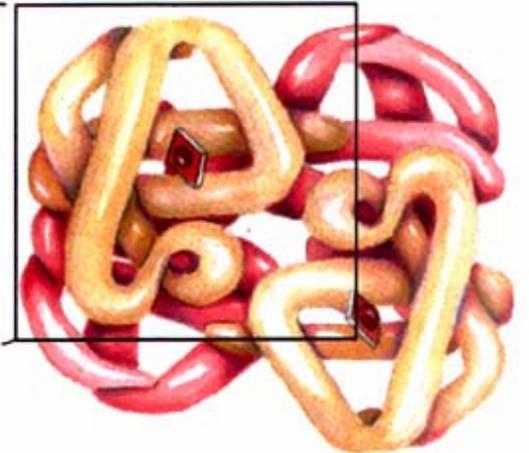
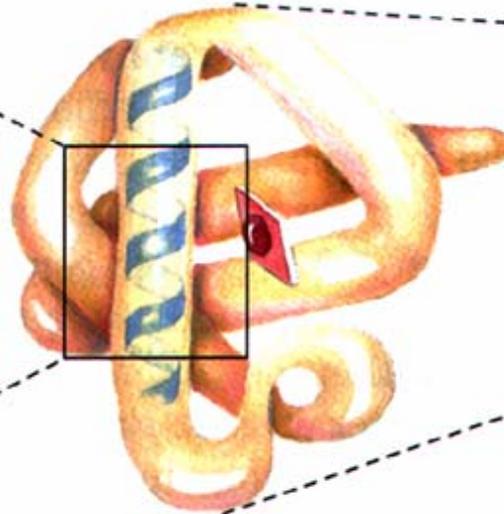
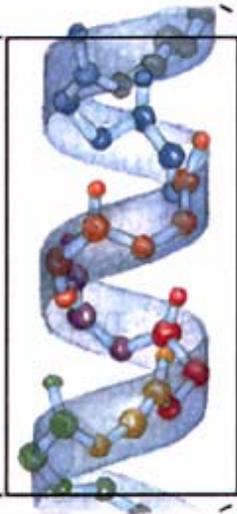


# 蛋白質的四級構造

胺基酸



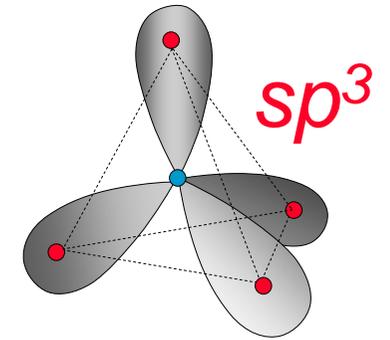
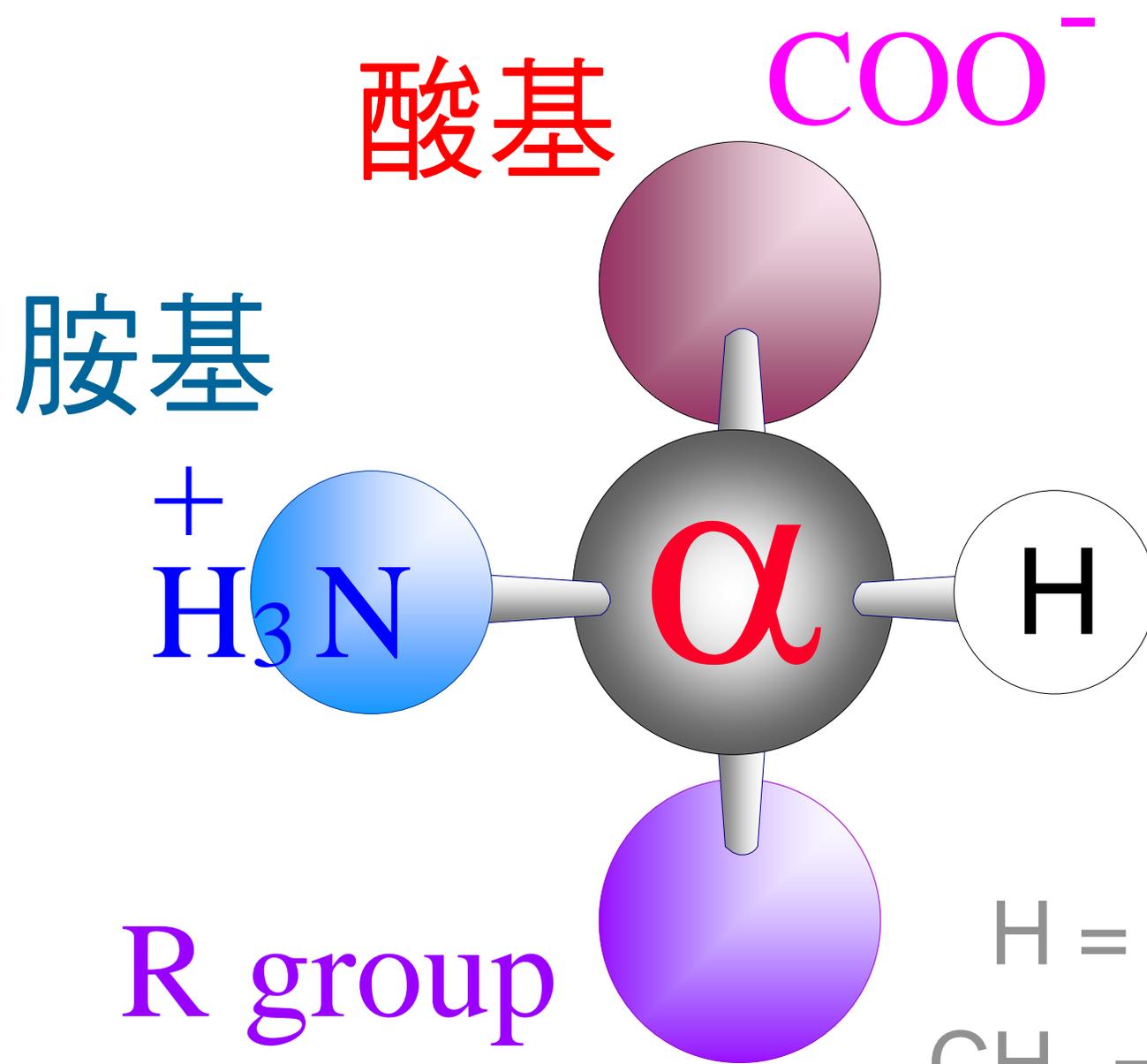
一級構造

二級構造

三級構造

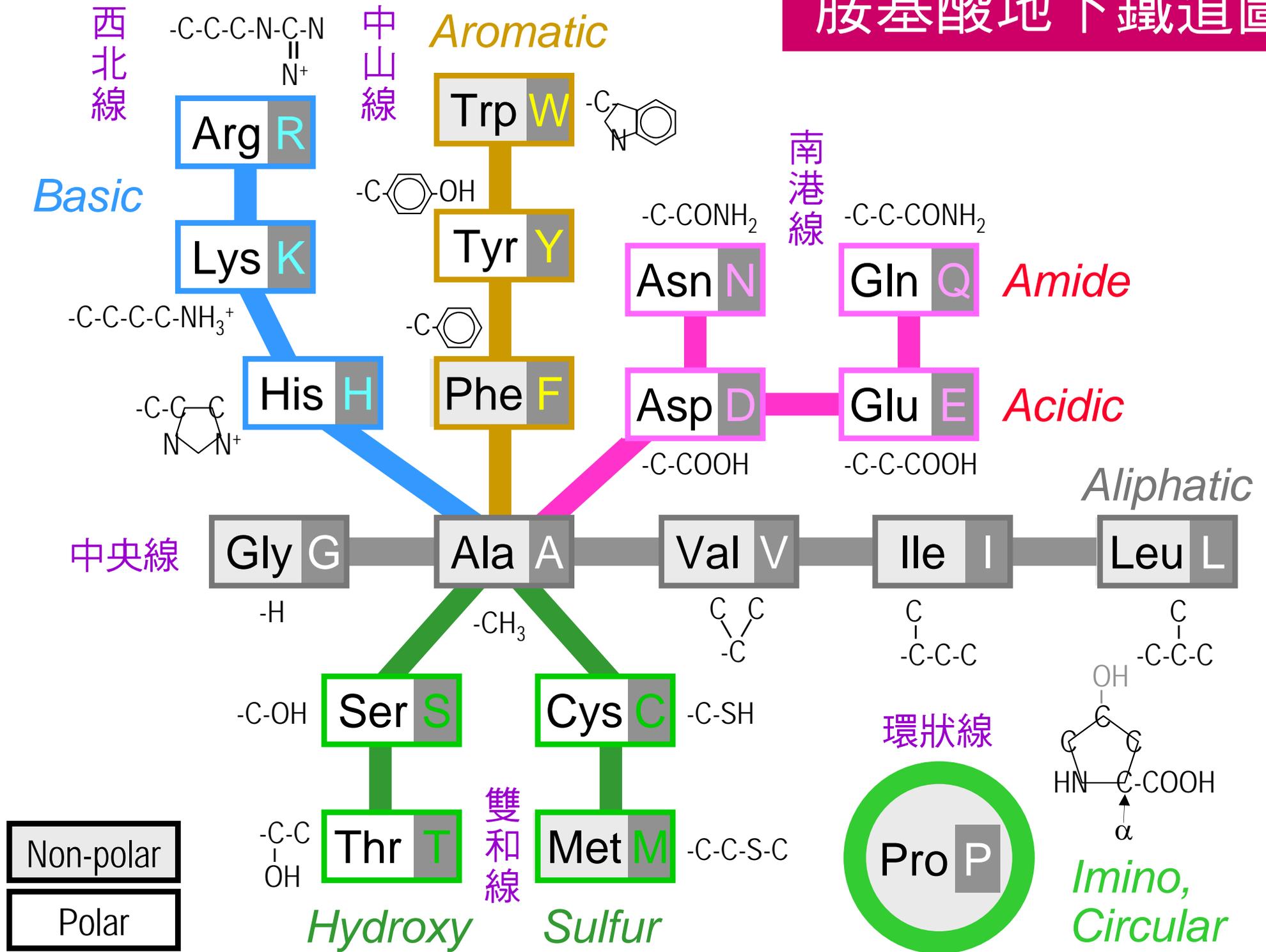
四級構造

# L-Form Amino Acid



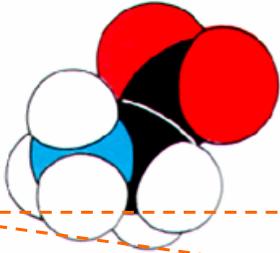
H = Glycine  
 $\text{CH}_3$  = Alanine

# 胺基酸地下鐵道圖

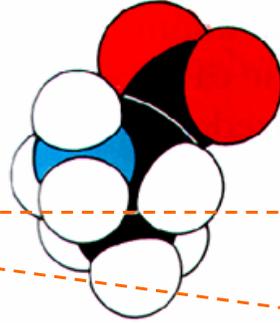


# 各式胺基酸的大小樣式齊全

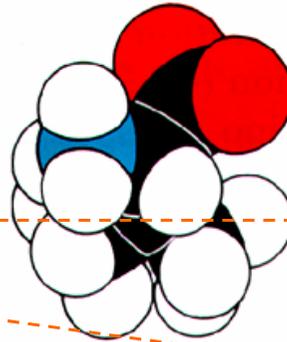
Gly



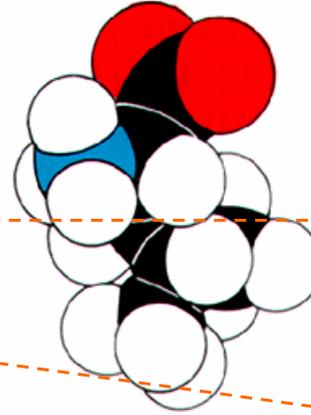
Ala



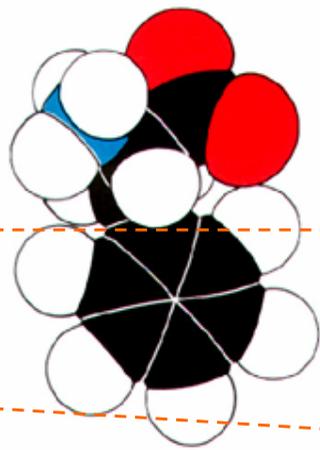
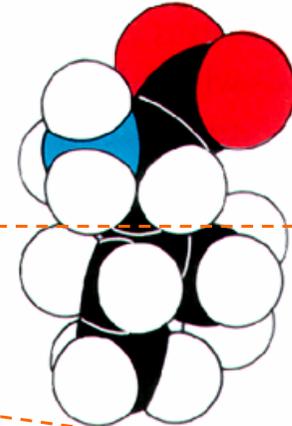
Val



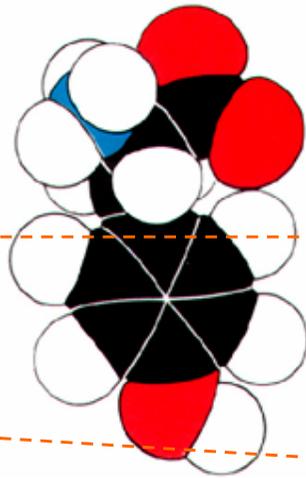
Leu



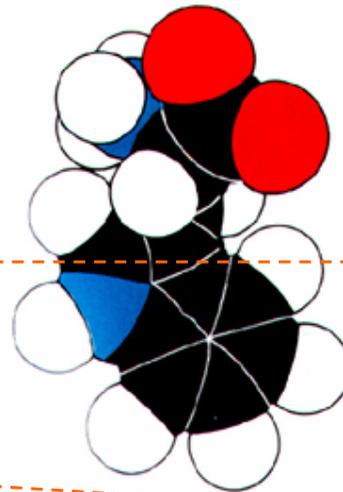
Ile



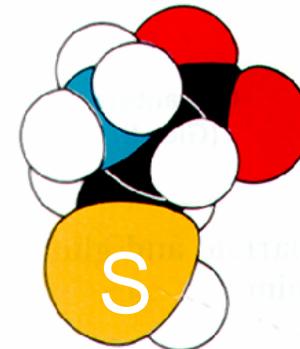
Phe



Tyr

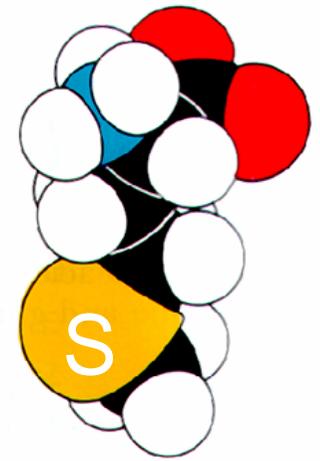


Trp



-SH

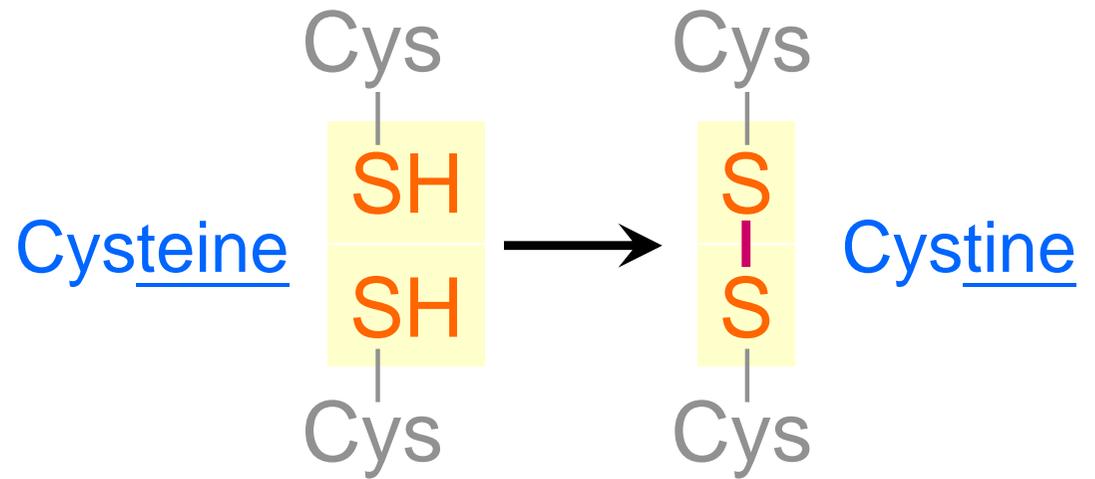
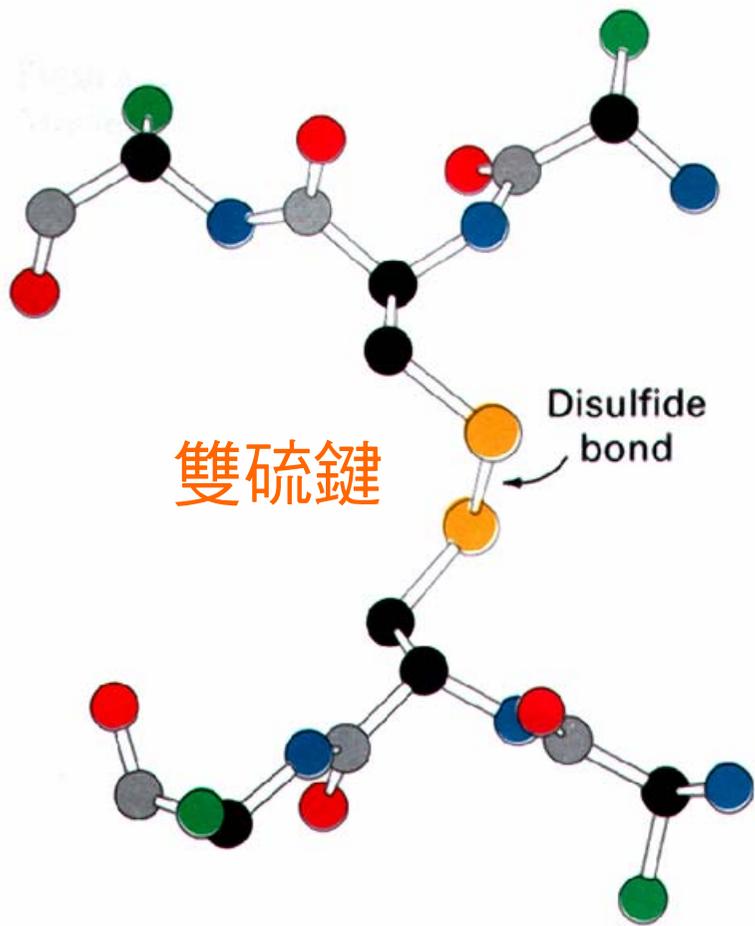
Cys



