

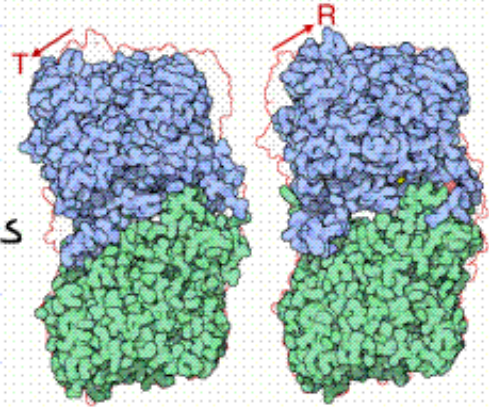
GCMB'07 Graph Rigidity for Modeling Allostery

Note Title

4/10/2007

Molecular Allostery
Glycogen phosphorylase

First-order rigidity of graphs
Laman conditions [L70]
Pebble games [JH95, 97]

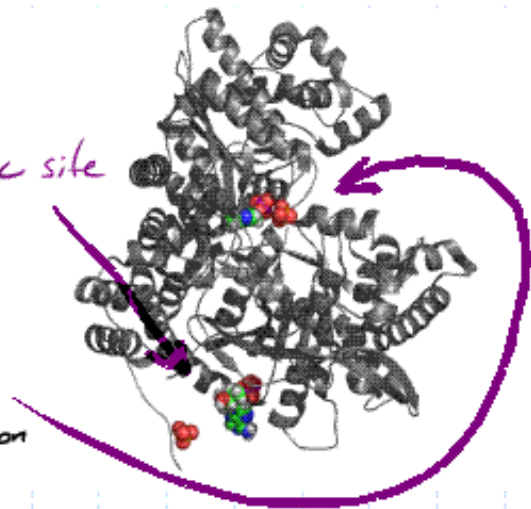


Applying to GP
Hydrogen bond strength

http://www.rcsb.org/pdb/static.do?p=education/discussion/molecule_of_the_month/pdb24_1.html

Allostery

Binding a molecule at an *allosteric site*
affects the ability to bind
at another *catalytic site*



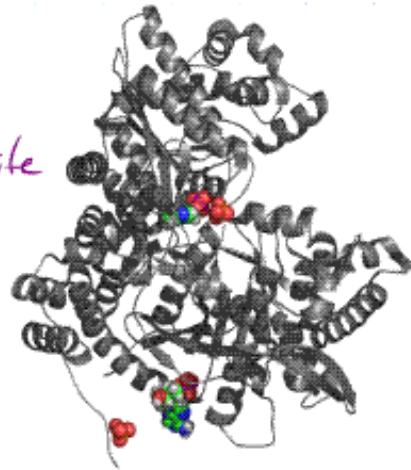
Is there a mechanical transmission
that can explain allostery?

A 2-d example:
struts transmit
1 degree of freedom



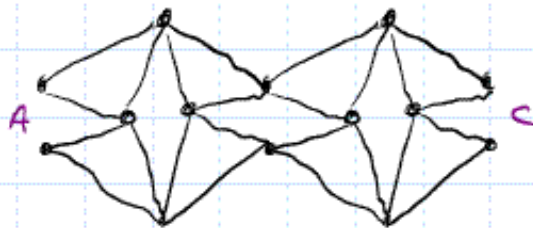
Allostery

Binding a molecule at an **allosteric site** affects the ability to bind at another **catalytic site**



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First-order Rigidity of Graphs

Let G be a graph with a straight-line embedding into \mathbb{R}^d with vertices in general position.

