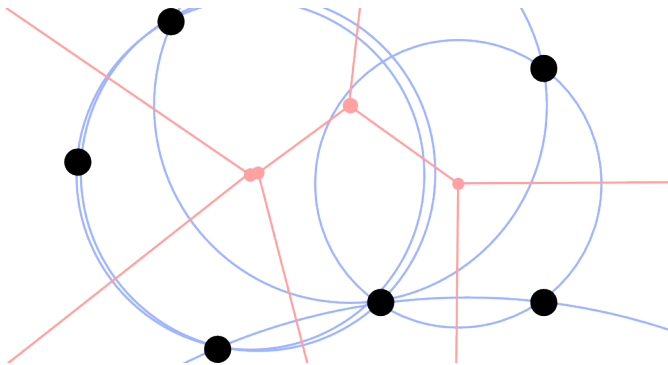
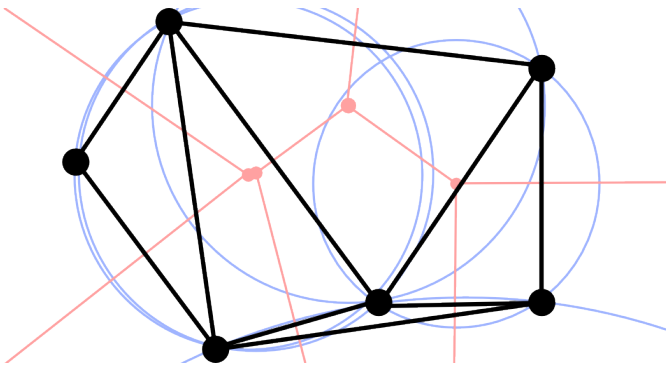


# Voronoi Diagram



- Given a set  $P$  of  $n$  points in the plane (general position), its **Voronoi diagram** is a partition of the plane into  $n$  cells, each containing one point of  $P$  and everything closer to it than to any other point.

