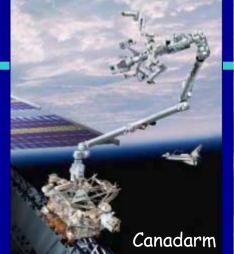
# Folding & Unfolding: Linkages

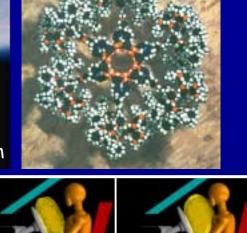
# Erik Demaine M.I.T.

edemaine@mit.edu http://theory.lcs.mit.edu/~edemaine/folding

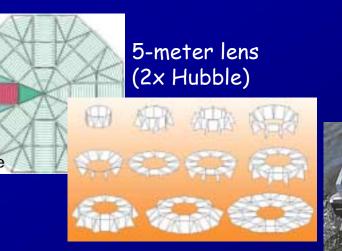
# Folding and Unfolding in Science

Linkages Robotic arms Proteins Paper Airbags Space deployment Polyhedra Hyde Sheet metal







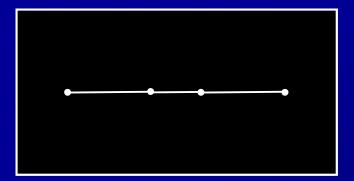




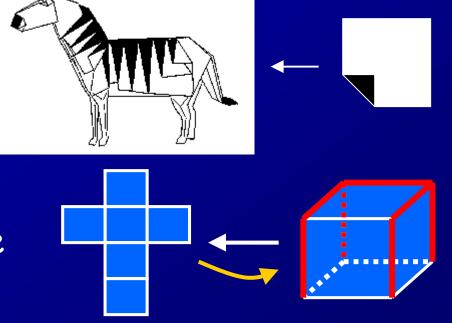
Touch-3D

#### Folding and Unfolding in Computational Geometry

Linkages
Preserve edge lengths
Edges cannot cross



Paper
Preserve distances
Cannot cross itself
Polyhedra
Cut the surface while keeping it connected



# Folding and Unfolding Talks

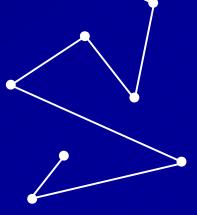
Linkage folding	Today	Erik Demaine
Paper folding	Tomorrow	Erik Demaine
Folding polygons into convex polyhedra	Friday	Joe O'Rourke
Unfolding polyhedra	Saturday	Joe O'Rourke

# **Outline: Linkages**

Definitions and History Rigidity Locked chains in 3D Locked trees in 2D No locked chains in 2D Algorithms Connections to protein folding