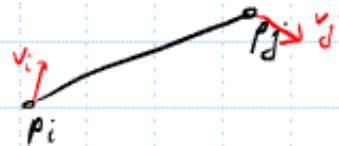
Consider assigning a vector v_i to each vertex i such that $\forall (p_i, p_j) \in E, (p_i - p_j) \cdot (v_i - v_j) = 0$

Preserve length

$$d(p_i + \epsilon v_i, p_j + \epsilon v_j) = d(p_i, p_j)$$

$$\begin{aligned} (p_i - p_j + \epsilon(v_i - v_j)) \cdot (p_i - p_j + \epsilon(v_i - v_j)) &= (p_i - p_j) \cdot (p_i - p_j) \\ &= \cancel{(p_i - p_j) \cdot (p_i - p_j)} + 2\epsilon(p_i - p_j) \cdot (v_i - v_j) + \epsilon^2 \cancel{(v_i - v_j) \cdot (v_i - v_j)} \end{aligned}$$

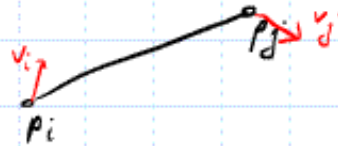
$$(p_i - p_j) \cdot (v_i - v_j) = 0$$



First-order Rigidity of Graphs

Let G be a graph with a straight-line embedding into \mathbb{R}^d with vertices in general position.

Consider assigning a vector v_i to each vertex i such that $\forall (p_i, p_j) \in E, (p_i - p_j) \cdot (v_i - v_j) = 0$



G is rigid iff the only assignments are rigid motions of \mathbb{R}^d .

Actually, the rigidity of generic embeddings is a property of the graph - e.g. all triangles are rigid.

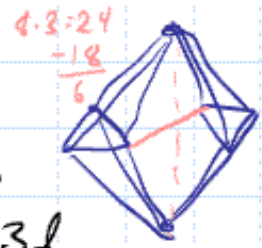
Degree of Freedom Counting

For a graph G with v vertices & e edges

| | 2d | 3d |
|--------------------|------------------------|------------------------|
| Vertices have | $2v$ dof | $3v$ dof |
| Edges remove | $- e$ dof | $- e$ dof |
| \mathbb{R}^d has | 2 dof | 3 dof |
| | $2v - e = 3$ for rigid | $3v - e = 6$ for rigid |

Degree of Freedom Counting

For a graph G with v vertices & e edges



| | | |
|---------------|-------------------------|-----------|
| | 2d | 3d |
| Vertices have | $2v$ dof | $3v$ dof |
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| R^d has | 3 dof | 6 dof |
| | $9 = 12 - 3 \checkmark$ | |



Laman [70] G is minimally rigid in 2d iff
 $e = 2v - 3$ and any vertex-induced subgraph
 (V', E') with $|V'| \geq 2$ satisfies $|E'| \leq 2|V'| - 3$.

Equivalent condition in R^3 , $e \leq 3v - 6$, is necessary, but not sufficient.

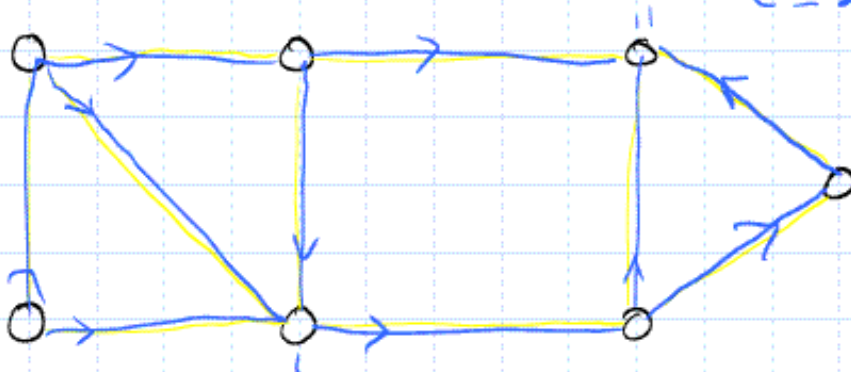
A Pebble Game to verify Laman's condition (2D)

Every vertex has two pebbles it can place on edges

Every edge has a pebble.

