



INTERNATIONAL ATOMIC ENERGY AGENCY
UNITED NATIONS EDUCATIONAL, SCIENTIFIC AND CULTURAL ORGANIZATION
INTERNATIONAL CENTRE FOR THEORETICAL PHYSICS
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H4.SMR/642 - 11

College on Methods and Experimental Techniques in Biophysics

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Protein Folding
(Transparencies)

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These are preliminary lecture notes, intended only for distribution to participants.

Protein Folding - Lecture 1

①

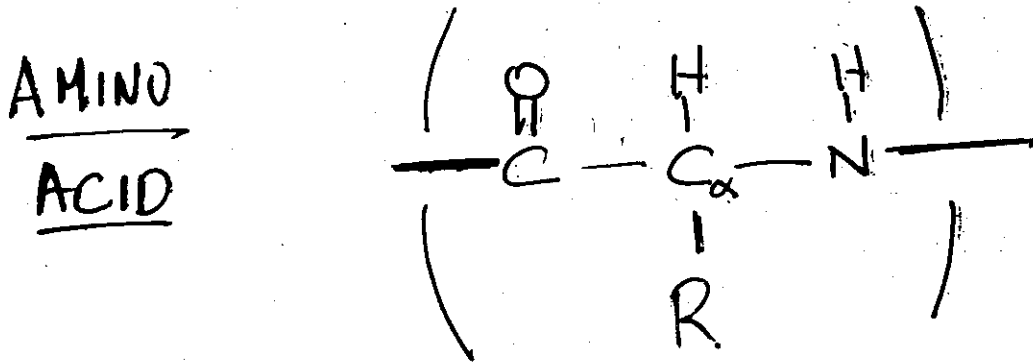
"PROTEIN'S"

MAIN TOPIC OF STUDY DURING
THIS COLLEGE

- Several techniques will be described - Most of the applications to proteins

WHAT ARE PROTEINS?

Proteins are heteropolymers
of Amino-Acids



R characterizes the Amino-Acid

A protein can be described by its
sequence of Amino-Acids!!!

Looks obvious!!!

But we will elaborate on
this statement during our lectures

Why we care about proteins?

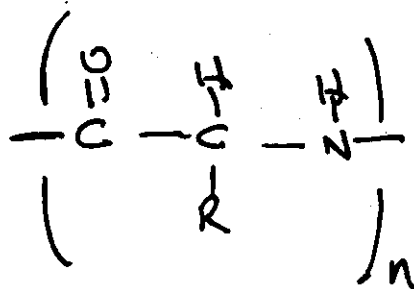
Biological Functions — Proteins
do the work!

For example !

- Energy Conversion
- Oxygen and CO_2 transport
- Catalysis (Enzymes)
- Immune Recognition
- Information Process
-

However

Function (and dynamics)
depends on the 3-D
structure of the protein
- Not the sequence



————→ D
proper
conditions



UNIQUE STATE
(FOLDED STATE)

ENERGETICS AND GEOMETRY OF

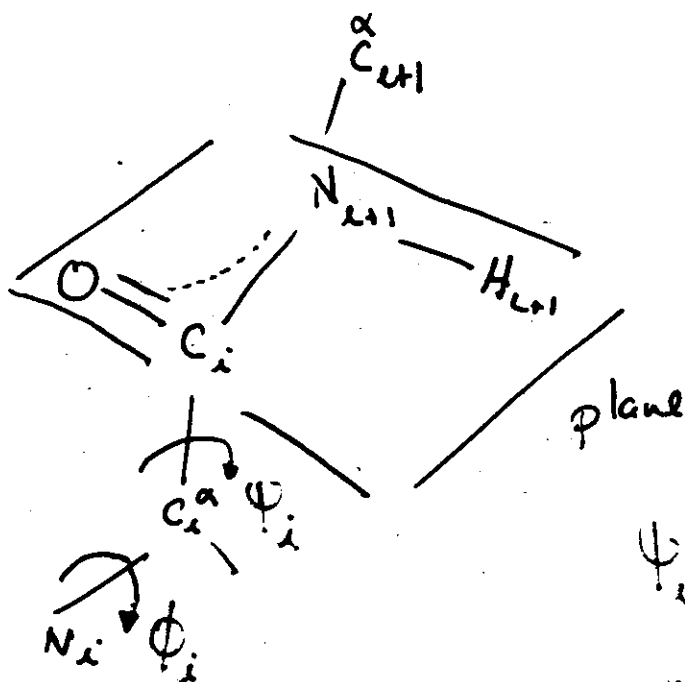
HOW PROTEINS ORGANIZE THEMSELVES 3D

- Lubert Stryer, "Biochemistry", chapter 2, W.H. Freeman, New York, third edition 1988
- T.E. Creighton, "Proteins: Structures and Molecular Properties", W.H. Freeman, New York 1984

(People without a reasonable knowledge of this subject should look for me.)