## Linkages / Frameworks

# Bar / link / edge = line segment Vertex / joint = connection between endpoints of bars



Open chain / arc Closed chain / cycle / polygon

Tree

General

## Configurations

Configuration = positions of the vertices that preserves the bar lengths

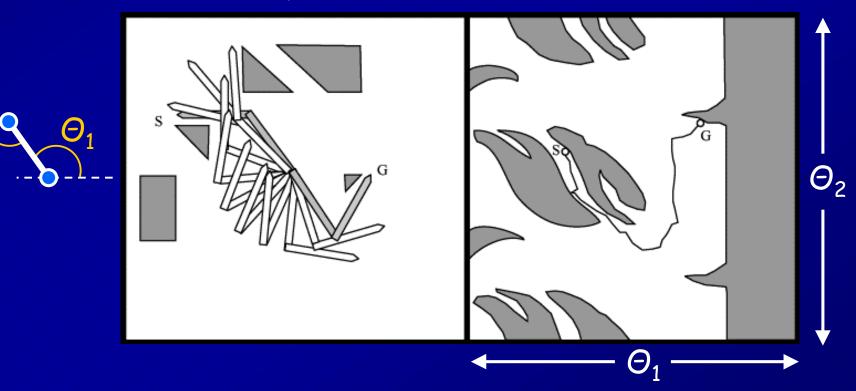
Non-self-intersecting = No bars cross

Non-self-intersecting configurations

Self-intersecting

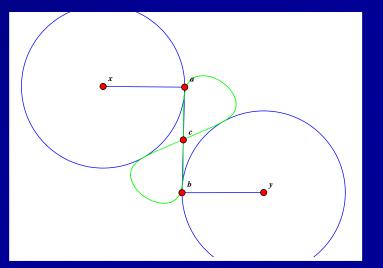
## **Configuration Space**

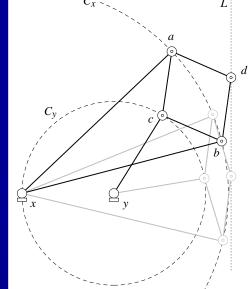
Configuration space: One point per config.
Free space: Only non-self-intersecting configs.
Paths in these spaces = Motions of linkage



#### Some History

# An early quest: Converting circular motion into linear motion





Watt parallel motion (1784) Peaucellier linkage (1864)

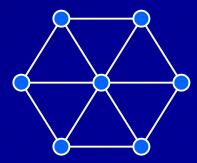
Many more in Kempe's How To Draw A Straight Line (1877)

# **Outline: Linkages**

Definitions and History Rigidity Locked chains in 3D Locked trees in 2D No locked chains in 2D Algorithms Connections to protein folding

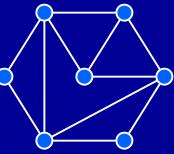


# Starting question: When can a config. of a linkage move at all? [excluding rigid motion] Yes ⇒ Flexible; No ⇒ Rigid



Rigid

(all triangulations)



Rigid

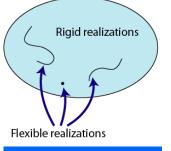


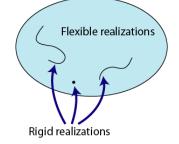
Flexible

Rigid

(all "pseudotriangulations")

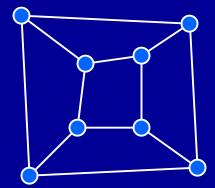
# **Generic Rigidity**



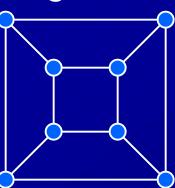


Whether a linkage configuration is rigid is almost always a combinatorial property of the underlying graph structure:

- Generically flexible = almost all realizations of the graph are flexible
- Generically rigid = almost all realizations rigid



Generically flexible



Rarely rigid



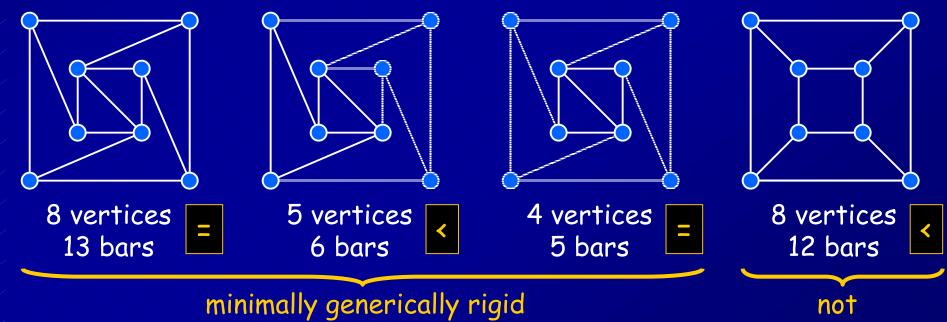
Generically rigid Rarely flexible

#### Laman's Characterization of Generic Rigidity

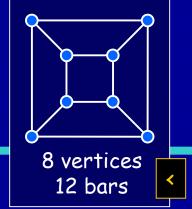
#### Theorem: [Laman 1970]

A graph with n joints and exactly 2n - 3 bars is generically rigid in 2D precisely if every induced subgraph on k joints has at most 2k - 3 bars