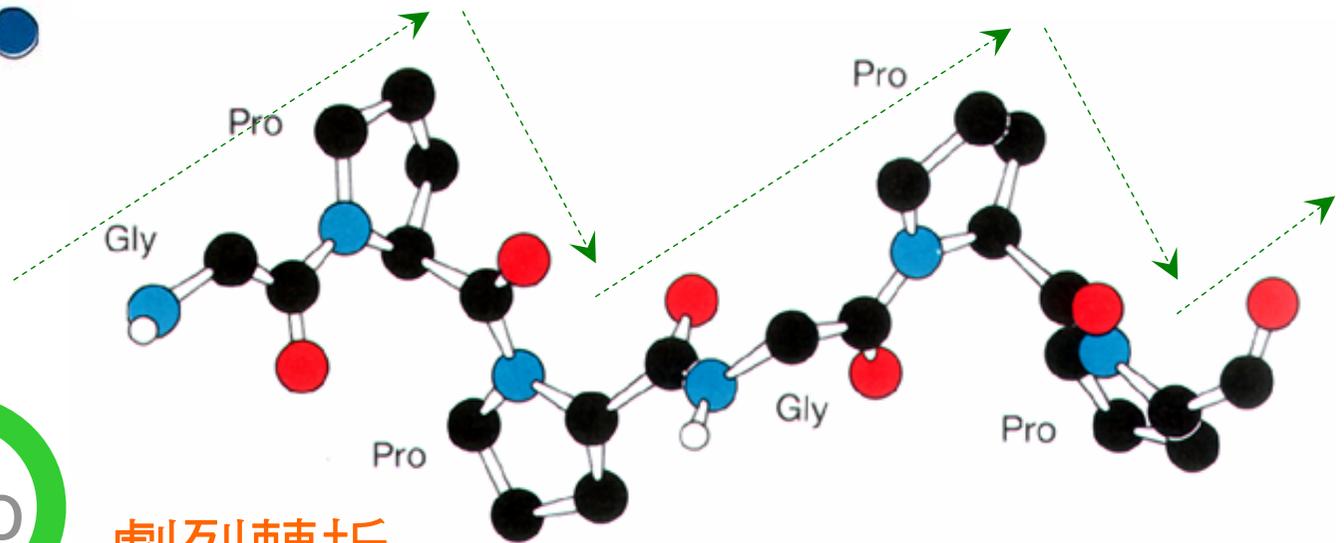
-S-CH<sub>3</sub>

Met

# 兩種胺基酸對蛋白質構造影響很大



劇烈轉折



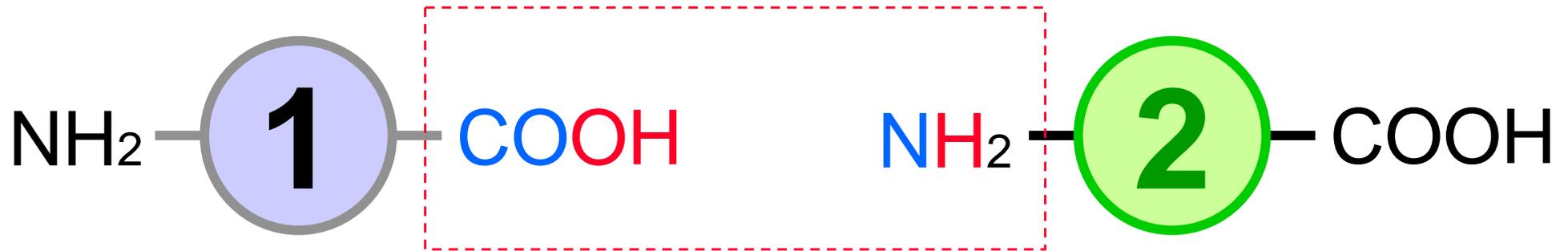
# 形形色色的胺基酸側基



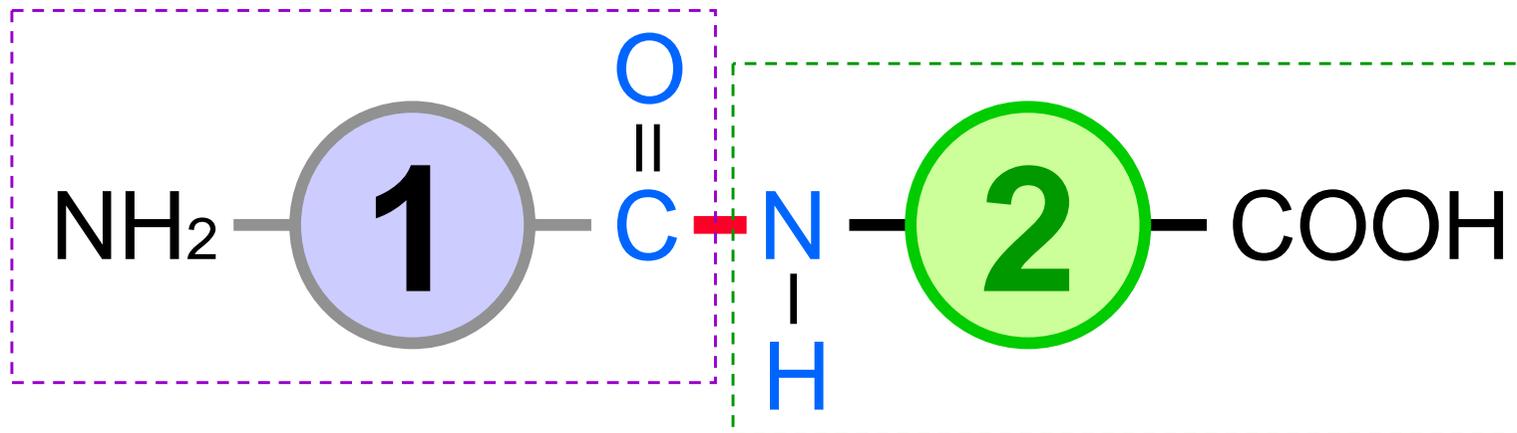
有大有小  
有正有負  
有極性  
非極性

# 胜肽鍵的形成

兩個胺基酸分子頭尾連接起來

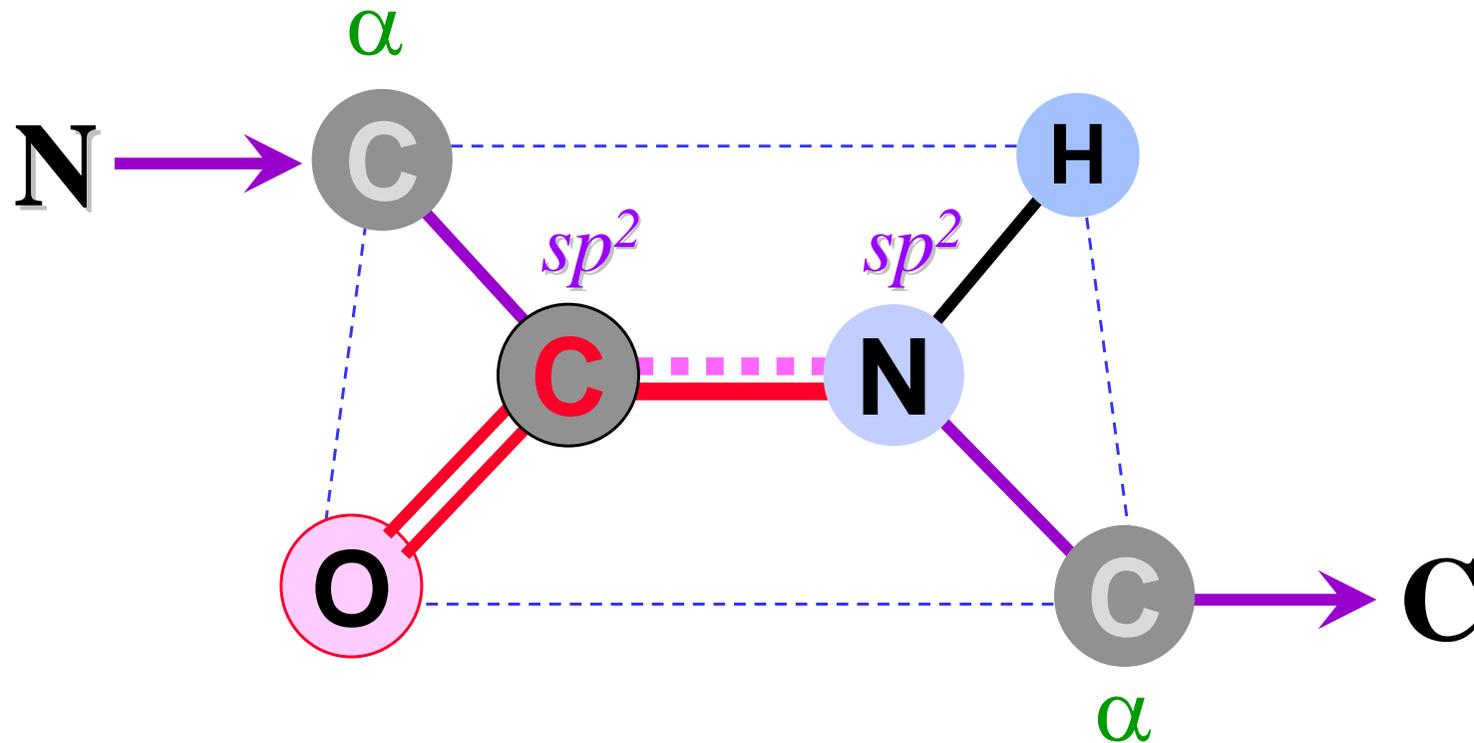


Carbodiimide ↓ 脫水

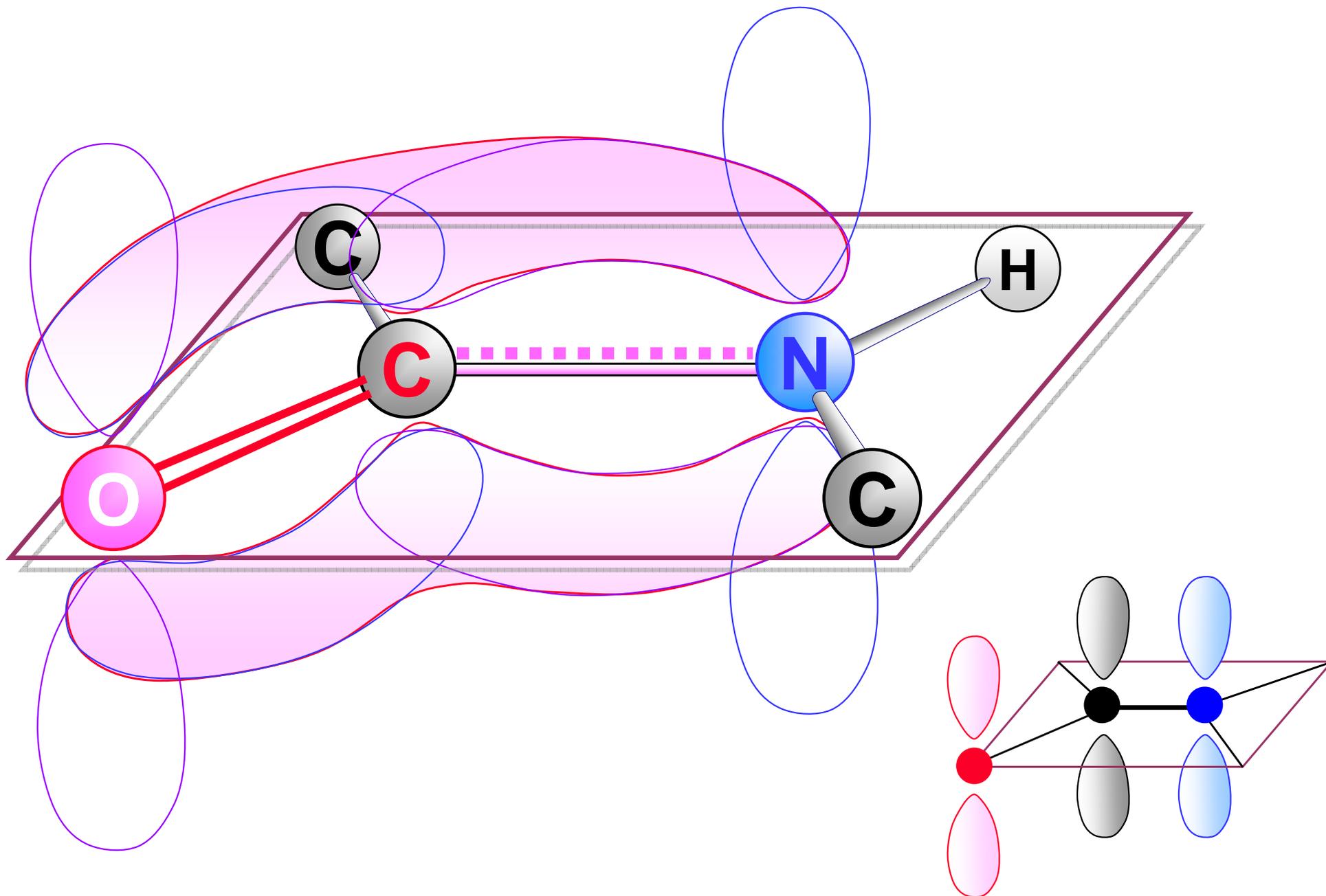


# 胜肽鍵的特性

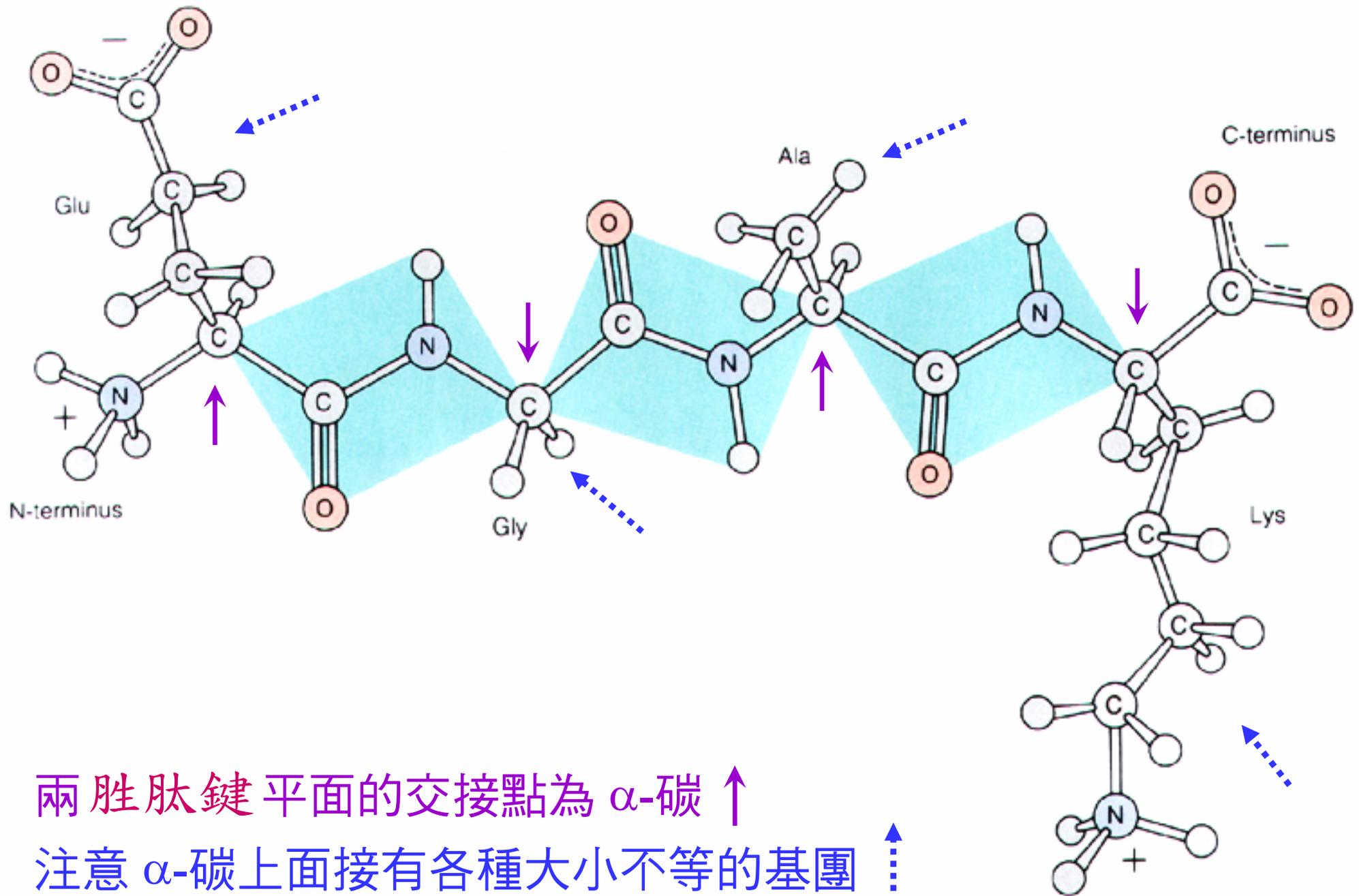
- 胜肽鍵 雖是單鍵卻有雙鍵性質
- 胜肽鍵 周邊六個原子在同一平面上
- 前後兩個胺基酸的  $\alpha$ -carbon 在對角 (trans)



