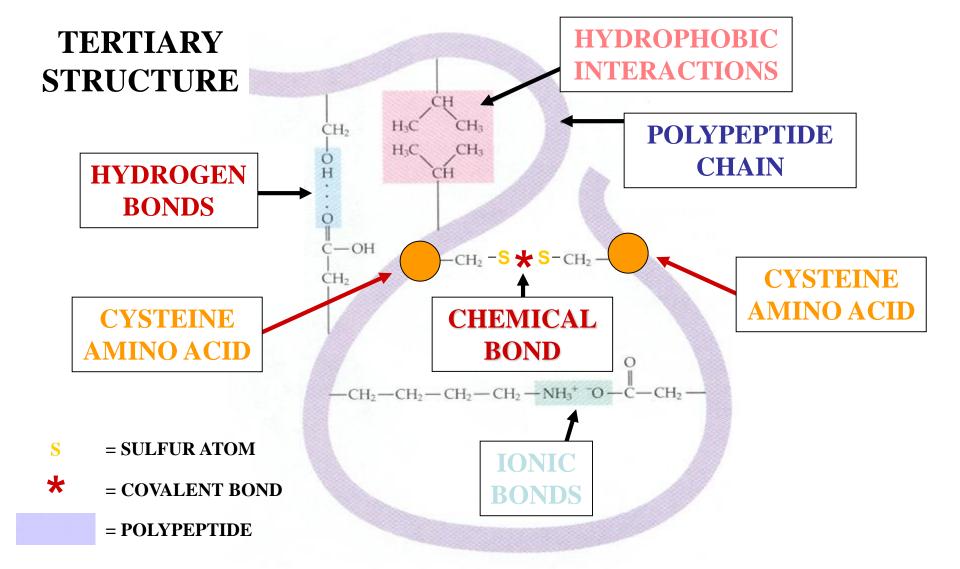
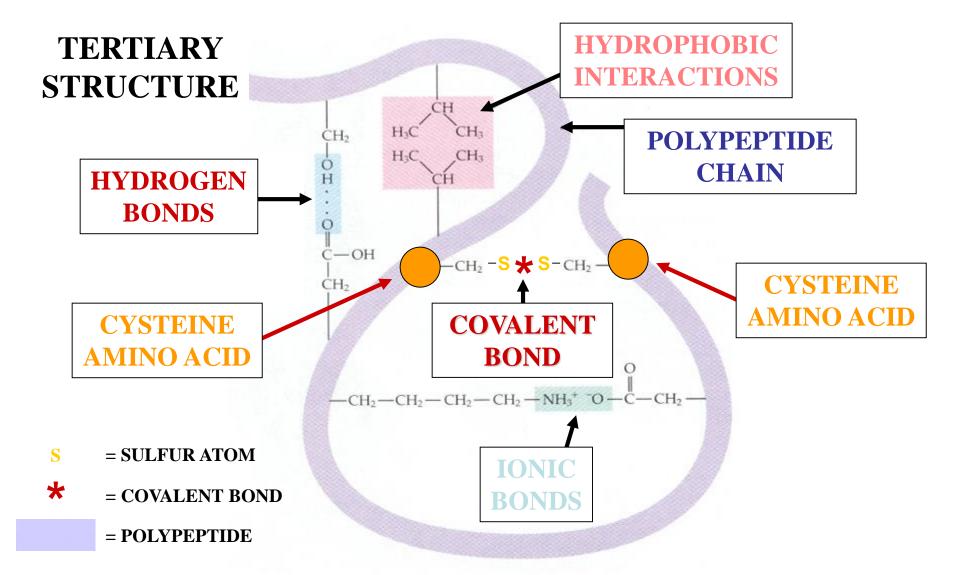
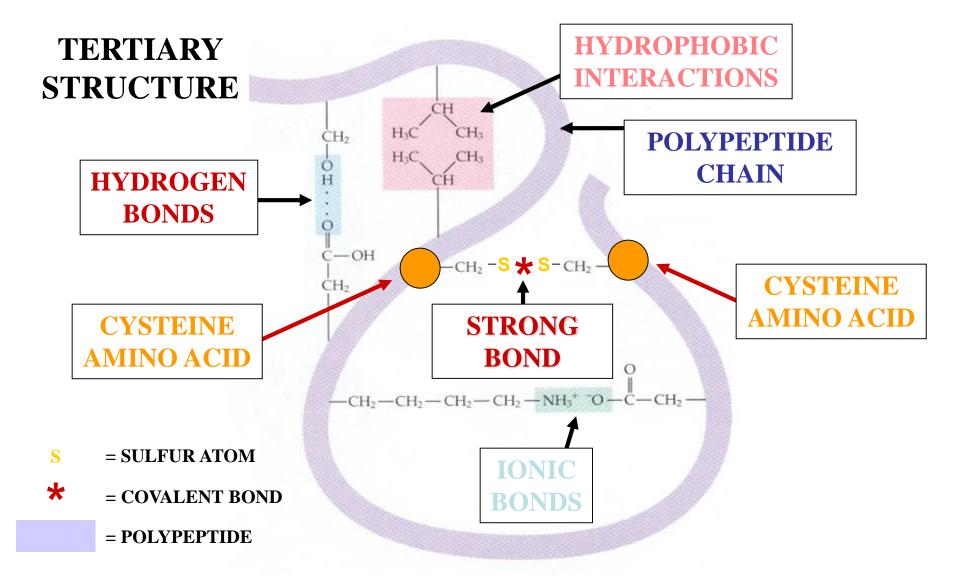


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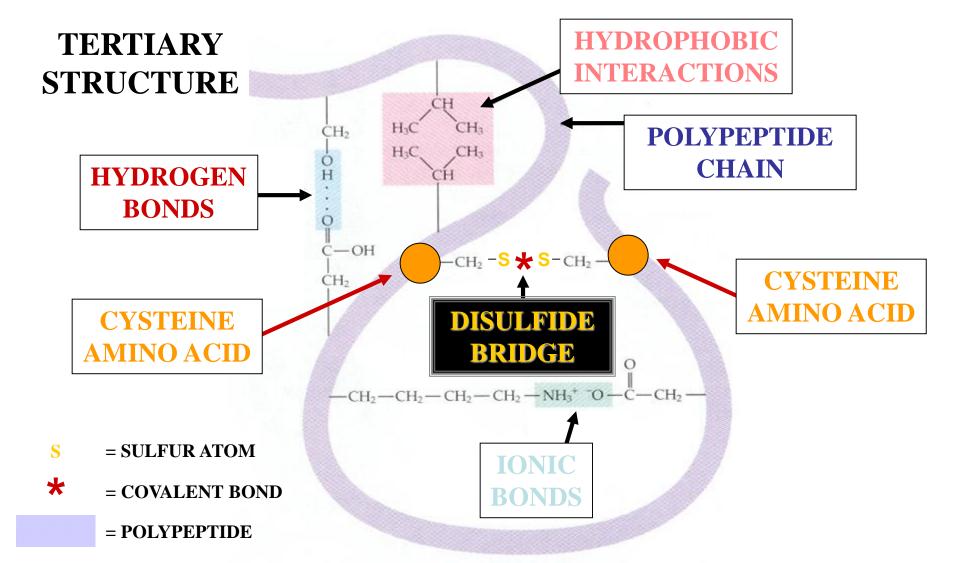






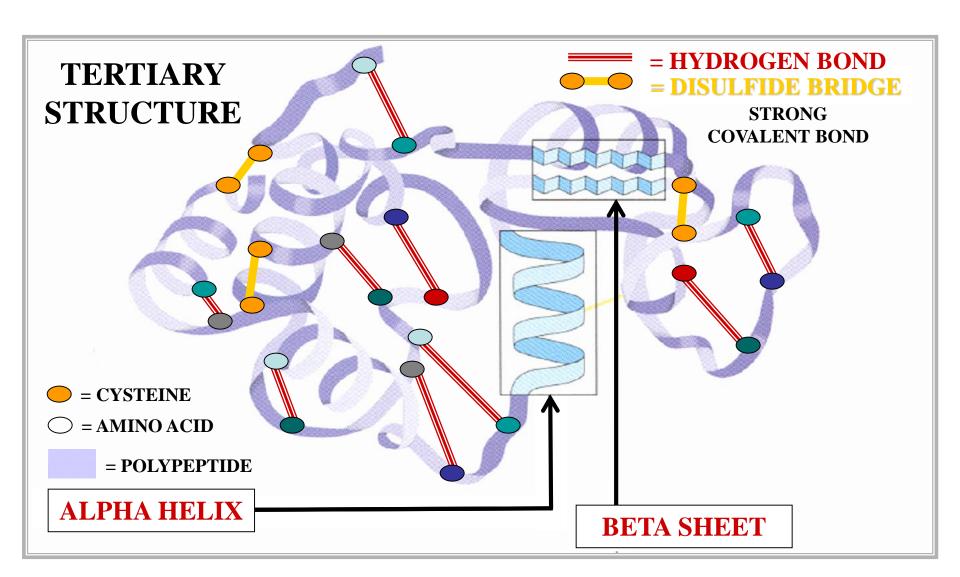






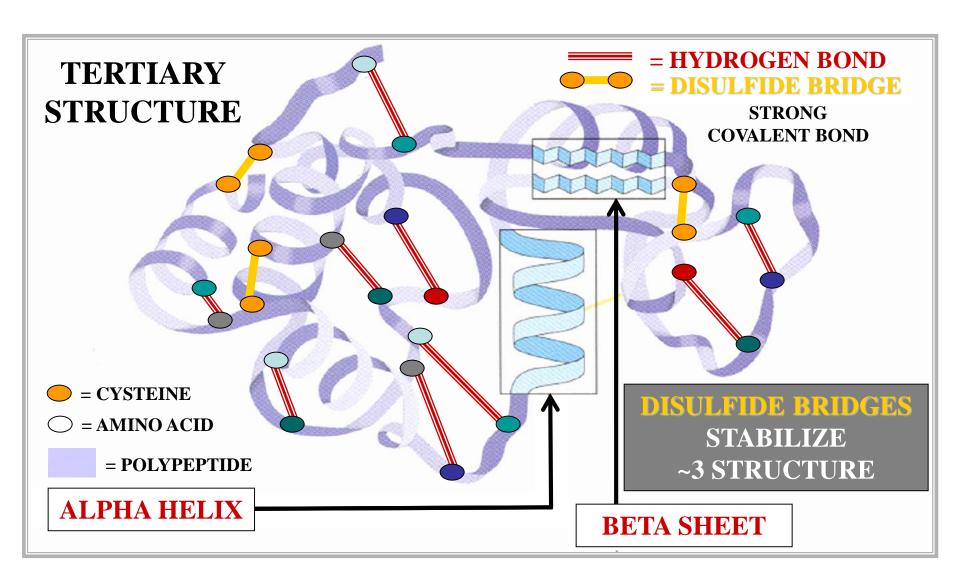
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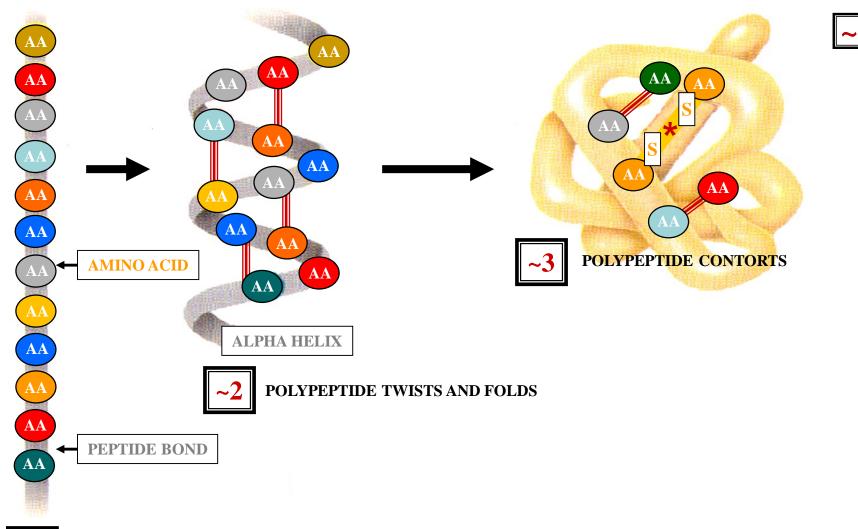
PROTEIN STRUCTURE



