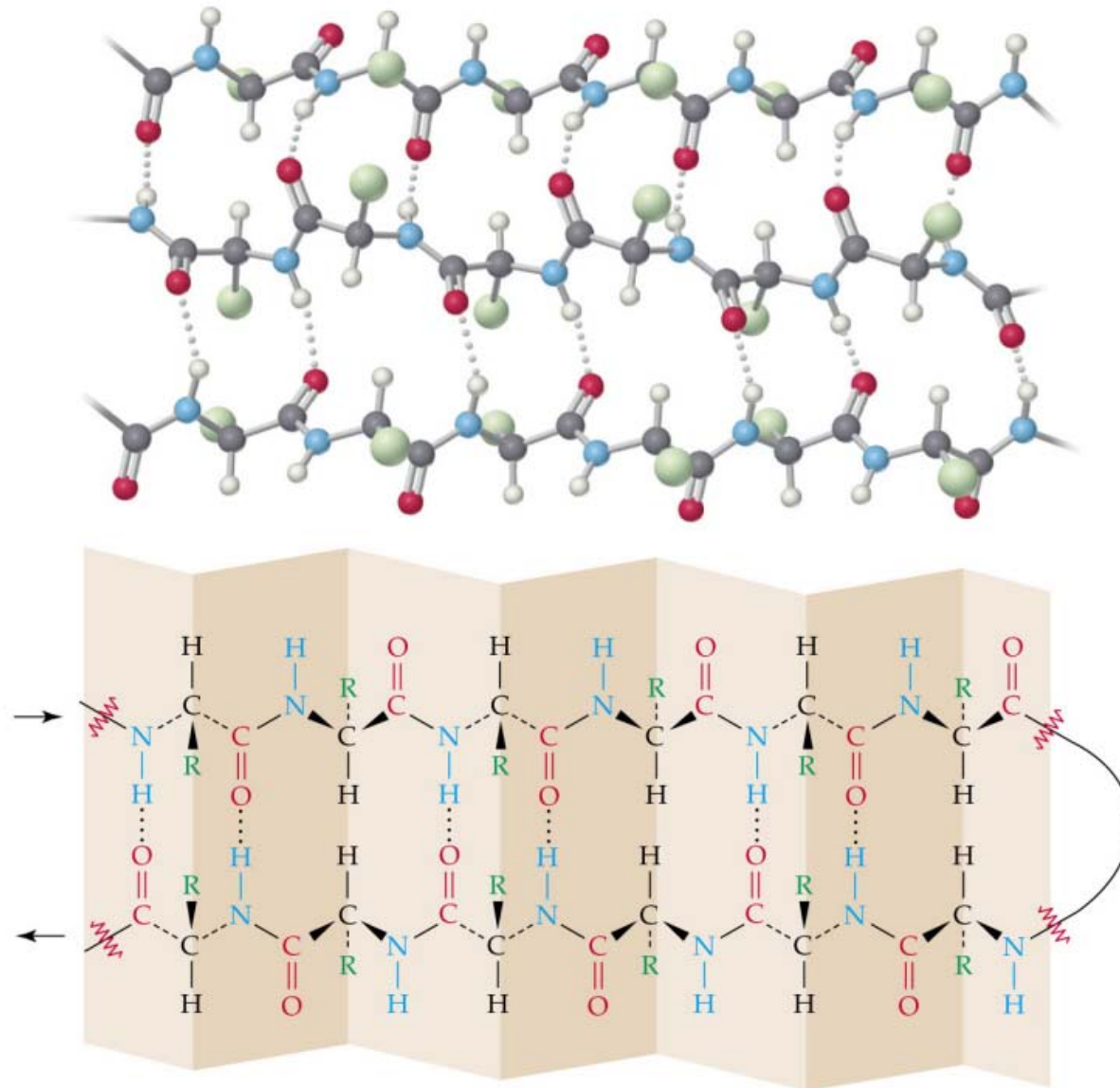
Prinzip der Polarimetrie



Sekundärstruktur von Proteinen

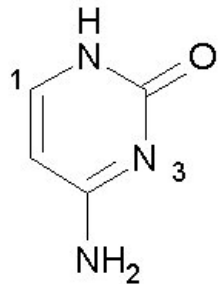


Sekundärstruktur von Proteinen: „Faltblattstruktur“

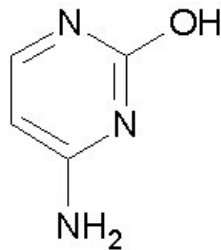


Die Pyrimidin-Basen

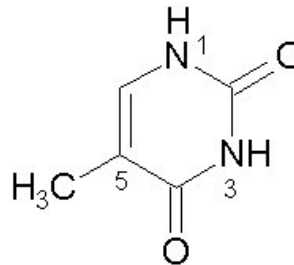
(Uracil in RNA, Thymidin in DNA)