# $p$ 軌道電子共振使胜肽鍵具雙鍵特性



# 以胜肽鍵平面連接成多肽長鏈

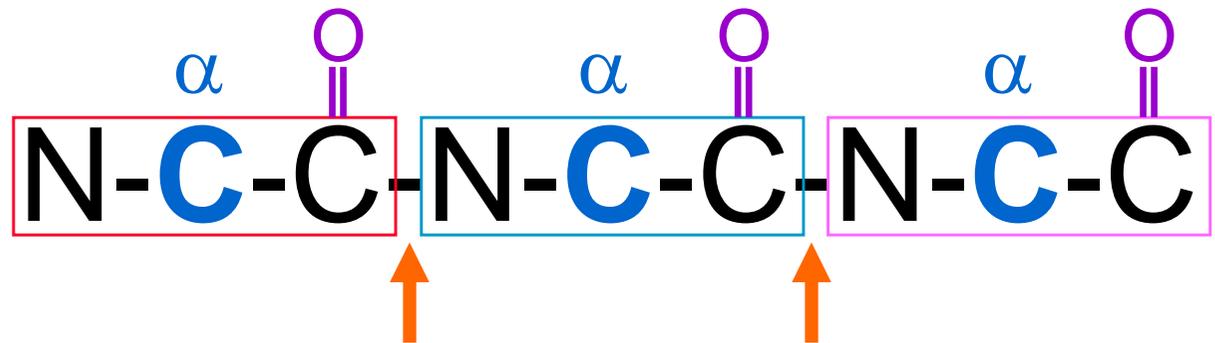
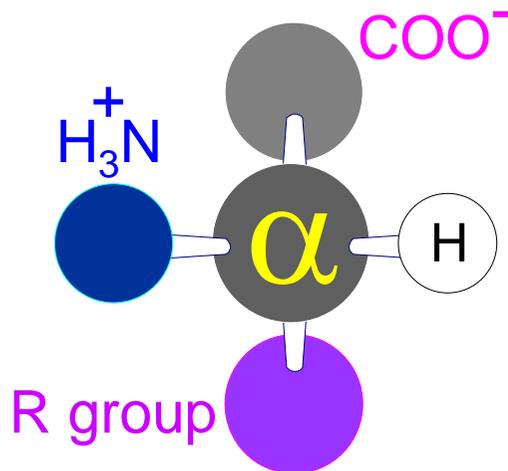
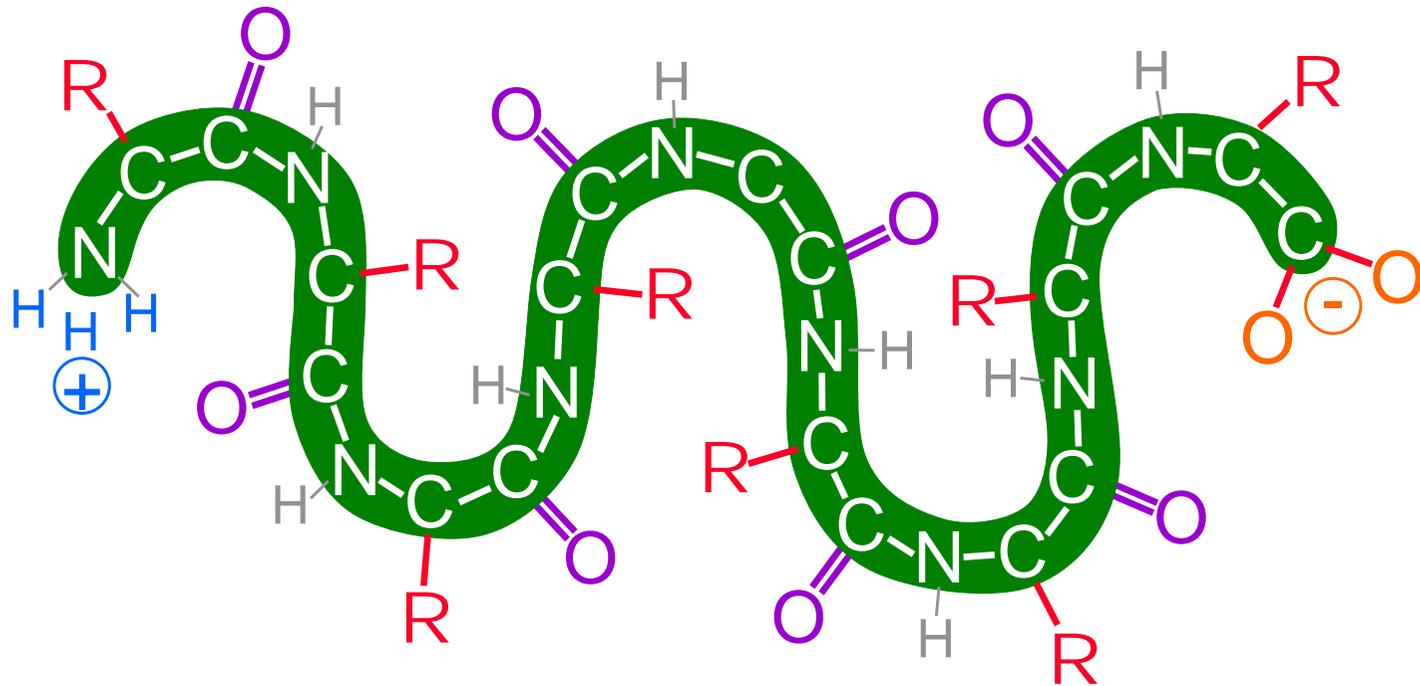




# 蛋白質的骨架 (Backbone)

Constant

Variable





# 蛋白質的胺基酸序列

ΑΡΧΙΕΡΕΥΣ ΤΩΝ ΣΕΒΑΣΤΩΝ ΔΙΑΒΙΟ  
ΚΛΑΥΔΙΟ ΚΑΙ ΣΑΡΟΣ ΣΕΒΑΣΤΟΥ Ε  
ΕΤΑΜΕΙΝΩΝ ΔΟΥΕΙΤΕΝ ΤΡΟΒΕΒΟΥ

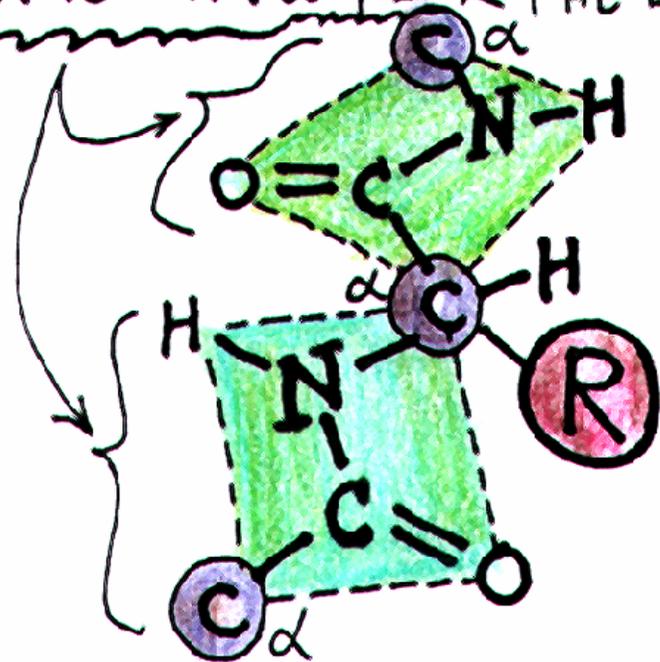
MSRLSGITPRARDDRSQFQNPRLLEIAYVDPRTAGLQRTKRTLKVLCVLDDETKQTIQHV  
VTEKNEGTLLEDAASIASSIKYHAEFSPAFSPERFELPKAYFATAQSV **RDALIVNNA**  
TYDYYEKLNMKQAYYLSMEFLOG **RALLNAIGNLE** LEITCEVAEALNKLGHNLENVAS  
KEPDAALGNGGLGRLASCFLDSLATINYPAGYGI **destroy me** GQEEVAE  
DLELGNPEIHRMDVSYVPVKFFGKVITGSDGKKHIGGEDILAYVADYVITGFKTRTTISI  
RLSTKVPSSEDFDLYSFNAGEHTKACEAQANA EKICYILYPGDESI EGKRLKQQYT  
LCSASLQDIARFERRSGEYVKWEEFPEKVAVQMN DTHPTLCIPELIRILIDLKGLSW  
KEAWNITQRTVA YTNHTVLP EALEKWSYELMEKLLPRHIEHEMIDEQLINEIVSEY G  
TSDLDMLEKKLNDMRILENFDIPSSIANLFTK **PKETSIVDPSEEVEVSGKVVTESVEV**  
DKVVTESSEKDELEEKDTELEKDEDPVFAPIPPKMVRM ANI CVV/GGHAYNGVAEIH  
DIVKEDVFNDFYQLWP EK FQNK TNGVTPRRWRFCNP **Cut me** EDWV LNTEI  
LAELRKFADNEDLQIEWRAAKRSNKVKVASFLKERTGIDSDINAMDIQVKRIHEY  
KRQLLNILGIVYRYKQMKEMSAREREAKFYPRVCIFGGKAFATYVCAKRIAKFITD  
VGATINHDP EIGDILKVI FVPDYNVSAAE LLIPASGLSQHISIA GMEASGQSNMKFAI  
NGCILIGTLDGANVEIRQEVG EENFFLFGAEAEHEIAGLRKERAE GKFVPDERFEEV  
EFIKRGVVEGSNTYDELLGSLGNEGEGRGDYFLV GKDFPSYIECQEKVDEAYRDOK

ΧΟΥ ΣΙΚΑΙ ΣΥΝΕΛΡΟΙΣ ΚΑΙ ΤΟ ΔΗΜΟ  
ΤΑΤ ΟΤΙ ΑΡΟΝΤΟ  
ΓΡΑΦΟΝΤΑΣ ΔΙΕΝΕΤΟ ΕΡΓΟΝ  
ΤΑ ΕΝ ΤΩΝ ΑΩΤΟΥ ΑΠΟΛΛΩΝΟΣ ΤΟΥ  
ΔΡΙΟΝ ΤΑΣ ΤΟΙΣ ΠΑΤΡΙΟΙΣ ΘΕΟ

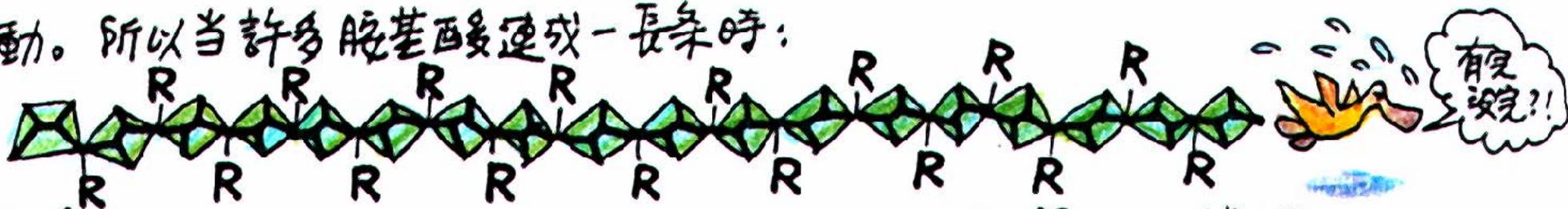
許多有意義的序列被各物種間所通用

# 胜肽平面連成蛋白質但不會展開成一直線

Peptide bond 的平面是不能扭曲的



這兩個平面又因為 R group 的關係，只能在一定範圍的角度內活動。所以當許多胺基酸連成一長條時：



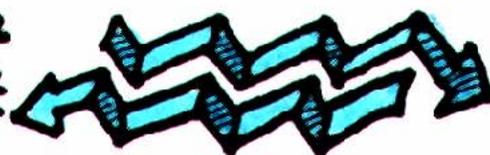
因為↑的關係，會自動捲曲成一定的構造 (Secondary structure),

大略說來有：  
兩種形式

Helix



或 Sheet



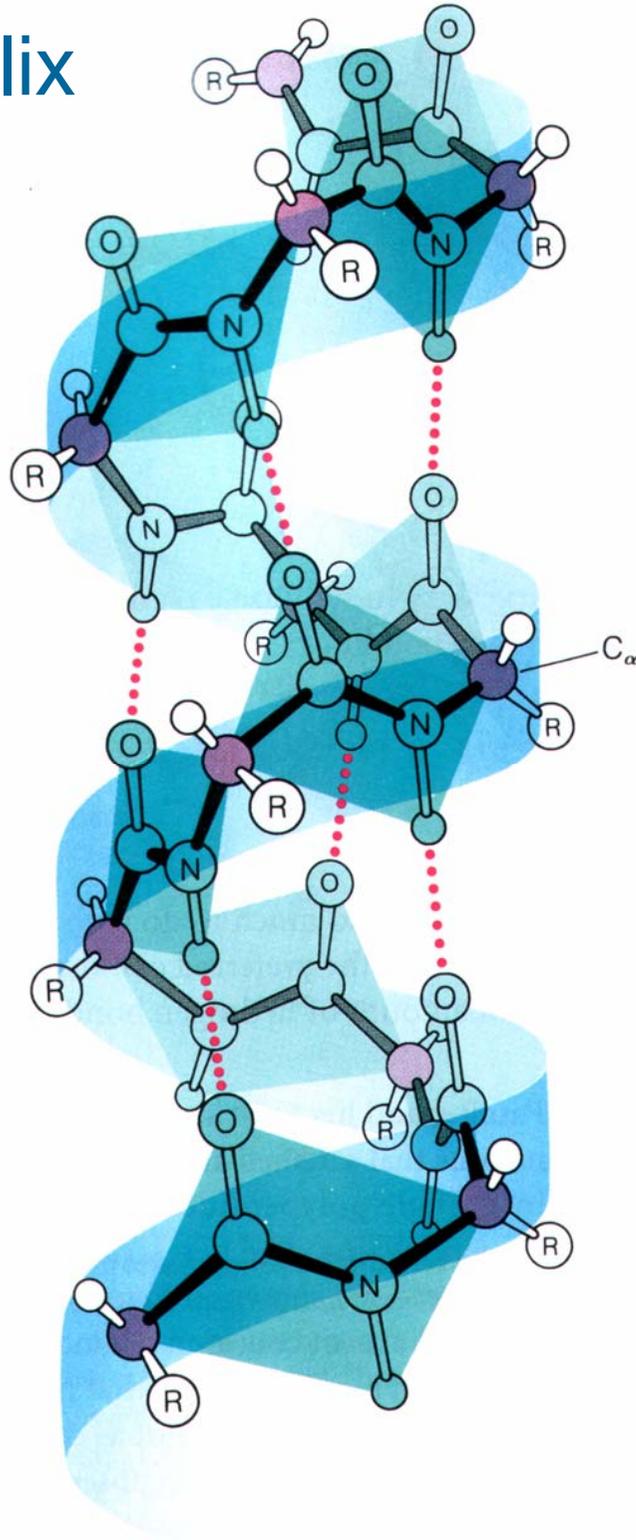
# 二級構造的成因

## $\alpha$ Helix, $\beta$ Sheet

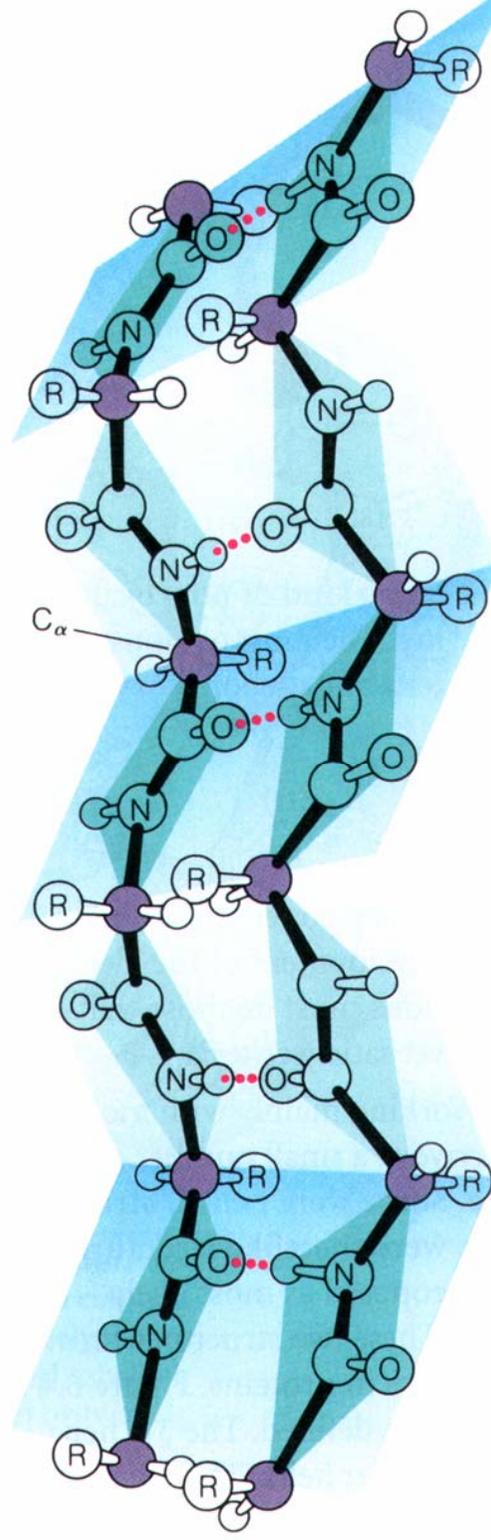
- (1) Peptide bond 不能轉動  $\rightarrow$  Peptide bond 平面
- (2) 一個胺基酸 R 基團 與前後 R 基團的限制  $\rightarrow$  Peptide bond 平面不能任意轉動  $\rightarrow$
- (3) R 基團的大小、電荷限制  $\rightarrow$  只做 規律摺疊  $\rightarrow$   $\alpha$  Helix,  $\beta$  Sheet Ramachandron plot
- (4) 穩定二級構造的力量： 氫鍵