PROTEIN STRUCTURE





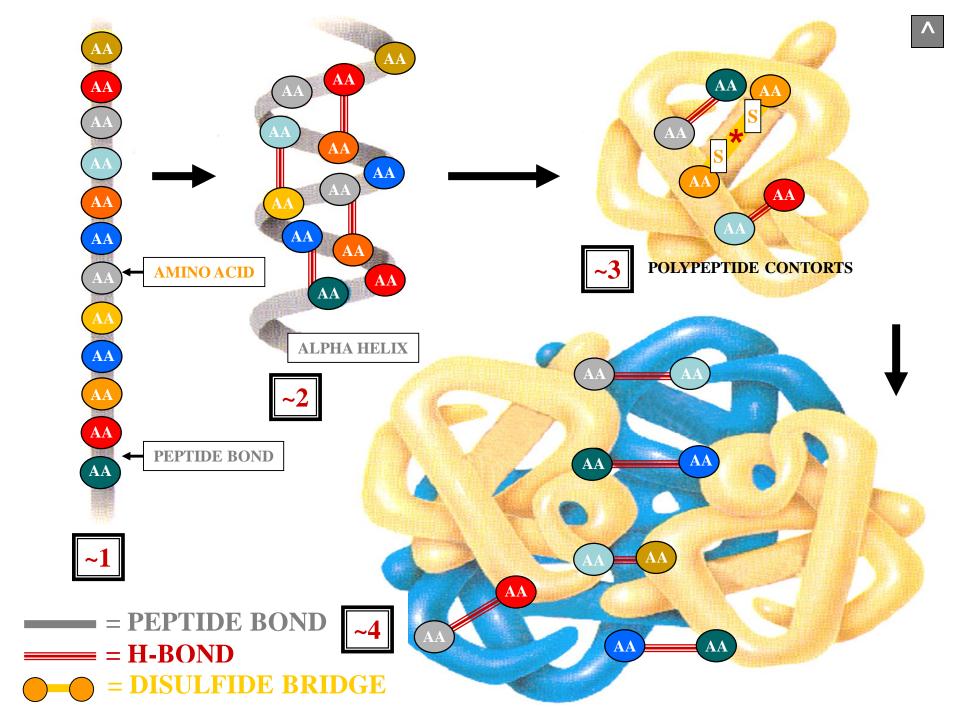


= PEPTIDE BOND

= = H-BOND

= DISULFIDE BRIDGE

|S| = SULFUR ATOM * = COVALENT BOND



PROTEIN QUATERNARY STRUCTURE

PROTEIN QUATERNARY STRUCTURE



PROTEIN ~4 STRUCTURE

SPACIAL ARRANGEMENT 2 OR MORE ~3 POLYPEPTIDES

PROTEIN ~4 STRUCTURE

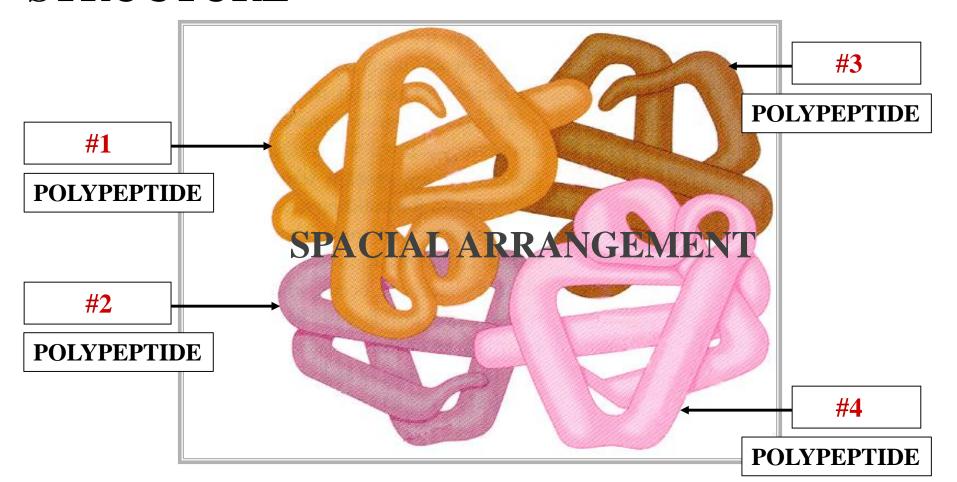


PROTEIN STRUCTURE

QUATERNARY STRUCTURE

QUATERNARY 1-2-3-4 = **POLYPEPTIDE**

IN~3 STRUCTURE

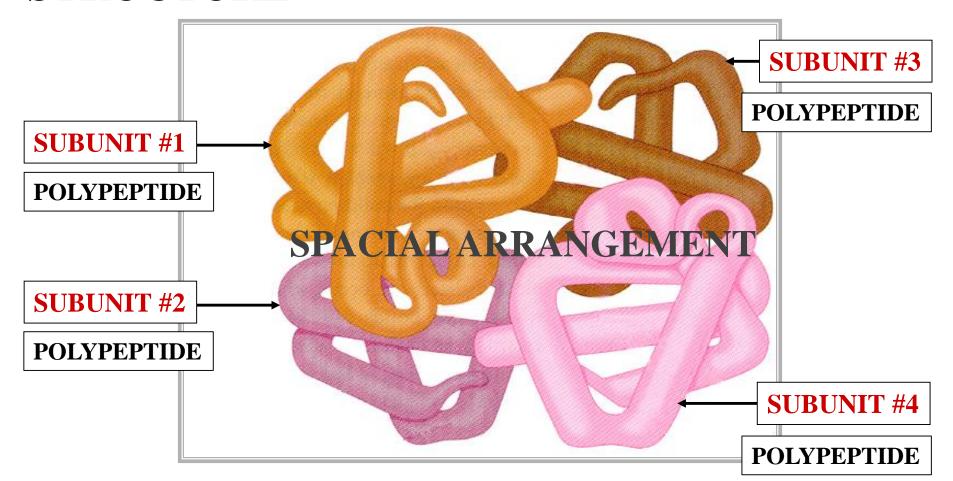


PROTEIN STRUCTURE

STRUCTURE

QUATERNARY SUBUNIT = POLYPEPTIDE

IN~3 STRUCTURE



SUBUNIT



~4 STRUCTURE SUBUNIT

DIFFERENT POLYPEPTIDE

ASSOCIATED WITH ~4 STRUCTURE

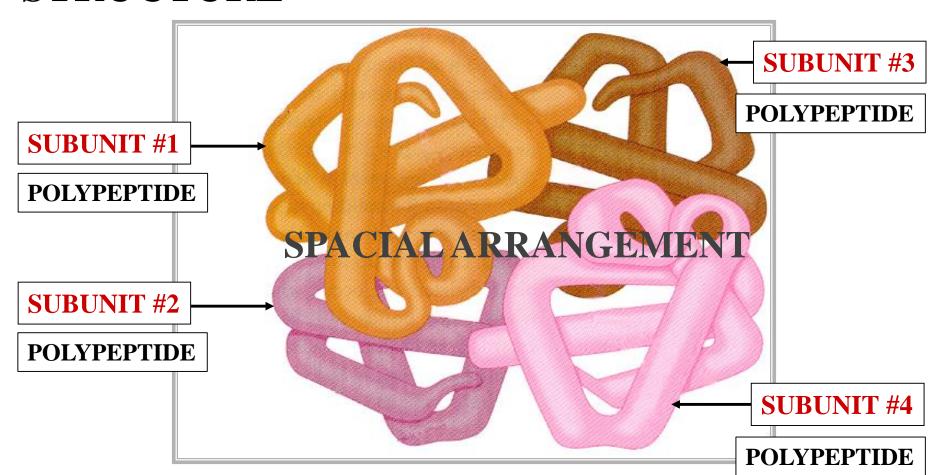
~4 STRUCTURE SUBUNIT

PROTEIN STRUCTURE

QUATERNARY STRUCTURE

QUATERNARY SUBUNIT = POLYPEPTIDE

IN ~3 STRUCTURE



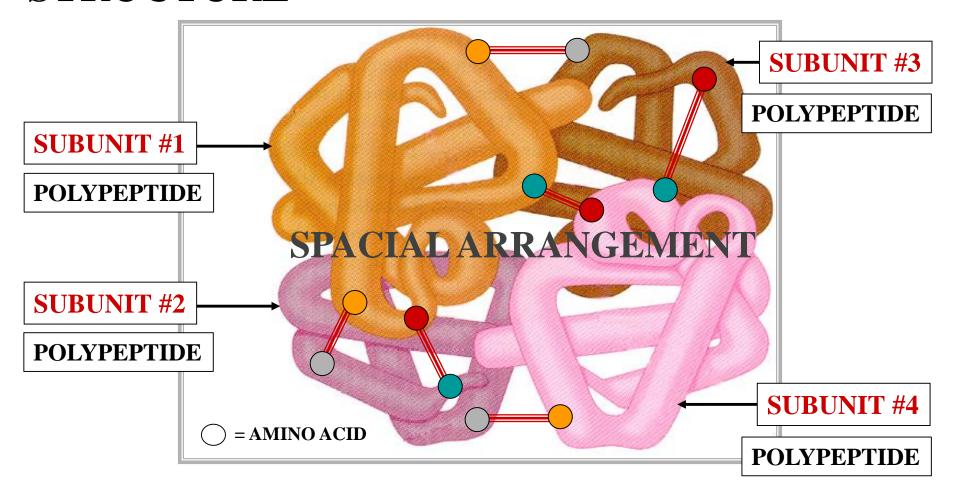
PROTEIN STRUCTURE

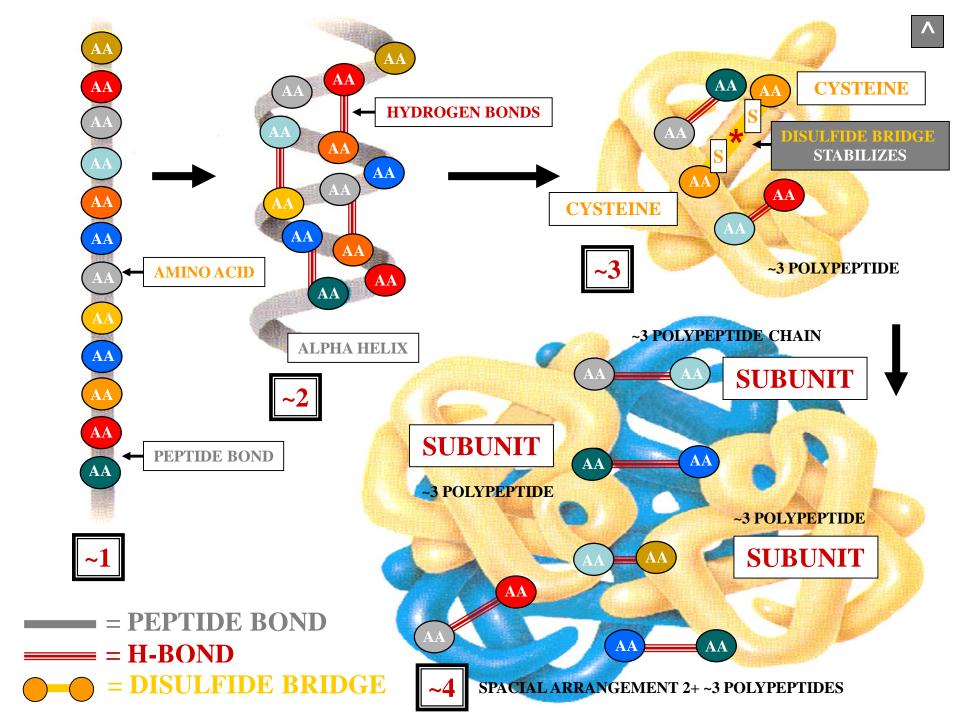