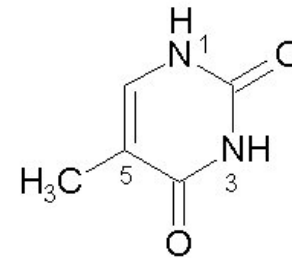
bzw.



Cytosin

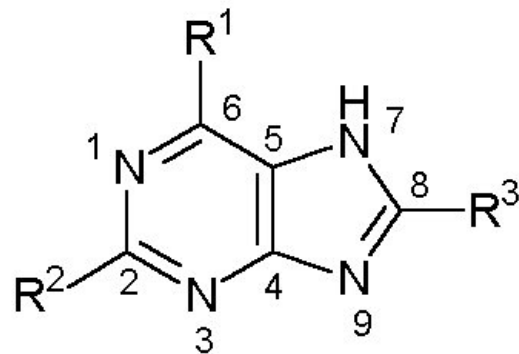


Uracil



Thymidin

Die Purin-Basen



Purin : $R^1 = R^2 = R^3 = H$

Adenin : $R^1 = NH_2, R^2 = R^3 = H$

Guanin : $R^1 = OH, R^2 = NH_2, R^3 = H$

Harnsäure : $R^1 = R^2 = R^3 = OH$

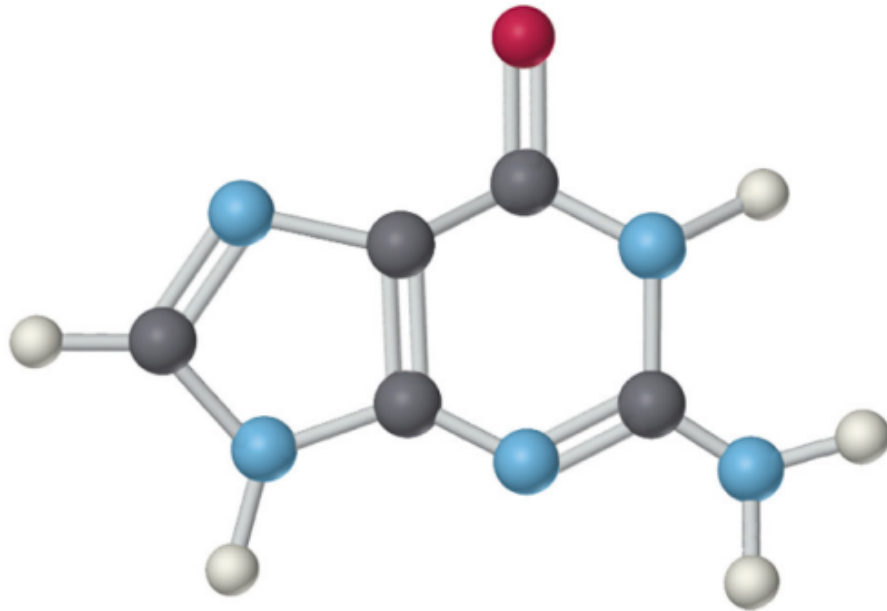
Hypoxanthin : $R^1 = OH, R^2 = R^3 = H$

6-Purinthiol : $R^1 = SH, R^2 = R^3 = H$

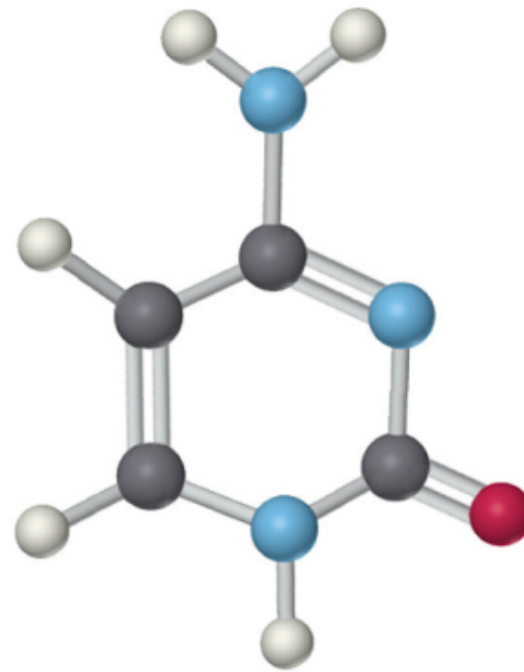
6-Thioguanin : $R^1 = SH, R^2 = NH_2, R^3 = H$

Xanthin : $R^1 = R^2 = OH, R^3 = H$

Um welche Purin-Base (a) bzw. Pyrimidin-Base (b)
handelt es sich?

