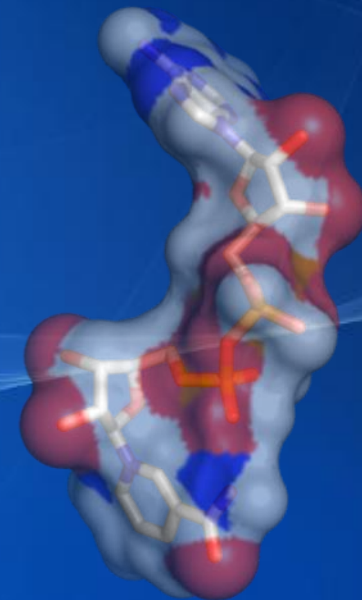
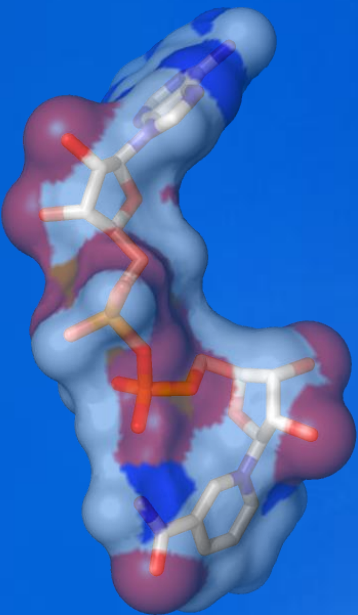


# FFI0750 – Biologia Molecular Estrutural

Prof. Rafael V. C. Guido  
[rvcguido@ifsc.usp.br](mailto:rvcguido@ifsc.usp.br)

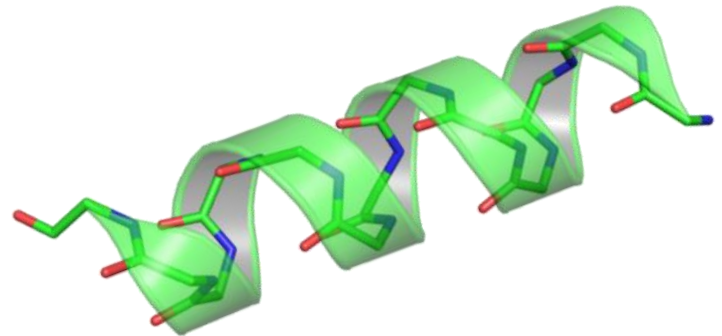
Aula 05

Bacharelado em Ciências Físicas e Biomoleculares  
Instituto de Física de São Carlos - USP

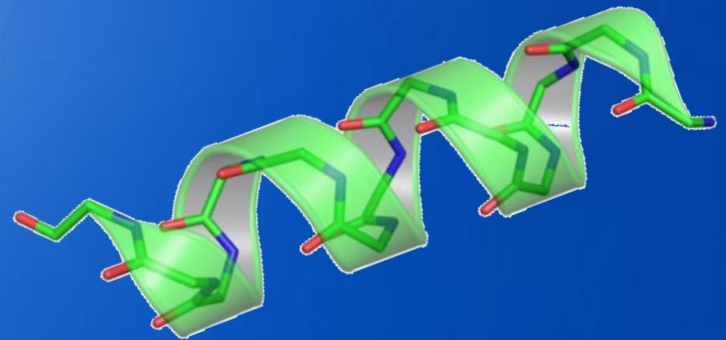




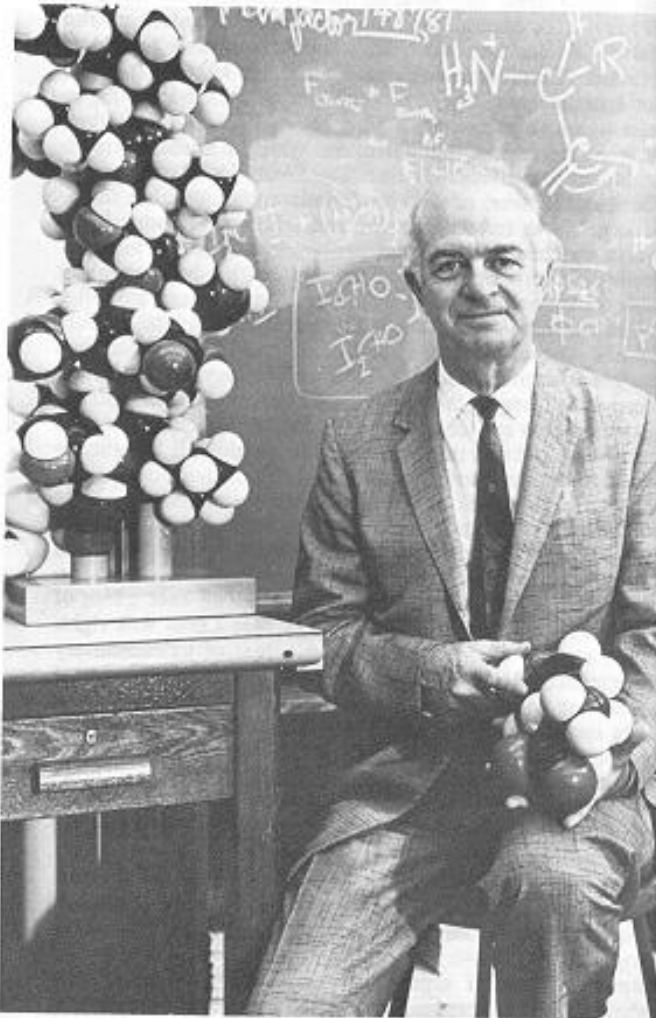
- Hélices em proteínas
  - Aspectos históricos
  - Características estruturais



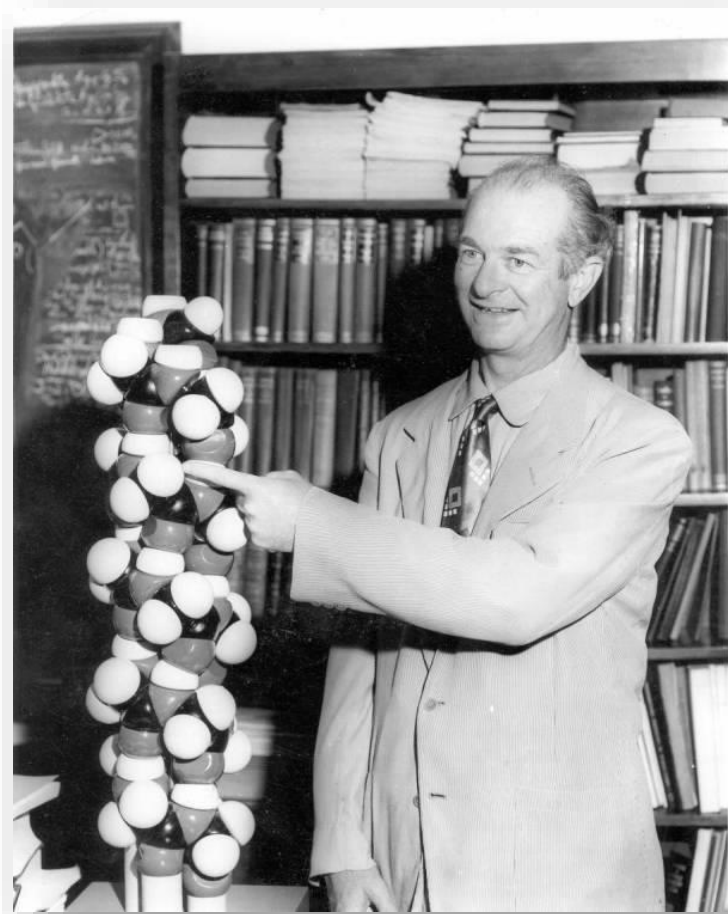
# Estrutura de Proteínas – Alfa-hélices



# Linus Pauli e alfa-hélice (1951)



Linus Pauling with his atomic models.







# Jonh Kendrew – Mioglobina (1958)

662

NATURE

March 8, 1958 VOL. 181

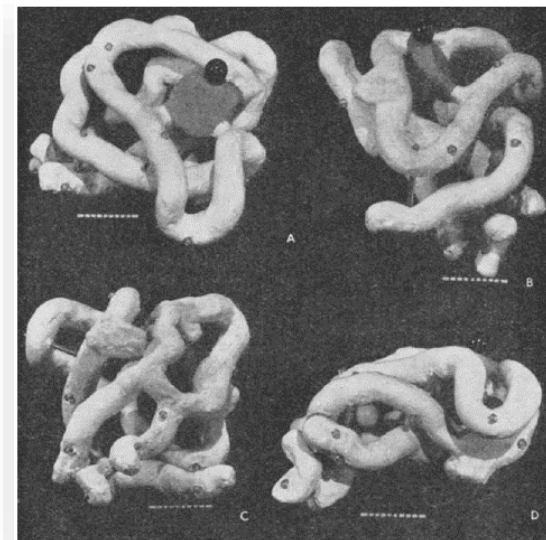
## A THREE-DIMENSIONAL MODEL OF THE MYOGLOBIN MOLECULE OBTAINED BY X-RAY ANALYSIS

By Drs. J. C. KENDREW, G. BODO, H. M. DINTZIS, R. G. PARRISH and H. WYCKOFF  
Medical Research Council Unit for Molecular Biology, Cavendish Laboratory, Cambridge

AND

D. C. PHILLIPS

Davy Faraday Laboratory, The Royal Institution, London



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NATURE

February 13, 1960 VOL. 185

## STRUCTURE OF MYOGLOBIN

A THREE-DIMENSIONAL FOURIER SYNTHESIS AT 2 Å. RESOLUTION

By Drs. J. C. KENDREW, R. E. DICKERSON, B. E. STRANDBERG, R. G. HART  
and D. R. DAVIES\*

Medical Research Council Unit for Molecular Biology, Cavendish Laboratory, Cambridge

AND

D. C. PHILLIPS and V. C. SHORE

Davy Faraday Laboratory, The Royal Institution, London





# Alfa-hélice: Definição

Alfa hélice ( $\alpha$ -hélice) é uma estrutura secundária cuja conformação em espiral posiciona o grupo C=O do resíduo  $n$  da cadeia polipeptídica em geometria favorável para receber ligação de hidrogênio do grupo NH do resíduo localizado quatro posições a frente ( $n + 4$ ) na cadeia polipeptídica.