QUATERNARY STRUCTURE

SUBUNIT = POLYPEPTIDE

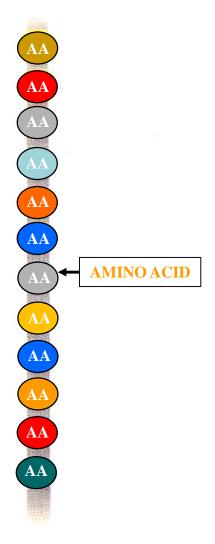
= HYDROGEN BOND



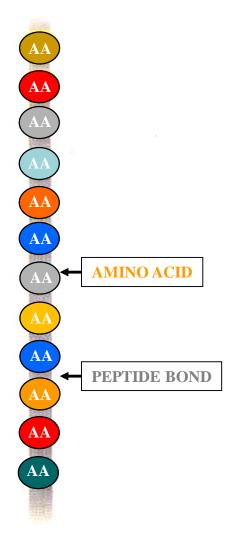




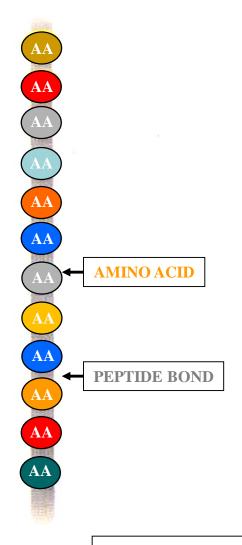
PROTEIN STRUCTURE SUMMARY



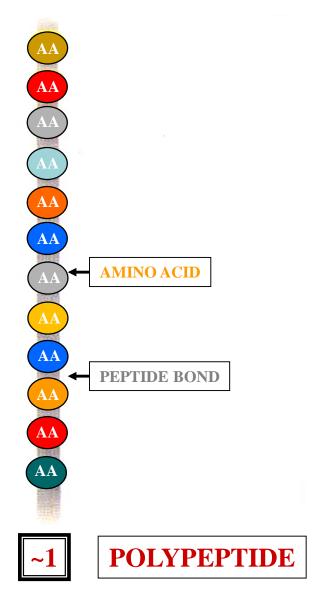
LINEAR AMINO ACID SEQUENCE

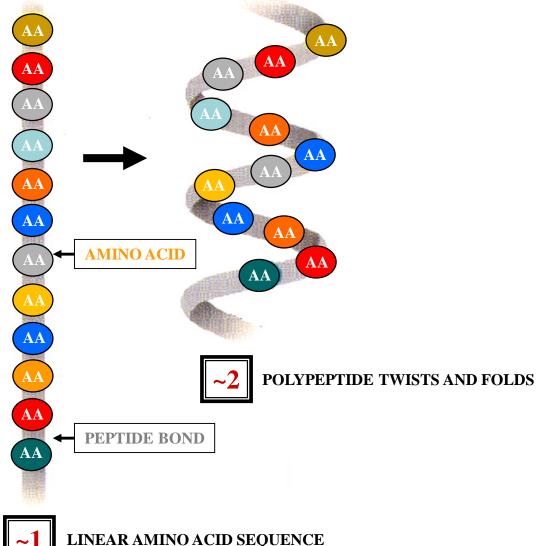


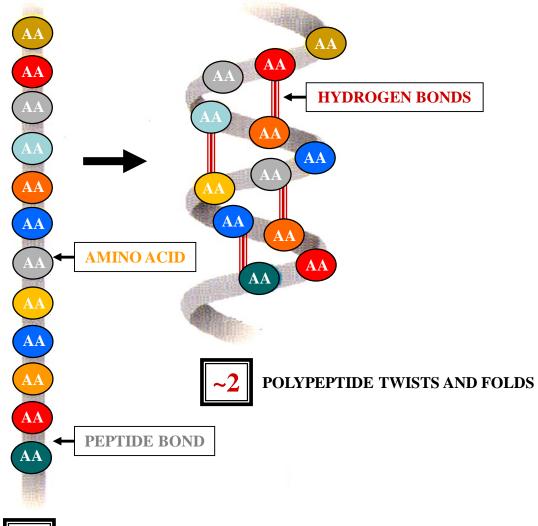
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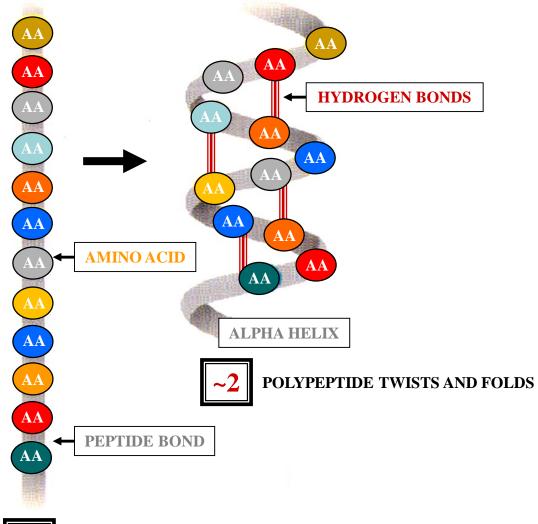
POLYPEPTIDE



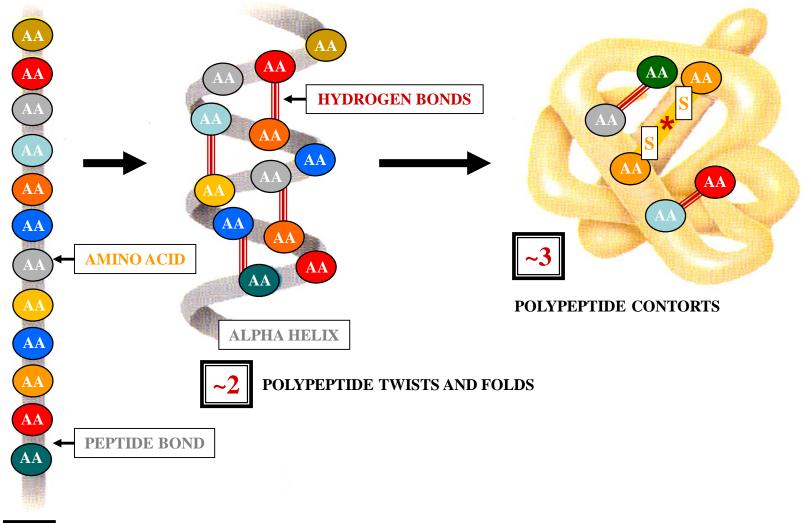




= PEPTIDE BOND = H-BOND

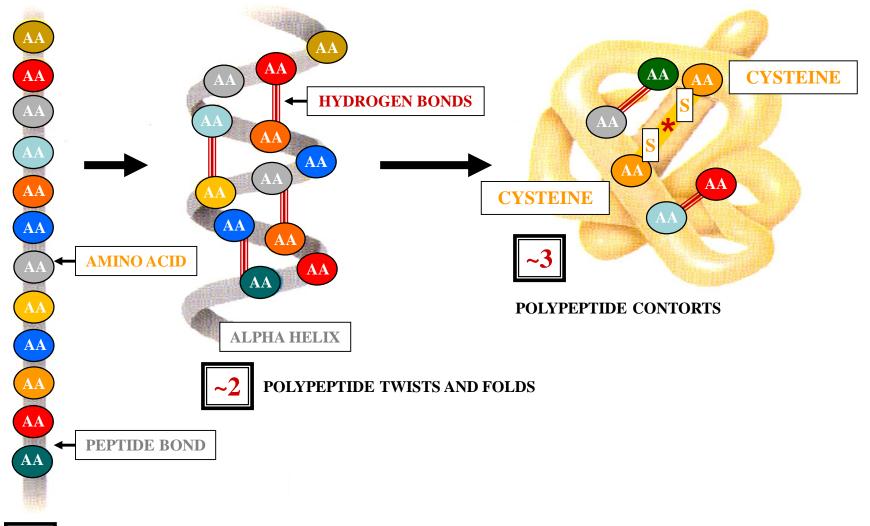


= PEPTIDE BOND = H-BOND

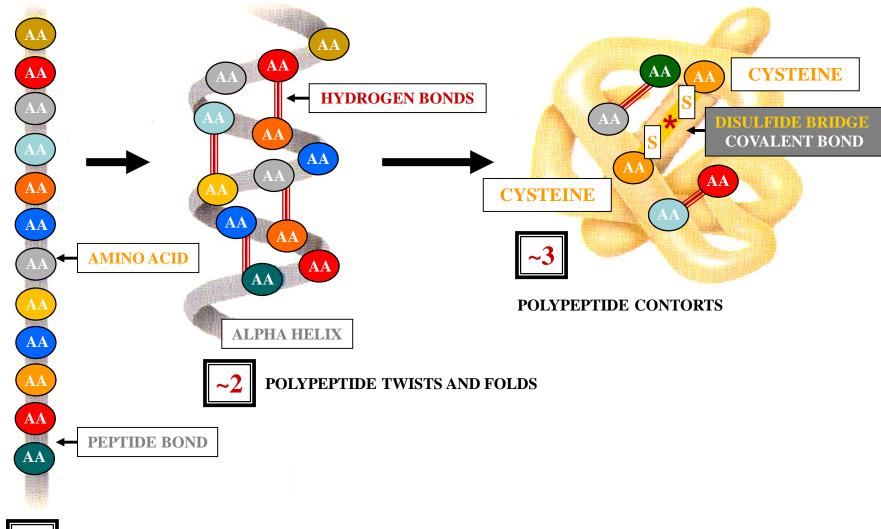


= PEPTIDE BOND

=== H-BOND



= PEPTIDE BOND

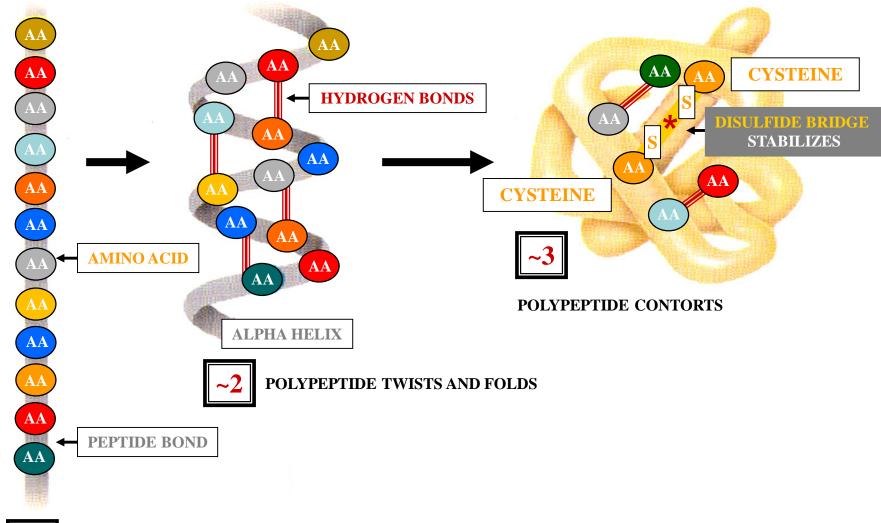


= PEPTIDE BOND

= = H-BOND

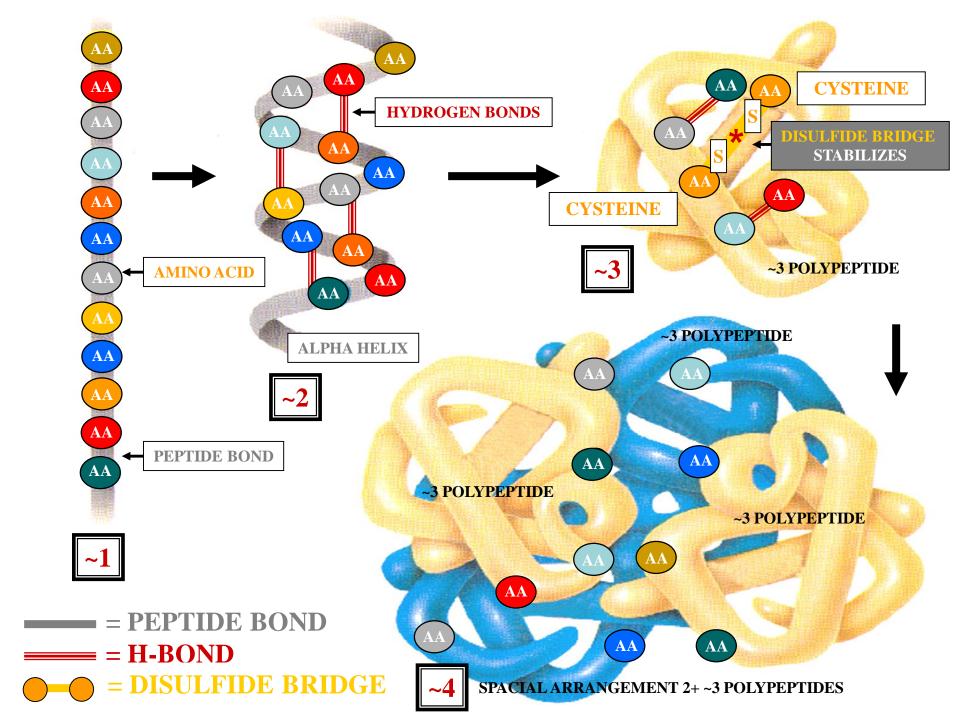
= DISULFIDE BRIDGE

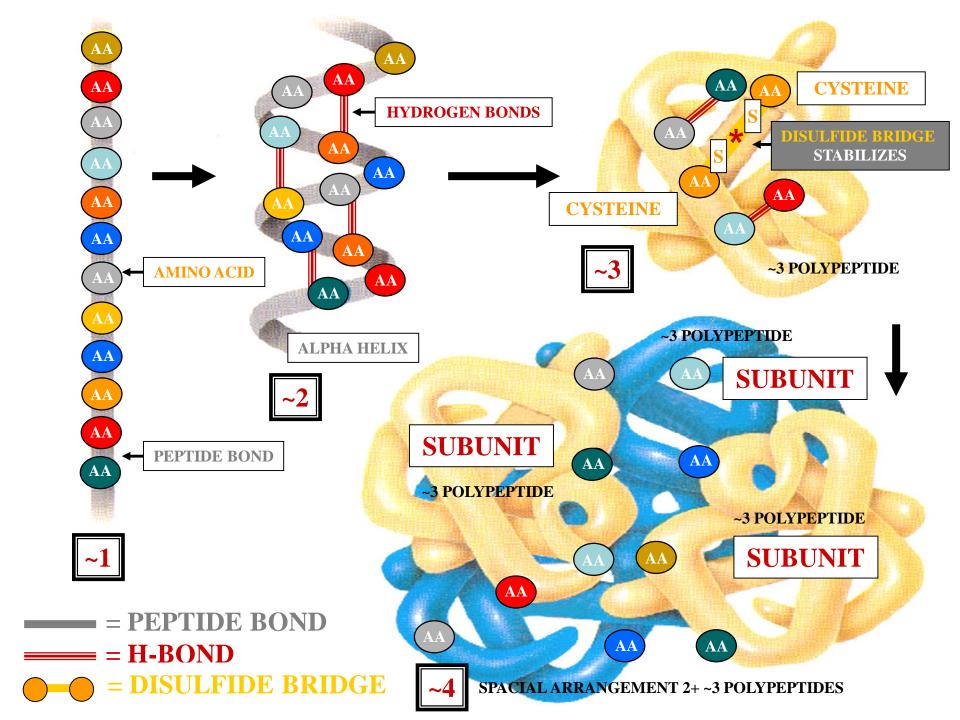
 $|S| = SULFUR ATOM <math>\star = COVALENT BOND$