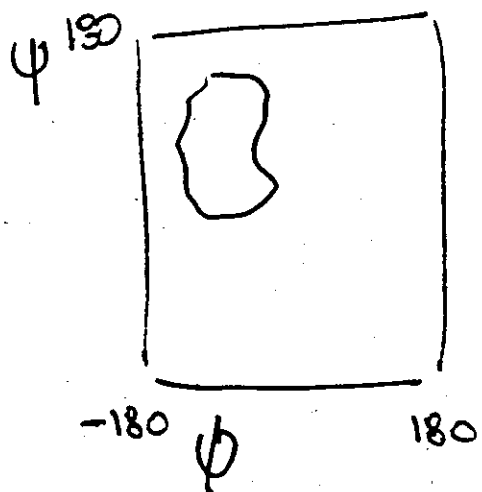
Highlight 3 points

①



ψ_i, ϕ_i define the
"local" 3-D structure
of the protein.

Rama Chandra Plot



Possible values of
 ψ and ϕ
 "Geometrical Constraints"
 Possible local geometries

② Hydrogen Bonds $\text{C}=\text{O}$ and $\text{H}-\text{N}$ of AA's
 like to H.B.

→ Two possible periodic structures
 (ϕ, ψ) are periodic
 α -Helix
 β -Sheet

③ Solvent Effects - hydrophobic interaction

any

the

the

STRUCT

Coc

for

Coupling

2.55E-01

4.00E-02

9.55E-01

1.06E-01

3.61E-02

Coup 117

1.12E+01

2.25E+00

4.55E+01

8.95E+02

1.80E+02



Primary Structure - Sequence (6)

Secondary Structure - local geometry (α helix, β sheet)

Tertiary Structure - global geometry

(quaternary structure - more than 1 sequence)

Look the importance
of the effects described before

SEQUENCE-STRUCTURE RELATIONSHIP

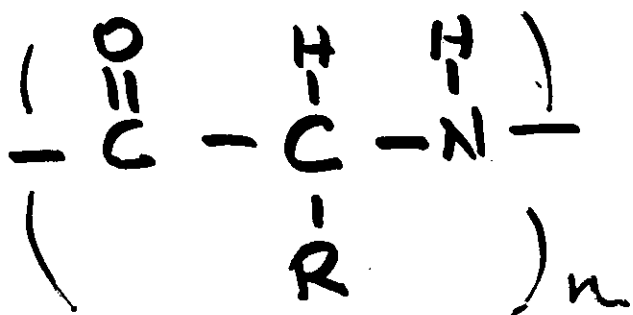
However, for Biological
Proteins, one sequence
defines one structure

Second Genetic Code

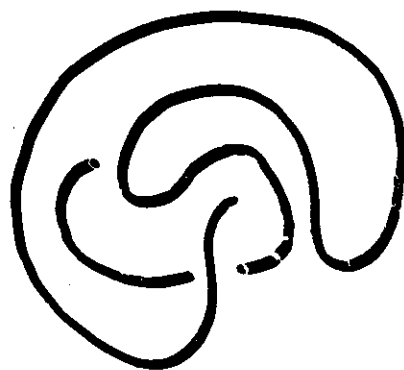
"All the information is on the sequence"

Proteins

Amino Acid Polymer



proper conditions



UNIQUE
STATE

(FOLDED
STATE)

R - defines the AA.

NOT ALL SEQUENCES FOLD !!!

Protein
Folding
Problem

CAN YOU PREDICT THE 3-D
STRUCTURE BASED ON THE
AMINO ACID SEQUENCE ?

CAN YOU PREDICT IF A
STRUCTURE WILL FOLD ?

FOLDING MECHANISM

(4)(4)

Ribonuclease

Anfinsen '60's (Ribonuclease)

SMALL PROTEINS (GLOBULAR) FOLD (REFOLD)
IN THE ABSENCE OF ANY CATALYTIC MOLECULE

• THERMODYNAMIC HYPOTHESIS (GLOBAL STABLE MINIMUM)

Levinthal '68

TOO MANY STATES TO PROBE TO FIND THE
RIGHT ONE!!! PARADOX

each AA. 2 positions 2^N states $N=100$
 10^{30}
? PROTEIN PROBE 10^{12} $\frac{\text{states}}{\text{sec}}$

$\searrow$
 10^{18} s

To save the T.H.

Baldwin '83 $\rightarrow$ Folding Pathway

FACT Not All Proteins fold
only biological ones

∴
Not all sequences meet the
thermodynamic and kinetic criteria

Theoretical Efforts → MECHANISM?

Analytical
Mean Field
Spin Glass
Polymer
Mean Field
} thermodynamics - Dill
folding times - Shakhnovich
- Bryngelson
and Wolynes

Lattice Models
Preserved
Sequence
Information
} sequence-energy-structure
exact summation
Lattice kinetic { Folding Pathways??
simulations
Dill, Shakhnovich, Skolnick, Je

THERMODYNAMICS

6

① Folded State Unique

- Two state system

② First Order Phase Transition

ORIGIN?

- ② If this is correct - we need
some energy gap

between the lowest state (folded)

and the

continuum of other (un-folded)
states

DYNAMICS - How to beat Levinthal?

① Diffusion-Collision Model

FIRST SECONDARY STRUCTURES
SECOND TERTIARY STRUCTURE

② Collapse-Reconfiguration Model

FIRST COLLAPSE -
SECOND Boundary Structure

③ Framework Model

single secondary structure
nucleates the process

→ This dynamical views influence even the

Results

66

- Folded Structures - stable
(thermodynamical stability)

??

- Principle of Minimal
Frustration

Band W

How?

- Lattice kinetics - Pathways

- Potentials Seq. - St.
 relationship

N- Letter Code