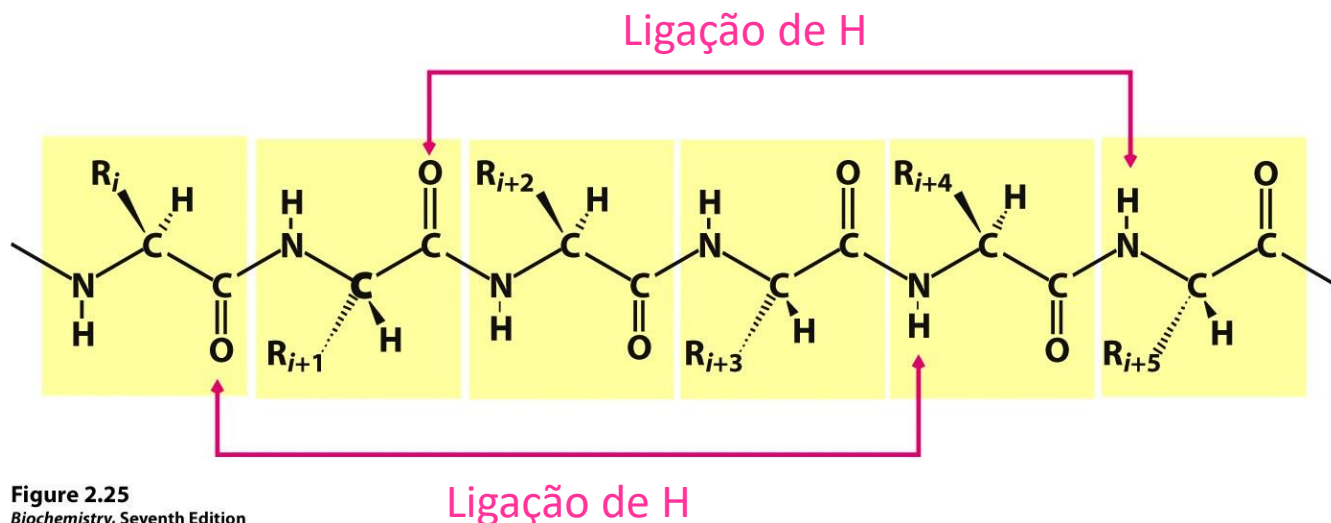
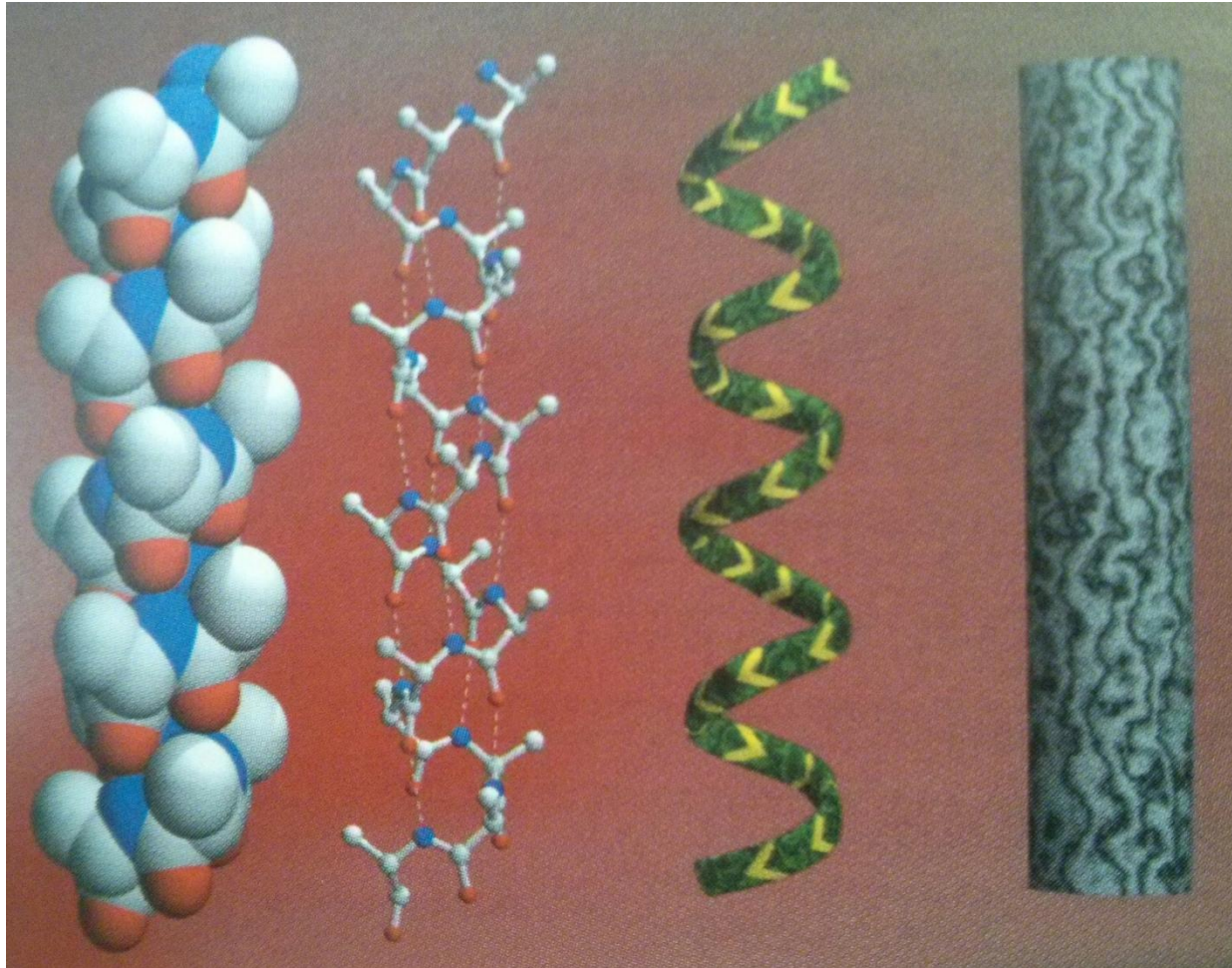


Figure 2.25  
Biochemistry, Seventh Edition  
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# Alfa-hélice

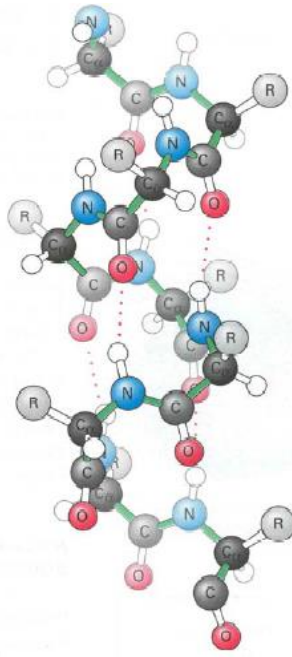
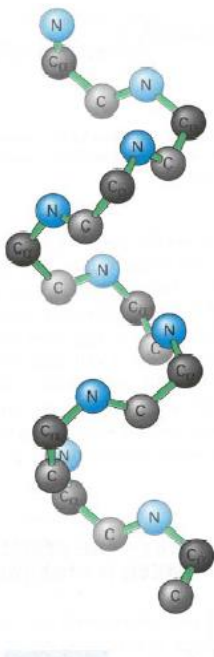
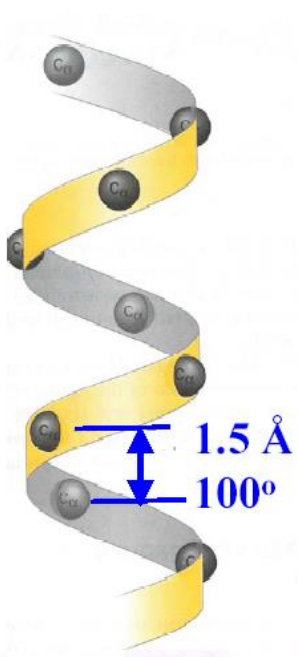