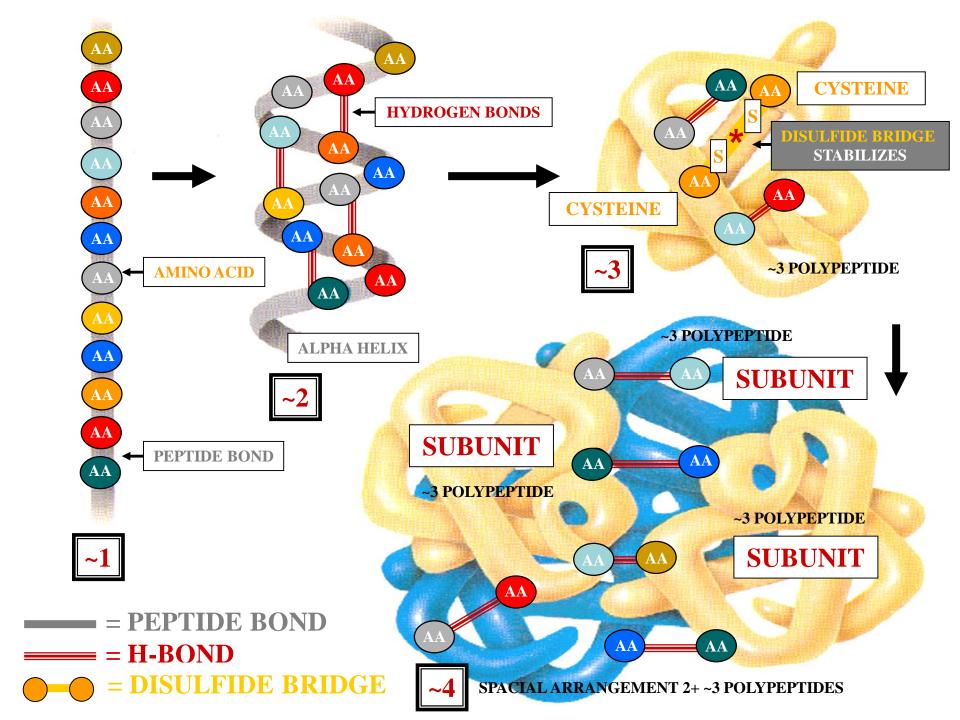


= PEPTIDE BOND

== H-BOND





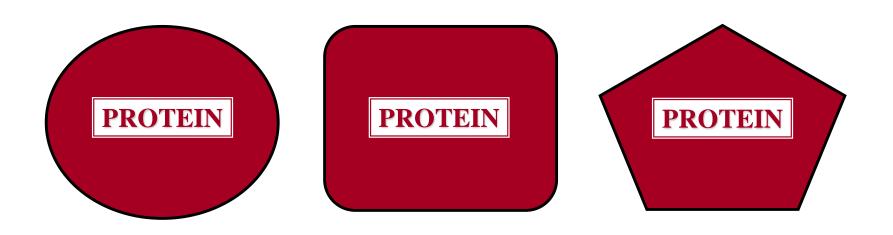




OVERALL PROTEIN STRUCTURE

PROTEIN CONFORMATION

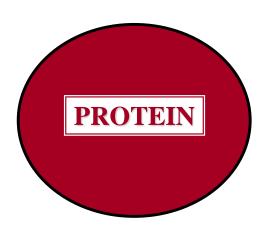


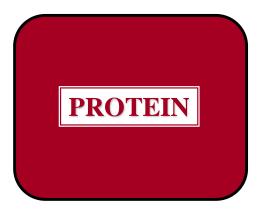


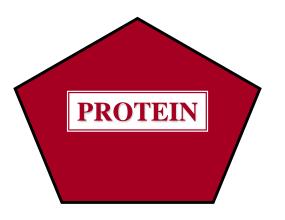
DIFFERENT PROTEINS







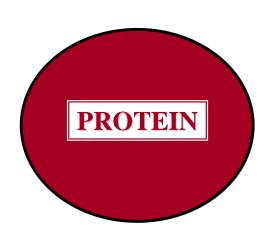


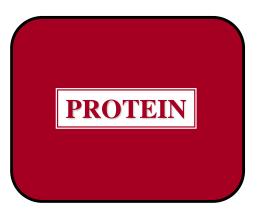


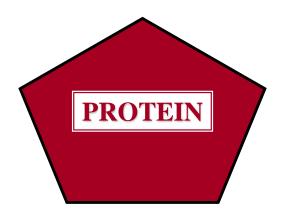
DIFFERENT PROTEINS

DIFFERENT STRUCTURE







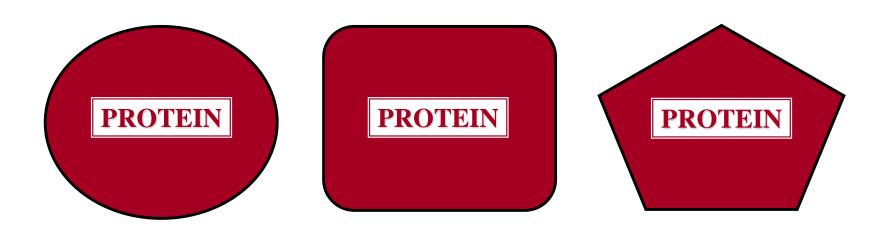


DIFFERENT PROTEINS

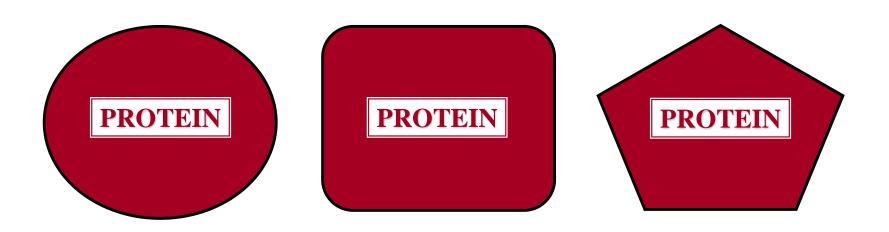
DIFFERENT STRUCTURE

DIFFERENT CONFORMATION





DIFFERENT FUNCTION



PROTEIN CONFORMATION
DICTATES
PROTEIN FUNCTION

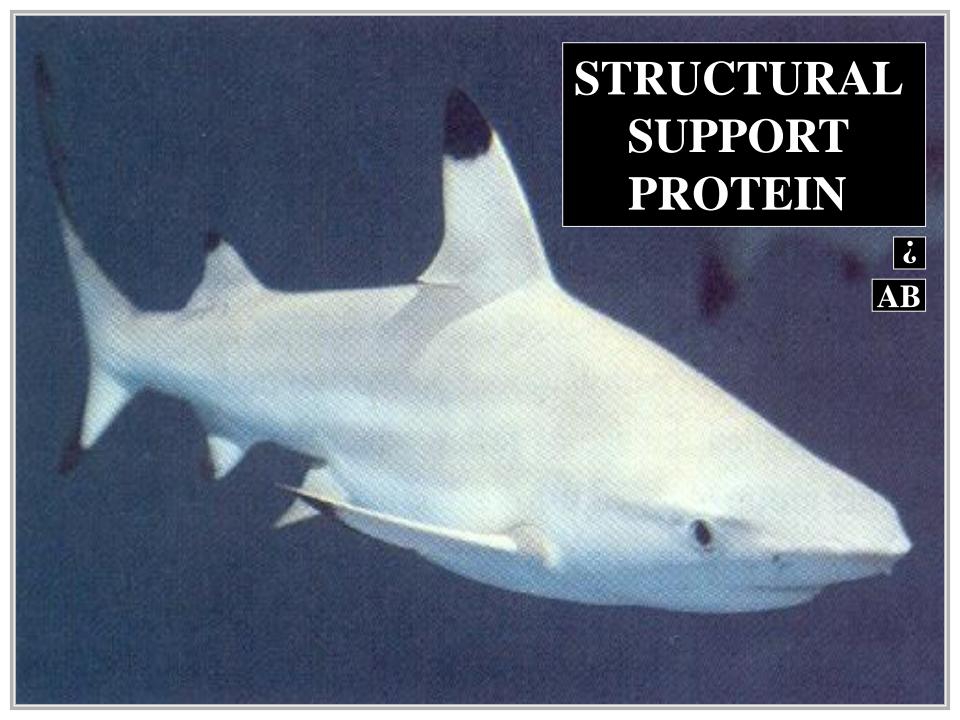
PROTEIN CONFORIVIATION DICIES IPIROTEDIN E.OV. C.L.OV

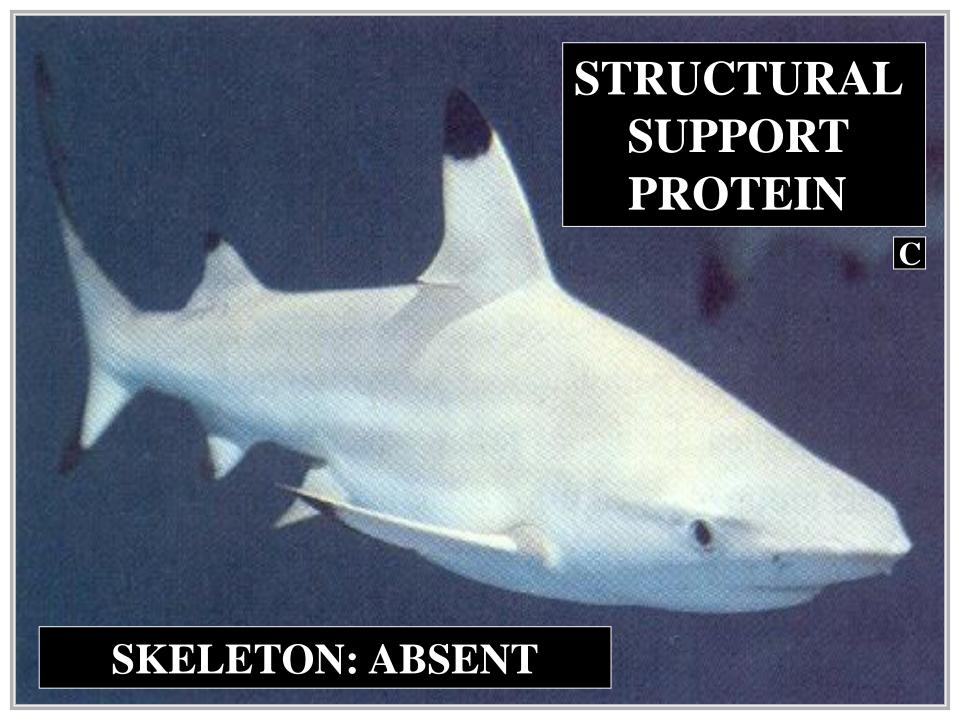
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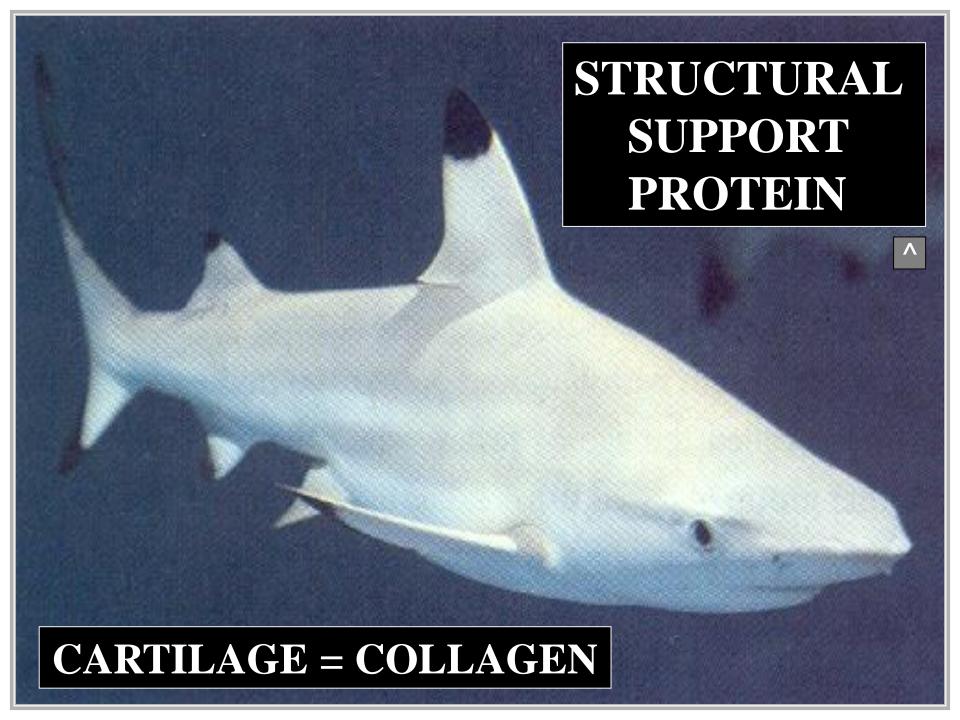
PROTEIN FUNCTION