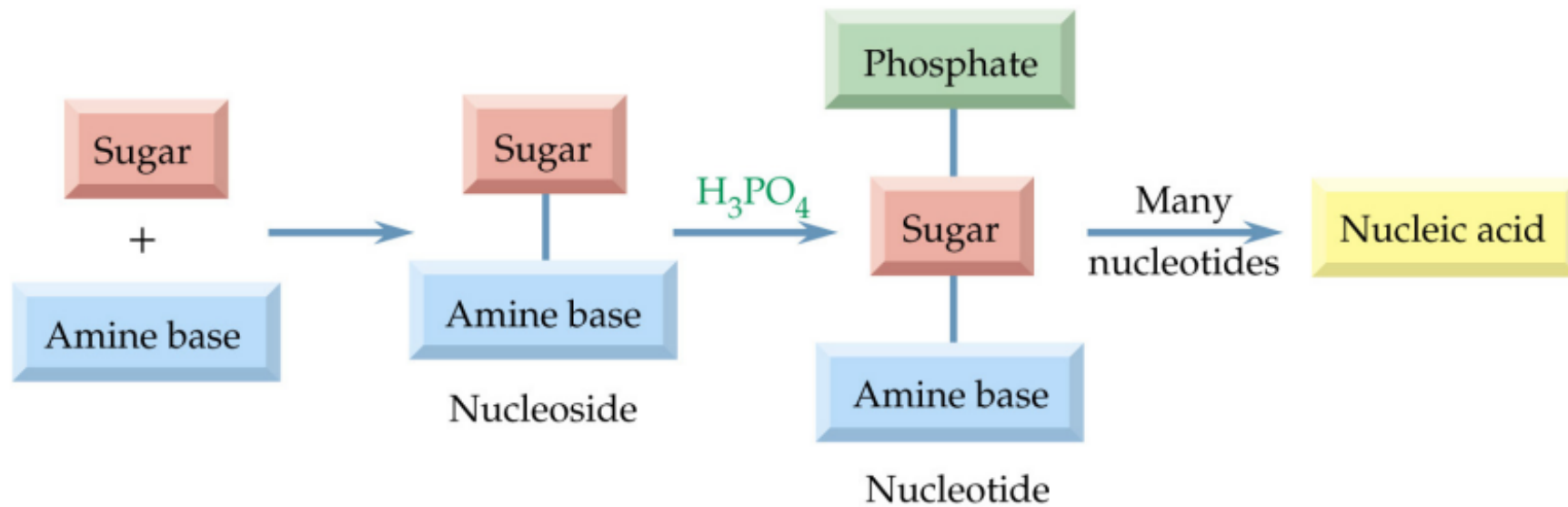


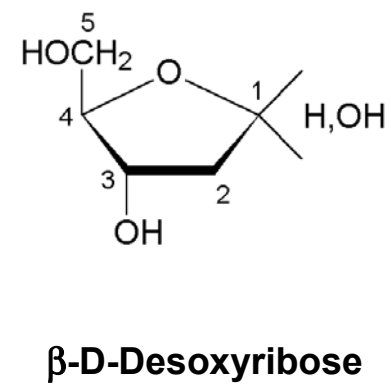