# Delaunay Triangulation

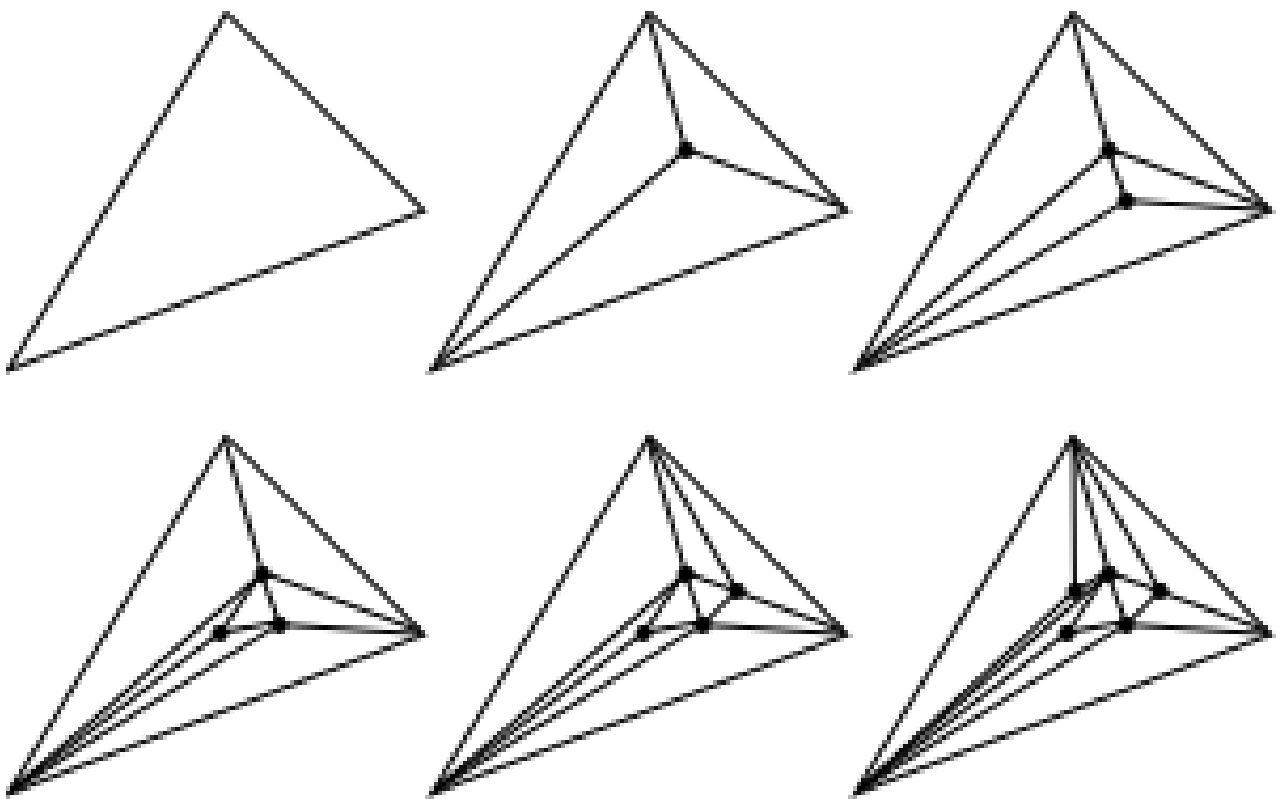


- 3 points  $p_i, p_j, p_k$  in  $P$  form a triangle in  $DT(P)$  iff the unique circle through  $p_i, p_j, p_k$  contains no other point from  $P$ .
- 2 points  $p_i, p_j$  in  $P$  form an edge in  $DT(P)$  iff there exists a circle through  $p_i$  and  $p_j$  that contains no other point from  $P$ .

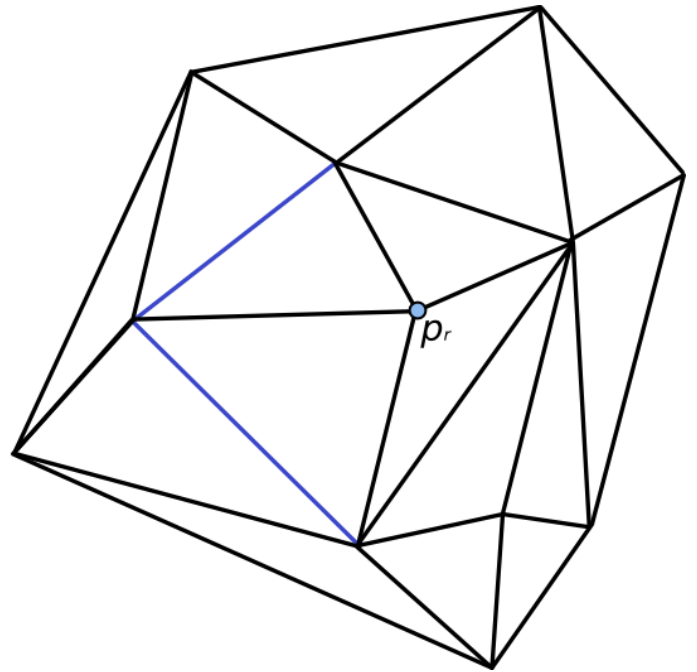
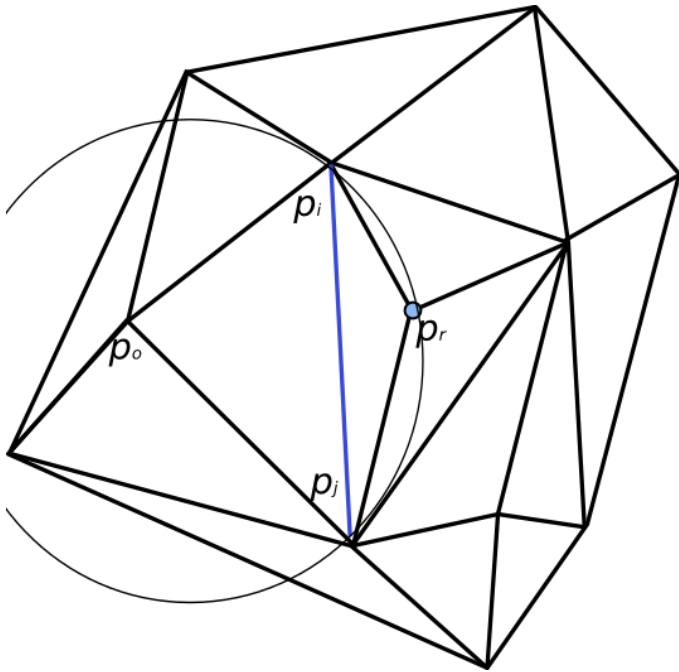
# Constructing DT

- Construct  $VD(P)$  in  $O(n \log n)$  time and  $O(n)$  space and convert it to  $DT(P)$  in  $O(n)$  time.
- $VD(P)$  algorithms are complex and can cause serious precision problems.
- $DT(P)$  can be determined directly in  $O(n^2)$  time.
  - Randomized version  $O_e(n \log n)$ .
  - Generalizes to 3D.