PROTEIN STRUCTURAL FUNCTION

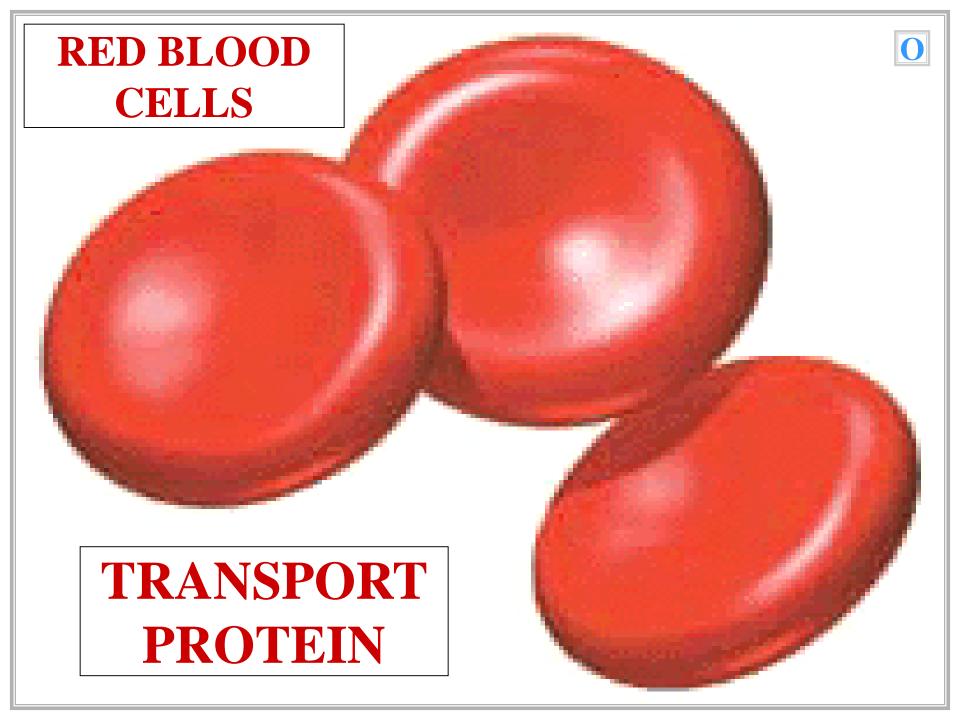


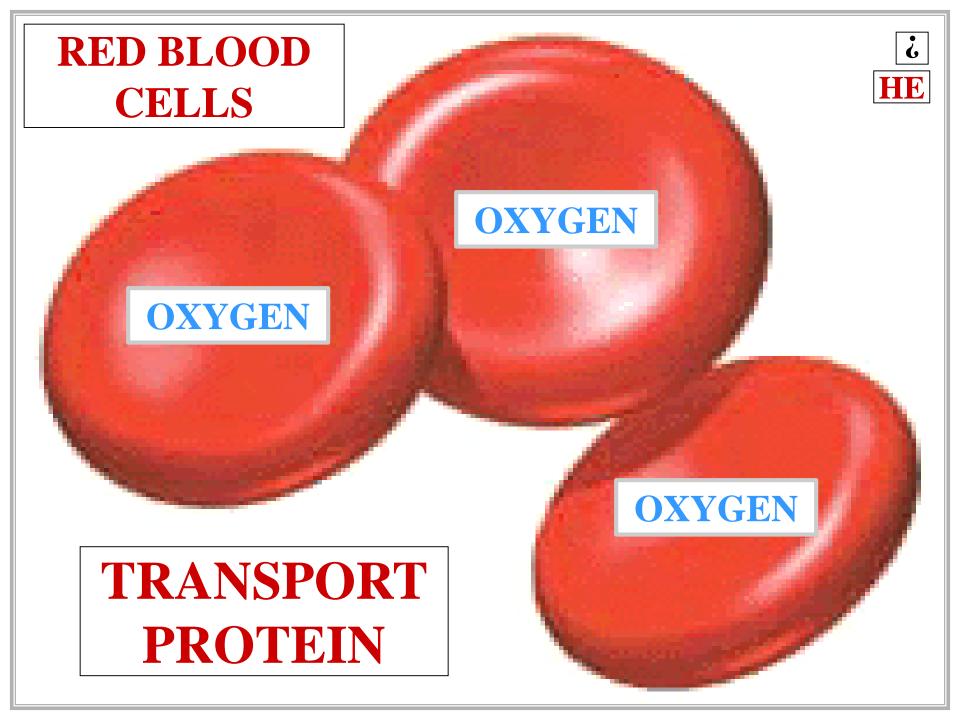


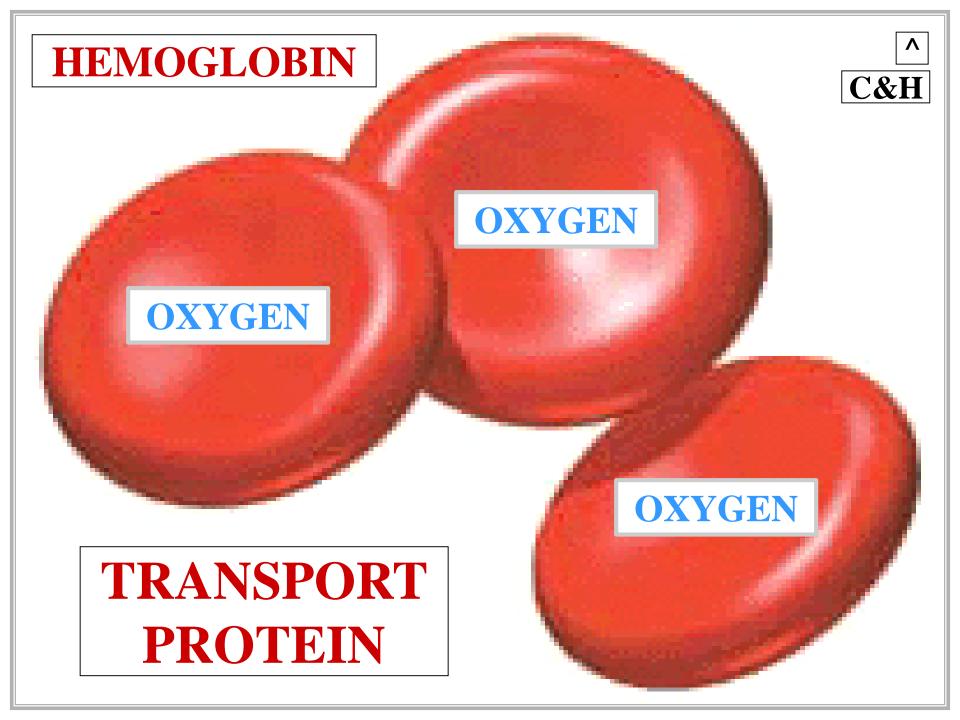




PROTEIN TRANSPORT FUNCTION



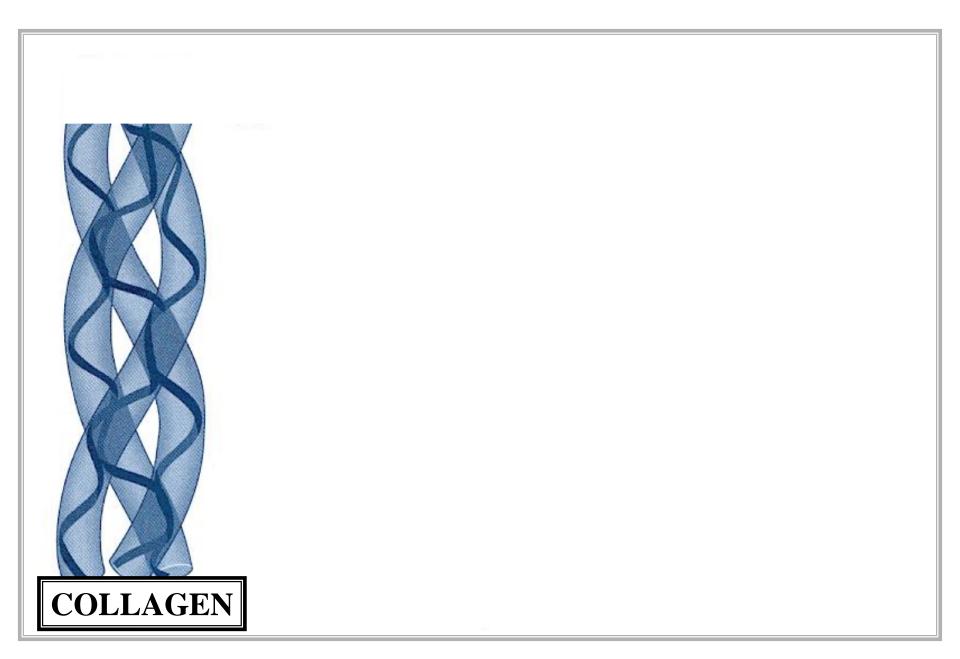




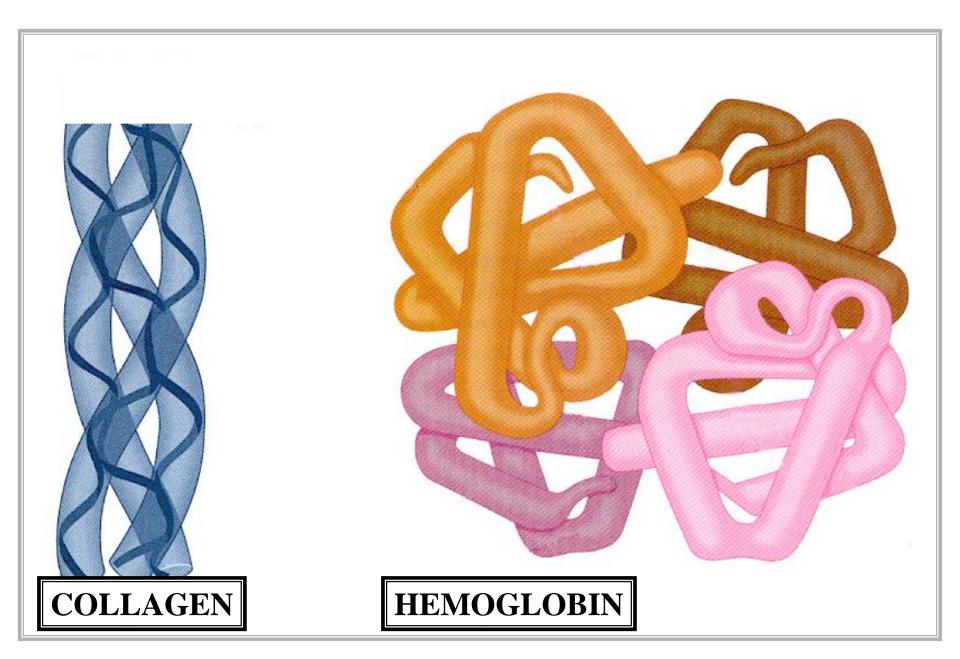


COLLAGEN VS HEMOGLOBIN CONTRAST

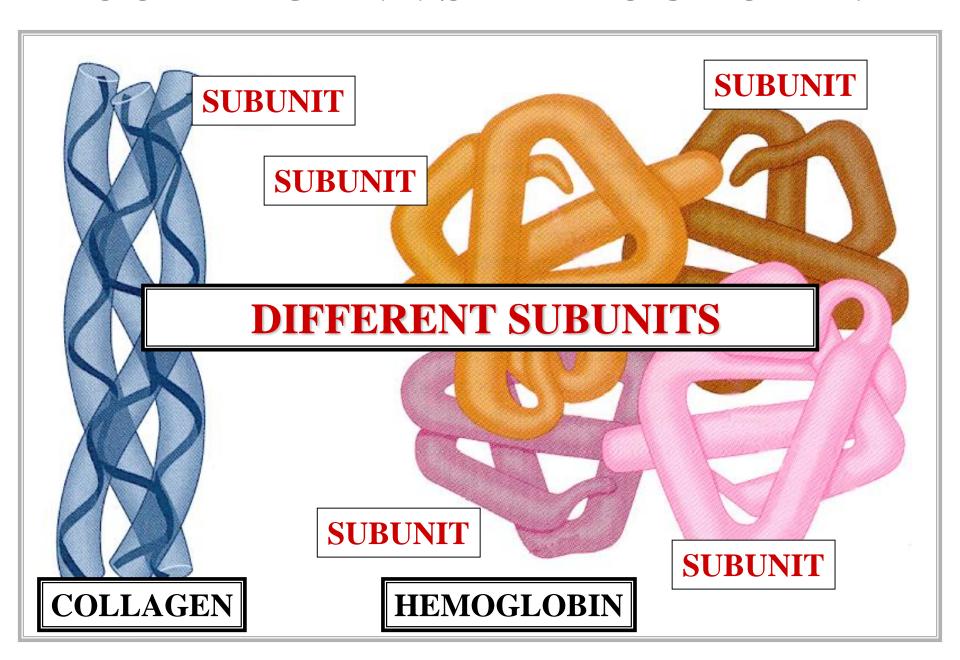




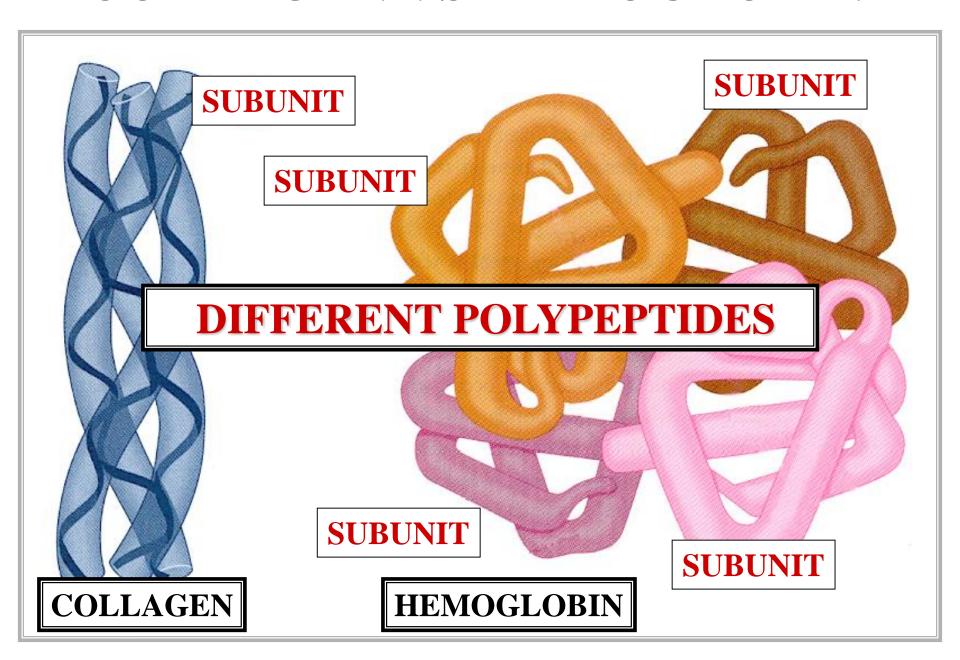