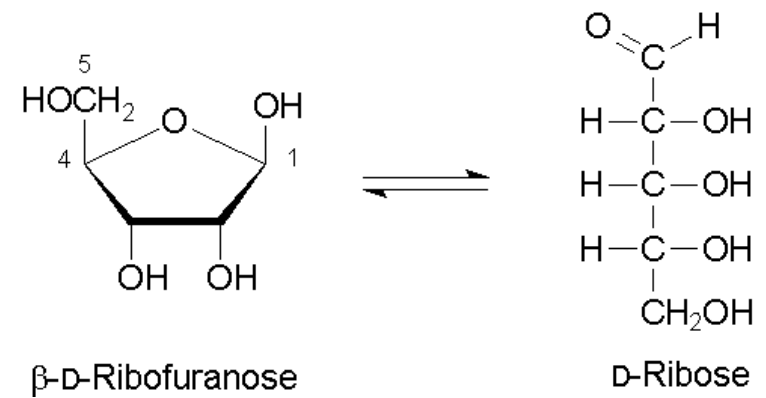
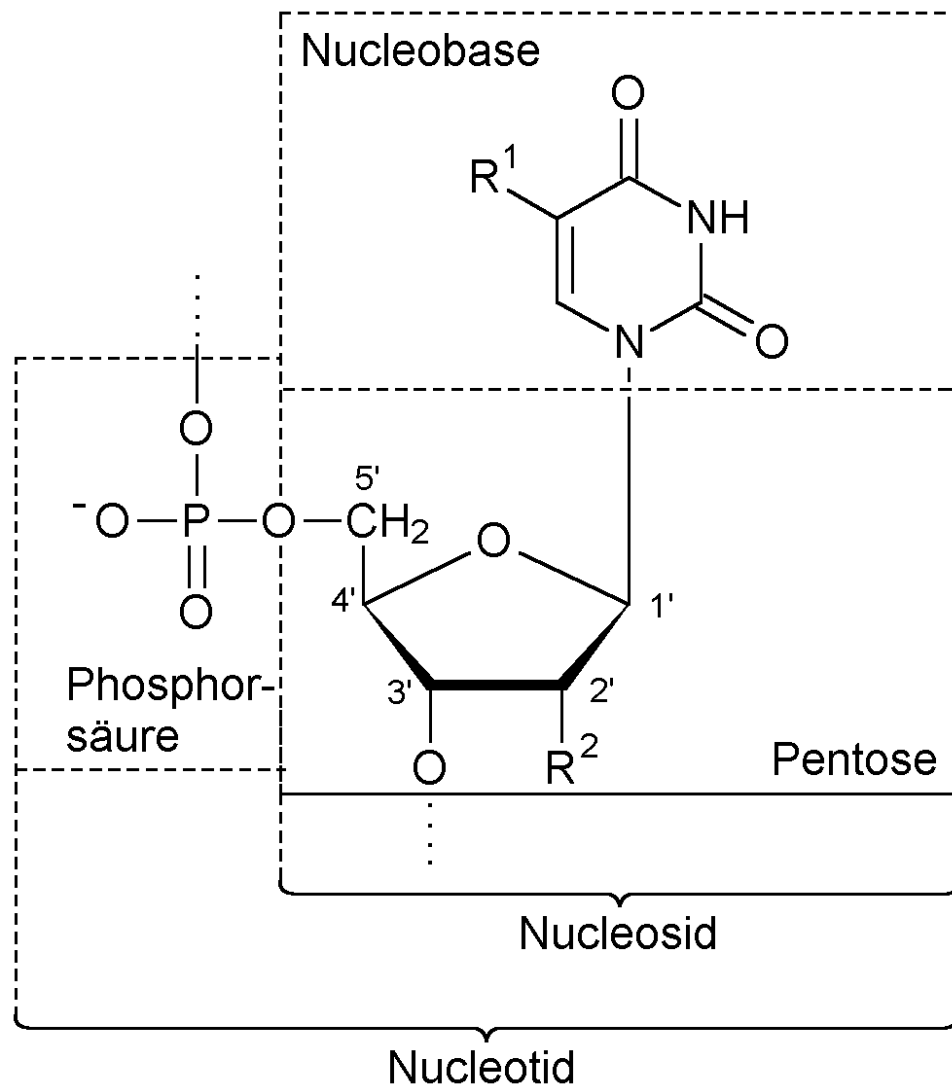
(a)



(b)