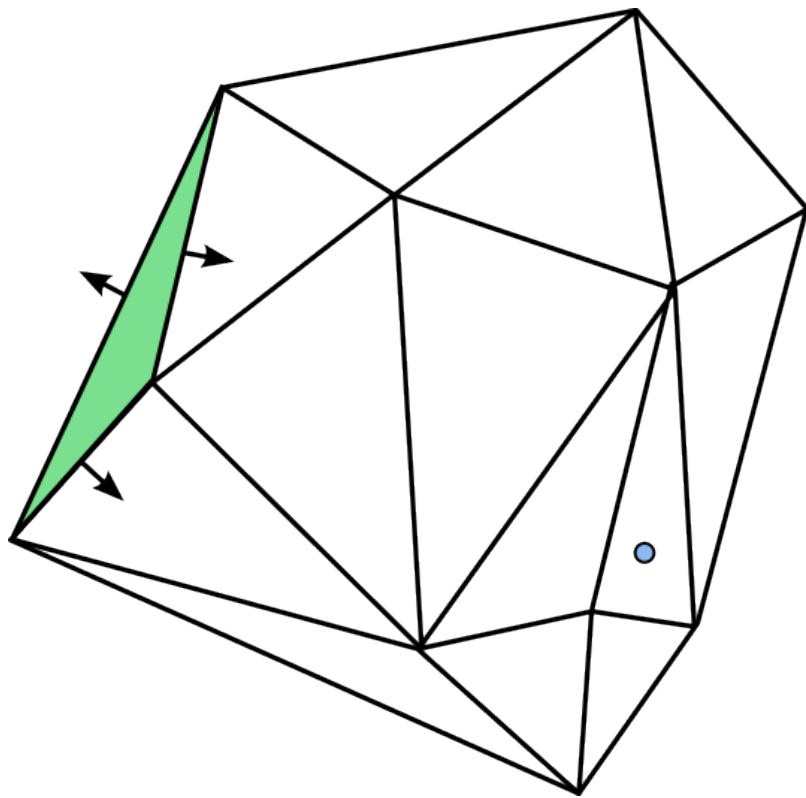
# Constructing DT



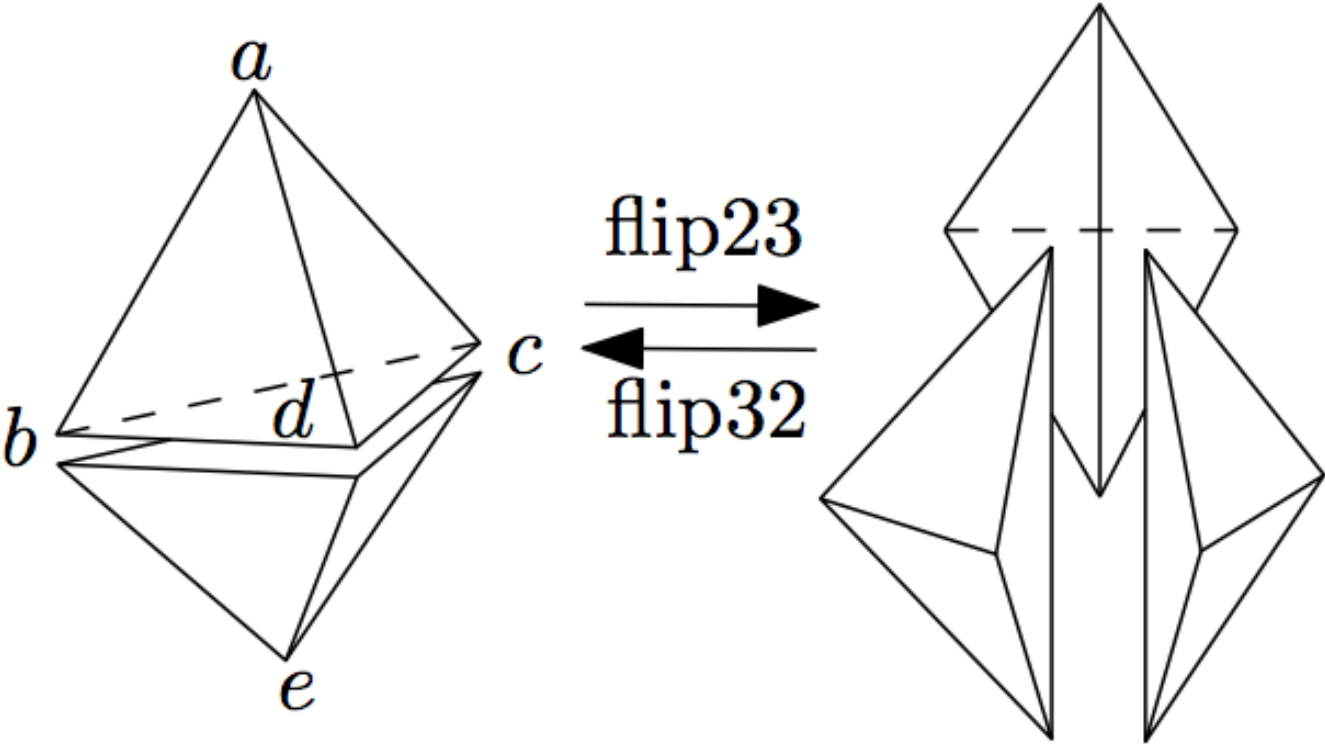
$O(n)$  Flips pr. Added Point



# Locating a Triangle in $O(n)$ Time



Flips in  $R^3$



# Kinetic Delaunay Triangulation in $\mathbb{R}^2$

- $n$  points with (piecewise) linear trajectories.
- Events:
  - Insertion and deletion (disregarded in the following).
  - Trajectory and/or speed change events (deletion followed by insertion).
  - Circle events: 4 points on a common circle.
  - Side events: 3 points on a common line where one half-plane contains no points. Can be considered as a special case of circle events (imagine everything inside a huge triangle with stationary corners).
