

생유기화학  
(*Bioorganic Chemistry*)

Amino Acids, Peptides, and Proteins-III  
(아미노산, 펩타이드, 단백질-3)

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순천향대

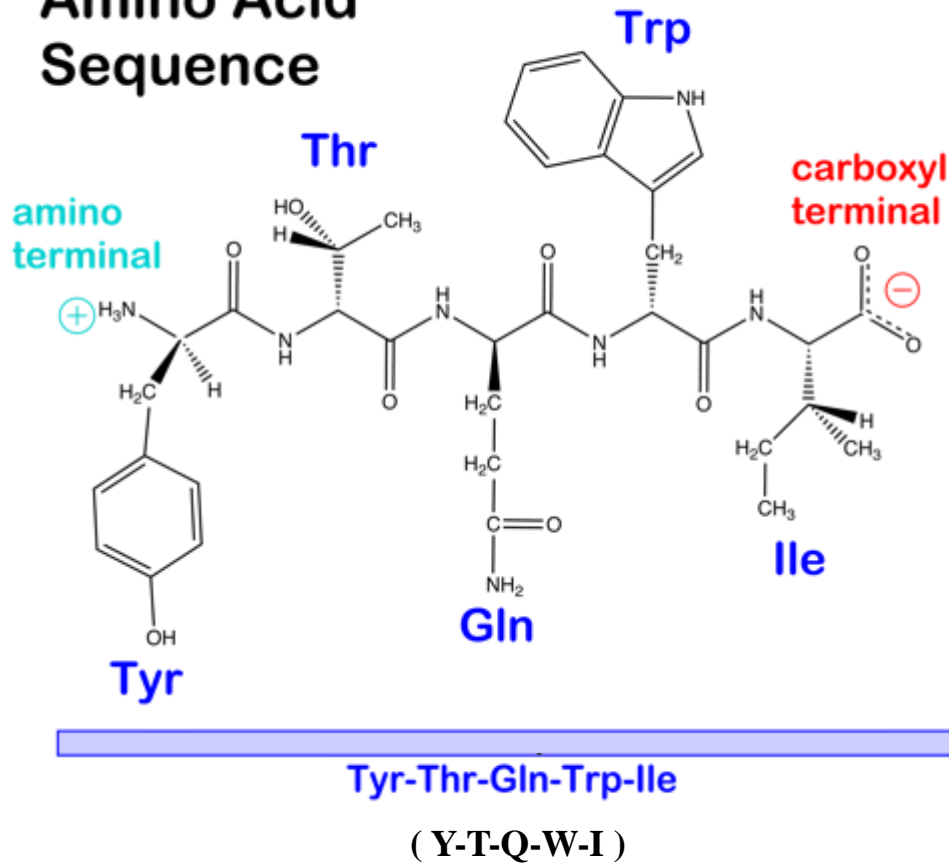
나노화학공학과

임정균 교수



## 9~10. Proteins, Structures

### Direction of an Amino Acid Sequence

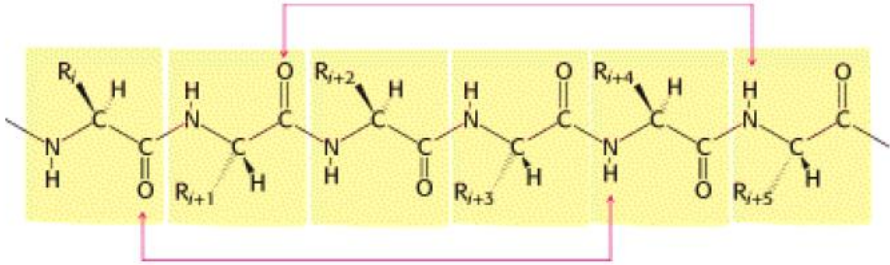


### Primary structure

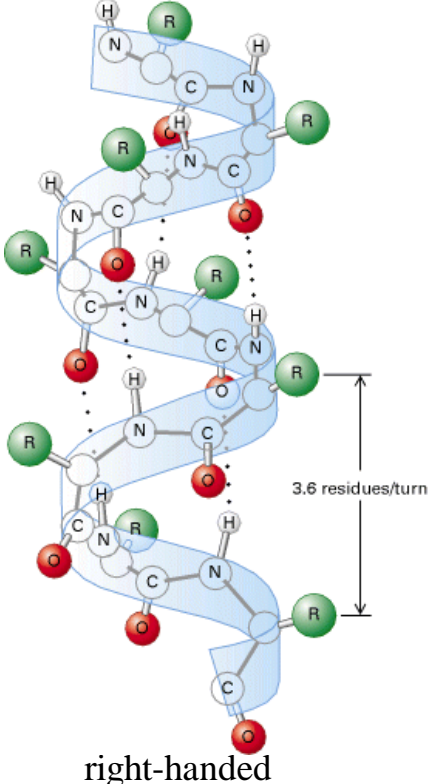