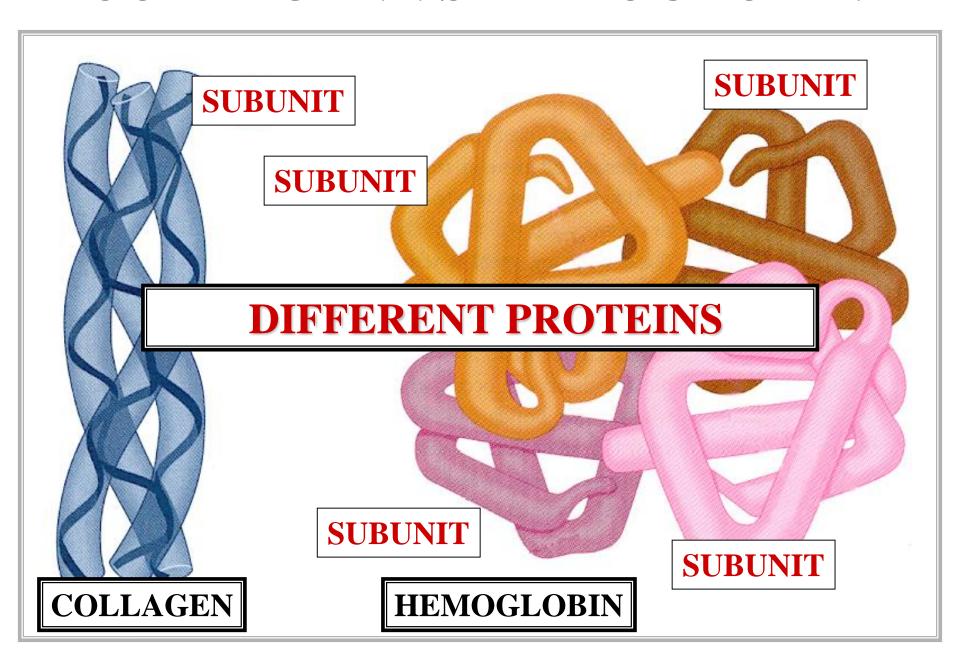




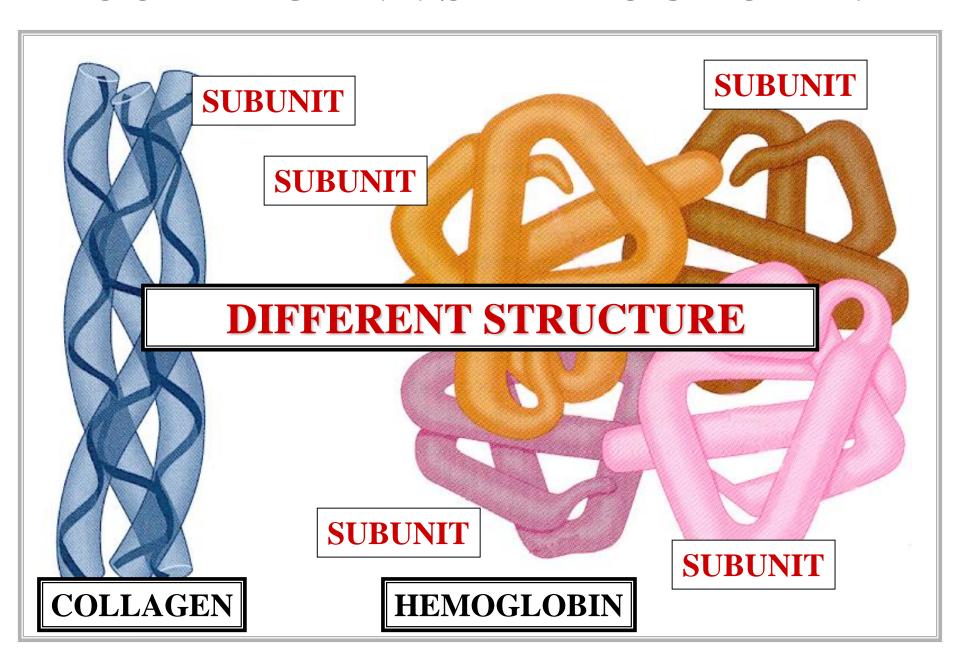




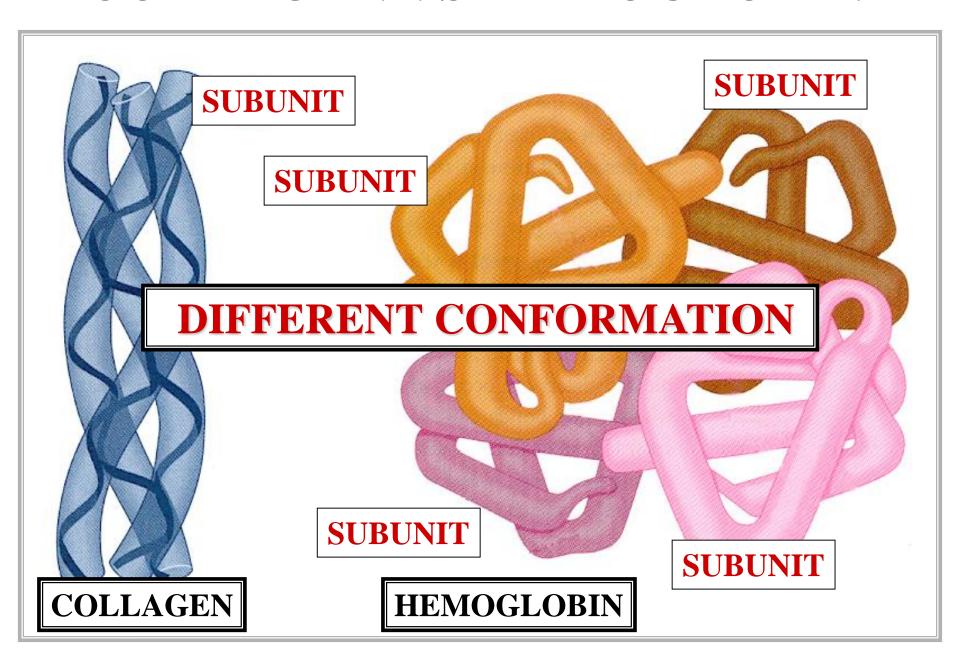




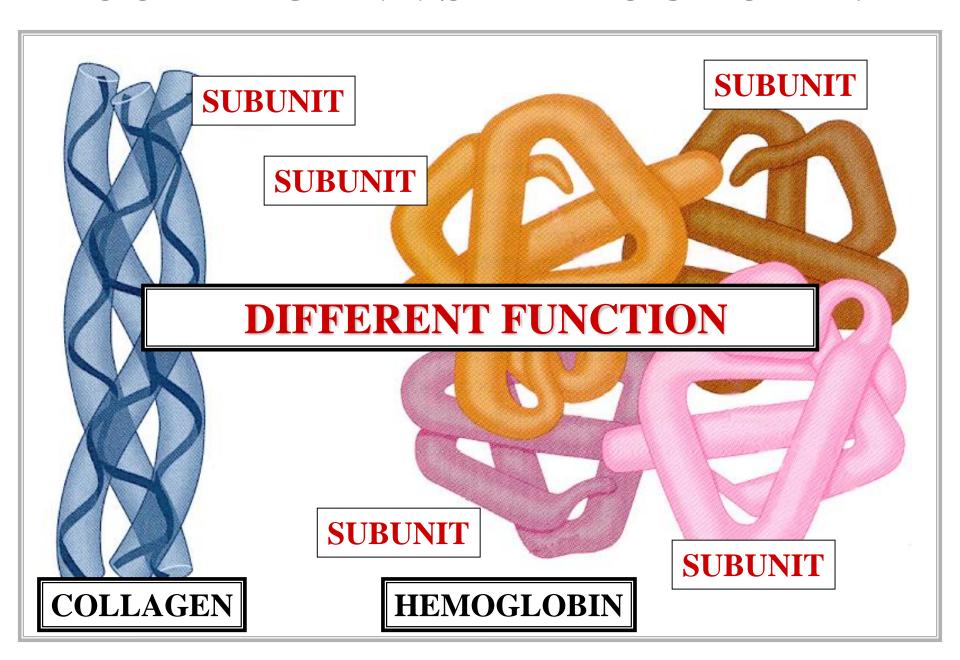




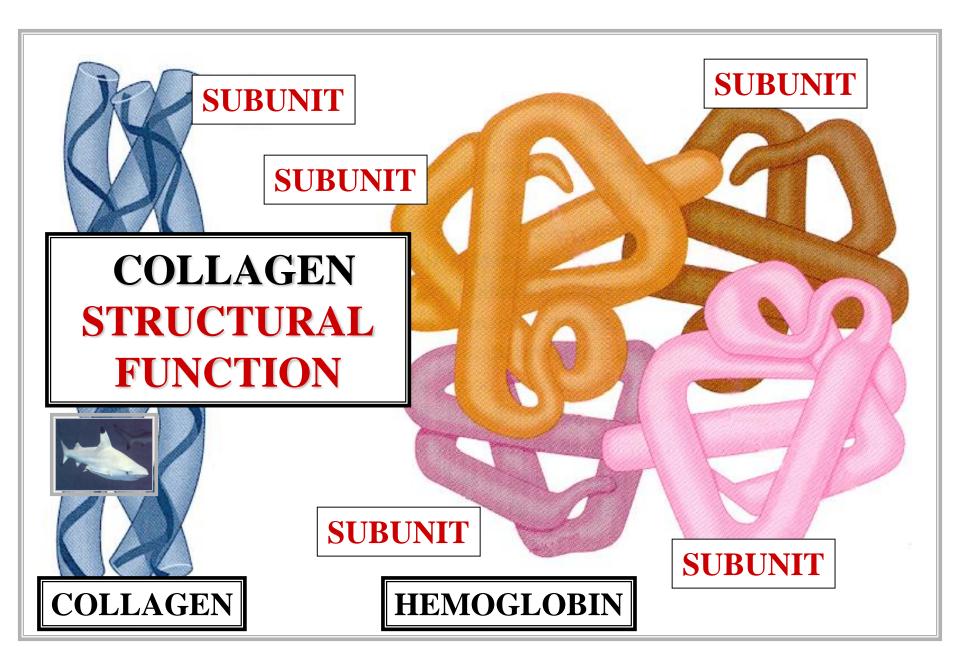


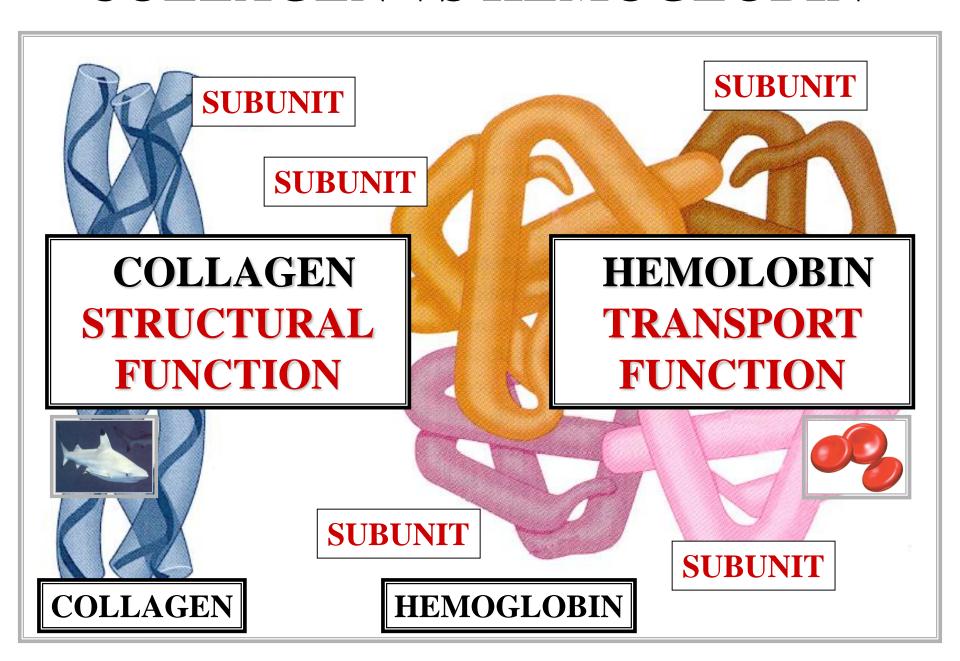












PROTEIN ENZYMATIC FUNCTION

ENZYME

ENZYME

PROTEIN BIOCHEMICAL CATALYST

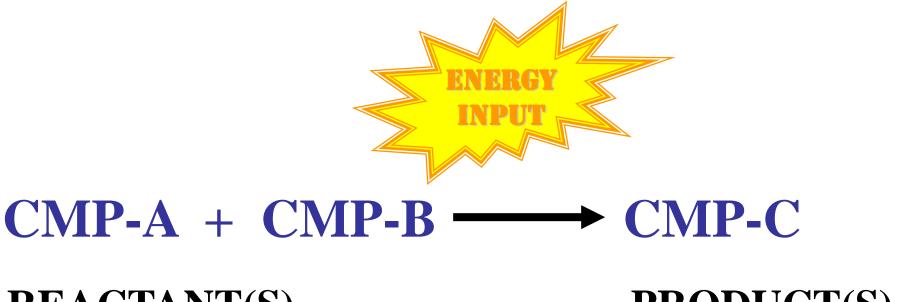
ENZYME



BIOCHEMICAL REACTION



BIOCHEMICAL REACTION