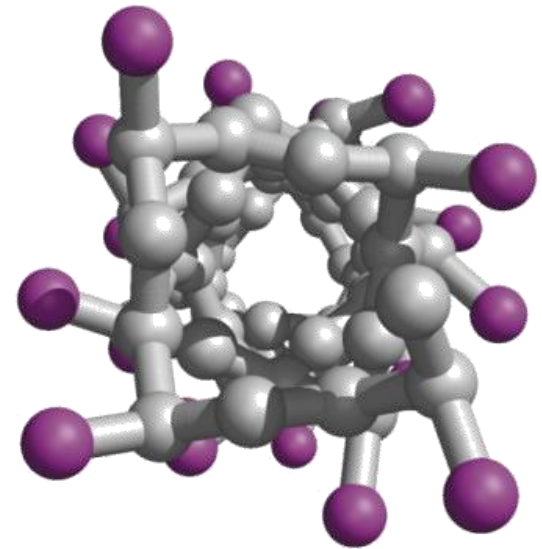


# Estrutura secundária – $\alpha$ hélices

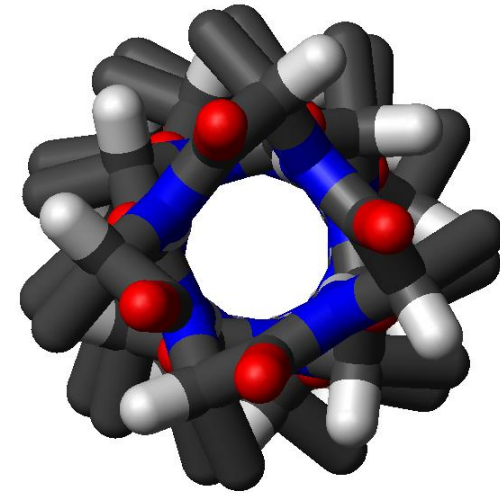
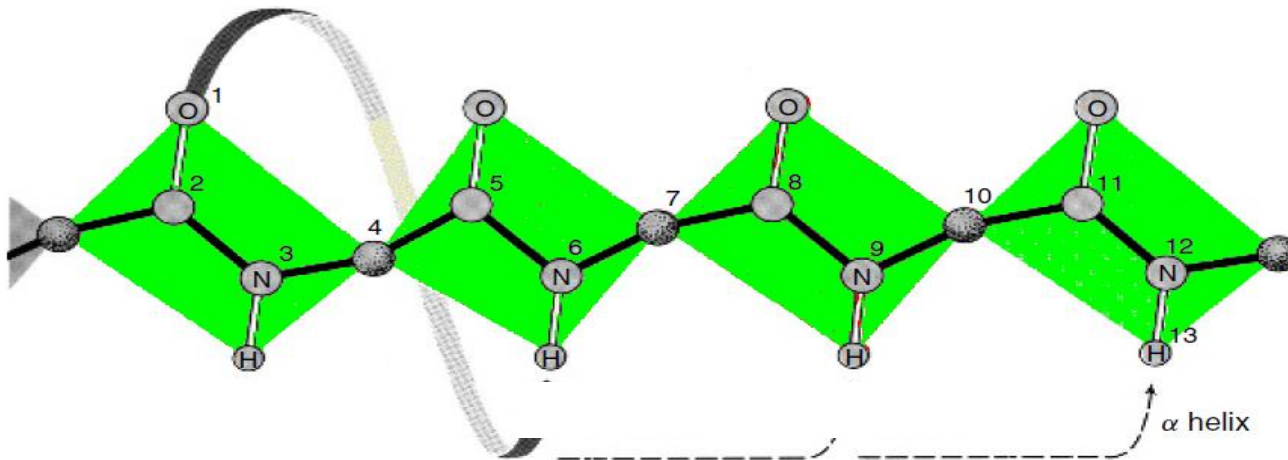
- Estrutura regular e energeticamente estável, pois todas as ligações de H possíveis entre a cadeia principal são estabelecidas ao longo do eixo da hélice;
- Em proteínas globulares o comprimento das  $\alpha$ -hélices varia consideravelmente (desde 4,5 até > 40 resíduos)
- O comprimento médio das  $\alpha$ -hélices é 10-15 resíduos (12-22 Å)



$$\psi = -47^\circ$$
$$\phi = -57^\circ$$







## α hélice

Ligação de H =  $n - n+4$

Período = 3,6 aa/volta

**Mão direita**

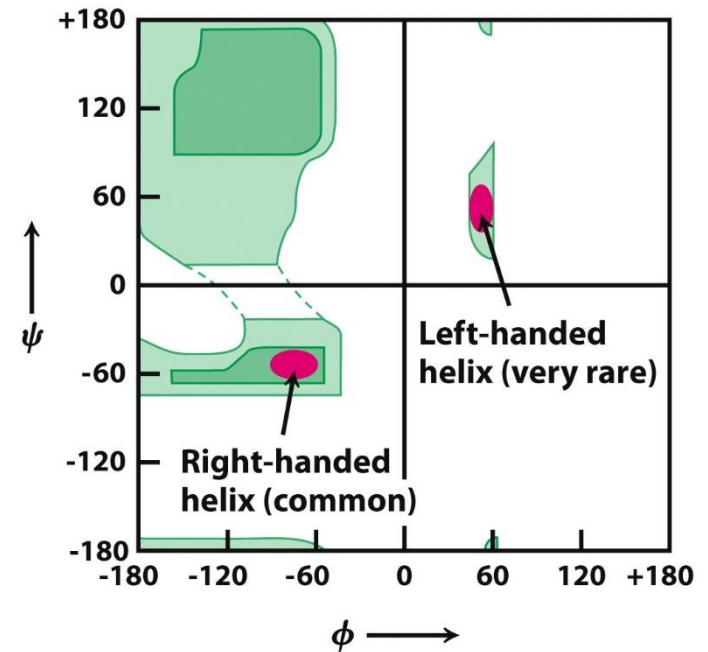
**Mão Esquerda**

$\Phi = -57^\circ$

$\Phi = 60^\circ$

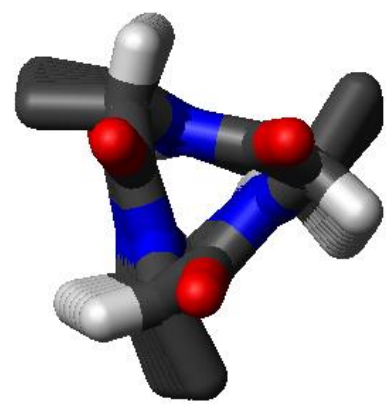
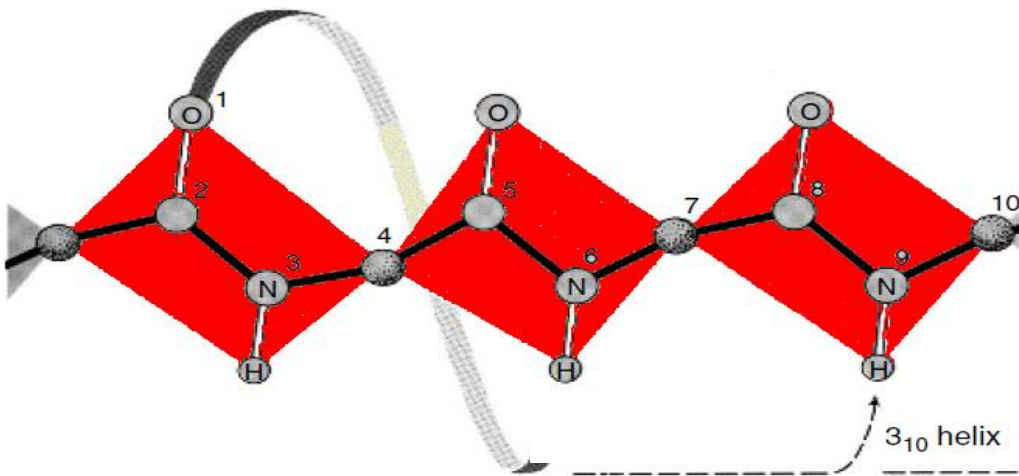
$\Psi = -47^\circ$