# 蛋白質二級構造

$\alpha$  helix

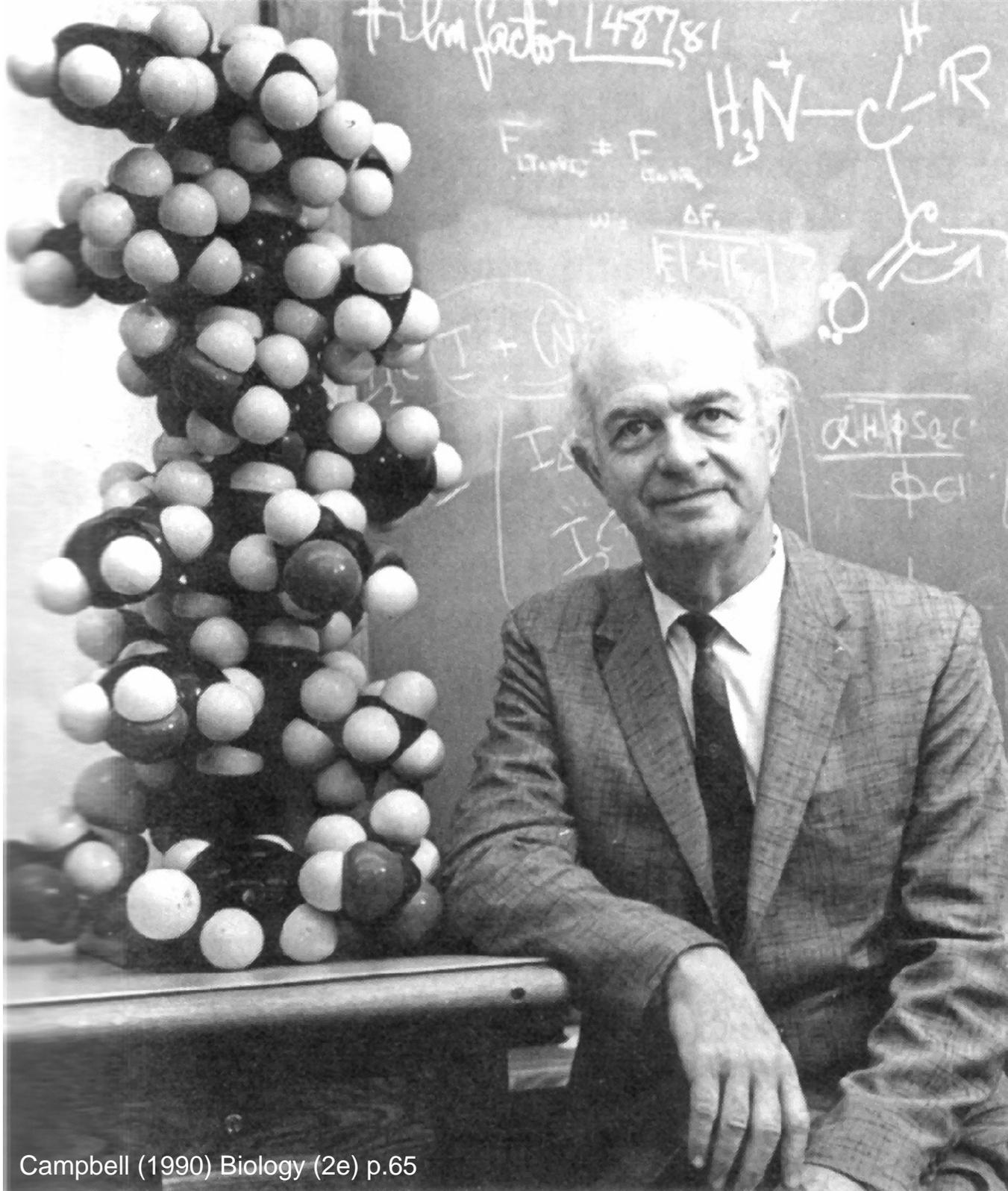


$\beta$  sheet

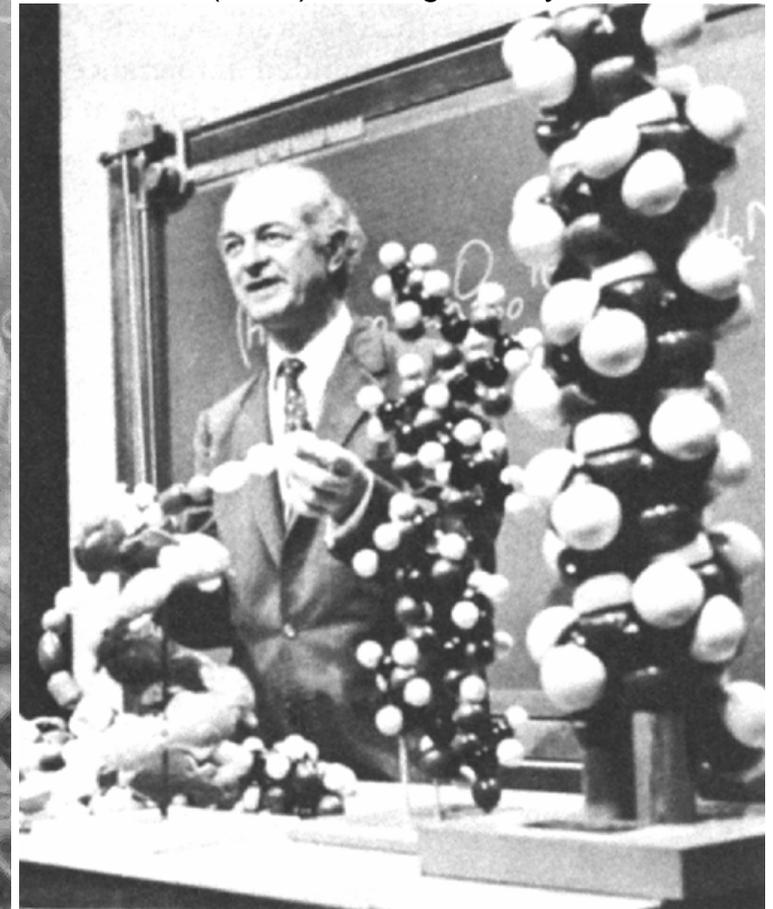


兩者都由  
氫鍵組成

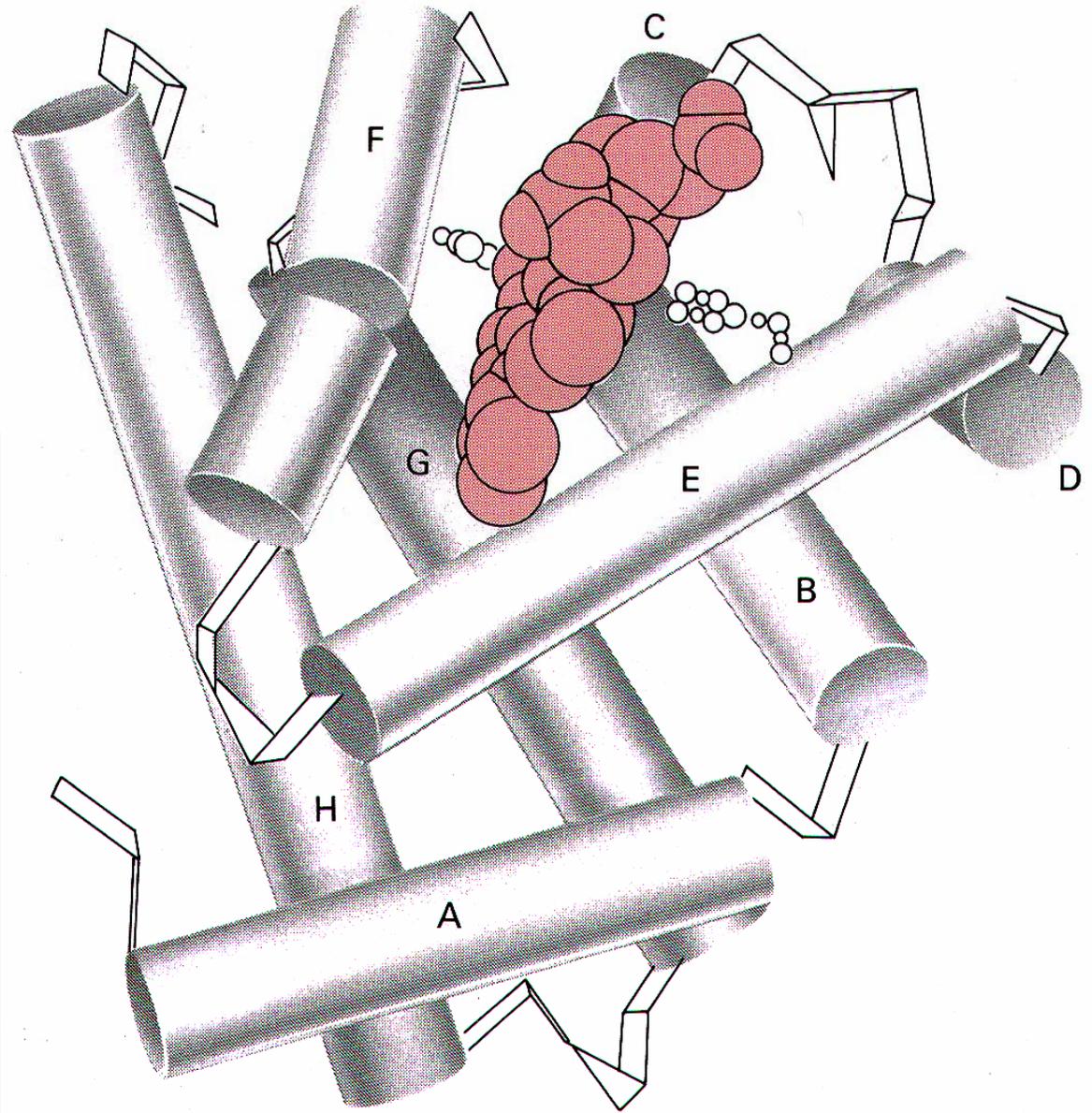
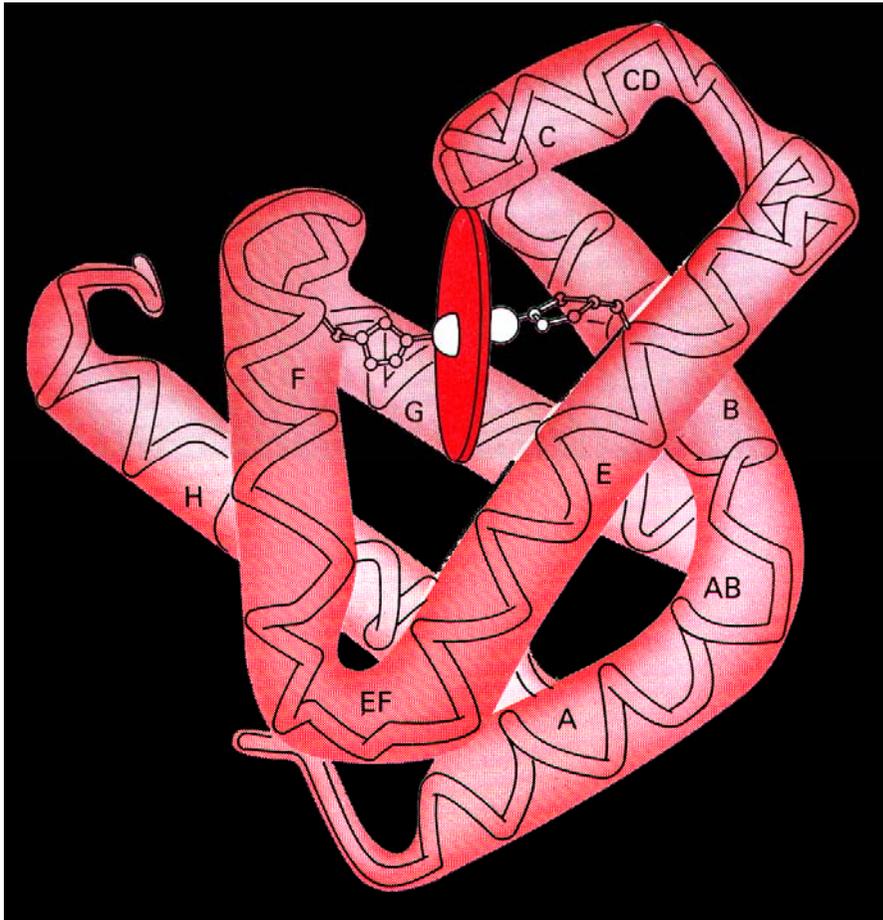
# Linus Pauline & $\alpha$ helix



Judson (1996) *The Eighth Day of Creation*

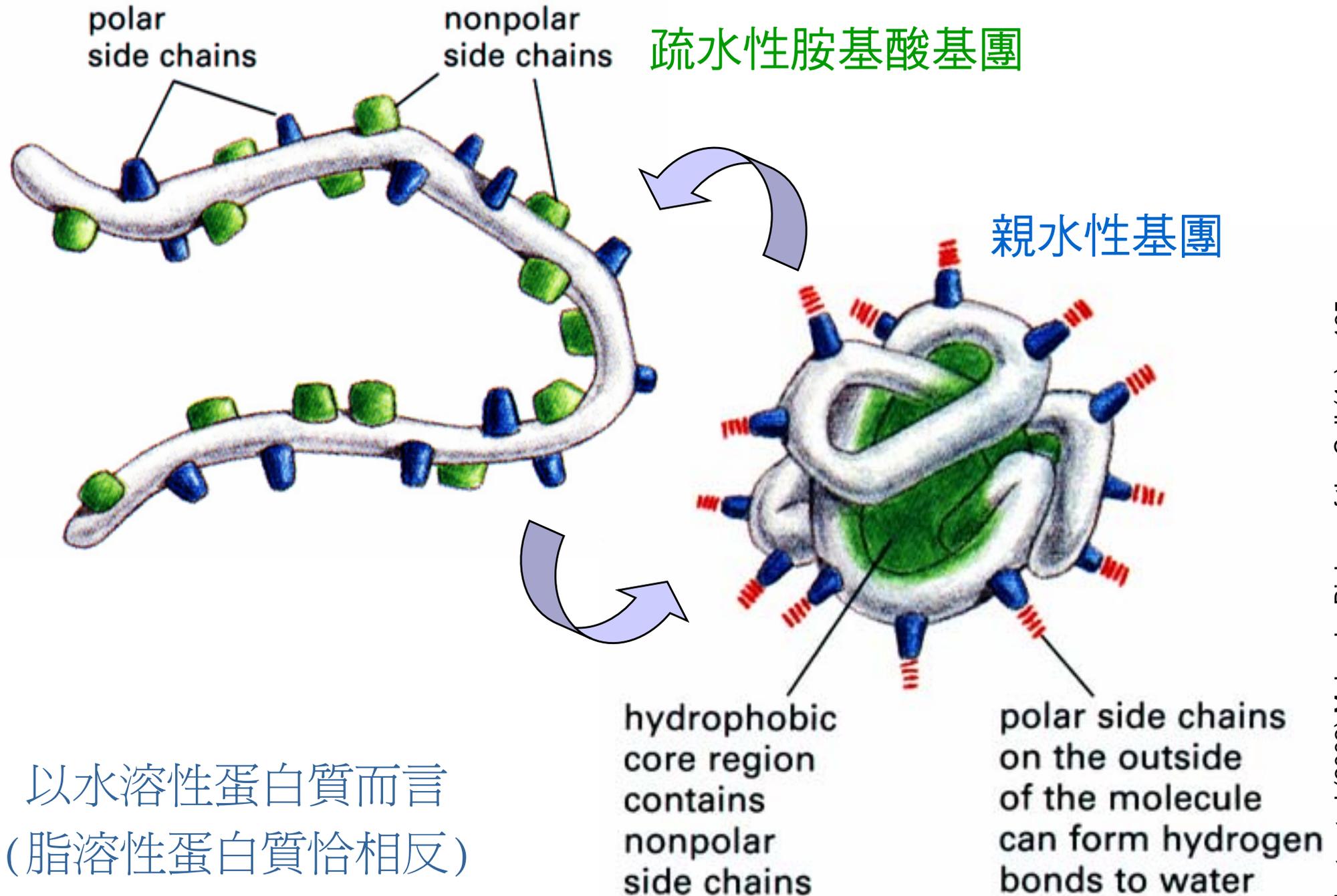


# 肌紅蛋白 myoglobin



用  $\alpha$  helix 組成堅固的立體構造

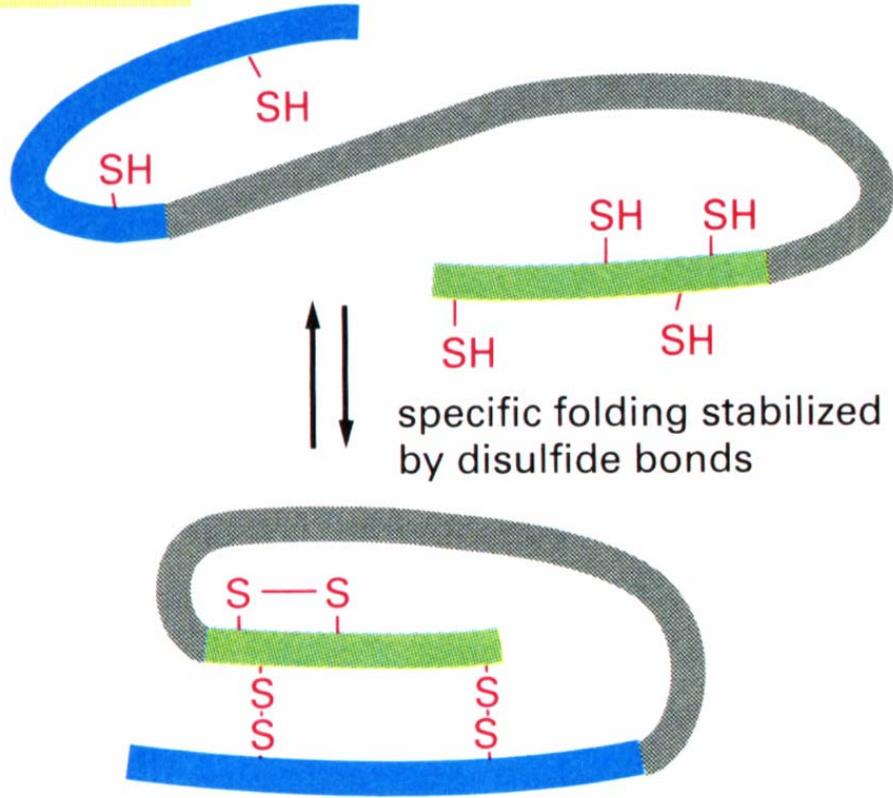
# 三級構造中的疏水性作用力



以水溶性蛋白質而言  
(脂溶性蛋白質恰相反)