To add edge (i, j) , first draw four pebbles to p_i, p_j .

$$e \leq 2v - 3$$

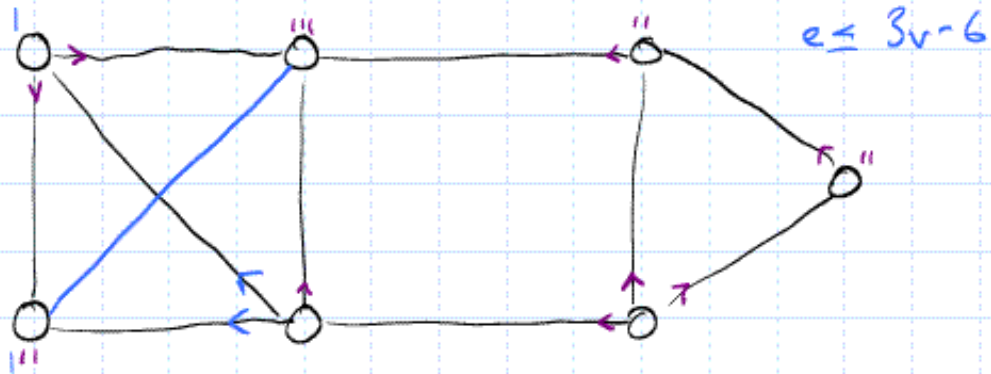


A Pebble Game to verify Laman's condition (3D)

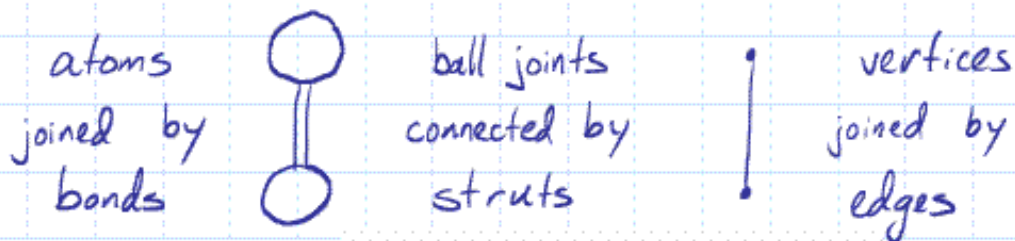
Every vertex has ~~two~~^{three} pebbles it can place on edges

Every edge has a pebble.

To add edge (i, j) , first draw ~~four~~^{six} pebbles to p_i, p_j , and one to each neighbor of p_i, p_j in turn



Allostery: Molecules \rightarrow rigid frameworks

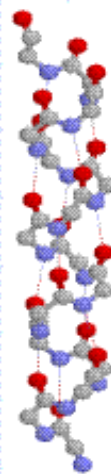
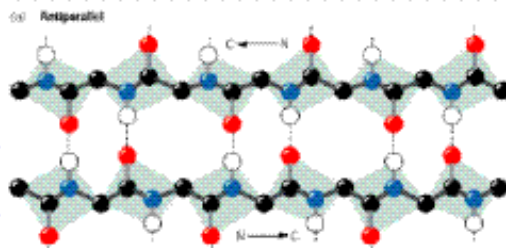
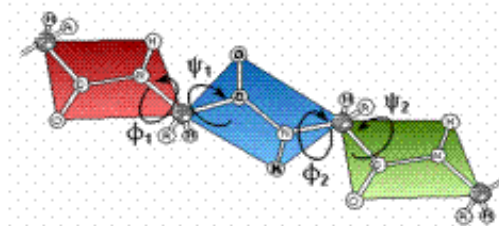


strong
peptide bonds
- angle rotations, so
add struts

disulfide bonds

hydrogen bonds

weak



Testing Allostery

1. Add edges for bonds to G
2. Test dof at catalytic site #
3. Add bonds to allosteric site A
4. Re-test dof at catalytic site #



Do this for all bond strengths