#### Such graphs are minimally generically rigid



#### Laman's Characterization of Generic Rigidity

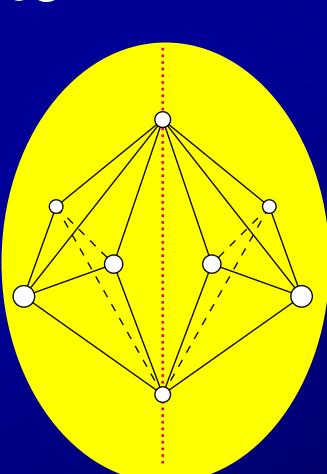


#### Theorem: [Laman 1970]

- A graph with n joints and exactly 2n 3 bars is generically rigid in 2D precisely if every induced subgraph on k joints has at most 2k – 3 bars
- A graph with n joints and less than 2n 3 bars is never generically rigid in 2D
- A graph with n joints and more than 2n 3 bars is generically rigid in 2D precisely if it has a generically rigid subgraph with 2n – 3 bars
- Intuitively, need 2n 3 bars to be rigid, but these bars cannot be too concentrated, else another subgraph would have too few edges

#### Generic Rigidity in Higher Dimensions

Open: Characterize the 3D generically rigid graphs Can these graphs be recognized in polynomial time? Laman's theorem generalizes to a necessary condition (3n - 6)



# **Infinitesimal Rigidity**

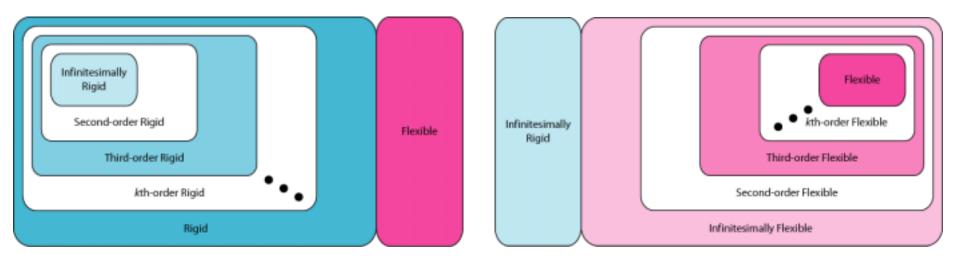
If a linkage is flexible, it is flexible by a smooth motion  $\Rightarrow$  can take first derivative Infinitesimal motion defines a motion to first order, from initial configuration x Suppose x<sub>i</sub> is the location of vertex Choose velocity vector v<sub>i</sub> for each i Lengths of each bar {i, j} must stay V: constant to the first order:  $(\mathbf{v}_i - \mathbf{v}_j) \cdot (\mathbf{x}_i - \mathbf{x}_j) = \mathbf{0}$ X Constraints form rigidity matrix

# **Rigidity and Infinitesimal Rigidity**

Flexibility ⇒ Infinitesimal flexibility
 Infinitesimal rigidity ⇒ Rigidity

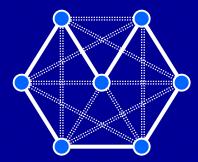
 So infinitesimal rigidity is a stronger condition

 There is also second-order, etc. rigidity

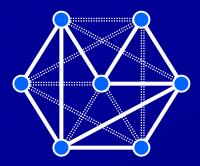


#### Tensegrities

 Tensegrity = generalization of linkage, where each edge can be one of
 Bar — length must stay equal
 Strut — length must stay equal or grow
 Cable — length must stay equal or shrink
 Plain/generic/infinitesimal rigidity similar

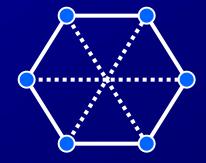


Flexible



Flexible

Flexible



Rigid

#### Infinitesimal Flexibility is Linear Programming

Infinitesimal flexibility can be expressed as a linear feasibility problem (special linear program)
Objective: Minimize 0
Constraints:

 $(v_i - v_j) \cdot (x_i - x_j) = 0$   $(v_i - v_j) \cdot (x_i - x_j) \ge 0$   $(v_i - v_j) \cdot (x_i - x_j) \le 0$   $v_i \text{ variable; } x_i \text{ given}$ 

for each bar {i, j}
for each strut {i, j}
for each cable {i, j}

#### Dual of Infinitesimal Motions: Equilibrium Stresses [Roth & Whiteley 1981]

Primal LP infeasible  $\Rightarrow$ dual LP infeasible or unbounded Equilibrium stress = assignment of weights  $W_{\{i,j\}}$  to edges  $\{i, j\}$  satisfying ■  $W_{\{i,i\}} \ge 0$  for all struts {i, j} w<sub>{i,j</sub>} ≤ 0 for all cables {i, j} Equilibrium: For each vertex i,  $\sum \{ w_{\{i,j\}} (x_i - x_j) \mid edge \{i, j\} \} = 0$ Any infinitesimally rigid framework has an equilibrium stress that's not everywhere 0