- Simplifying assumption: No collisions, no pair of events occurs at the same time.



## Incircle test

$$\begin{vmatrix} a_x & a_y & a_x^2 + a_y^2 & 1 \\ b_x & b_y & b_x^2 + b_y^2 & 1 \\ c_x & c_y & c_x^2 + c_y^2 & 1 \\ d_x & d_y & d_x^2 + d_y^2 & 1 \end{vmatrix}$$

- If this determinant is 0 then point  $d$  is inside the circle through points  $a$ ,  $b$ , and  $c$ ?
- Why?
- Assuming linear trajectories, roots of a 4-th degree polynomial need to be determined to find time  $t$  where these 4 points are cocircular.

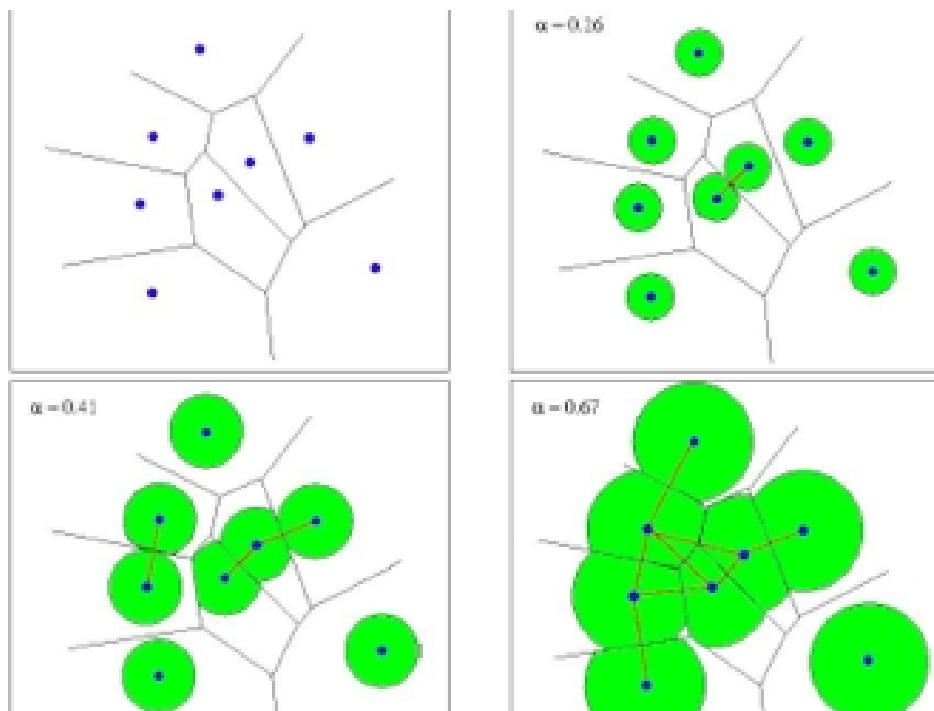
## Events – Kinetic DT in $\mathbb{R}^2$