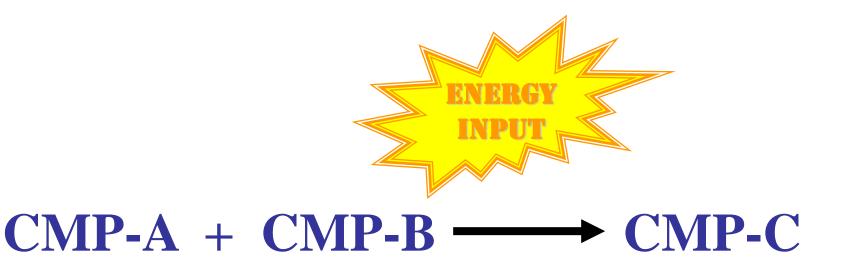
REACTANT(S)

PRODUCT(S)

?



BIOCHEMICAL REACTION

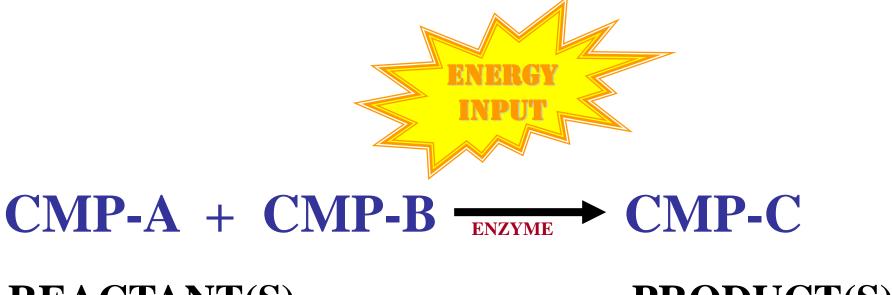


REACTANT(S)

PRODUCT(S)

ACTIVATION ENERGY



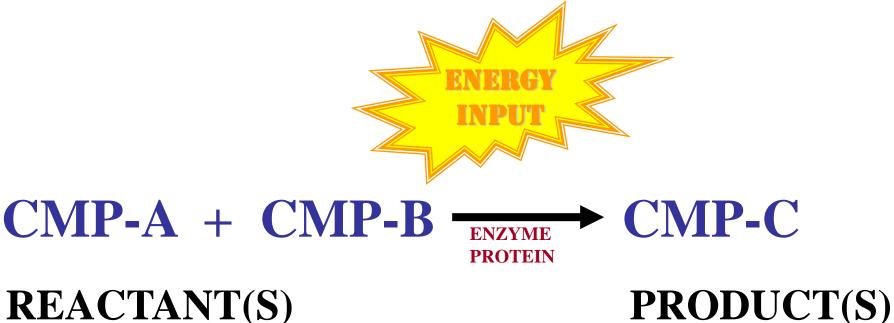


REACTANT(S)

PRODUCT(S)

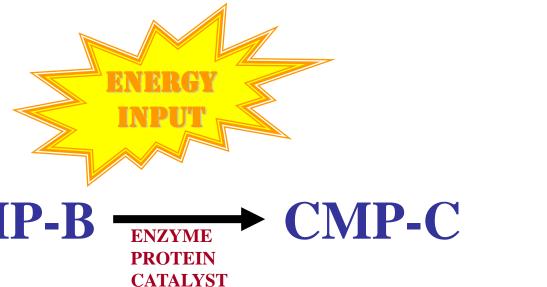
ACTIVATION ENERGY





ACTIVATION ENERGY





CMP-A + CMP-B

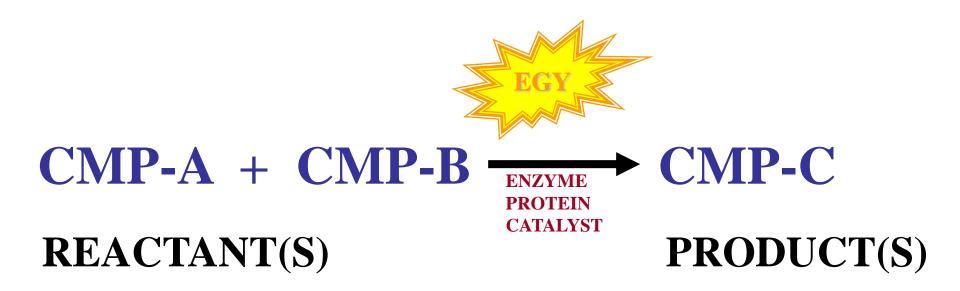
REACTANT(S)

PRODUCT(S)