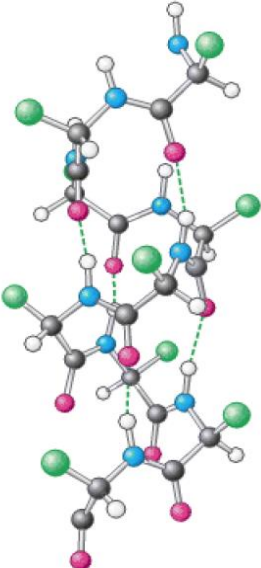
the sequence of amino acids in the peptide chain

# Alpha helix

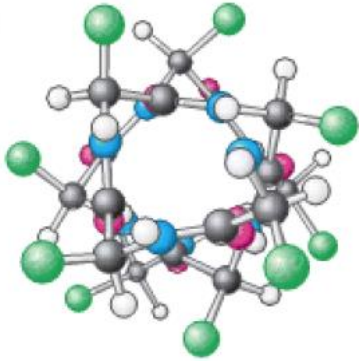
a coiled structure stabilized by ( ) hydrogen bonds



Hydrogen bonding between n and n+4 residue

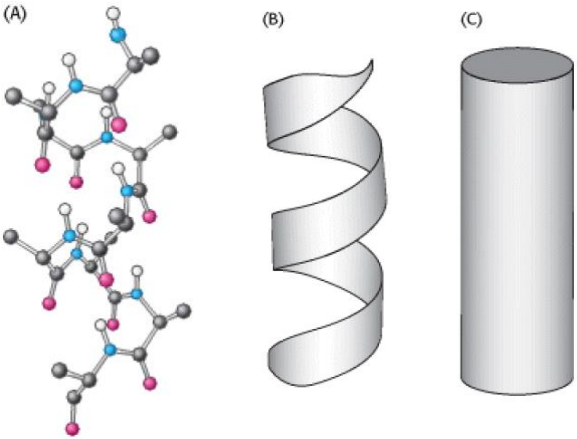


top view



R groups are pointing out

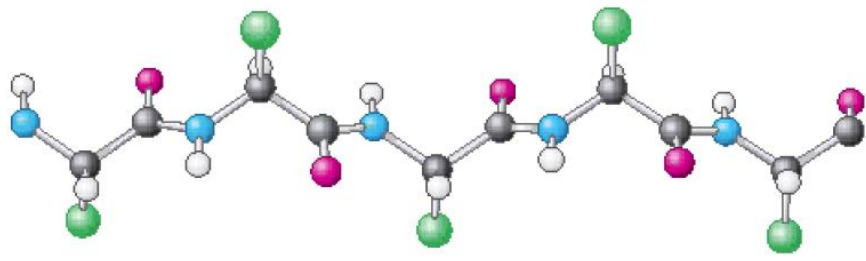
단백질에 따라서  $\alpha$ -helix의 분포는 0%에서 100%까지 다양하다.