$\Psi = 60^\circ$





# Hélice 3<sub>10</sub>



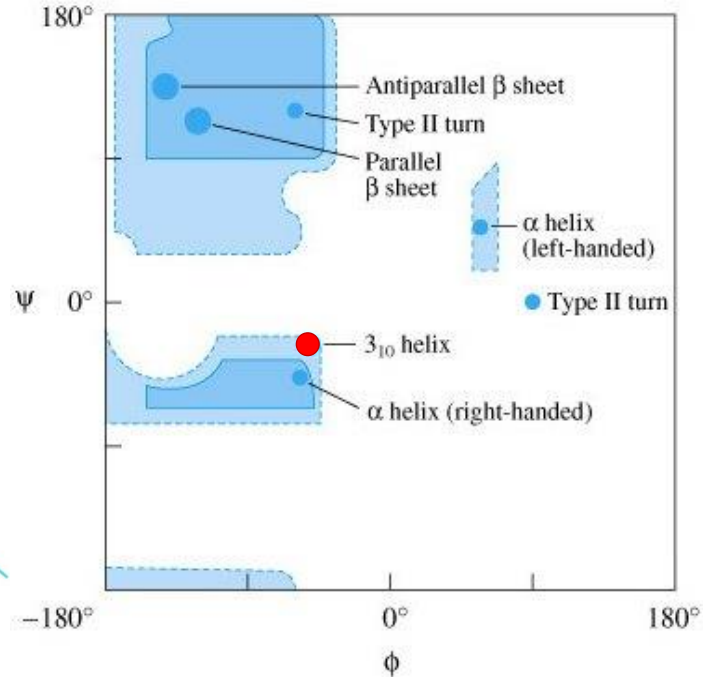
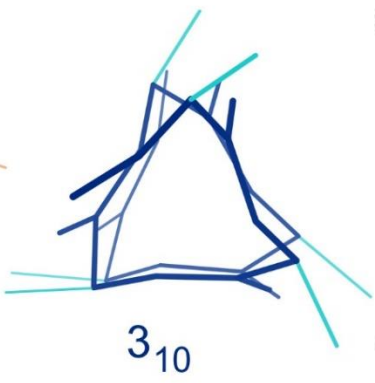
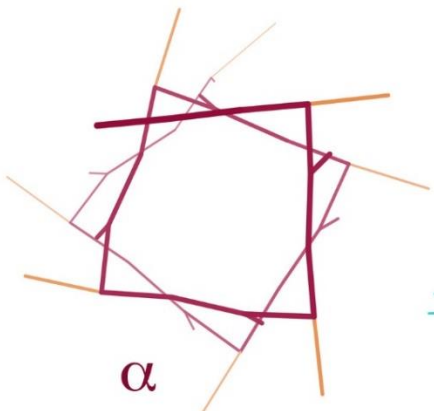
## hélice 3<sub>10</sub>

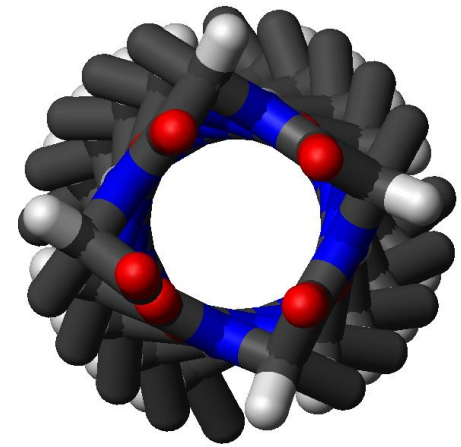
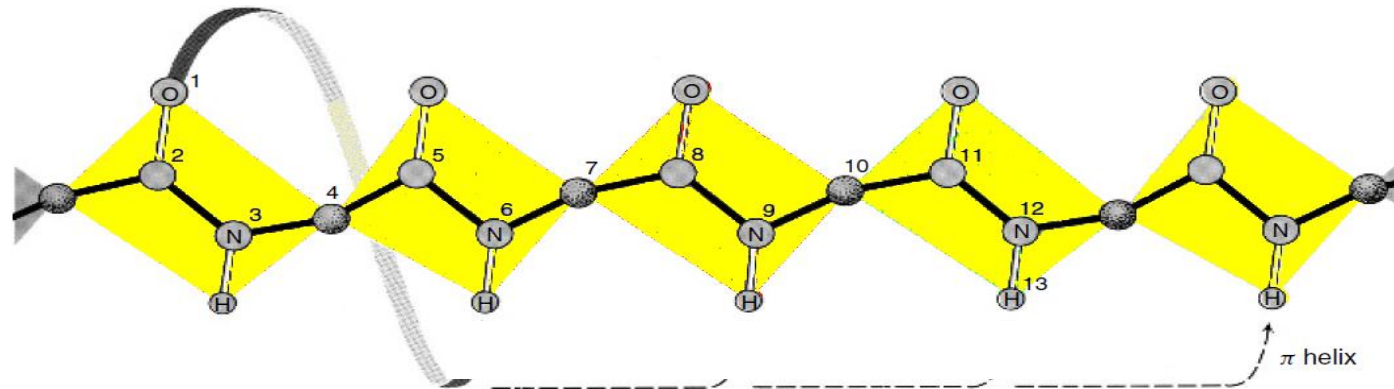
Ligação de H =  $n - n+3$

Período = 3,0 aa/volta

$\Phi = -49^\circ$

$\Psi = -26^\circ$





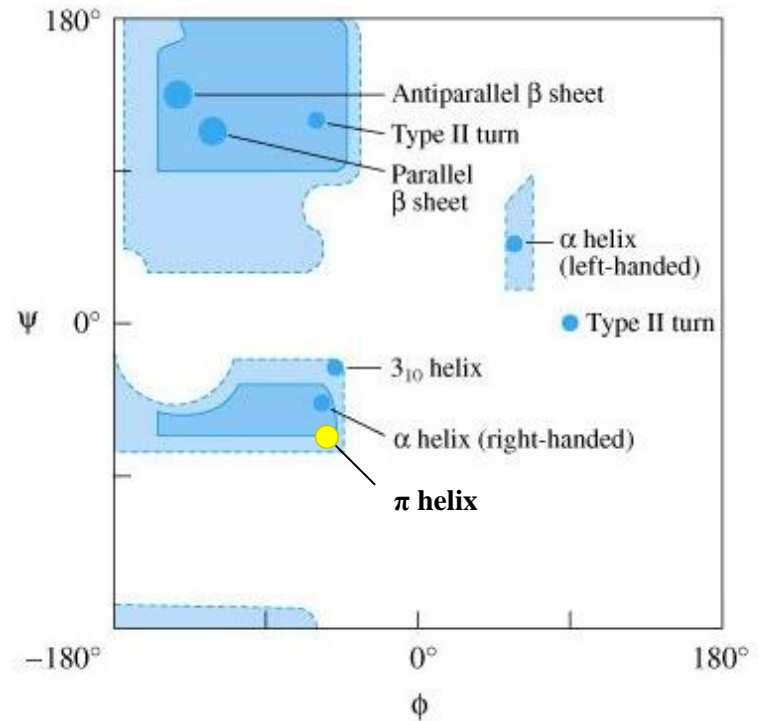
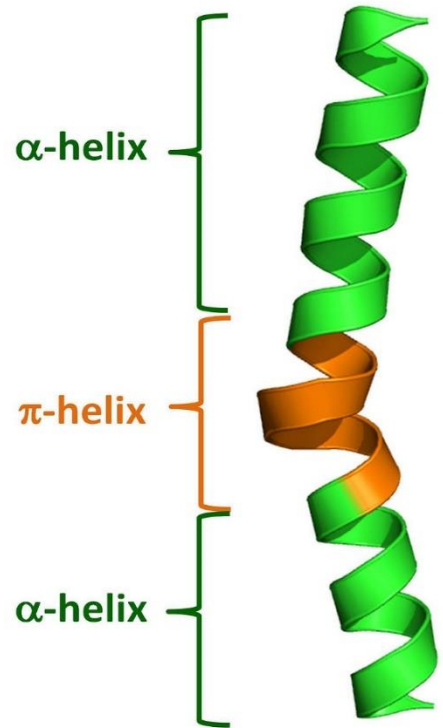
## π hélice

Ligação de H =  $n - n+5$

Período = 4,4 aa/volta

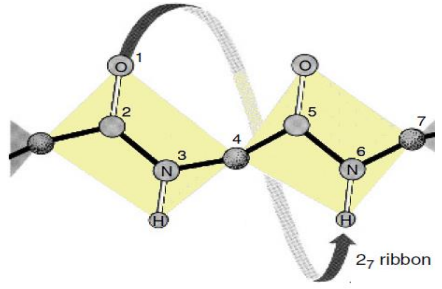
$\Phi = -57^\circ$

$\Psi = -70^\circ$

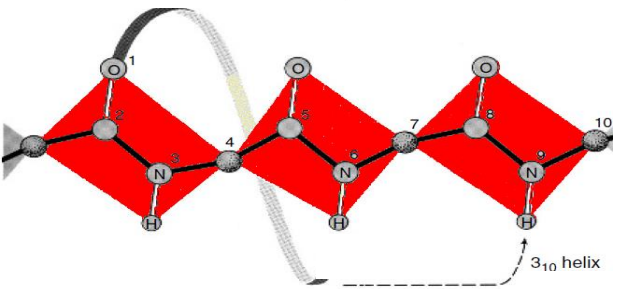




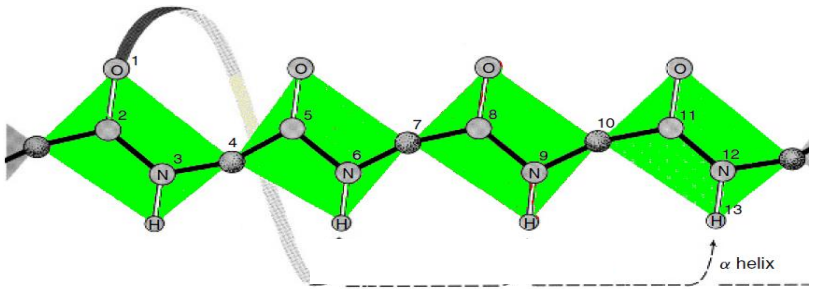
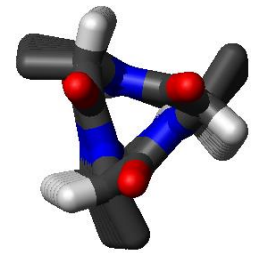
# Hélices em proteínas: Resumo