Bausteine der Bildung von Nucleinsäuren

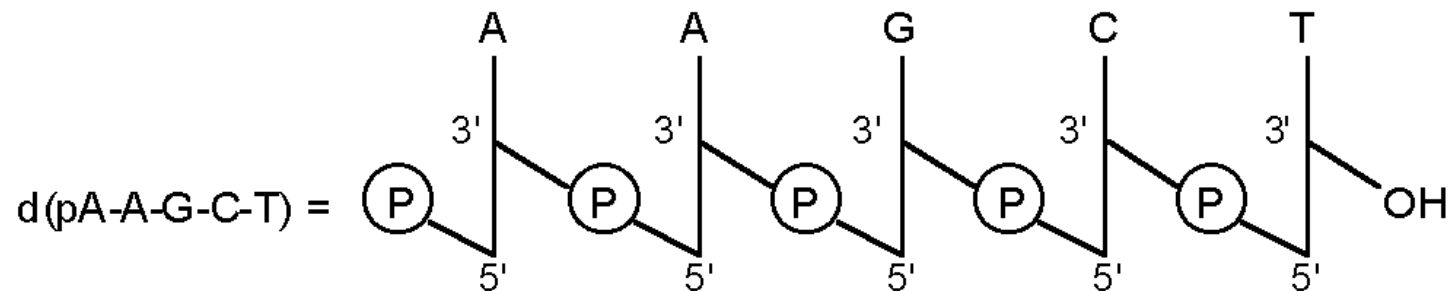




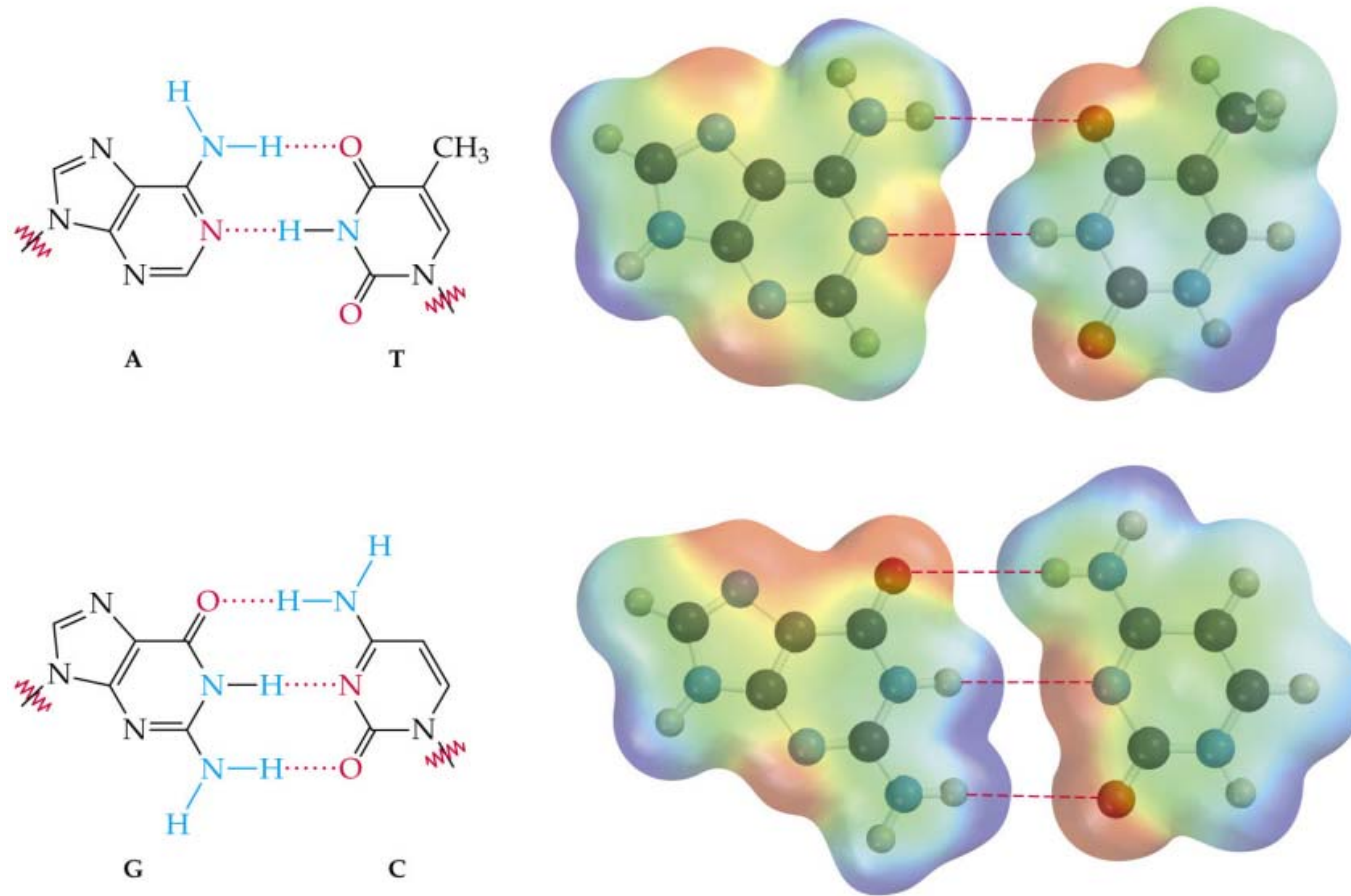
Kurze DNA-Sequenz:

Desoxyribose und Phosphat bilden das „Rückgrat“

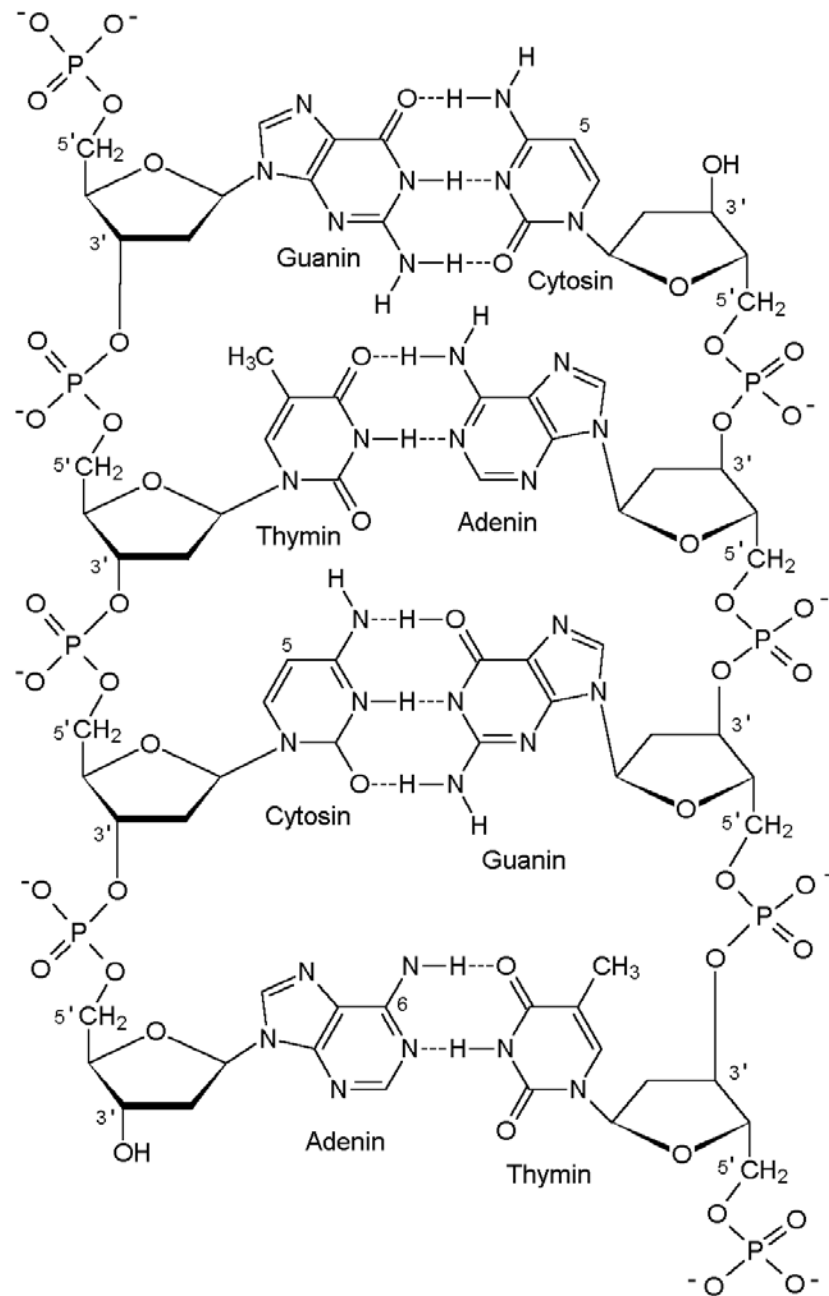
In der Abfolge der Basen A, C, G, T ist die genetische Information verschlüsselt