- Determine DT at  $t = 0$ .
- Identify times for flip events for pairs of triangles sharing an edge and place these events on a heap. One certificate pr. edge.
- Repeat:
  - Remove top flip events from the heap and flip.
  - Delete future events involving one of the two deleted triangles (or use lazy deletion).
  - Identify times for up to 5 new flip events and place them on the heap.
- until the heap is empty or  $t > t_{\text{end}}$

## Kinetic DT in $\mathbb{R}^2$

- Responsive: Well, flips require  $O(1)$  time, 5 new certificates need to be added. Computing their failure time depends on trajectories. Linear trajectories involve finding roots of polynomials of at most 4-th degree.
- Compact: Yes,  $O(n)$  certificates at any time since DT is planar.
- Local: No, but expected degree of a vertex in DT is 6, so this is the expected number of certificates involving it.
- Efficient: Unsure. Some scheduled events can become inactive due to neighboring lines

# $\alpha$ -Complexes via Voronoi Diagrams



# $\alpha$ -Complexes via Delaunay Triangulations

- 0-simplex=vertex, 1-simplex=edge, 2-simplex = triangle, 3-simplex = tetrahedron (in  $\mathbb{R}^3$ )
- $k$ -simplex is **short** iff its smallest circumcircle has radius at most  $\alpha$ .
- $k$ -simplex is **Gabriel** iff its smallest circumcircle contains no other points.
- $k$ -simplex of DT is in  $\alpha C$  iff it short and Gabriel, or it is a face of another DT-simplex that is short and Gabriel.
- A face of a short simplex is short but a face of a Gabriel simplex is not necessarily Gabriel.

## Events – Kinetic $\alpha C$ in $R^2$

- Determine DT at  $t = 0$ .
- Short triangles of DT are automatically Gabriel. They and their faces are in  $\alpha C$ .
- If an edge is not already in  $\alpha C$ , it is checked if it is short and Gabriel. If so, it is added to  $\alpha C$ .
- Times for flip and radius events are determined and stored in a heap.
- What about Gabriel events for edges? They are actually redundant.

# Redundancy of Gabriel Events

- Any triangle of  $\alpha C$  in  $R^2$  is automatically Gabriel (otherwise it would not be in DT).
- Consider an edge  $ab$  of DT that is about to change from Gabriel to non-Gabriel or vice versa.
- At the time of transition, its smallest circumcircle goes through  $a$  and  $b$  and through a third vertex  $c$ .
- $\Delta abc$  is short and Gabriel and therefore in  $\alpha C$ . Hence, all faces of  $\Delta abc$  (including  $ab$ ) are in  $\alpha C$ .