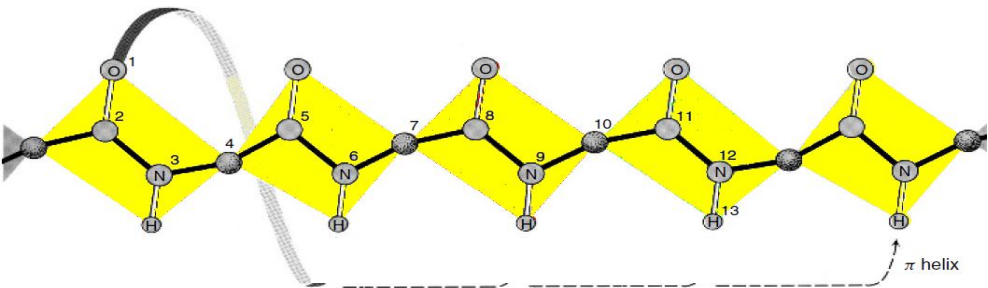
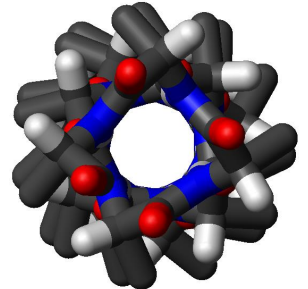
**2<sub>7</sub> ribbon**  
Ligação de H =  $n - n+2$   
Período = 2,0 aa/volta



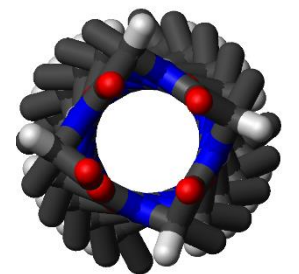
**hélice 3<sub>10</sub>**  
Ligação de H =  $n - n+3$   
Período = 3,0 aa/volta



**α hélice**  
Ligação de H =  $n - n+4$   
Período = 3,6 aa/volta



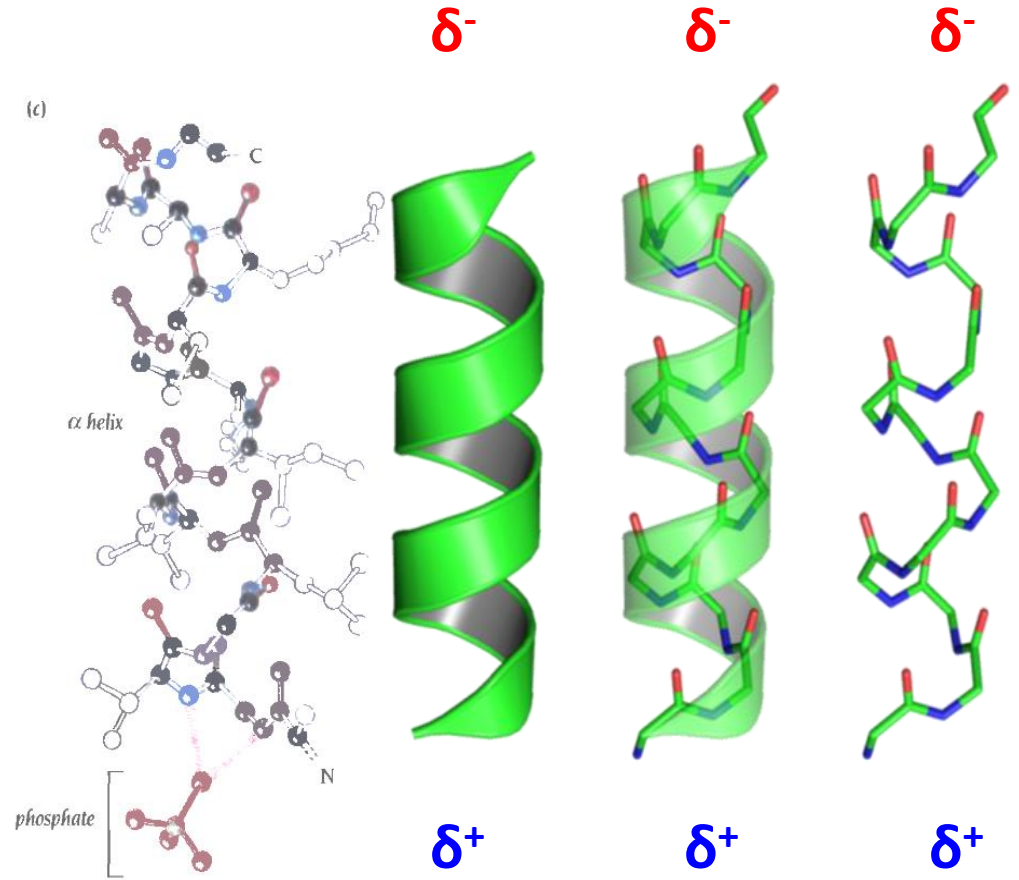
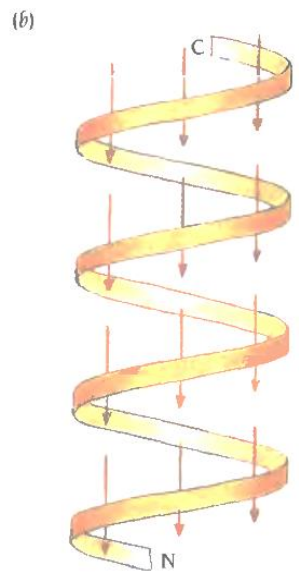
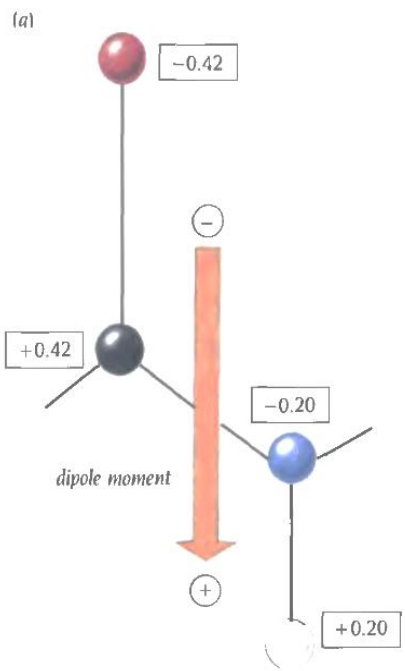
**π hélice**  
Ligação de H =  $n - n+5$   
Período = 4,4 aa/volta







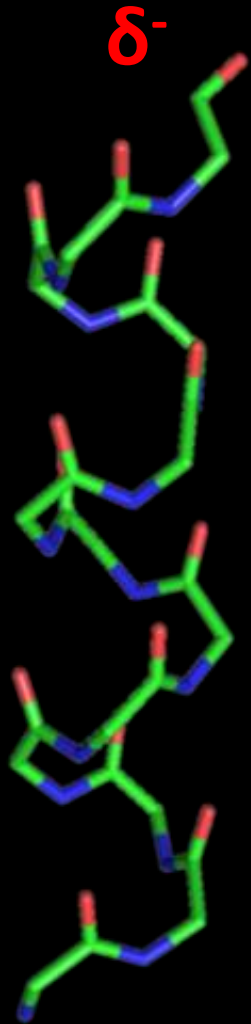
# Alfa-hélice: dipolos





# Alfa-hélices: dipolo

```
all A S H L C  
  
Mouse Mode  
Buttons L M R Wheel  
& Keys  
  Shift  
  Ctrl  
  C+Sh  
SnglClk  
Db1Clk  
Selecting Residues  
State 1/ 1  
|< < << >> > > S > F
```

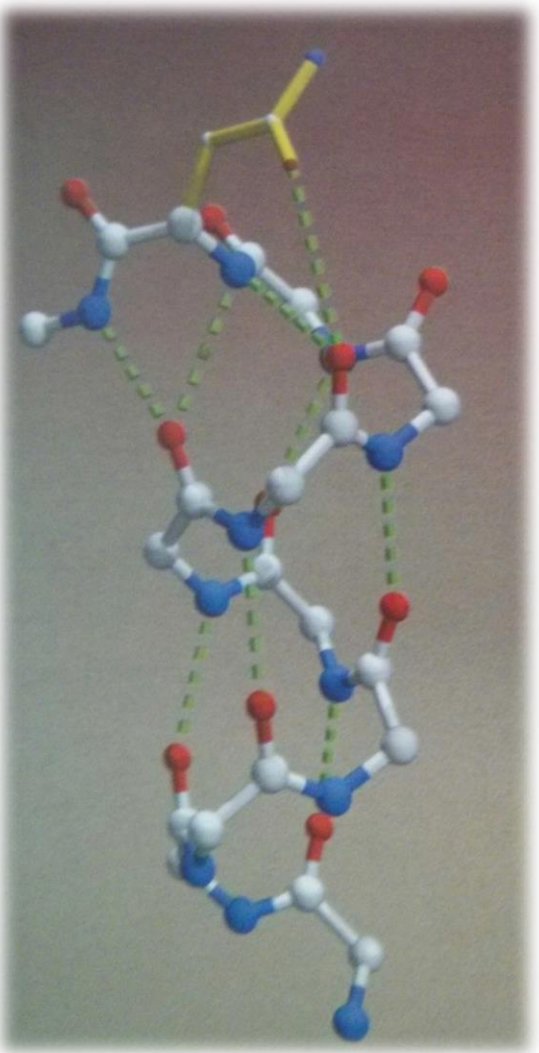


$\delta^-$

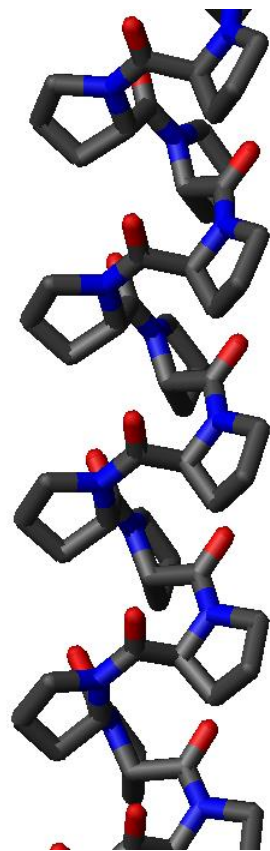
$\delta^+$



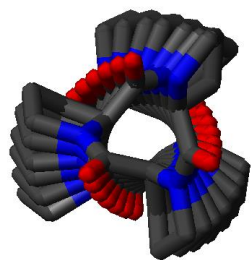
### Hélice "caps"