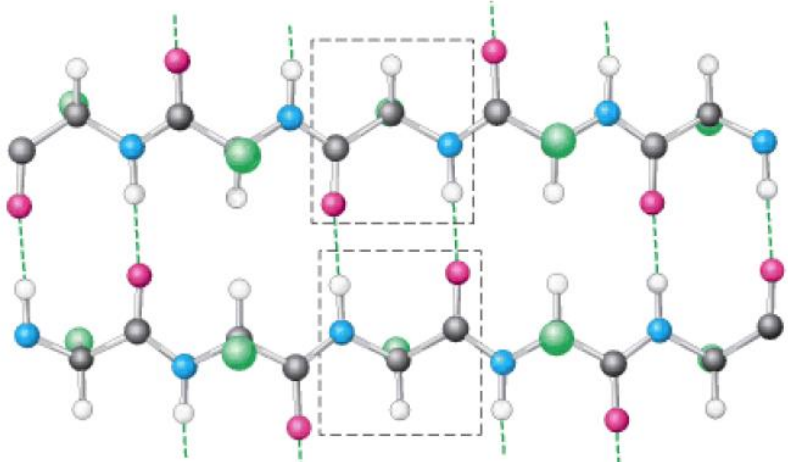
Schematic view of alpha helix: (a) ball-stick model, (b) ribbon, (c) cylindrical shape

# Beta sheet

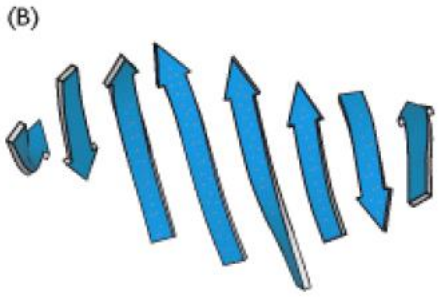
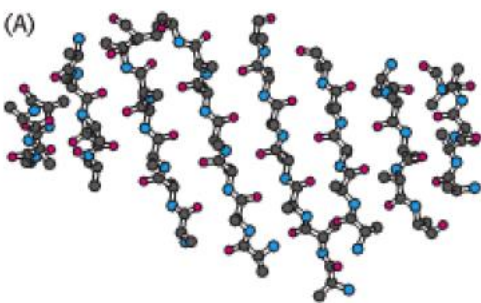
- stabilized by ( ) hydrogen bondings
- beta strand는 coil의 형태가 아닌 extended structure를 갖고있음