# Kinetic $\alpha C$ in $R^2$

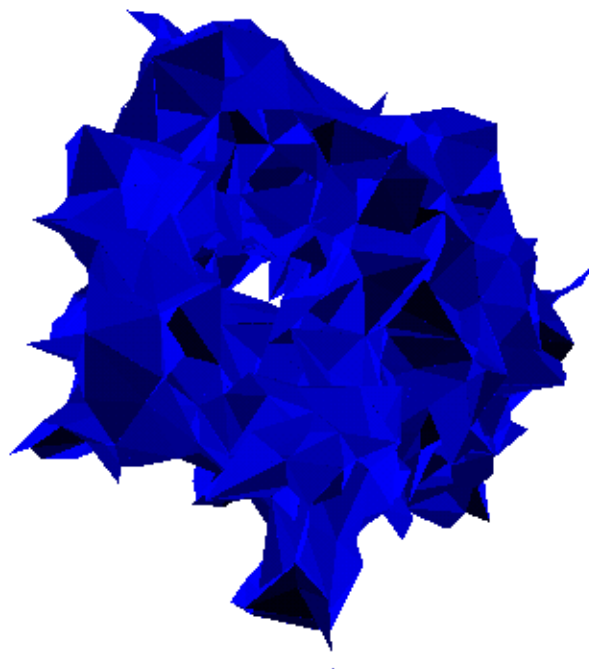
- Place flip events for pairs of edge-sharing DT-triangles and radius events for DT-triangles and DT-edges on a heap.
- Repeat:
  - If next is a flip event, flip. If the original pair of triangles was short so is the new pair. If it was non-short, check if the new edge is short. Create 5 new flip events.
  - If it is a radius event and its simplex  $\sigma$  is to become
    - short: If  $\sigma$  is Gabriel,  $\sigma$  and all its faces are added to  $\alpha C$ .
    - non-short: all faces of  $\sigma$  are short. Remove  $\sigma$  from  $\alpha C$  (if it is in  $\alpha C$ ). If a face of  $\sigma$  is not Gabriel and has no other coface in  $\alpha C$ ,  $\sigma$  has to be removed from  $\alpha C$ .
- until the heap is empty or  $t > t_{end}$



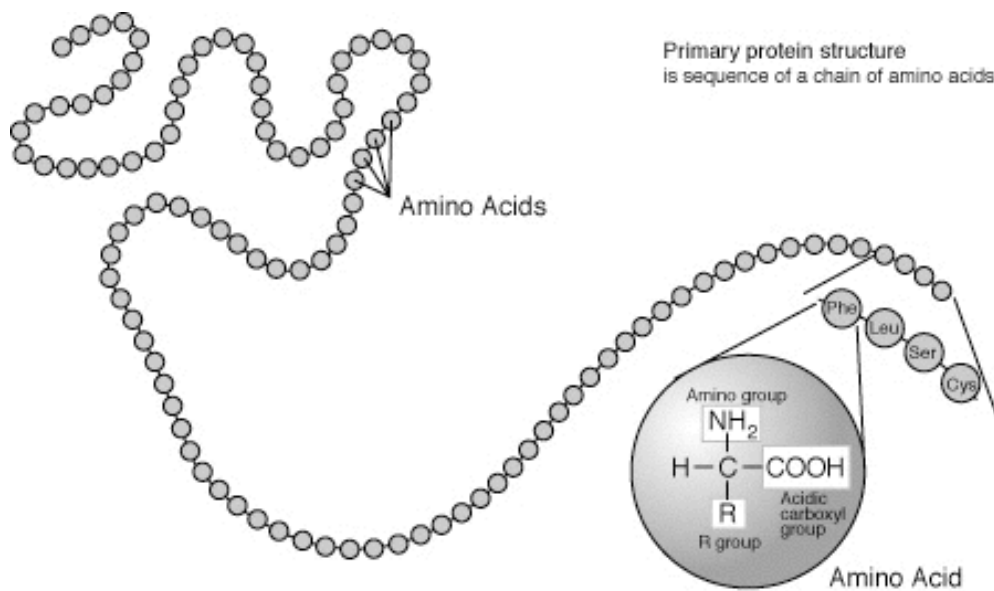
# Certificate Failures

- Assuming piecewise-linear trajectories, finding the time when a radius certificate of an
  - Edge-sharing pair of DT-triangles fails, requires finding a root of a polynomial of degree 2.
  - triangle fails, requires finding a root of a polynomial of degree 5.
  - tetrahedron fails, requires finding a root of a polynomial of degree 8 (this is  $R^3$  case).