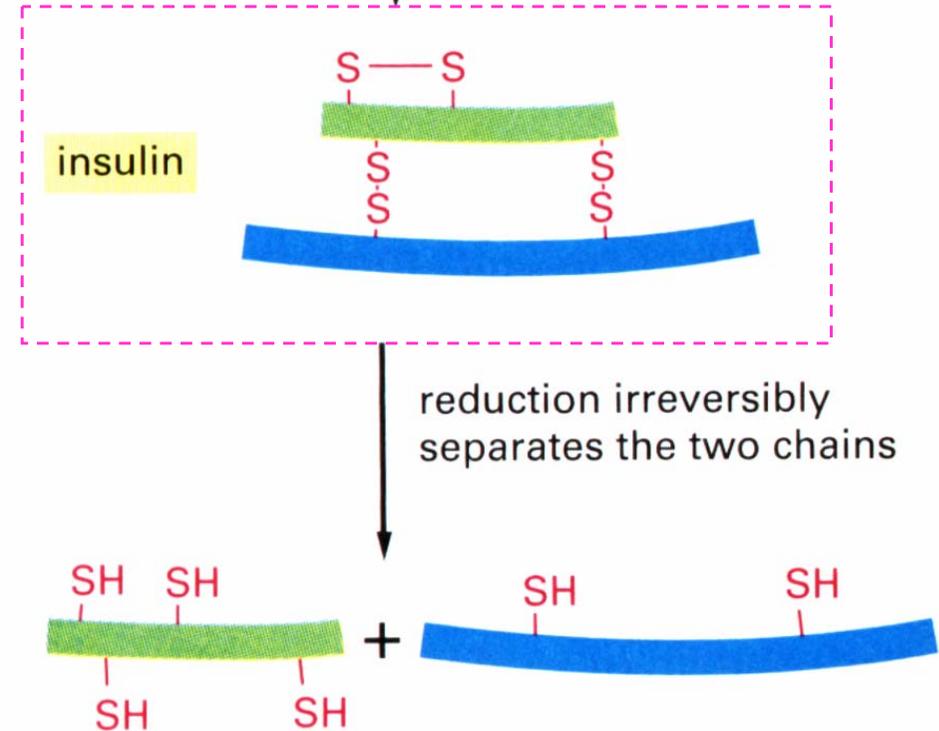
# 裂解是調節蛋白質功能的手段之一

proinsulin



connecting peptide removed, leaving complete two-chain insulin molecule

insulin

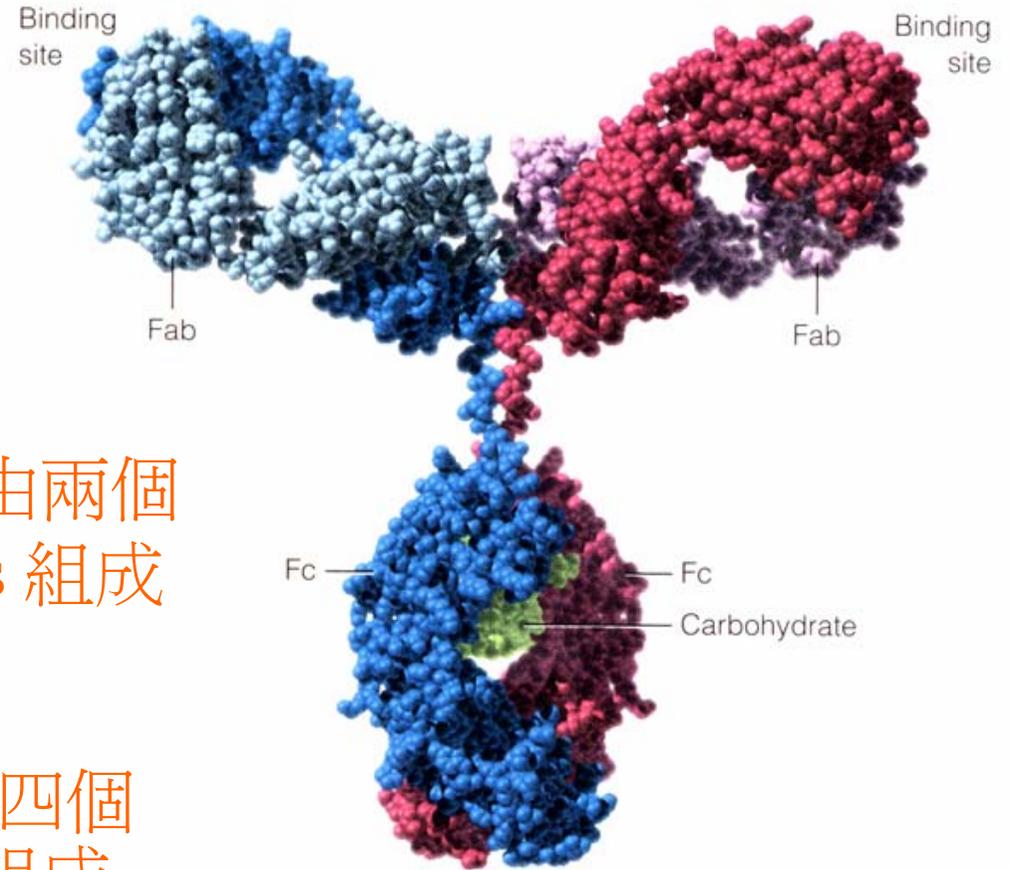
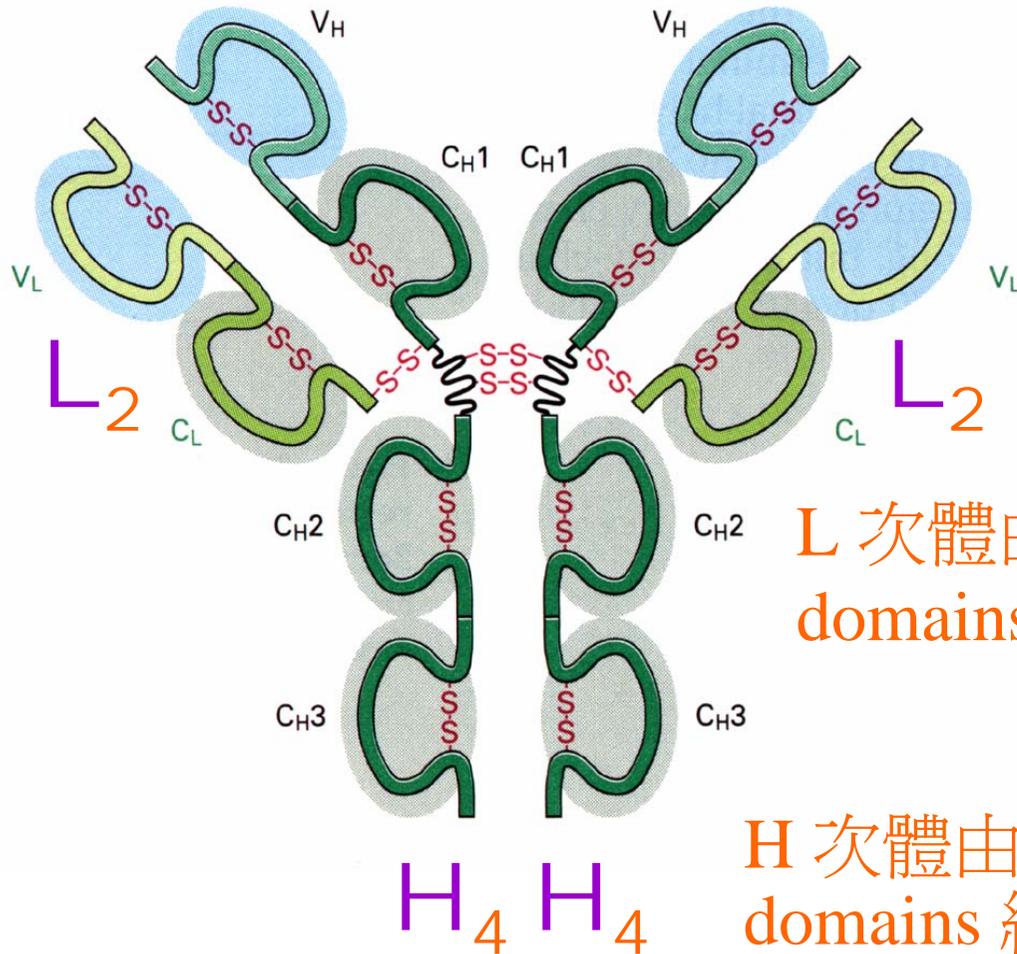


# 抗體具有十二個以上功能區塊



Edelman, Porter (1972)

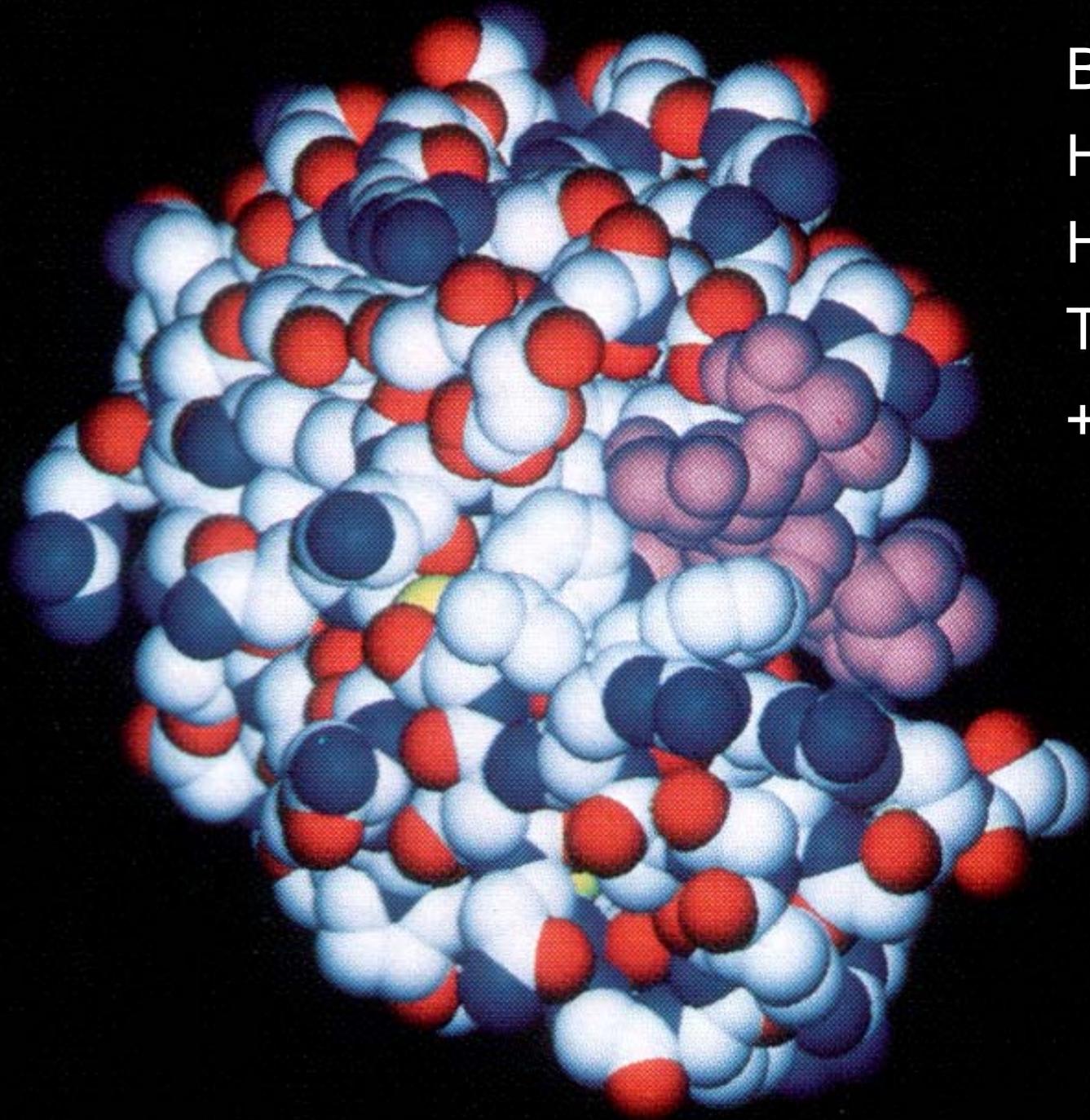
每一抗體分子有四條次體 (subunits) = 2H (heavy) + 2L (light)