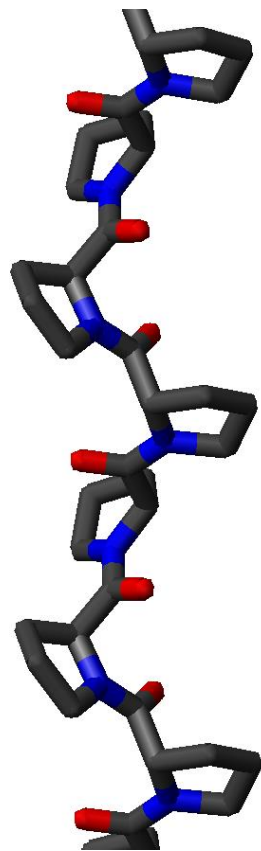
### Hélice poliprolina I



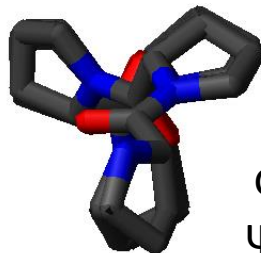
cys  
 $\Phi = -75^\circ$   
 $\Psi = 160^\circ$



### Hélice poliprolina II



trans  
 $\Phi = -75^\circ$   
 $\Psi = 150^\circ$





Resíduos de **Ala, Leu, Glu, Lys, Arg e Met (MALERK)** são encontrados mais frequentemente em **hélices alfa**, enquanto **Pro, Gly e Asp** são encontrados com menos frequência que a média.

| Residue          | Helix propensity, $\Delta G$ (kJ mol <sup>-1</sup> ) | Residue | Helix propensity, $\Delta G$ (kJ mol <sup>-1</sup> ) |
|------------------|--|---------|--|
| Ala              | 0  | Ile     | 0.41   |
| Arg              | 0.21   | Leu     | 0.21   |
| Asn              | 0.65   | Lys     | 0.26   |
| Asp <sup>0</sup> | 0.43   | Met     | 0.24   |
| Asp <sup>-</sup> | 0.69   | Phe     | 0.54   |
| Cys              | 0.68   | Pro     | 3.16   |
| Gln              | 0.39   | Ser     | 0.50   |
| Glu <sup>0</sup> | 0.16   | Thr     | 0.66   |
| Glu <sup>-</sup> | 0.40   | Tyr     | 0.53   |
| Gly              | 1.00   | Trp     | 0.49   |
| His <sup>0</sup> | 0.56   | Val     | 0.61   |
| His <sup>+</sup> | 0.66   |         |  |







# Diagrama Wenxiang

<http://www.jci-bioinfo.cn/wenxiang2>

**Wenxiang:** A web-server to draw Wenxiang diagram

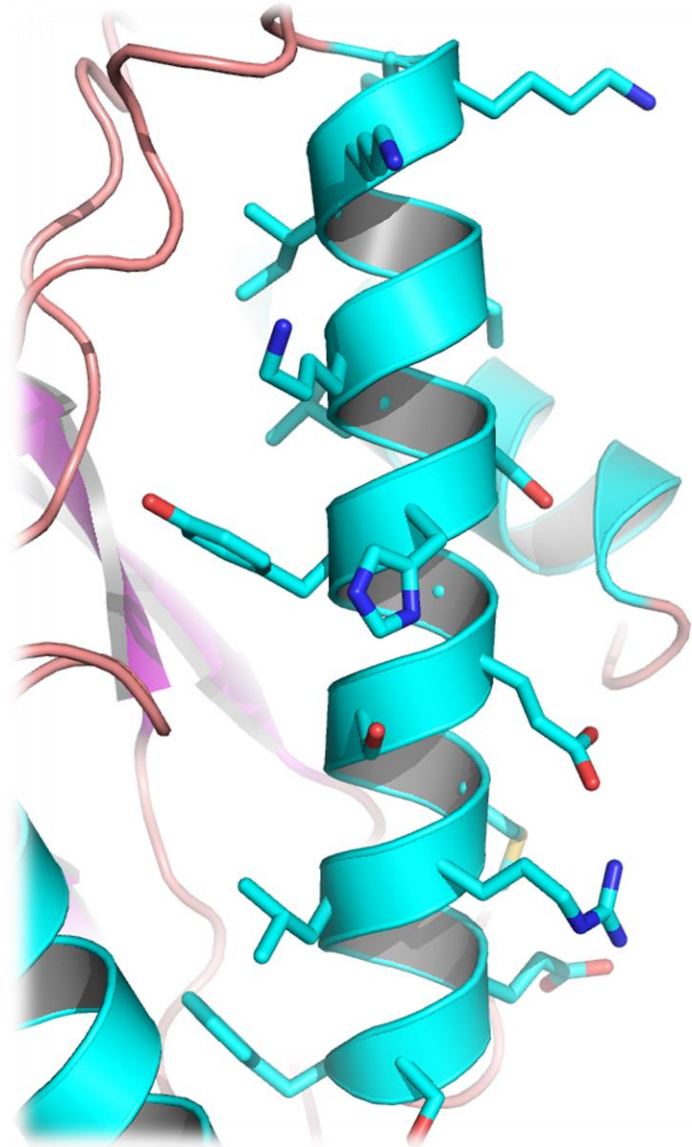
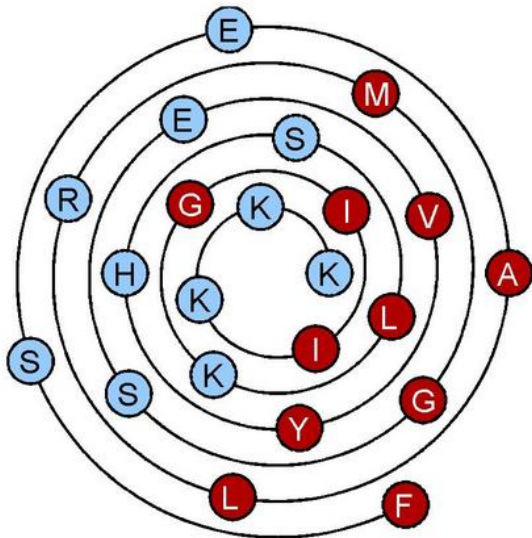
[Read Me](#) | [Citation](#)

Input the sequence of an  $\alpha$ -helix (Example)

```
FSEALRMGSEVYHSLKGIKKK
```

[Black&White](#) [Color](#) [Cancel](#)