# A-Complexes of Proteins



# Amino Acid Chain

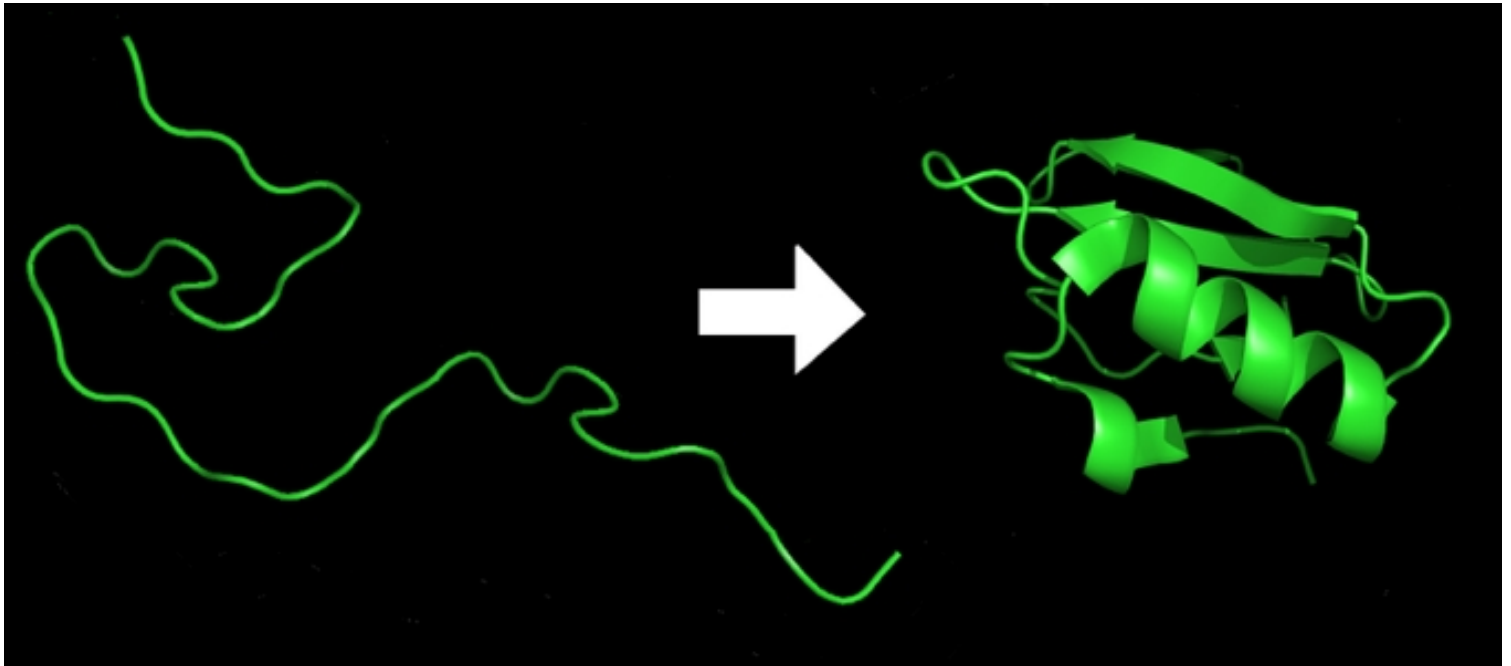


**Backbone:** disregard side chains and hydrogens.

From wikipedia

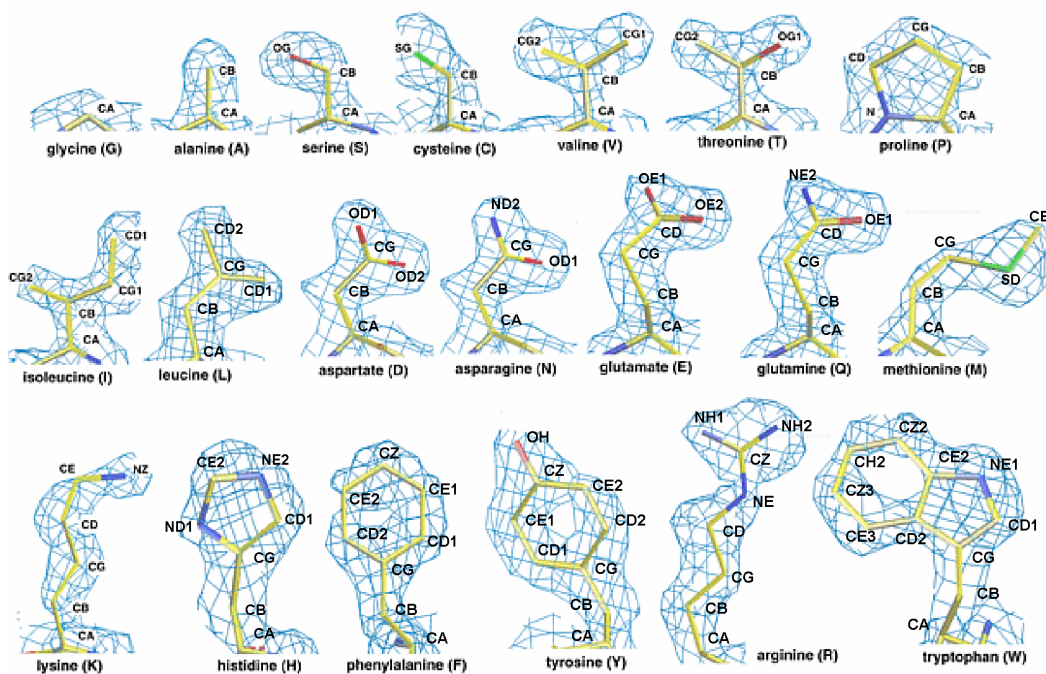
**$\text{C}_\alpha$ -trace:** disregard all but  $\text{C}_\alpha$ -atoms.

# Folding



From Wikimedia Commons

# 20 Different Side Chains



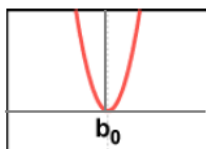
0-5 degrees of freedom pr. side chain

From <http://www.doe-mbi.ucla.edu/Legacy/CHEM153BH/info2.htm>

# Potential Energy Function

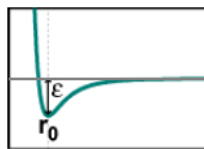
- $U = \text{Bond} + \text{Angle} + \text{Dihedral} + \text{van der Waals} + \text{Electrostatic}^{1,2,3}$

Bonded interactions



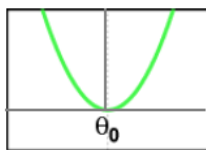
**Bond**

$$\sum_i^{\text{bonds}} K_{b,i} (b_i - b_{0,i})^2$$



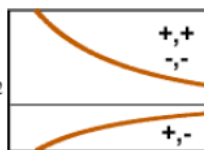
**van der Waals**

$$\sum_{\text{pairs } i,j} \left[ \epsilon_{ij} \left( \frac{r_{0,ij}}{r_{ij}} \right)^{12} - 2\epsilon_{ij} \left( \frac{r_{0,ij}}{r_{ij}} \right)^6 \right]$$



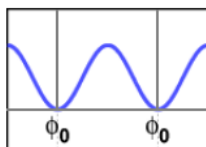