各個 domain 可能由同一原始基因所演化

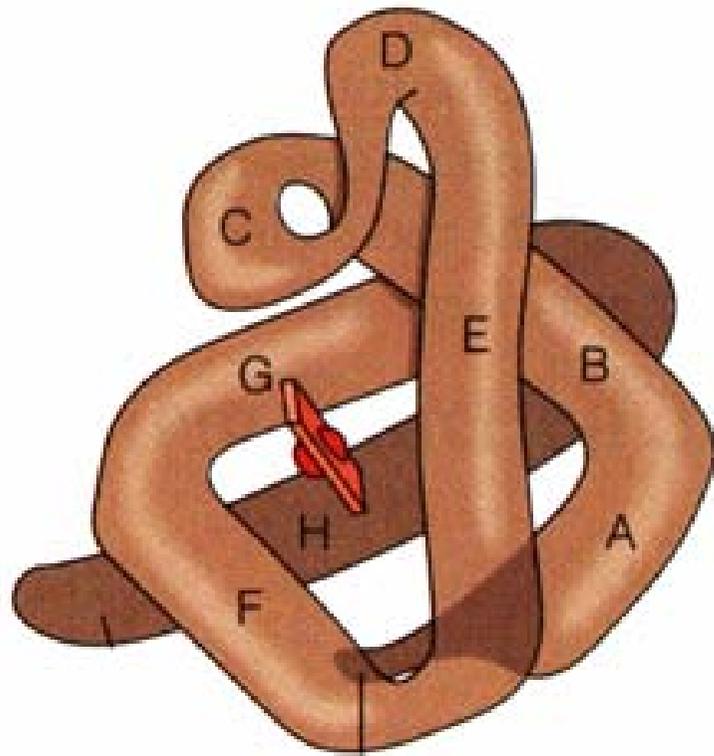
*Gene duplication*

# Lysozyme

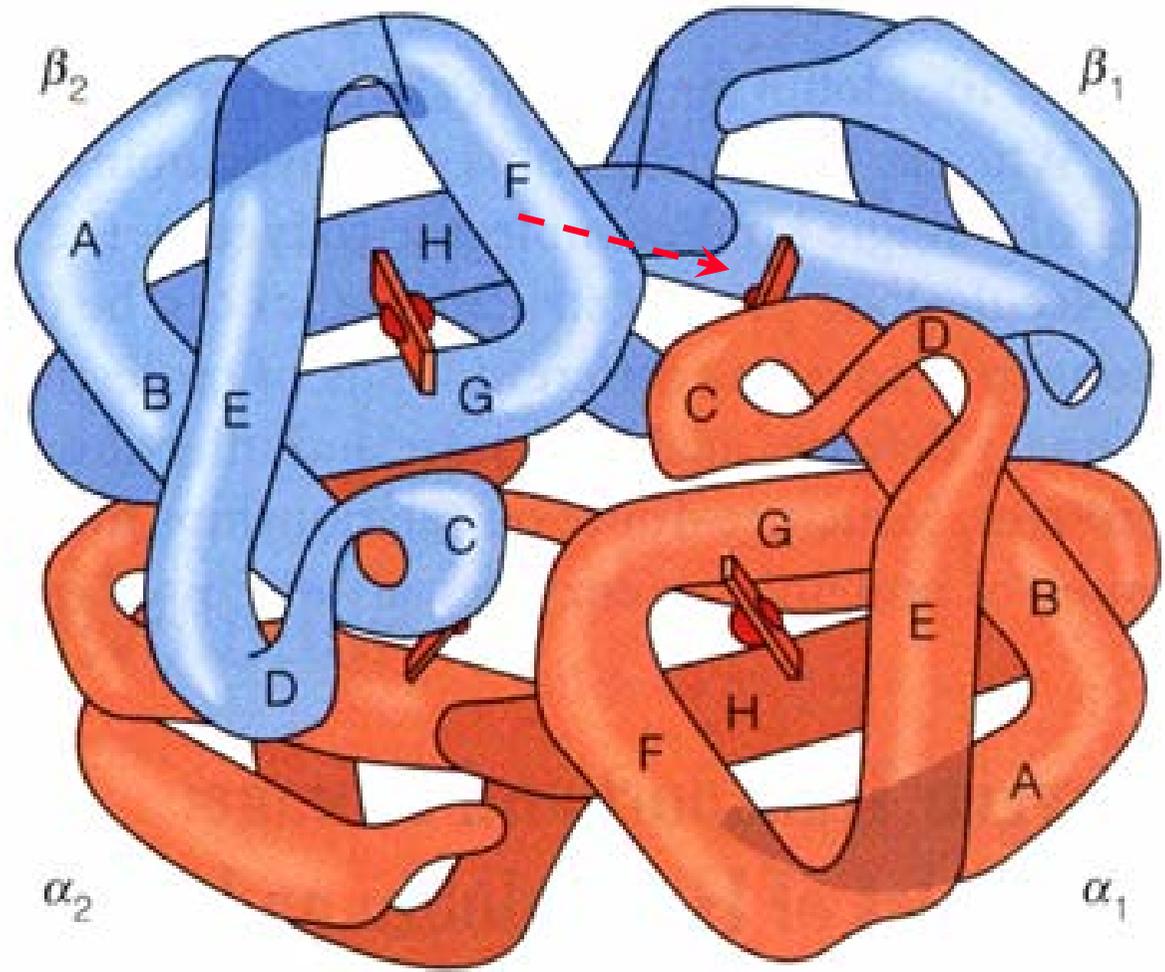


Backbone  
Hydrophobic  
Hydrophilic  
Total  
+ substrate

# 四級構造 (Quaternary structure)

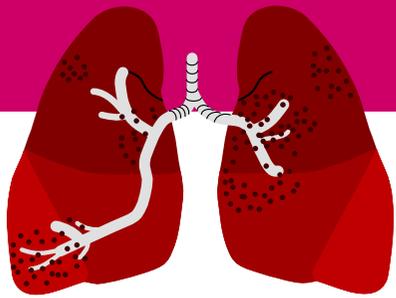


**Myoglobin**

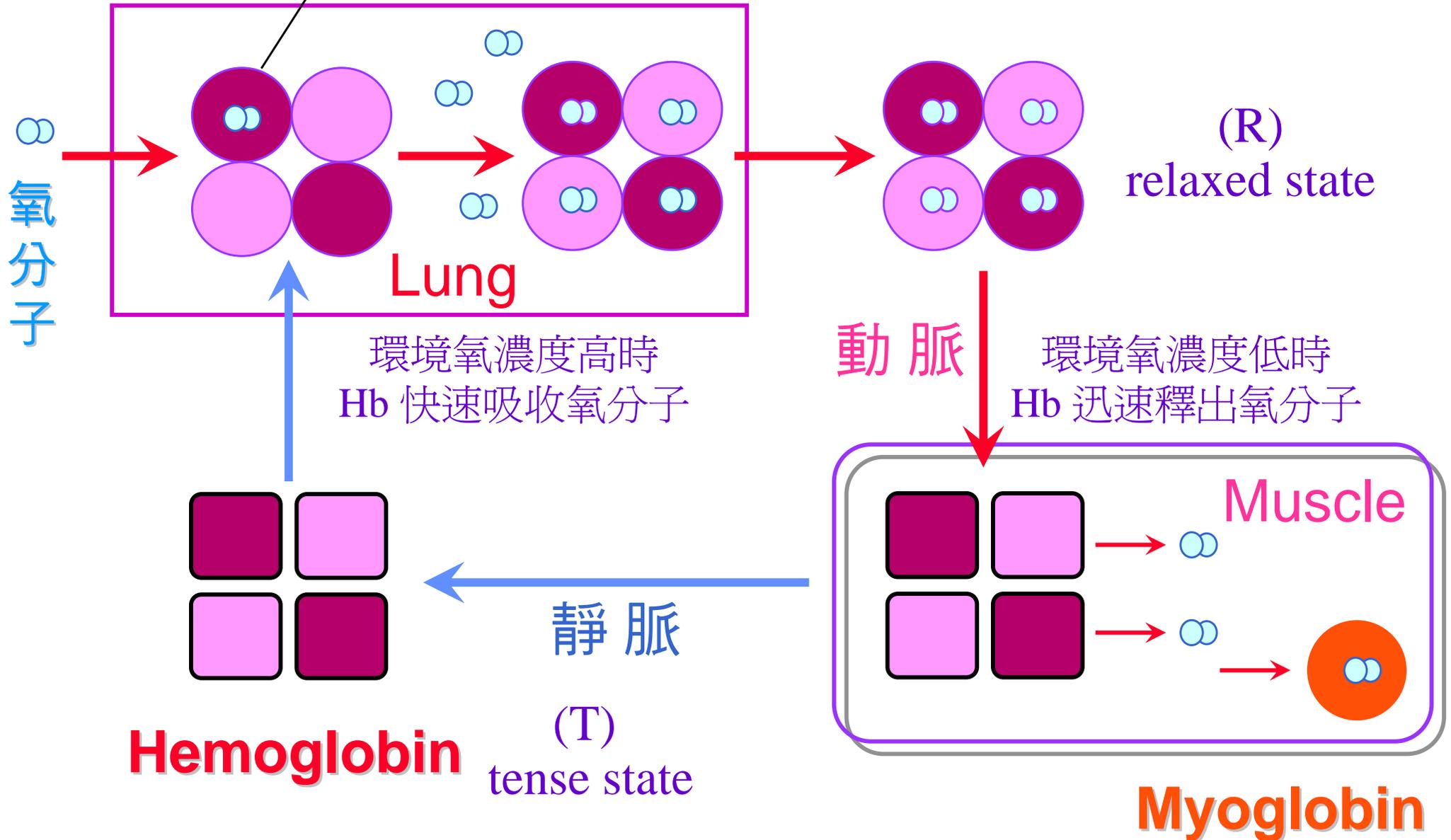


**Hemoglobin**

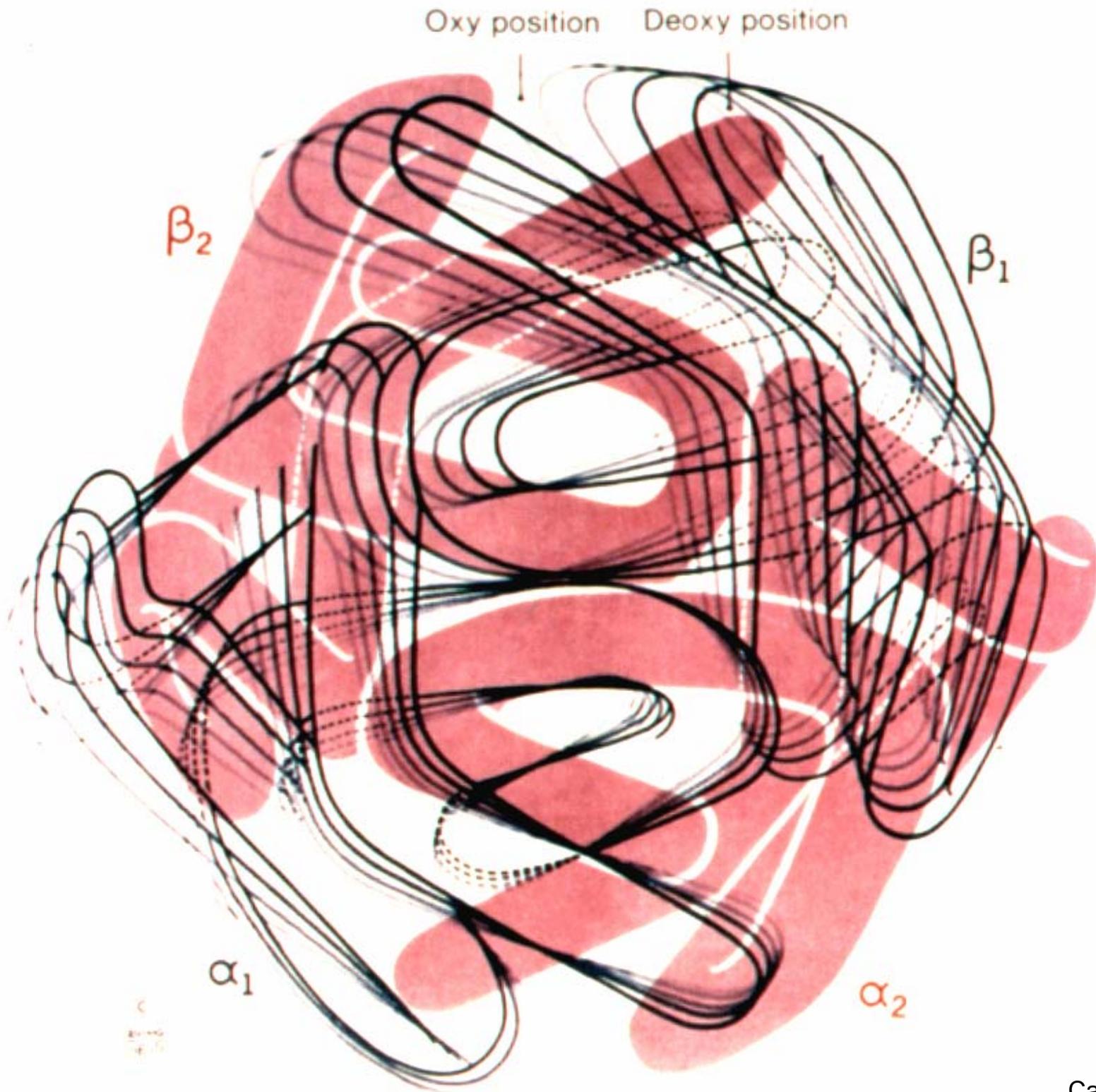
# 血中氧分子的運送



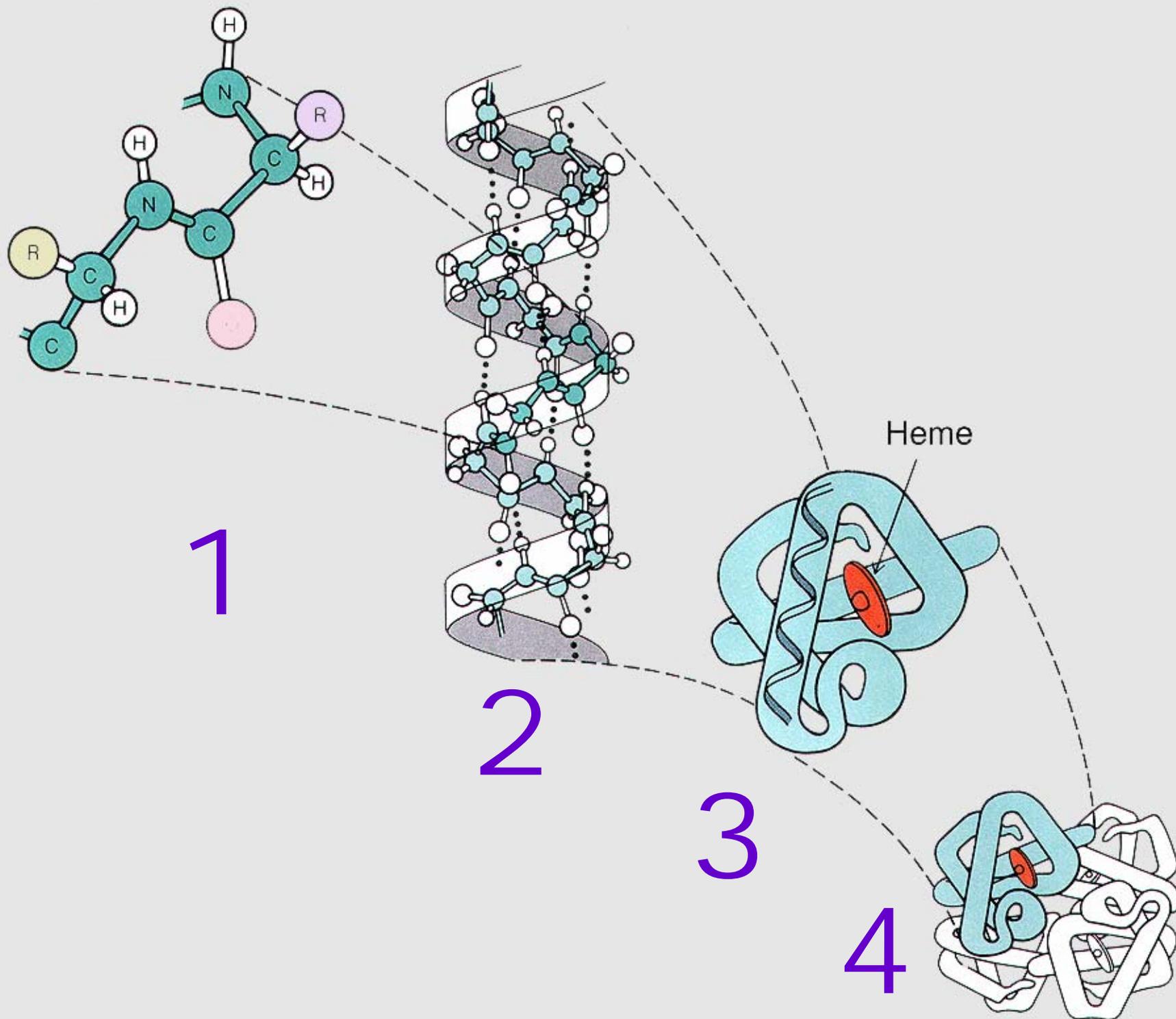
任何一個次體接受氧分子後，會增進其它次體吸附氧分子的能力



牽一髮動全身  
所有次體均受影響

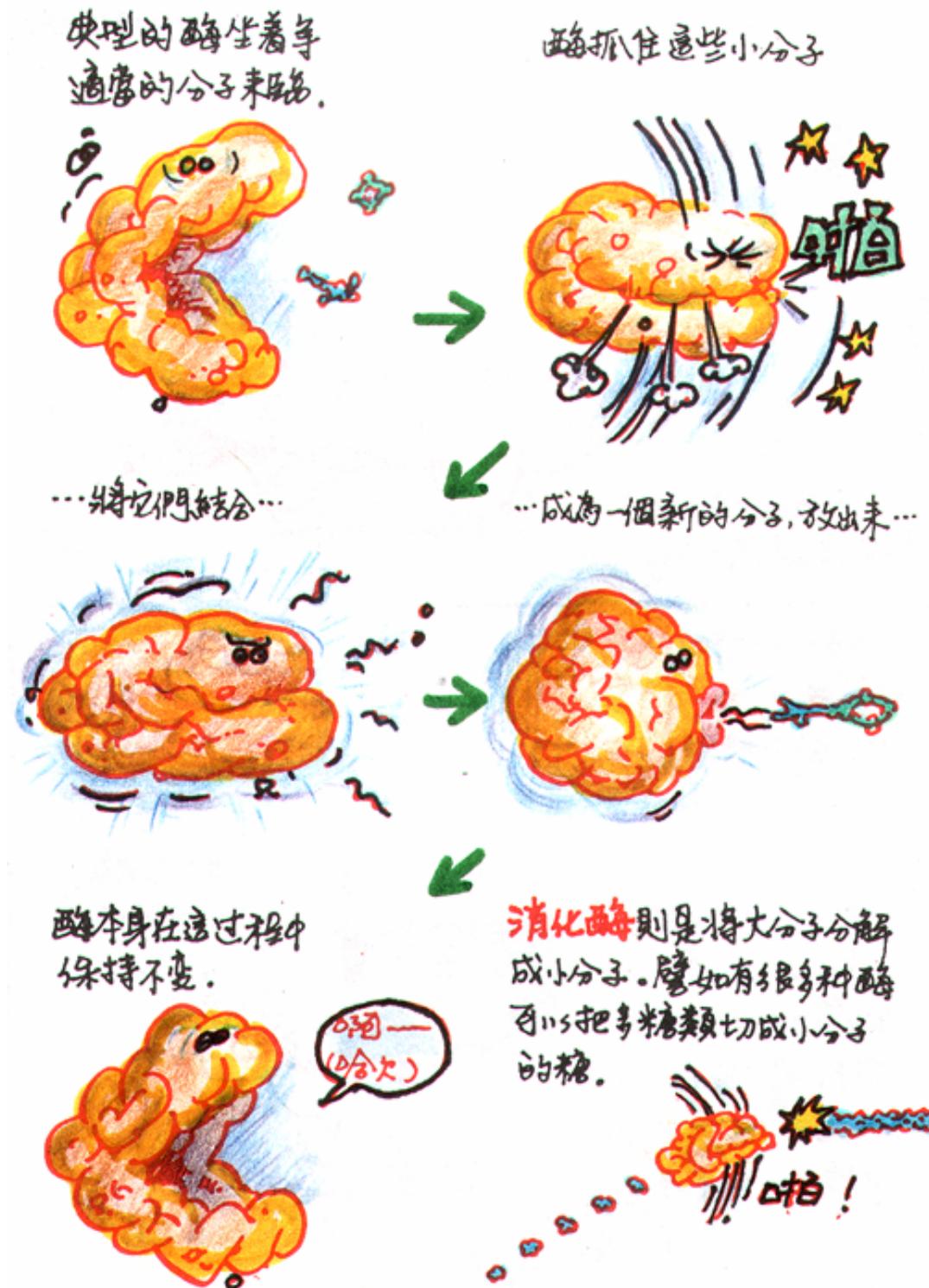


# 蛋白質的四級構造

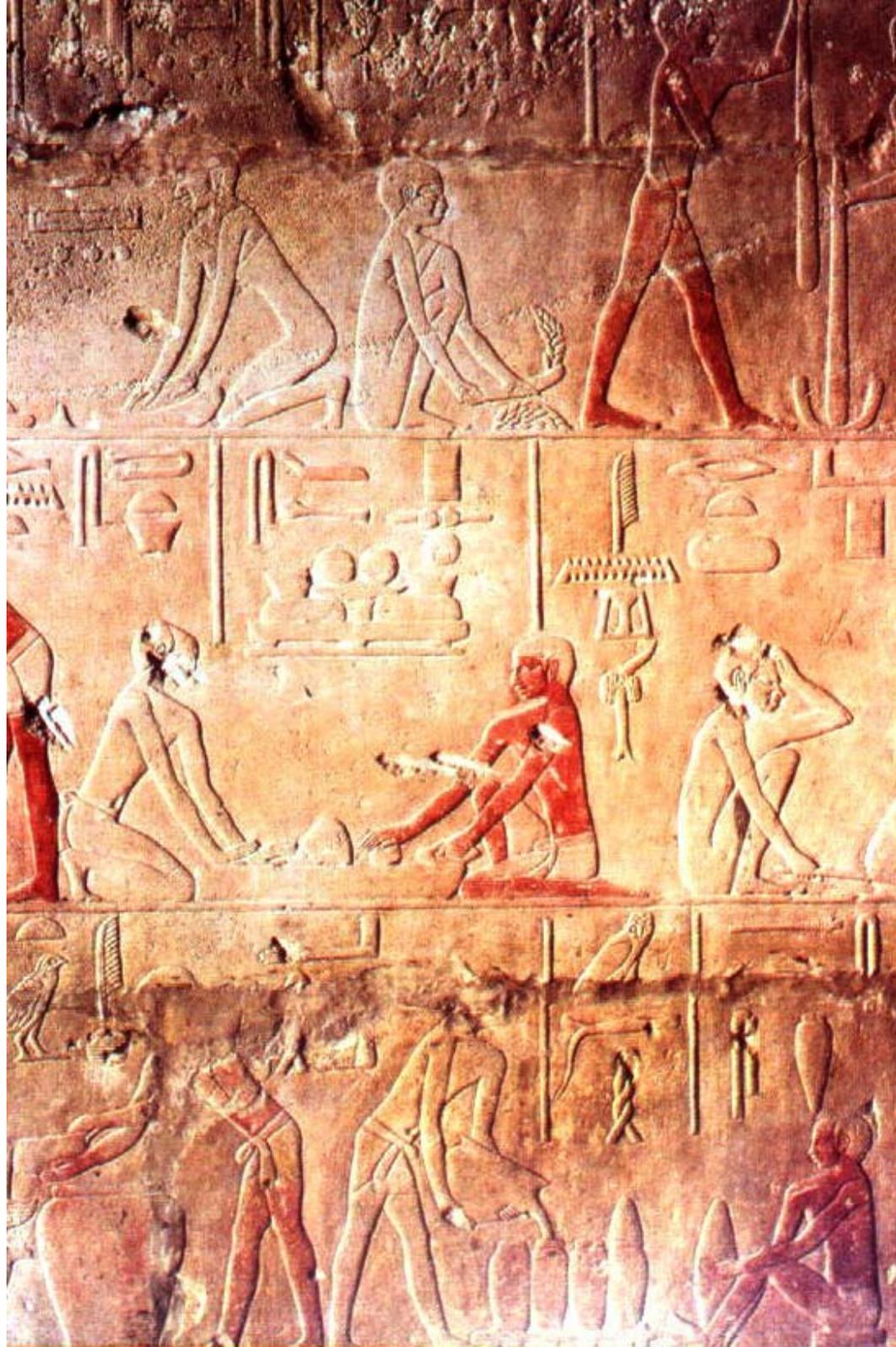


# 蛋白質的構形不是固定的

## Hexokinase



# 古埃及人用麥粉醱酵製造啤酒



磨粉 去糠 打碎

麥芽 萌發 浸潤

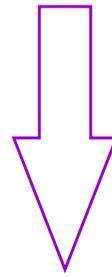
成酒 醱酵 裝瓶

# 酵素很早就為人類所利用

## 埃及壁畫



神奇轉化力量？