#### Dual of Infinitesimal Motions: Equilibrium Stresses [Roth & Whiteley 1981]

Equilibrium stresses do not imply rigidity Do imply "local rigidity" where nonzero: No infinitesimal motion can change the length of a strut or cable with nonzero weight in some equilibrium stress Tensegrity is infinitesimally rigid iff There is an equilibrium stress that is nonzero on all struts and cables • Underlying linkage (all edges  $\rightarrow$  bars) is infinitesimally rigid

# Maxwell-Cremona Relation: Stresses and Liftings

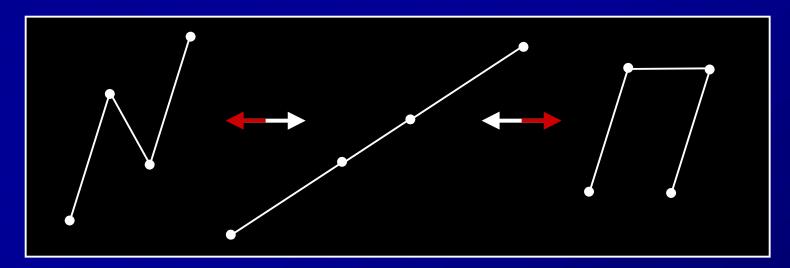
Polyhedral lifting = assignment of z coordinates to vertices such that faces of framework remain planar Can assume z = 0 on boundary face Maxwell-Cremona Theorem: A framework has a nonflat lifting precisely if it has a nonzero stress Valleys  $\leftrightarrow$  positive weights w (struts and bars) Mountains  $\leftrightarrow$  negative weights  $\omega$ (cables and bars)

# **Outline: Linkages**

Definitions and History Rigidity Locked chains in 3D Locked trees in 2D No locked chains in 2D Algorithms Connections to protein folding

#### Locked Question

- Can a linkage be moved between any two non-self-intersecting configurations?
- Can any non-self-intersecting configuration be unfolded, i.e., moved to "canonical" configuration?
  - Equivalent by reversing and concatenating motions

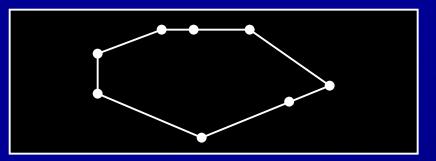


## **Canonical Configurations**

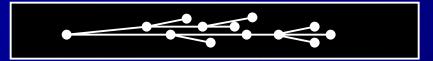
#### Arcs: Straight configuration



#### Cycles: Convex configurations



Trees: Flat configurations



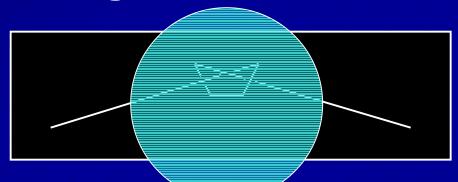
#### What Linkages Can Lock? [Schanuel & Bergman, early 1970's; Grenander 1987; Lenhart & Whitesides 1991; Mitchell 1992]

Can every arc be straightened?
Can every cycle be convexified?
Can every tree be flattened?

	Arcs	Cycles	Trees
2D	Yes	Yes	No
3D	No	No	No
4D & higher	Yes	Yes	Yes

Locked 3D Chains [Cantarella & Johnston 1998; Biedl, Demaine, Demaine, Lazard, Lubiw, O'Rourke, Overmars, Robbins, Streinu, Toussaint, Whitesides 1999]

#### Cannot straighten some chains



Sphere separates turns from ends

#### Idea of proof:

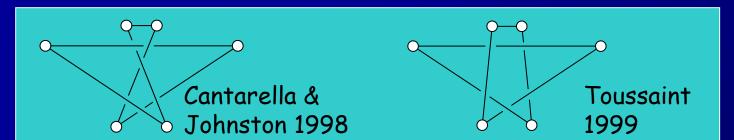
Ends must be far away from the turns
 Turns must stay relatively close to each other
 ⇒ Could effectively connect ends together
 Hence, any straightening unties a trefoil knot