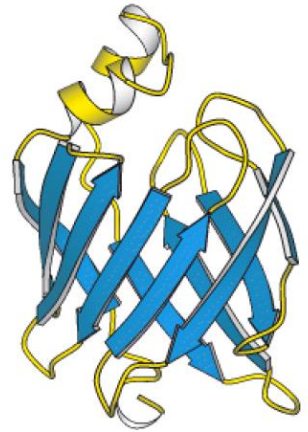
residue는 peptide plane에서 90°



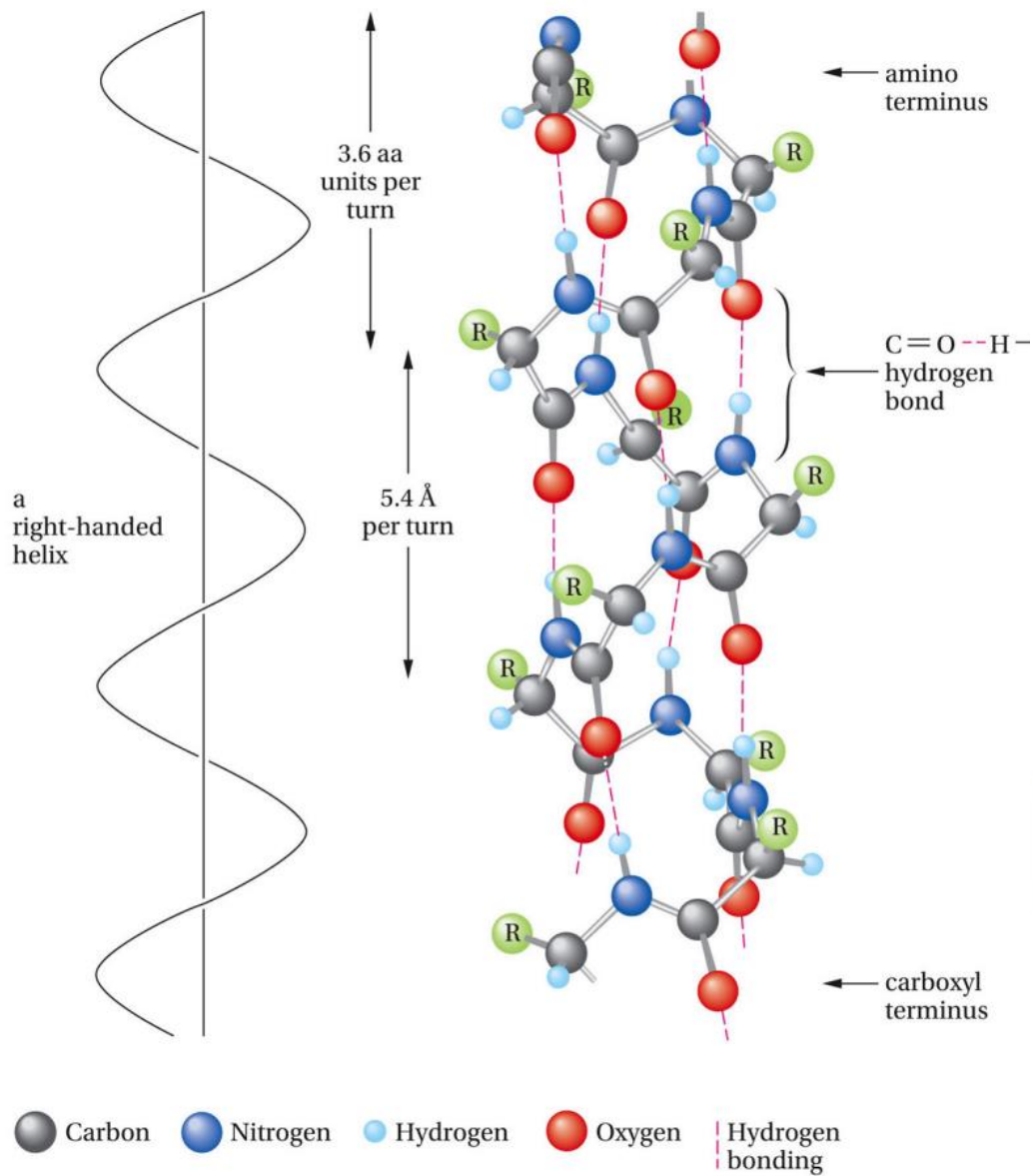
antiparallel beta sheet  
(parallel beta sheet도 가능)



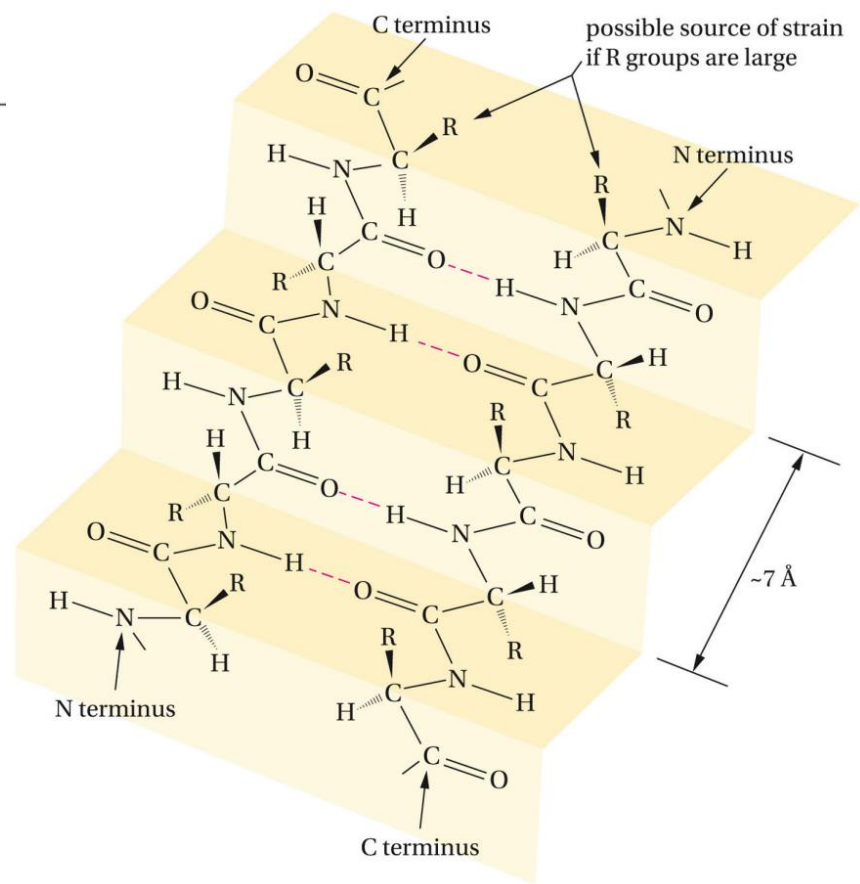
Schematic view of beta sheets: (a) ball-stick model, (b) schematic model