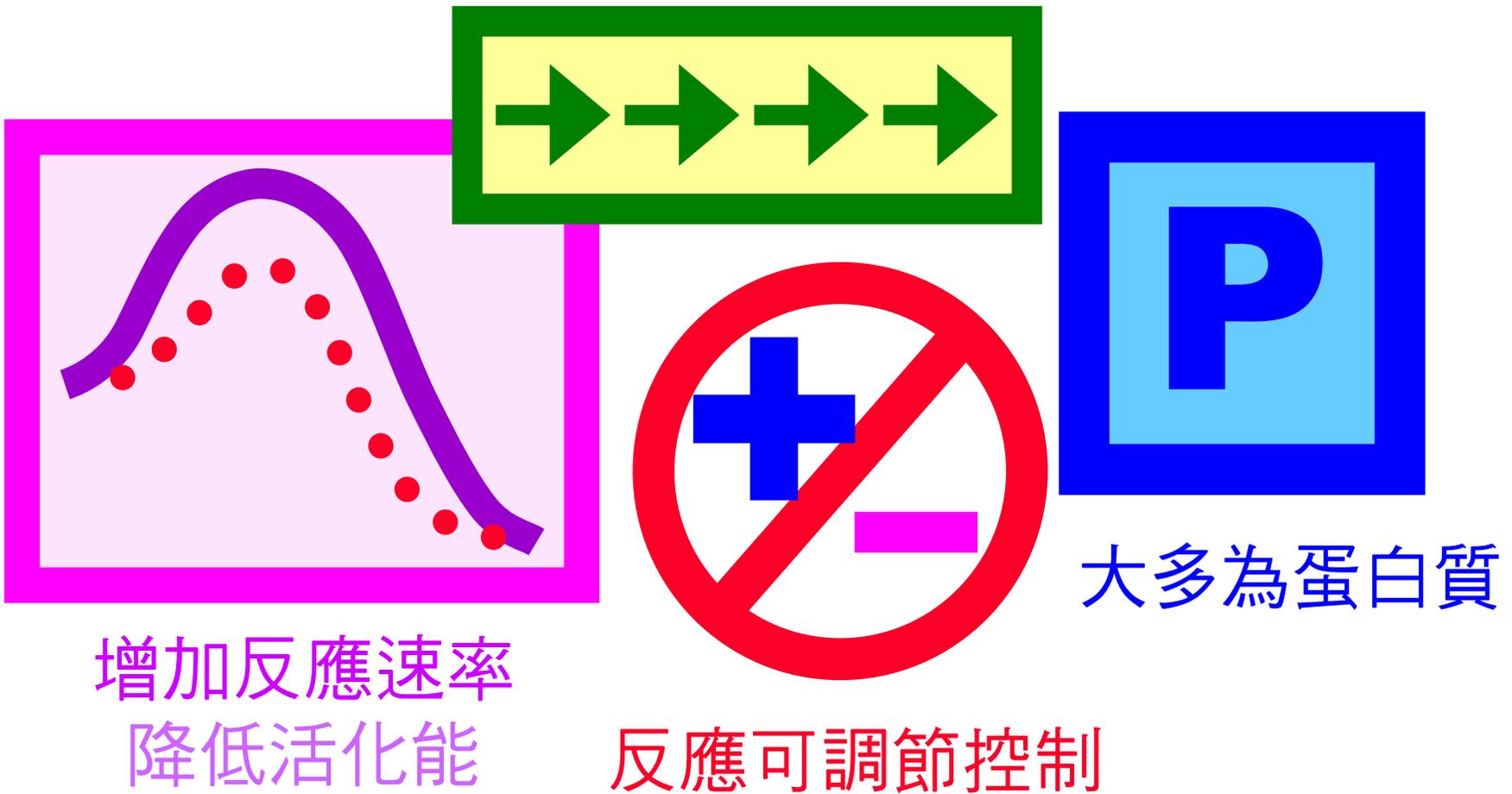
醱解作用產生酒精

■ 1926 *Sumner* 結晶出 **urease** 可分解尿素

酵素是一種蛋白質

# 酵素印象

催化專一性反應



# 酵素的催化效果



1

雙氧水裂解



1,000

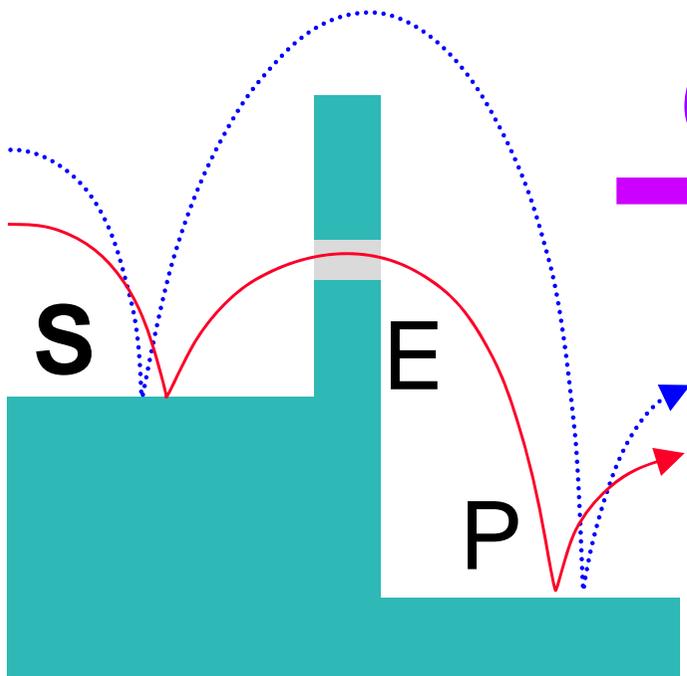


1,000,000

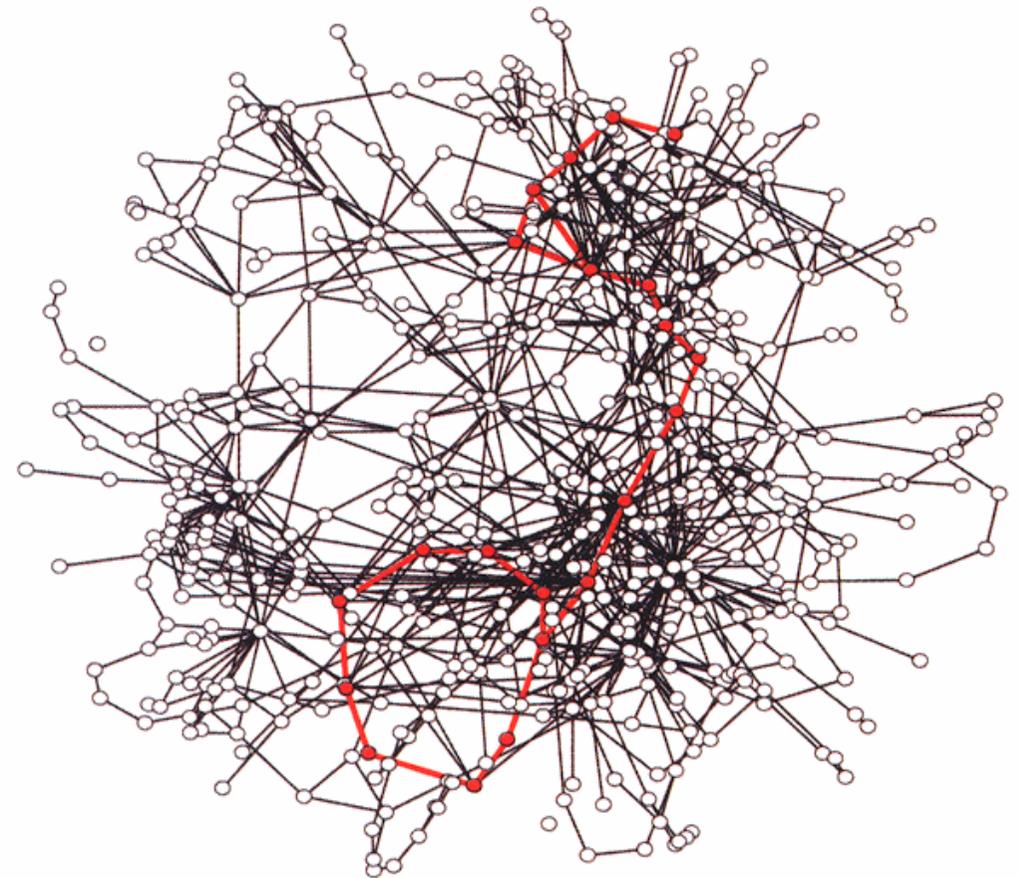
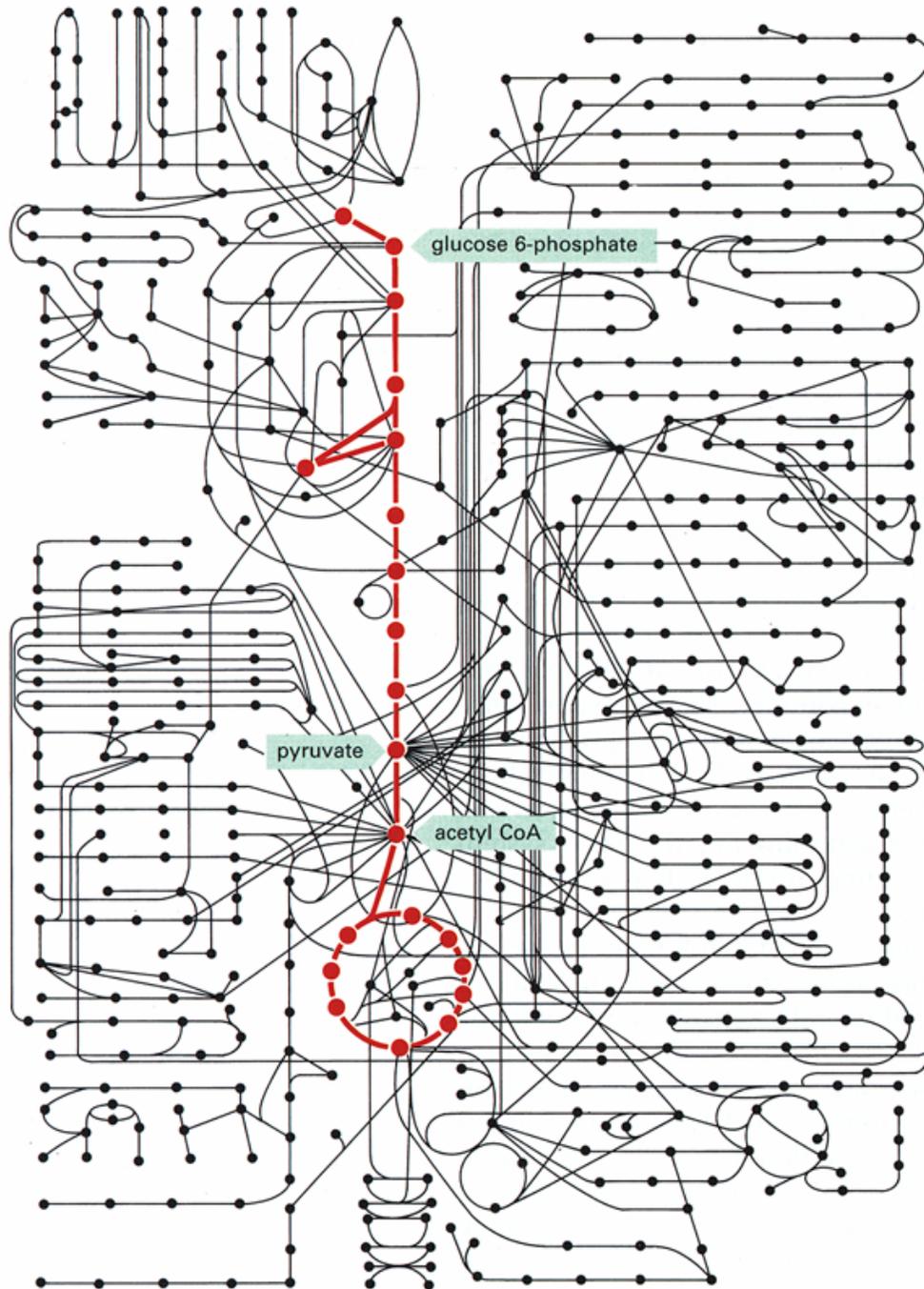


1,000,000,000

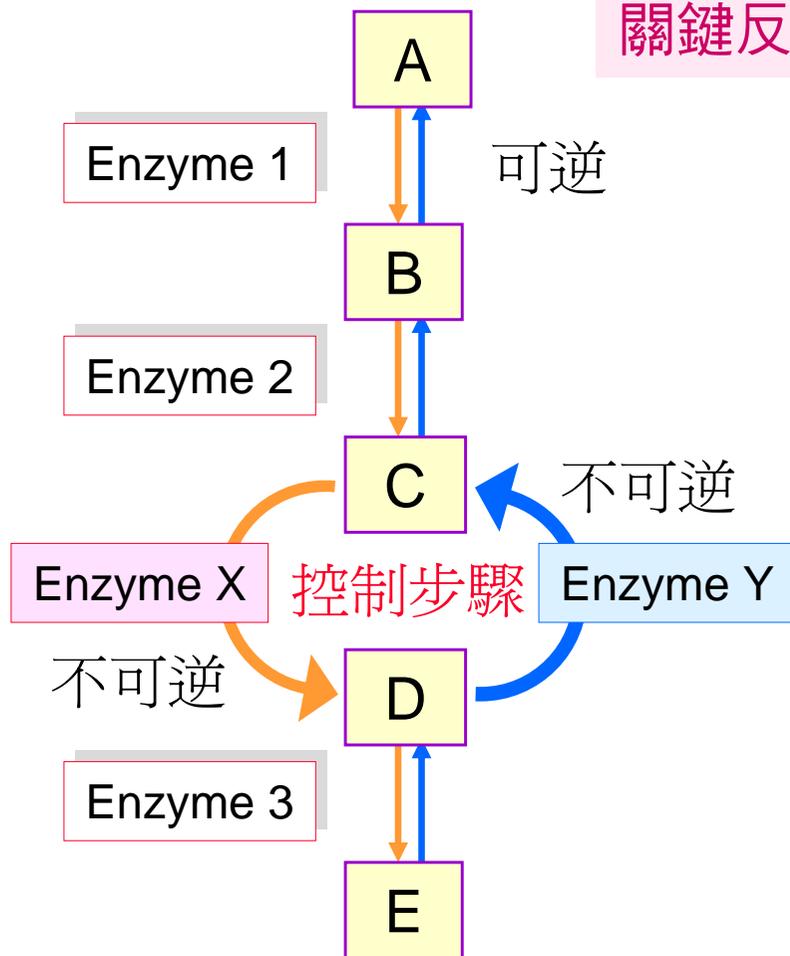
反應速率



# 酵素參與細胞所有代謝路徑



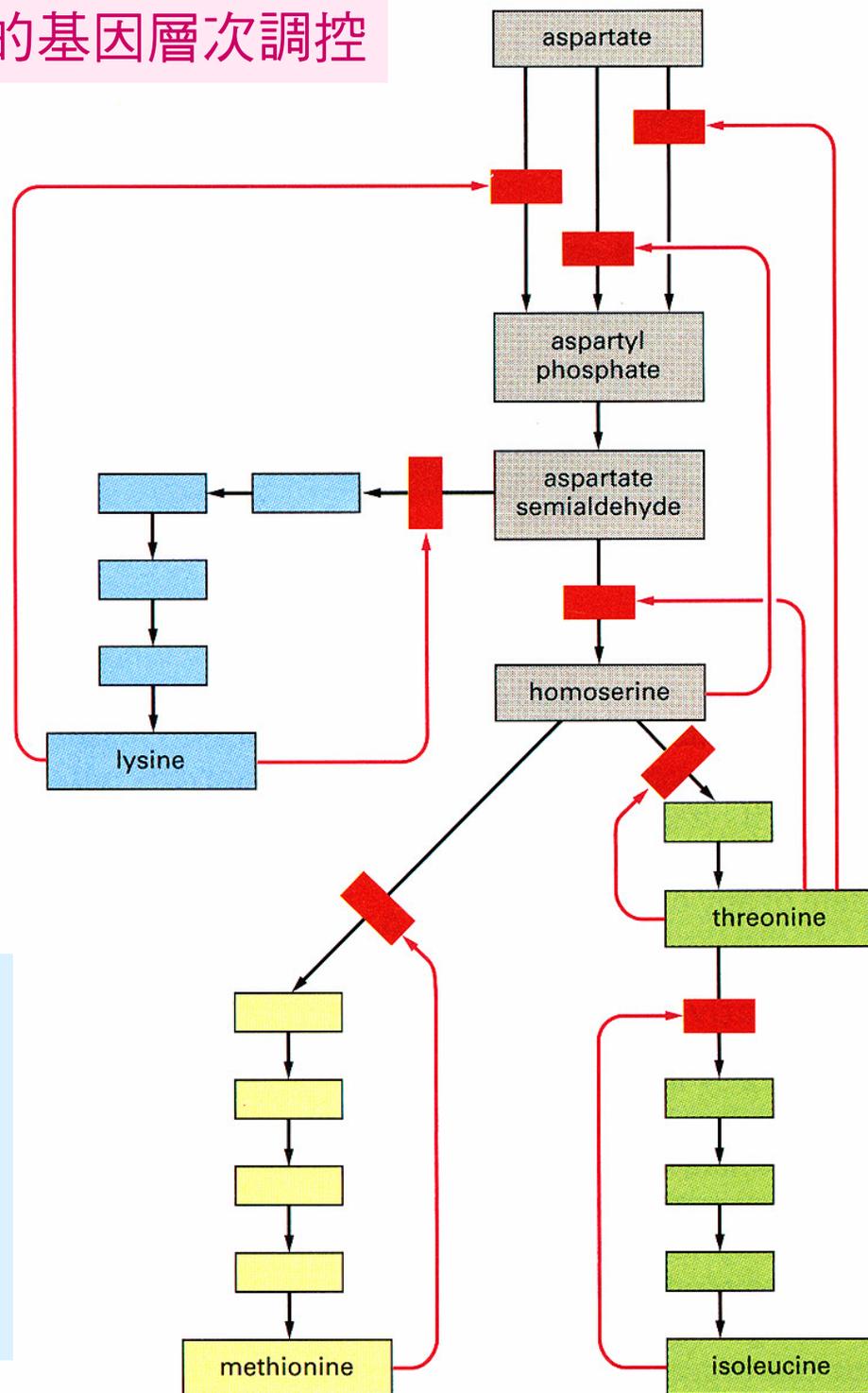
# 代謝路徑調控原則



- (1) 每一步反應均有酵素催化
- (2) 代謝路徑有速率決定步驟
- (3) 反應可能為可逆或不可逆
- (4) 各條代謝路徑之間可互通

Yang Cycle (楊祥發發現植物乙烯合成路徑)

## 關鍵反應的基因層次調控



# 蛋白質四級構造的意義

一級構造 (胺基酸序列)



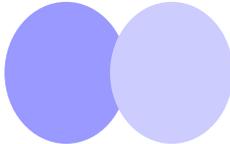
二級構造 ( $\alpha$  helix,  $\beta$  sheet)



*Domain* (區塊  $\alpha\alpha\beta\beta$ )

三級構造 (monomer) 



四級構造 (dimer) 

構形

有形体?