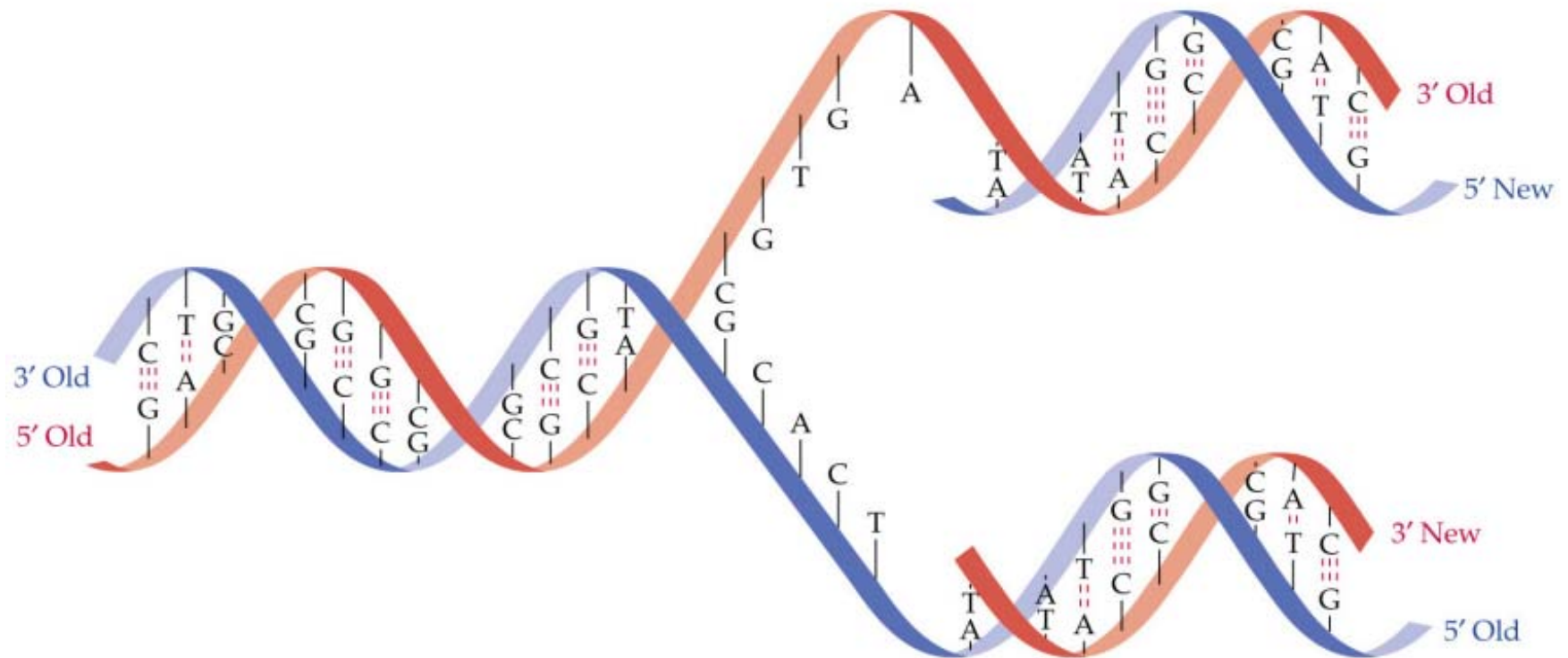
Die Basen C-G bzw. A-T bilden jeweils Paare
(Doppelstrang-Struktur)



Kurzer Ausschnitt aus
einem Doppelstrang
einer DNA



Duplikation (Replikation) eines DNA-Strangs



Replikation: siehe vorherige Folie = Verdoppelung der DNA

Transkription: RNA-Synthese aus dem Code der DNA (vermittelt durch RNA-Polymerase)

Translation: Übersetzung des Codes der RNA (Messenger-RNA) in Aminosäuren bzw. Peptide (Proteine)