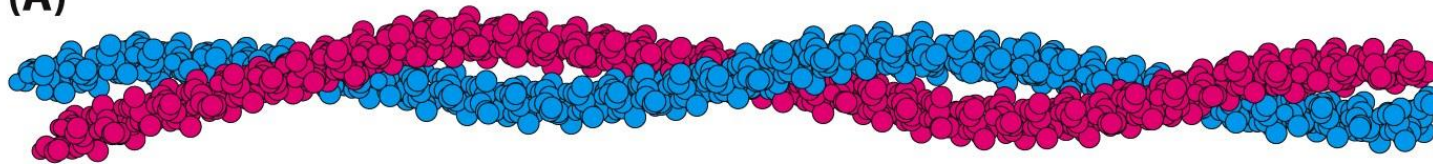
004287



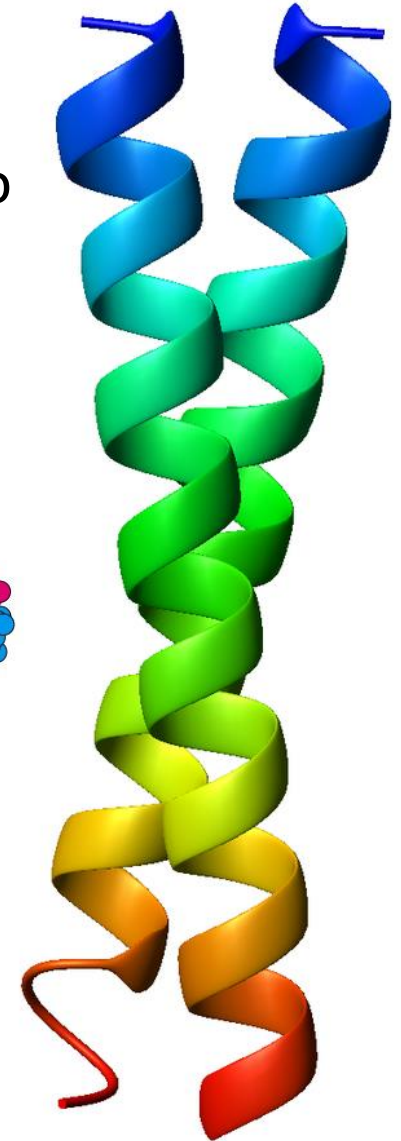
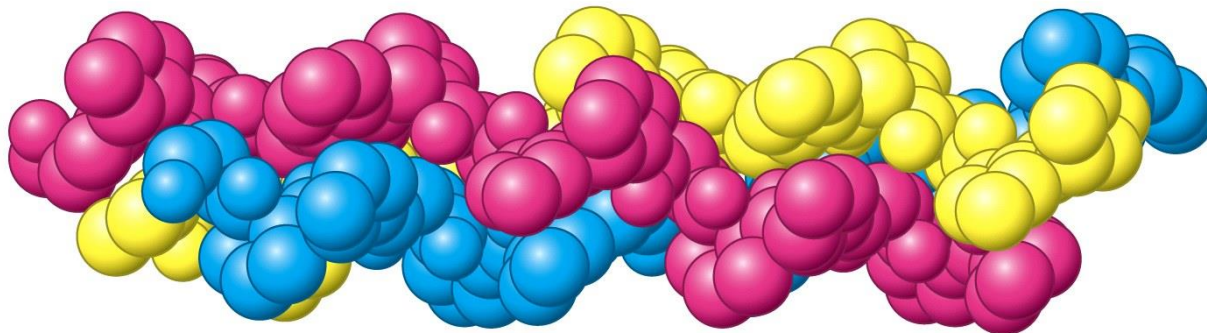


- *Coiled-coil* (“bobina em espiral”) é um motivo estrutural em proteínas em que 2 a 7 alfa-hélices são enroladas em conjunto como os fios de um cabo
- Dímeros e trímeros são os tipos mais comuns

(A)



(B)



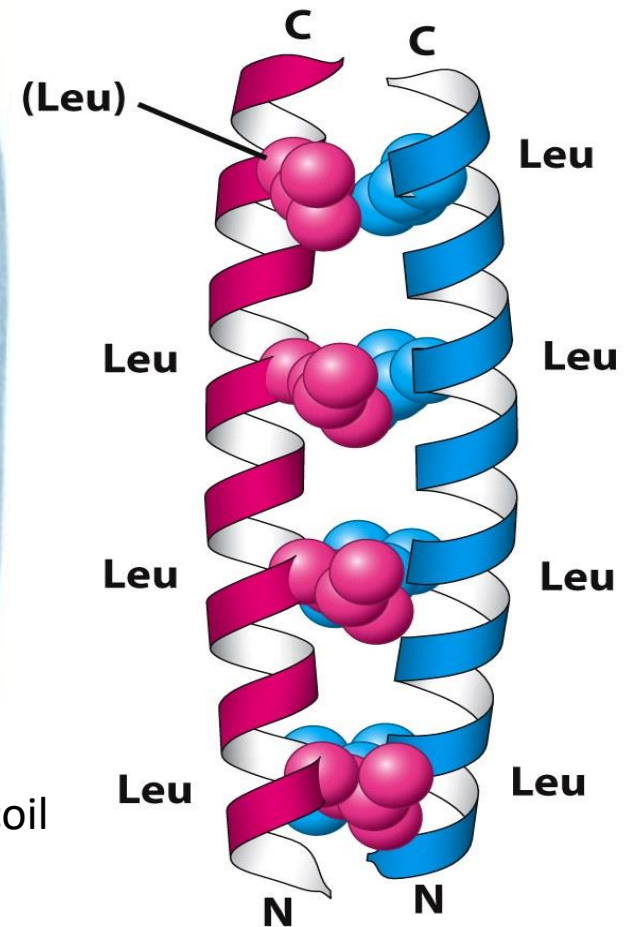
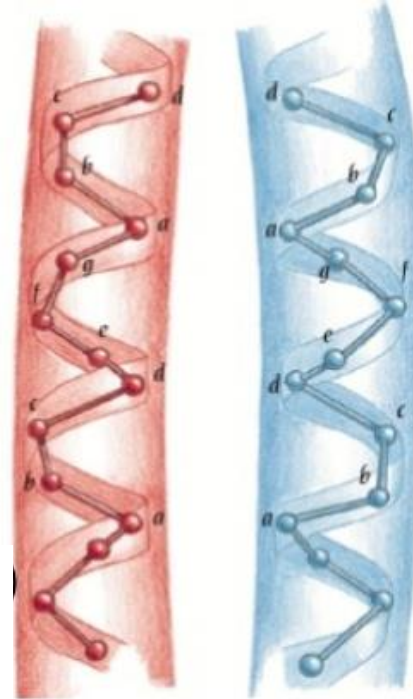




# Repetição Hepta “Heptad repeat”

## The heptad repeat

|    | a   | b   | c   | d          | e   | f   | g   |
|----|-----|-----|-----|------------|-----|-----|-----|
| 1  | Met | Lys | Gln | <b>Leu</b> | Glu | Asp | Lys |
| 8  | Val | Glu | Glu | <b>Leu</b> | Leu | Ser | Lys |
| 15 | Asn | Tyr | His | <b>Leu</b> | Glu | Asn | Glu |
| 22 | Val | Ala | Arg | <b>Leu</b> | Lys | Lys | Leu |



- d: frequentemente Leu ou Ile
- a: frequentemente hidrofóbico
- e,g: frequentemente carregado
- b,c,f : carregado ou polar
- essas características são utilizadas para predição de coiled-coil em estruturas de proteínas

### Por que hepta?

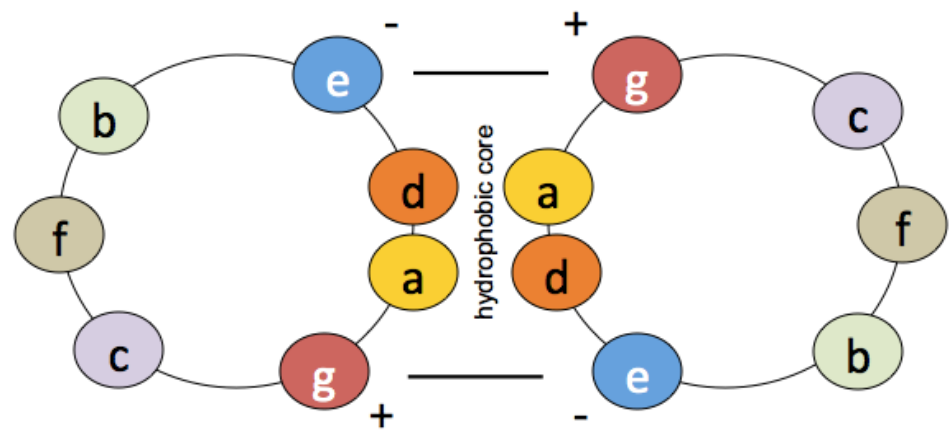
- hélice  $\alpha$ : 3,6 resíduos/volta
- hélice  $3_{10}$ : 3 resíduos/volta
- hélices  $\alpha$  em “coiled-coil” são ligeiramente distorcidas e apresentam 3,5 resíduos/volta
- $3,5 \times 2 = 7$ , portanto, a cada duas voltas forma-se uma repetição hepta



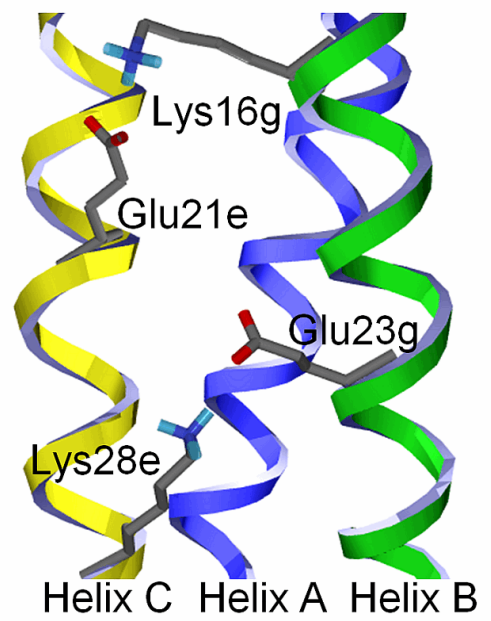


# Interações em Coiled-coil

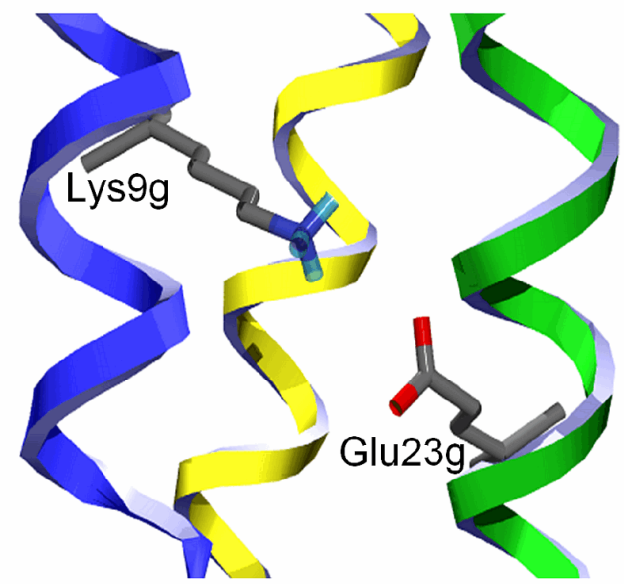
|    | a   | b   | c   | d          | e   | f   | g   |
|----|-----|-----|-----|------------|-----|-----|-----|
| 1  | Met | Lys | Gln | <b>Leu</b> | Glu | Asp | Lys |
| 8  | Val | Glu | Glu | <b>Leu</b> | Leu | Ser | Lys |
| 15 | Asn | Tyr | His | <b>Leu</b> | Glu | Asn | Glu |
| 22 | Val | Ala | Arg | <b>Leu</b> | Lys | Lys | Leu |