A protein rich in β sheets



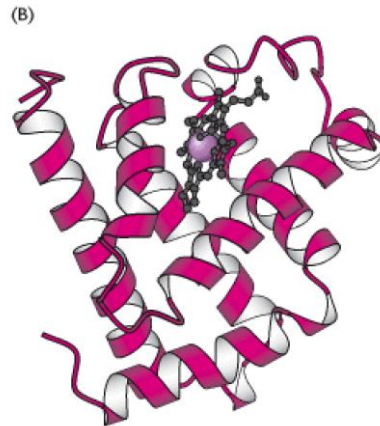
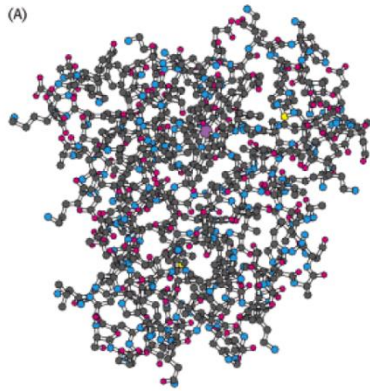
( $\alpha$ -helix)



( $\beta$ -sheet)

## Tertiary structure

Folding into compact structures



Myoglobin(oxygen carrier in muscle)의 구조

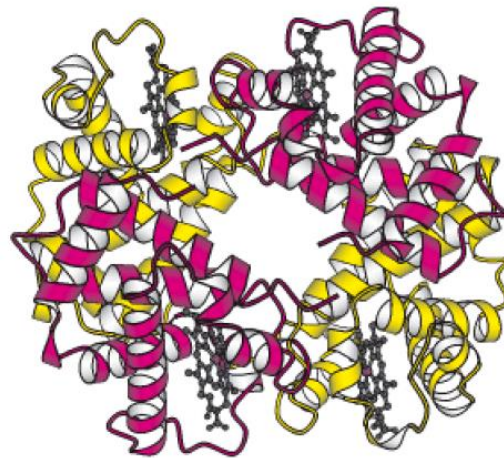
Protein의 안쪽은 대부분 nonpolar residue 존재,  
바깥 부분은 대부분 polar residue들이 존재 →  
driving force for protein folding