Locked 3D Chains [Cantarella & Johnston 1998; Biedl, Demaine, Demaine, Lazard, Lubiw, O'Rourke, Overmars, Robbins, Streinu, Toussaint, Whitesides 1999]

Double this chain:



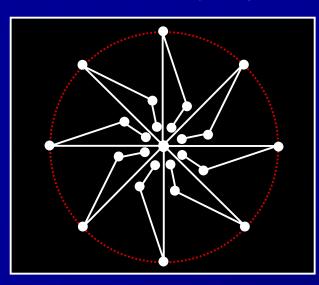
This unknotted cycle cannot be convexified by the same argument
 Several locked hexagons are also known



#### Locked 2D Trees

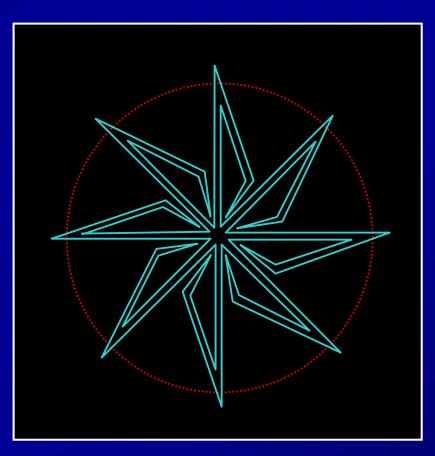
[Biedl, Demaine, Demaine, Lazard, Lubiw, O'Rourke, Robbins, Streinu, Toussaint, Whitesides 1998]

- Theorem: Not all trees can be flattened
  - No petal can be opened unless all others are closed significantly
  - No petal can be closed more than a little unless it has already opened



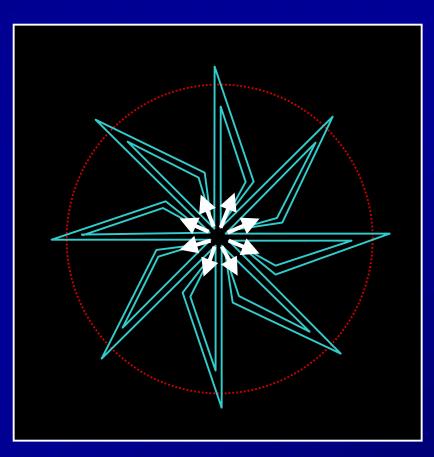
## **Converting the Tree into a Cycle**

#### Double each edge:



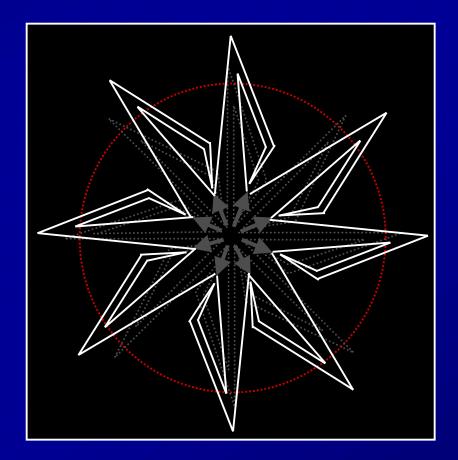
## **Converting the Tree into a Cycle**

#### But this cycle can be convexified:



## **Converting the Tree into a Cycle**

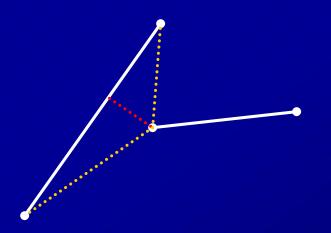
#### But this cycle can be convexified:



# One Key I dea for 2D Cycles: Increasing Distances

A motion is expansive if no inter-vertex distances decreases

Lemma: If a motion is expansive, the framework cannot cross itself

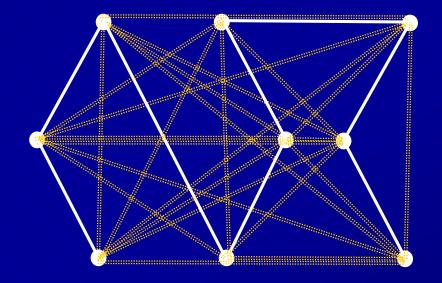


#### Theorem [Connelly, Demaine, Rote 2000]

- For any family of chains and cycles, there is a motion that
  - Makes the arcs straight
  - Makes the cycles convex
  - Increases most pairwise distances (and area)
- Except: Arcs or cycles contained within a cycle might not be straightened or convexified
- Furthermore: Motion preserves symmetries and is piecewise-differentiable (smooth)

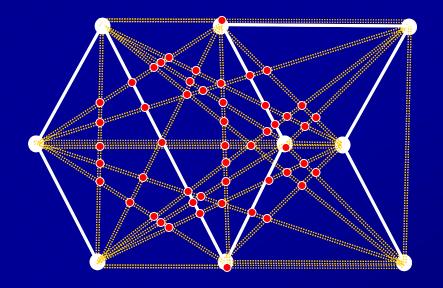
#### Transforming Linkage into Tensegrity

In addition to bars of the framework, add struts between each pair of vertices not in a common convex cycle



# Planarizing the Tensegrity

Subdivide edges at intersection points
Remove multiple overlapping edges
Replace with a bar if there is a bar
Replace with a strut otherwise

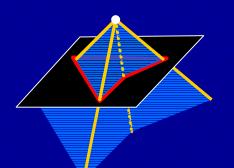


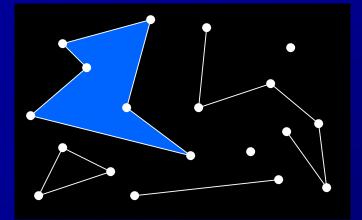
## Proof of Existence of an Infinitesimal Motion

- Original framework infinitesimally flexible
   (duality)
- Original framework has only the zero equilibrium stress
  - I ↑ (planarization lemma)
- Planar framework has only the zero equilibrium stress
  - I ↑ (Maxwell-Cremona Theorem)
- Planar framework has only the flat polyhedral lifting

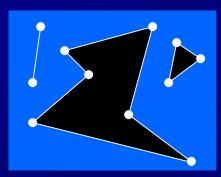
#### Proof of Existence of an Infinitesimal Motion

- Consider the top extreme M of some polyhedral lifting of the planar framework
  - One case: A vertex and none of its surroundings are at the top
    - Slice just below the vertex





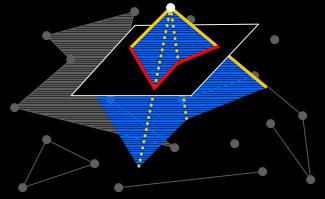
Red polygon has reflex vertices for valleys, convex for mountains
Must be at least three convex vertices

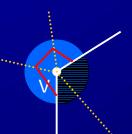


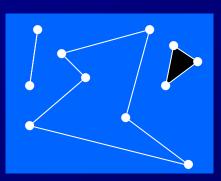
#### Proof of Existence of an Infinitesimal Motion

Consider the top extreme M of some polyhedral lifting of the planar framework

- Let v be a vertex of  $\partial M$
- Consider a small disk around v
- Suppose there is a reflex portion of the disk between two consecutive bars
- Claim this portion must be contained in M







#### **Existence of Global Motion**

Quadratic program defines a unique ordinary differential equation on an open subset of the configuration space, defined by the conditions: No vertex has angle 180° No vertex touches a bar Some vertex is reflex Continuity  $\Rightarrow$  There is a path to infinity or the boundary of the open set Path is bounded because of bars' limited reach

#### **Outline: Linkages**

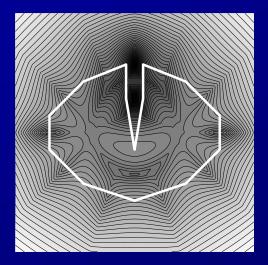
Definitions and History Rigidity Locked chains in 3D Locked trees in 2D No locked chains in 2D Algorithms Connections to protein folding