序列



構形



活性



調節

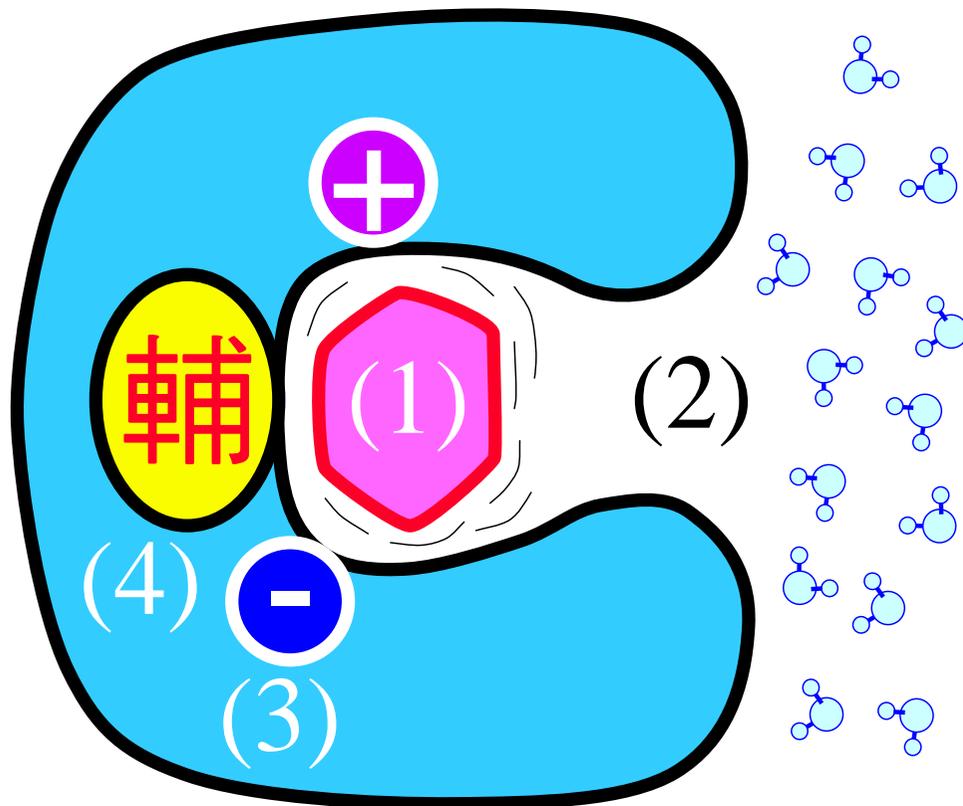
有生命?

有智慧?

# 酵素活性區是一個凹陷的口袋

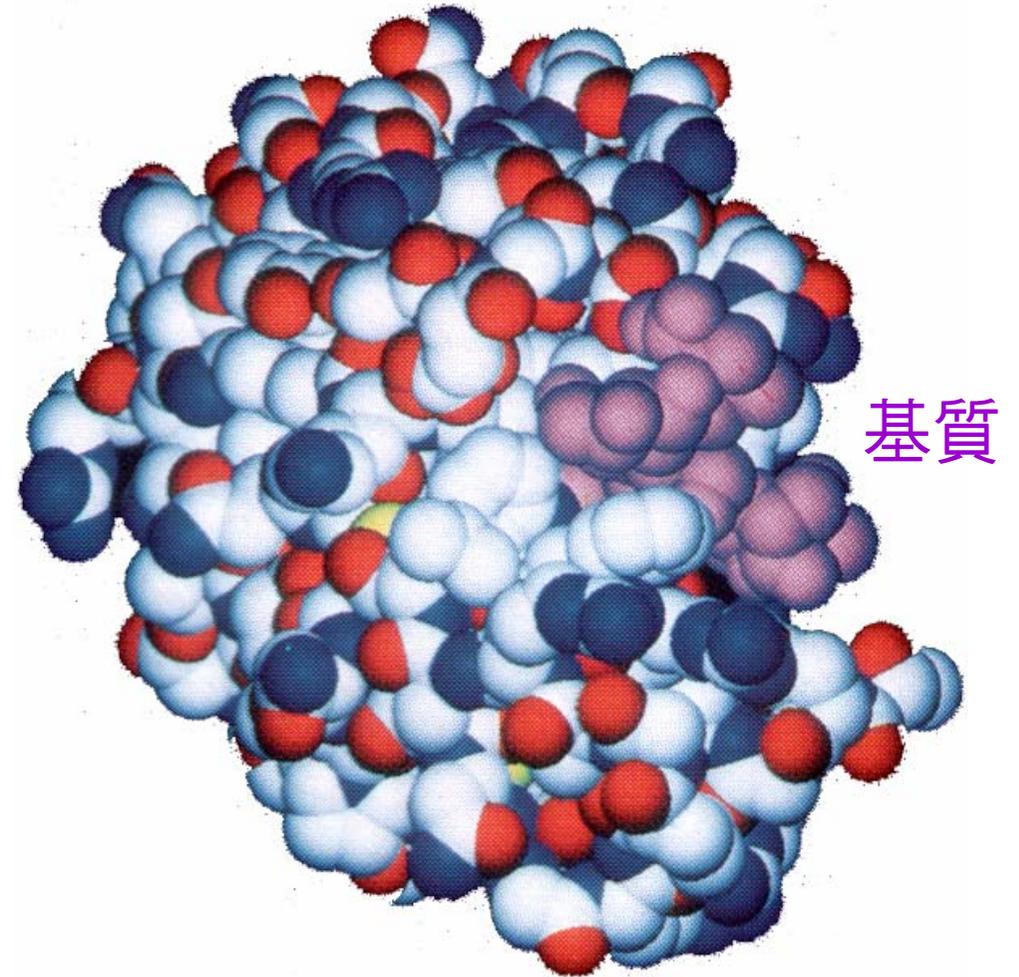
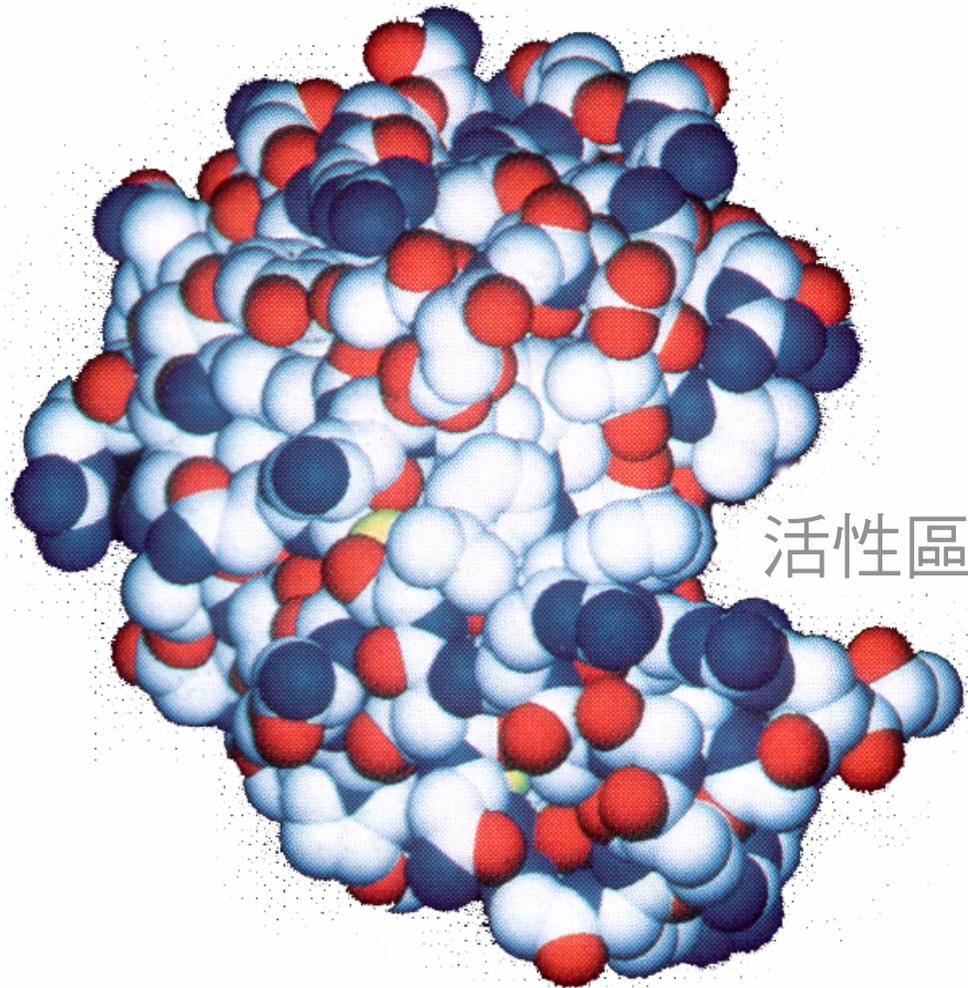
酵素活性區為何可降低活化能？

是一個魔術口袋



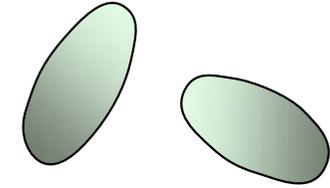
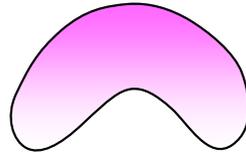
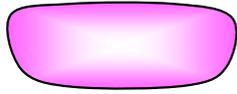
- (1) 可穩定過渡狀態
- (2) 防止水分子干擾
- (3) 具高反應性基團
- (4) 有輔酶幫助反應

# Lysozyme 活性區的口袋



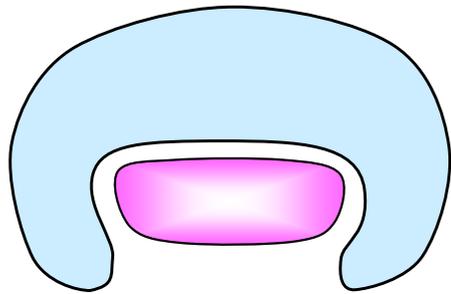
# 棒棒脢 Stickase

基質

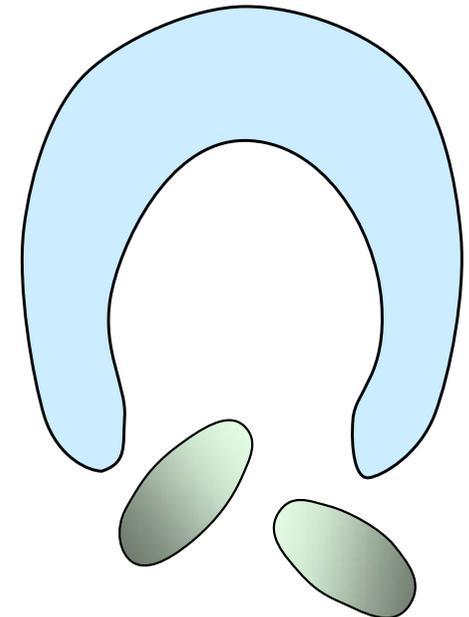
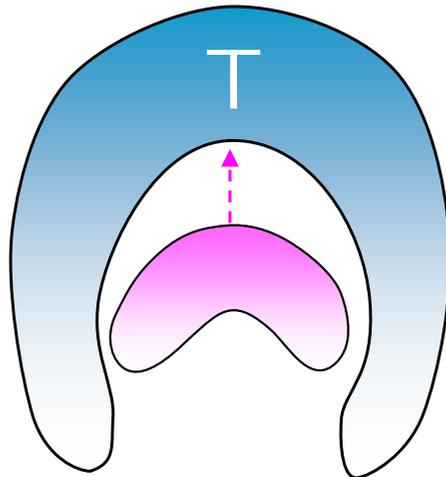
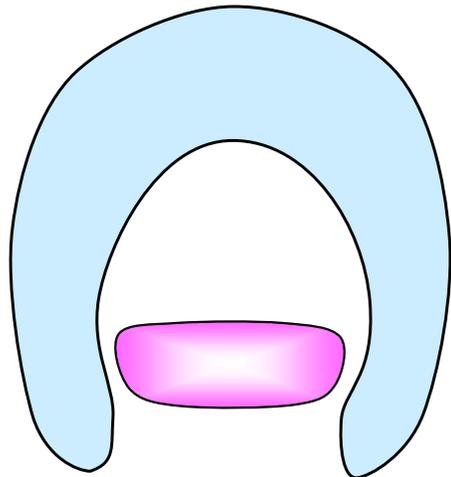


中間過渡狀態

生成物



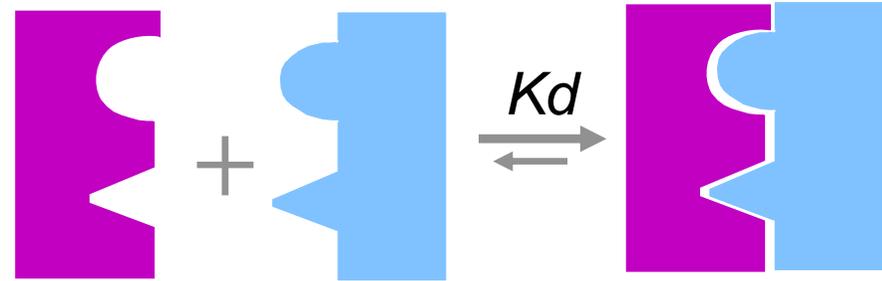
**X** 若只是與基質互補結合則無催化反應



酵素不但能基質結合 還會誘導過渡狀態生成

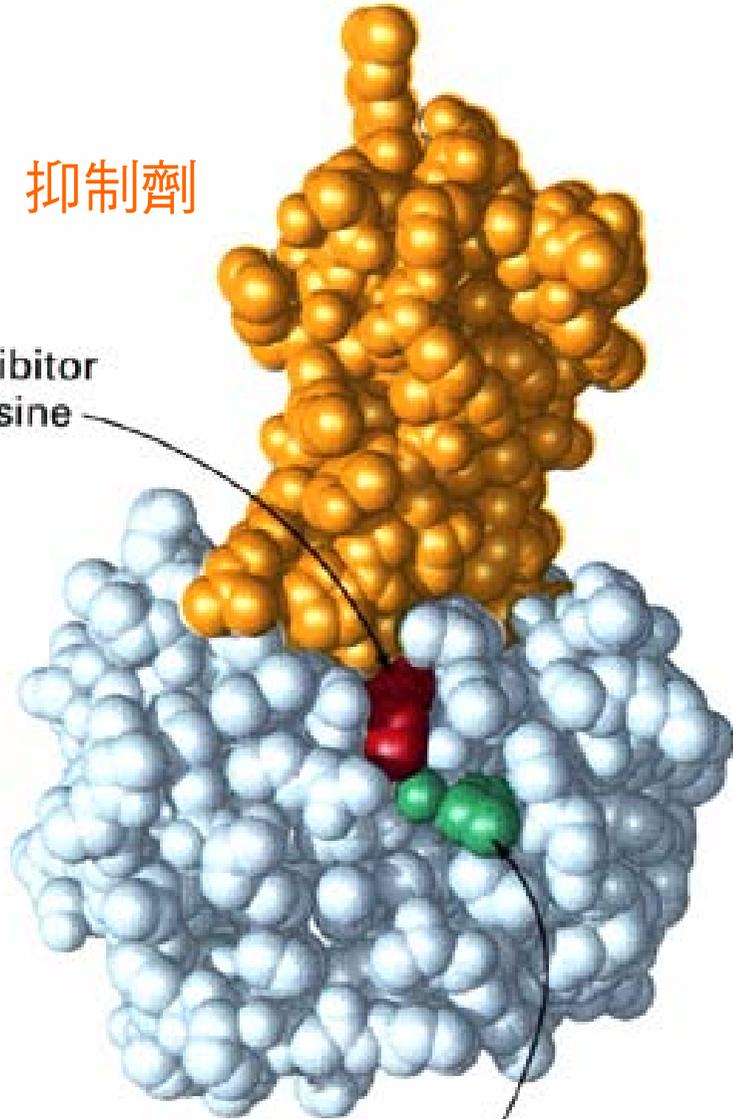
# 蛋白質間專一性的形成

## I. Conformational Match: Van der waals interaction



抑制劑

Inhibitor  
lysine



胰蛋白酶

Active site  
aspartate

## II. Interaction Forces:

- (1) Hydrogen bond
- (2) Hydrophobic interaction
- (3) Electrostatic interaction
- (4) Van der waals interaction