polar residue들은 hydrogen bonding, nonpolar  
residue들은 van der Waals attraction

## Quaternary structure

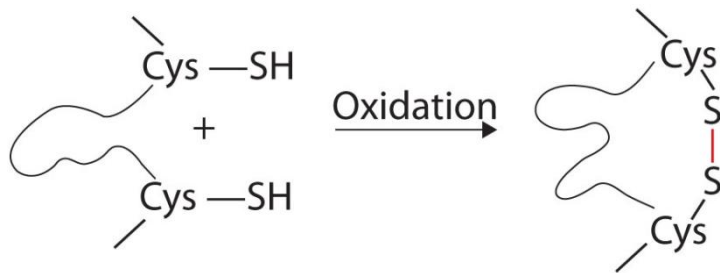
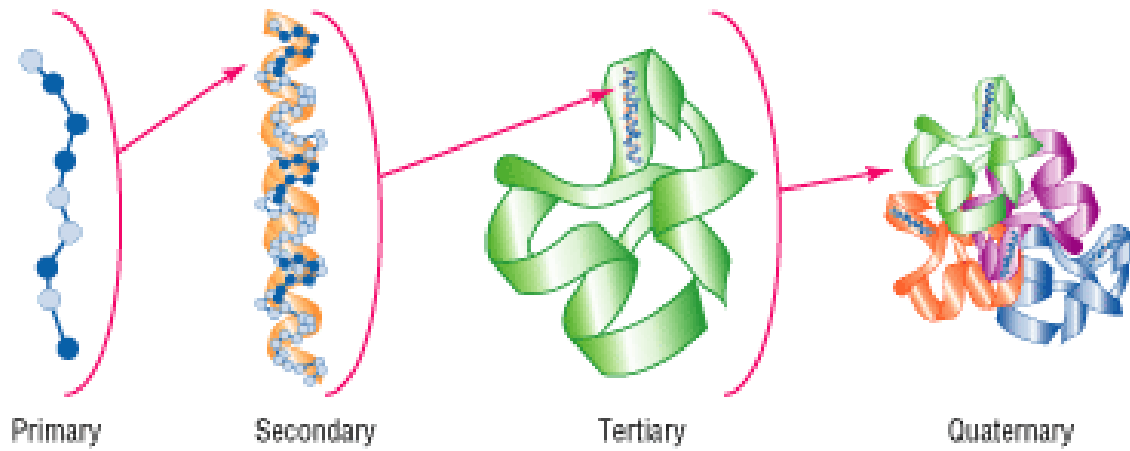
Spatial arrangement of subunits

heme group은 검은색

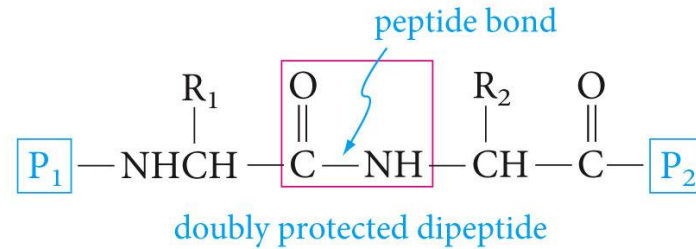
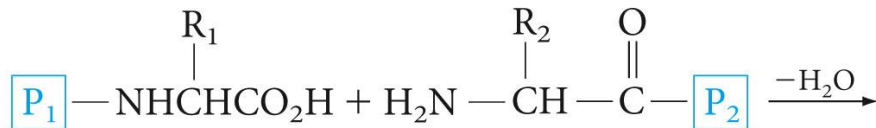
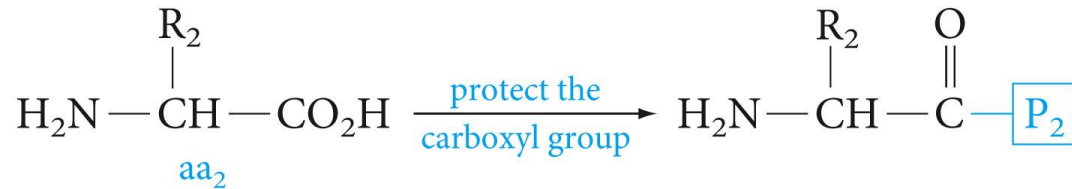
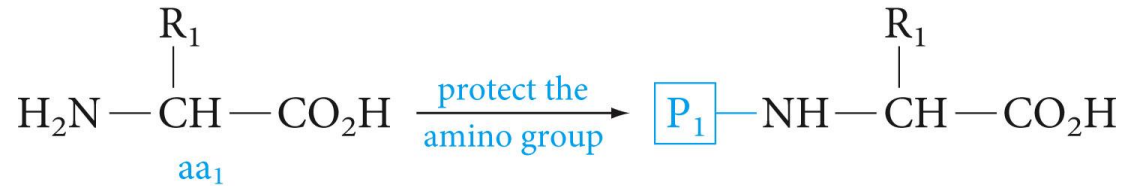