# **Algorithms for 2D Chains** Connelly, Demaine, Rote (2000) — ODE + convex programming Streinu (2000) — pseudotriangulations + piecewise-algebraic motions Cantarella, Demaine, Iben, O'Brien (2003) — energy



#### An Energy-Driven Approach to Linkage Unfolding

Jason Cantarella Erik Demaine Hayley Iben James O'Brien









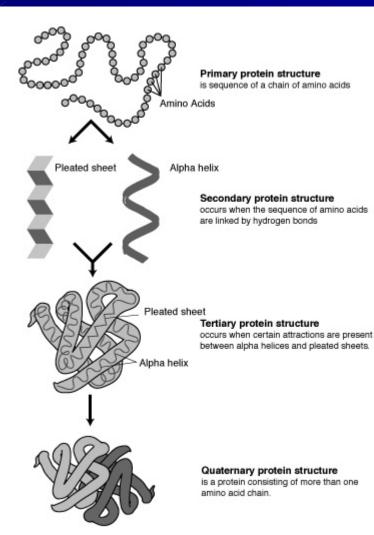


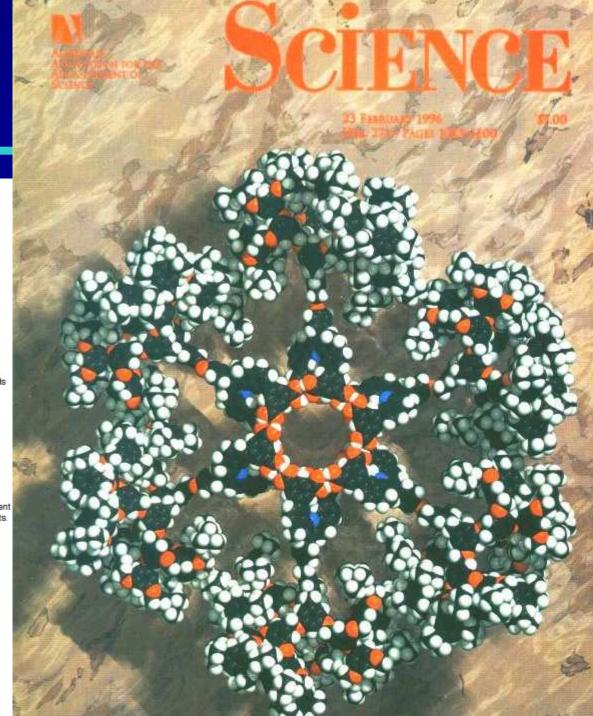


#### **Outline: Linkages**

Definitions and History Rigidity Locked chains in 3D Locked trees in 2D No locked chains in 2D Algorithms Connections to protein folding

#### Protein Folding





#### **Protein Folding**

#### Motivation

Geometry of a protein folding is an important aspect of its behavior Prediction of protein folding, and synthesis of proteins with desired foldings, are central problems in computational biology Drug design Preventing diseases (e.g., Alzheimer's, mad-cow disease, cystic fibrosis, some forms of cancer) Understanding genomes

## Linkage problems arising in protein folding

Fixed-angle linkages (in 3D): In addition to bar lengths, joint angles remain fixed Protein is roughly a fixed-angle tree When are all flat states connected via motions? [Aloupis, Demaine, Dujmović, Erickson, Langerman, Meijer, O'Rourke, Overmars, Soss, Streinu, Toussaint 2002x2] Nonacute chains; equal-angle acute chains Not general planar graphs **Open:** All chains? All trees?

## Linkage problems arising in protein folding

Equilateral chains All bar lengths (roughly) equal Protein backbone is roughly such a chain Open: Can all 3D equilateral chains be straightened? Open: Can all 3D equilateral trees be flattened?

Hydrophobic-Hydrophilic / H-P Model [Dill 1990]

Nodes (20 amino acids) categorized into two types:

Hydrophobic (H): Afraid of surrounding water Hydrophilic (P): Like surrounding water Model: Proteins fold on 2D or 3D lattice to maximize number of bonds = lattice connections between nonadjacent H nodes NP-hard to find optimal protein folding Open: Design protein with a desired shape as (roughly) its unique optimal folding