e/g in Parallel face



g/g in Antiparallel face





# Repetição Hepta "Heptad repeat"

**a**

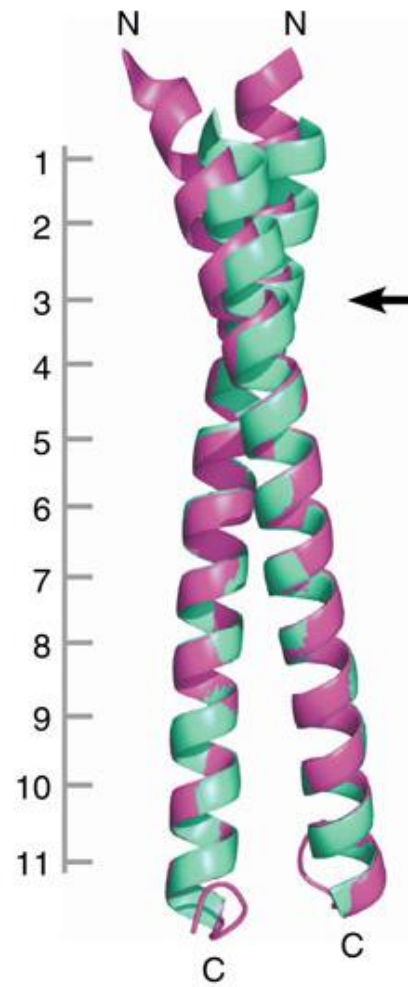
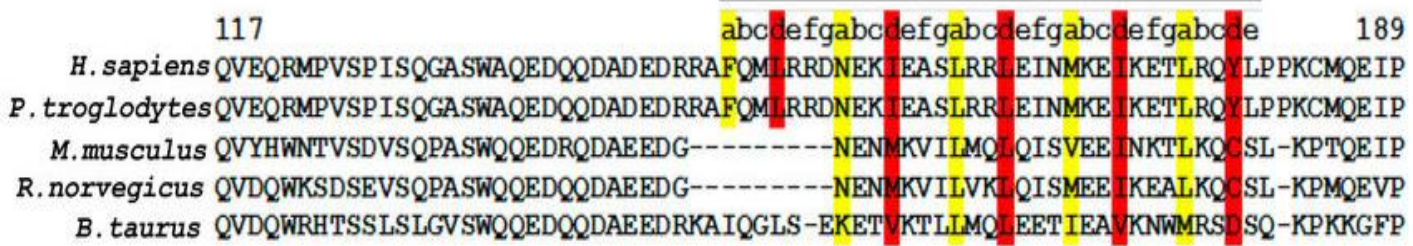
S4 Coiled-coil domain



Coiled-coil domain 1



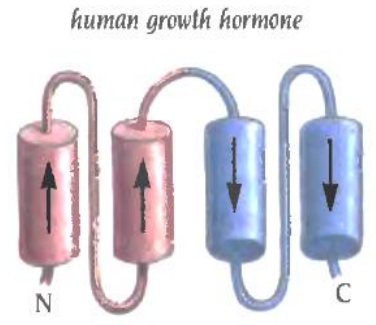
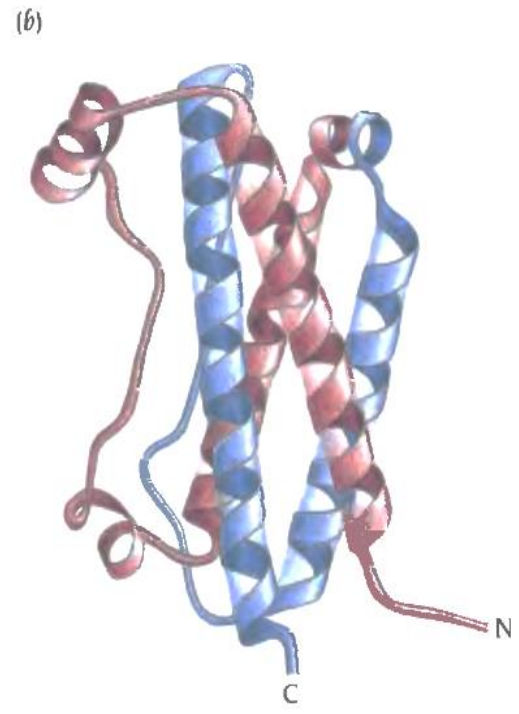
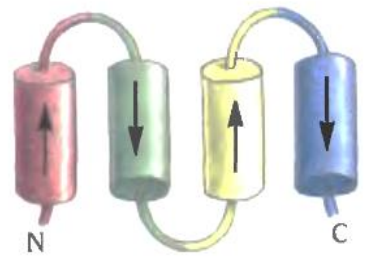
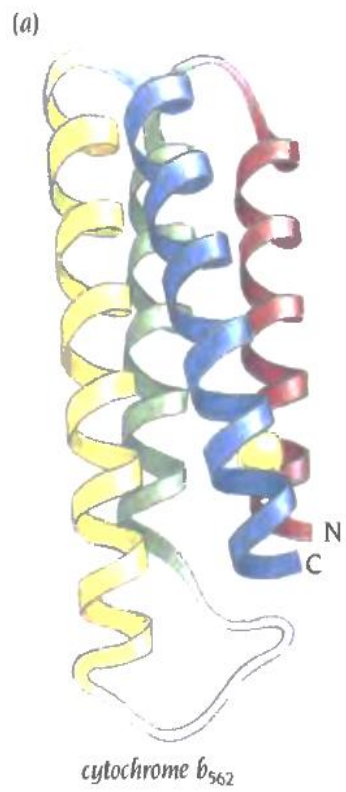
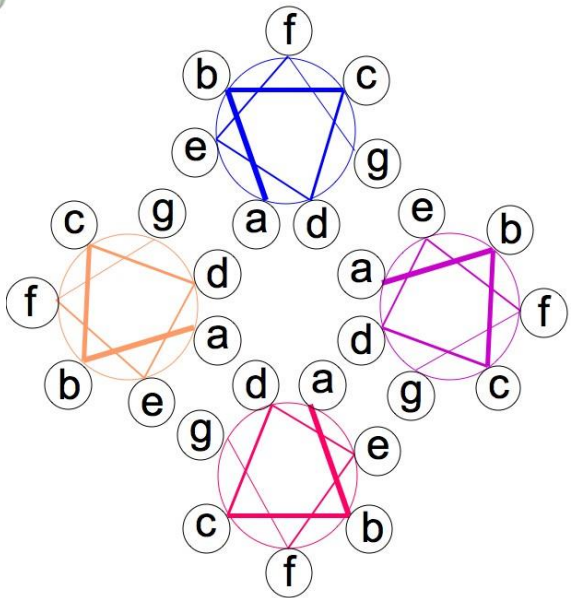
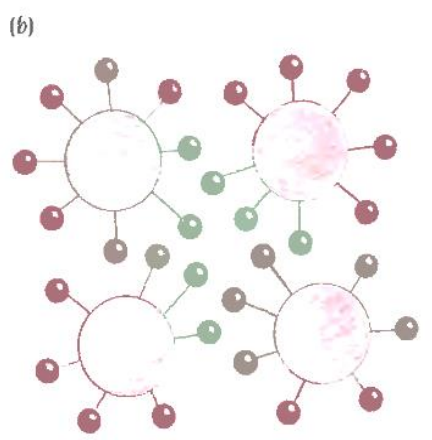
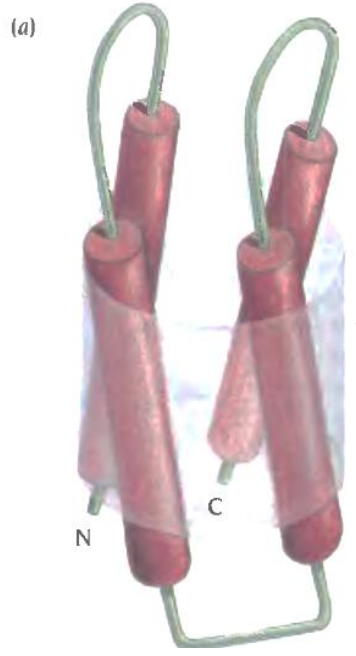
Coiled-coil domain 2







# Four-helix bundle “pacote de 4 hélices”





# Domínio Globina