$\alpha_2\beta_2$  tetramer

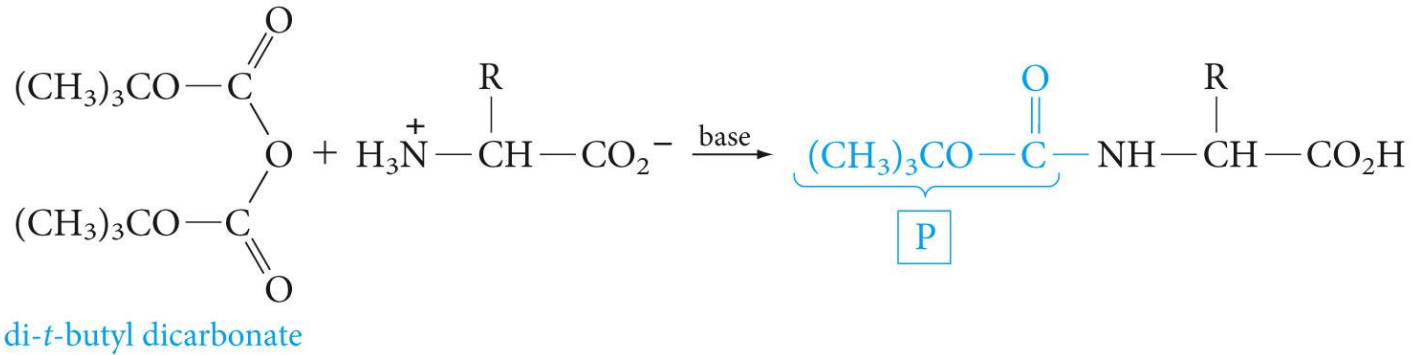
**Hemoglobin (Hb)**



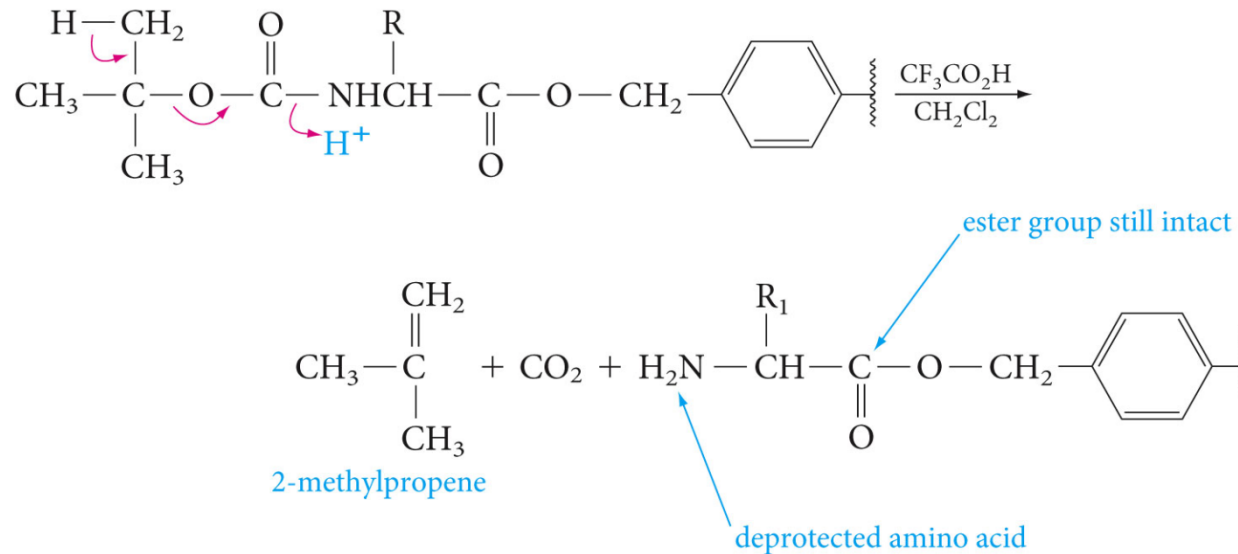
## 12. Peptide Synthesis







펩타이드 합성에 있어서 가장 많이 쓰이는 N-protecting group은 *t*-butoxycarbonyl (Boc) group이다.



Boc 보호기 제거는 산 조건 하에서 가능하다.

두가지 by products는 모두 가스이기 때문에 따로 정제과정이 필요없다.

펩타이드 연결반응(peptide coupling)시 가장 많이 쓰이는 coupling agent로서 DCC (dicyclohexylcarbodiimide)가 있다.