**Angle**

$$\sum_i^{\text{bond angles}} K_{\theta,i} (\theta_i - \theta_{0,i})^2$$



**Electrostatic**

$$332 \sum_{\text{pairs } i,j} \left( \frac{q_i q_j}{r_{ij}} \right)$$



**Dihedral**

$$\sum_i^{\text{torsion angles}} K_{\phi,i} \{1 - \cos[n_i(\phi_i - \phi_{0,i})]\}$$

Nonbonded interactions

1. Levitt M. Hirshberg M. Sharon R. Daggett V. Comp. Phys. Comm. (1995) 91: 215-231
2. Levitt M. *et al.* J. Phys. Chem. B (1997) 101: 5051-5061
3. Dynameomics: Protein Mechanics, Folding and Unfolding through Large Scale All-Atom Molecular Dynamics Simulations. David A. C. Beck. Valerie Daggett Research Group

# $\alpha$ -Complexes for Rotating $\mathbb{R}^2$ -Points

- Subset of points rotates around common point with the same circular velocity.
- Flip and radius events require solutions of second degree polynomials

# $\alpha$ -Complexes for Rotating $R^3$ -Points

- Subset of points rotates around common axis with the same circular velocity.
- Flip and radius events require solutions of fourth degree polynomials.
- Radius events for tetrahedra with 2 rotating and two stationary points still refuses to be solved analytically.