ACTIVATION ENERGY



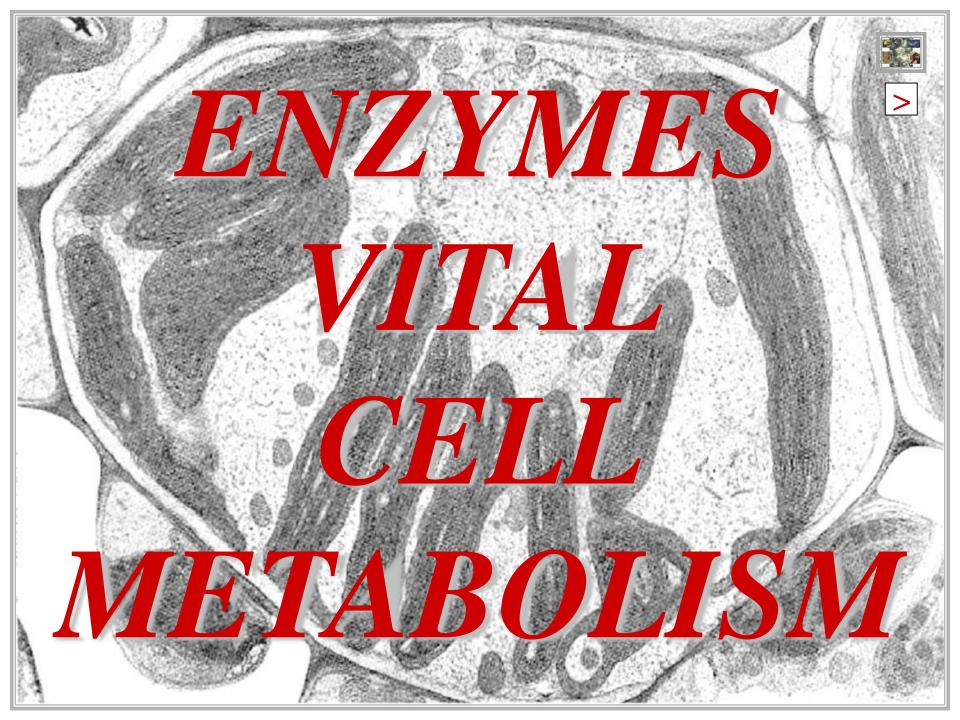


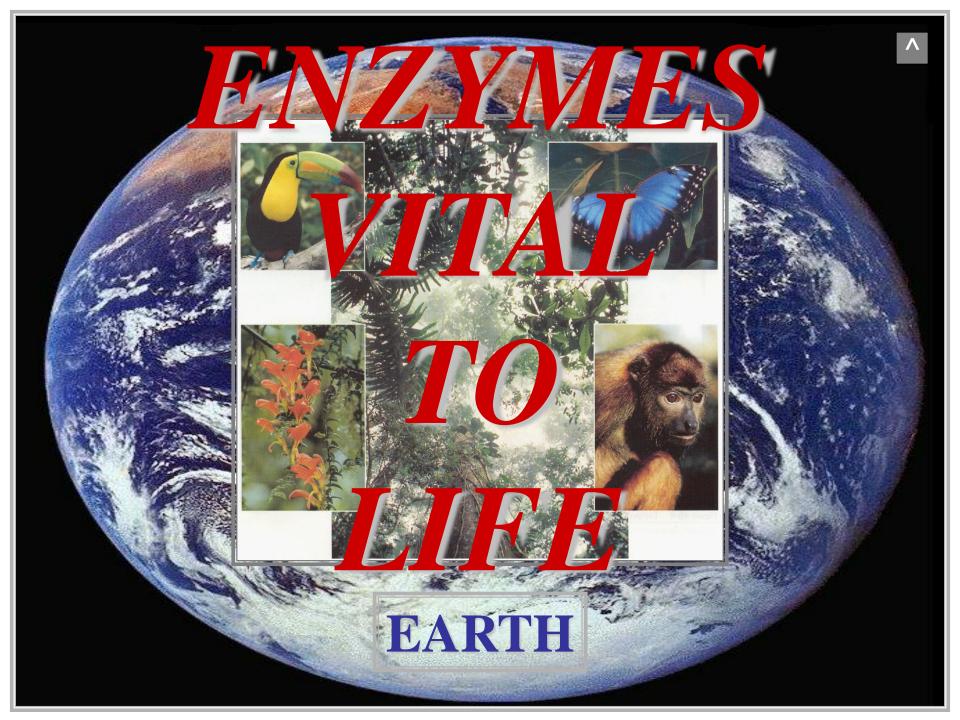


ACTIVATION ENERGY DECREASES







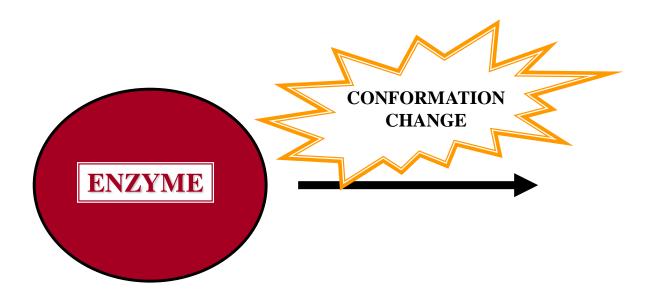


CONFORMATION CHANGE

ENZYME DENATURES



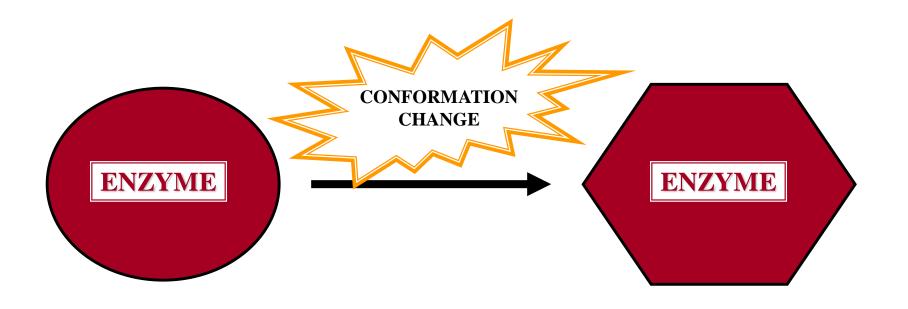




ENZYME CONFORMATION CHANGE

L

ENZYME DENATURES

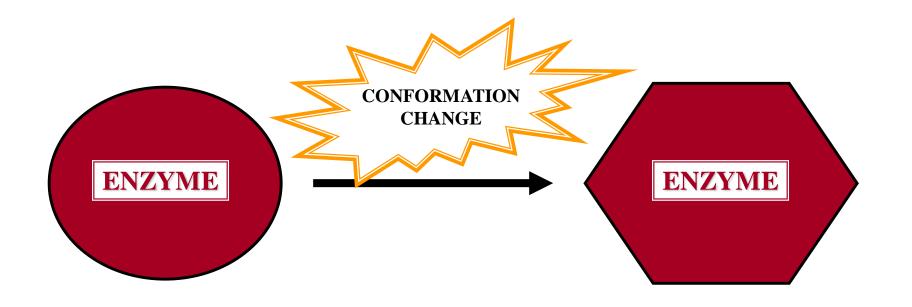


ENZYME STRUCTURE CHANGE



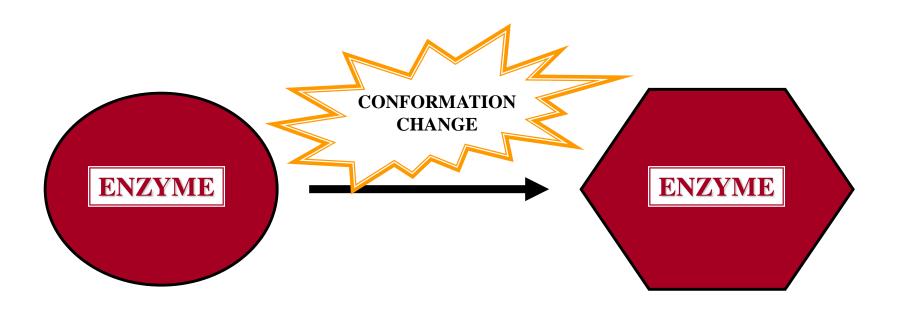




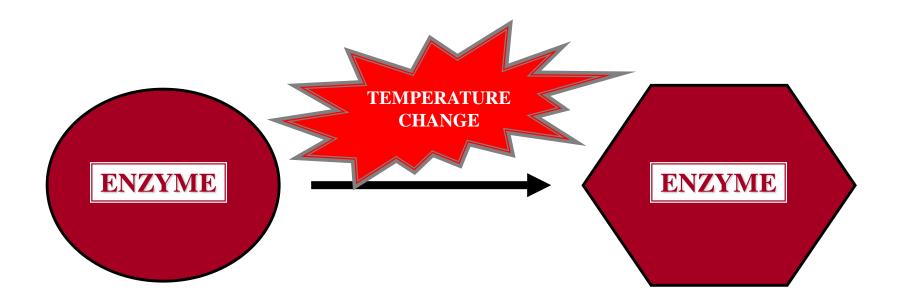


ENZYME LOST FUNCTION

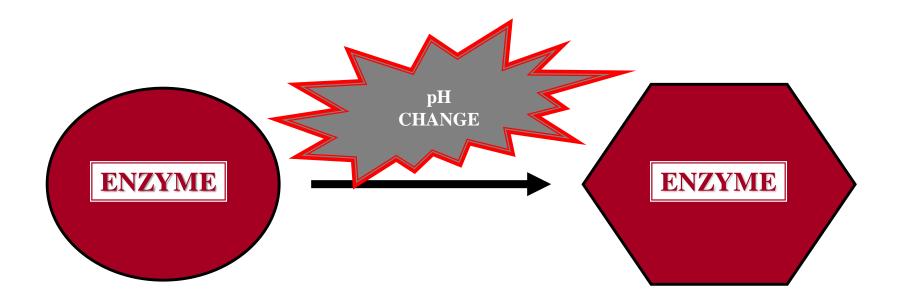




ENZYMES SENSITIVE:



ENZYMES SENSITIVE: TEMPERATURE CHANGE



ENZYMES SENSITIVE: pH CHANGE

ENZYME DENATURES TIMPERATURE CHANGE

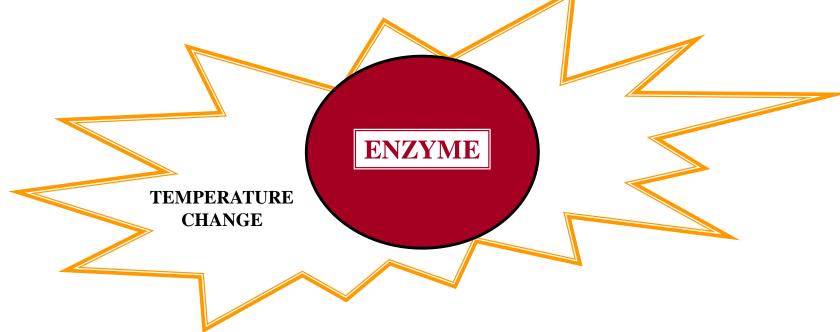




ENZYMES NARROW TEMPERATURE TOLERANCE

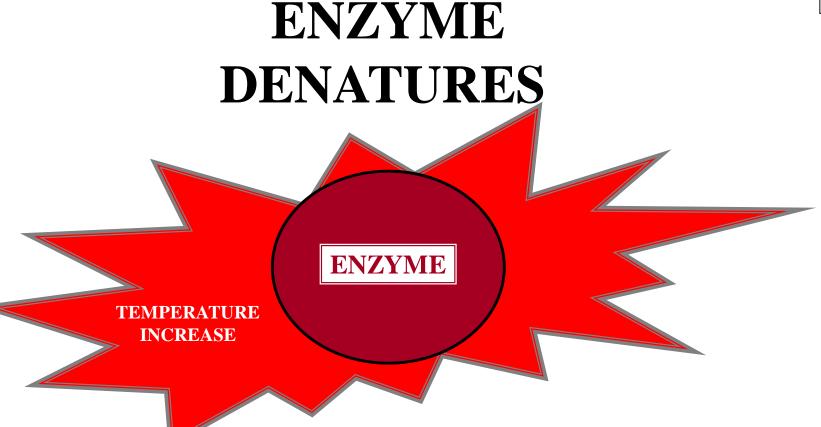






ENZYMES NARROW TEMPERATURE TOLERANCE

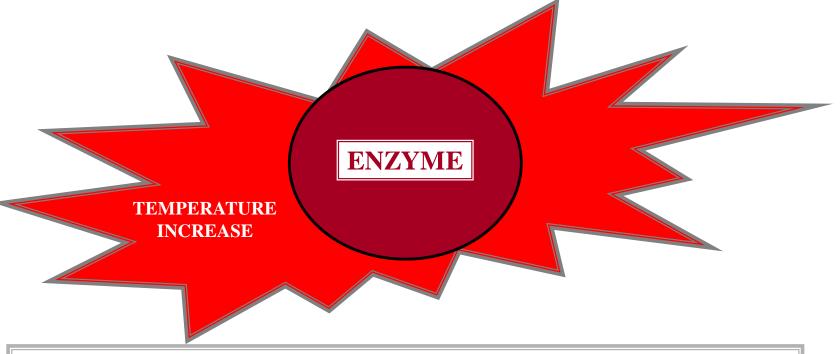




ENZYMES NARROW TEMPERATURE TOLERANCE







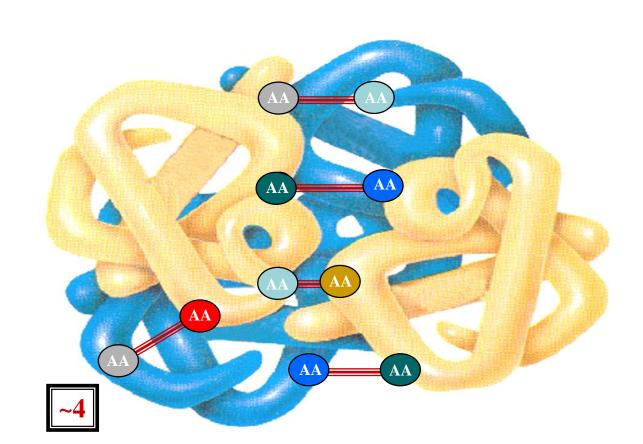
ENZYME INCREASES MOLECULAR MOVEMENT

WB

TEMPERATURE INCREASE INCREASES ENZYME MOLECULAR MOVEMENT

H-BONDS

ENZYME

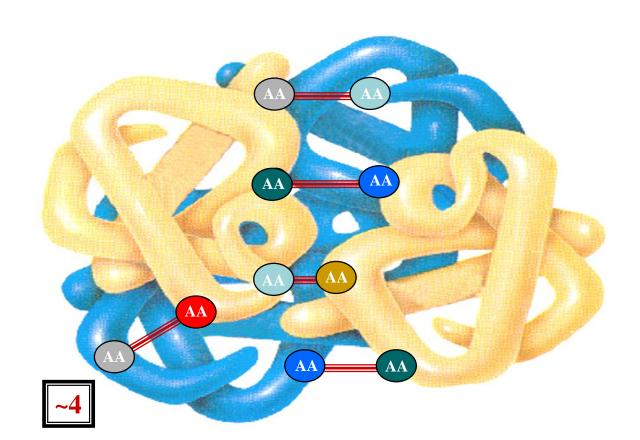


BD

TEMPERATURE INCREASE INCREASES ENZYME MOLECULAR MOVEMENT

H-BONDS WEAK BONDS

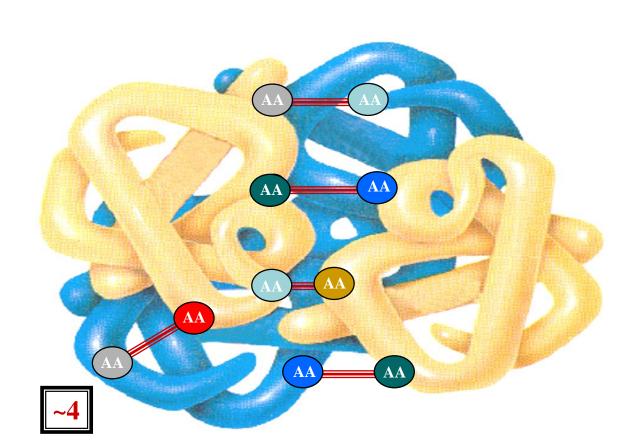
ENZYME



TEMPERATURE INCREASE INCREASES ENZYME MOLECULAR MOVEMENT

H-BONDS BREAK-DOWN

ENZYME

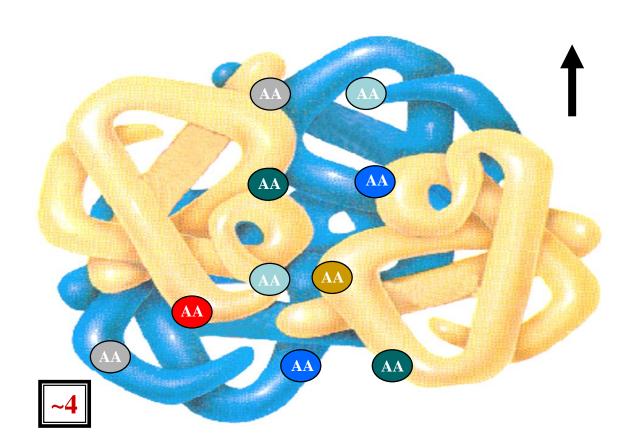


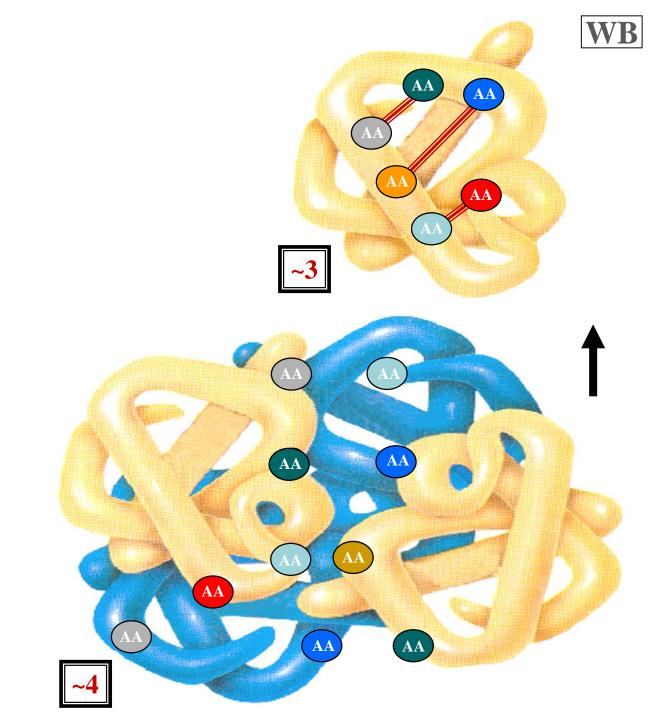


TEMPERATURE INCREASE INCREASES ENZYME MOLECULAR MOVEMENT

H-BONDS BREAK-DOWN

ENZYME





H-BONDS

ENZYME