

Note Title

# Problem set Cyclic Coord Descent

5/2/2007

Given ICDA residues 34-46 form loop

1. extract bond length, bond angle, & dihedral angle parameters from given coordinates
2. Choose random  $\phi, \psi$  dihedrals for loop,
3. Construct coords from parameters - note this breaks the loop at C46
4. Use Cyclic Coord Descent on loop  $\phi, \psi$ s to bring C46 back to correct position
  - adjust each angle in turn  $\phi_{34}, \psi_{34}, \phi_{35}, \psi_{35} \dots \phi_{46}, \psi_{46}$
  - bring C46 as close to position as possible by changing  $\phi_i$  (or  $\psi_i$ ).
5. Repeat from 2 to get 10 loop conformations.

## Coordinates to parameters

Bond length  $ab$   $\|a-b\| = \sqrt{(a_x-b_x)^2 + (a_y-b_y)^2 + (a_z-b_z)^2}$   
 MATLAB  $\text{norm}(a-b) = \text{sqr}(\text{sum}((a-b).^2, 2))$   
 $\uparrow$  works if  $a, b$  are  $n \times 3$  vectors.

Math review for 3-d vector geometry:

dot product  $u \cdot v = u_x v_x + u_y v_y + u_z v_z$

when  $u$  is a unit vector, i.e.  $\|u\| = \sqrt{u \cdot u} = 1$

$u \cdot v$  = the length of the projection of  $v$  onto  $u$ .

Thus if  $v$  is perpendicular to  $u$ ,  $u \cdot v = 0$ .

We can use perpendicular unit vectors  $u, v$  to give coordinates on the  $uv$  plane.

E.g. here is the angle to the projection of  $w$ :

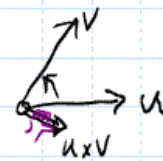
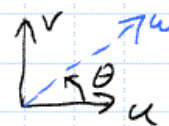
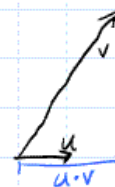
$$\theta = \text{atan2}(w \cdot v, w \cdot u)$$

cross product  $u \times v = (u_y v_z - u_z v_y, u_z v_x - u_x v_z, u_x v_y - u_y v_x)$

gives perpendicular vector in 3d. (0 if  $u, v$  colinear!)

order matters - curl fingers of right hand from  $u$  to  $v$

Don't forget to divide by length if you want to use for coordinates.



Bond angles & dihedrals will be most easily constructed if we agree on a standard coordinate frame:

For atom  $b$  in the sequence  $a \rightarrow b \rightarrow c$ ,  
 let  $Z = \frac{b-a}{\|b-a\|}$ , the unit vector  $a \rightarrow b$   
 $X = \frac{(a-b) \times (c-b)}{\|(a-b) \times (c-b)\|}$ , the unit vector  $\perp$  to plane  $abc$   
 Note: choose any  $\perp$  to  $Z$  if  $a, b, c$  colinear.  
 $Y = \frac{Z \times X}{\|Z \times X\|}$   $\leftarrow$  should already be 1.

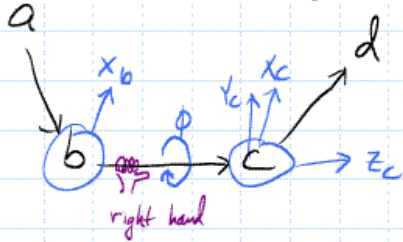
Result of this choice:

Bond angle: angle to rotate  $a$  around  $X$  to match  $c$ .

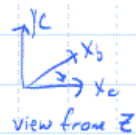


Since  $c$  is in the  $YZ$  plane,  
 $\theta = \pi - \text{atan2}((c-b) \cdot Y, (c-b) \cdot Z)$

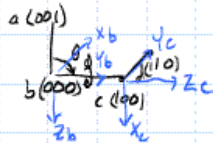
Dihedral angle: angle to rotate  $abc$  plane around  $bc$  to match  $bcd$  plane.



$\equiv$  Rotate  $X_b$  to  $X_c$ , since both are  $\perp$  to  $bc$   
 $\phi = -\text{atan2}(X_b \cdot Y_c, X_b \cdot X_c)$



Check signs for a small example.



$$\theta = \pi - \text{atan}(1, 0) = \frac{\pi}{2} \checkmark$$

$$\phi = -\text{atan}(1, 0) = -\frac{\pi}{2} \checkmark$$

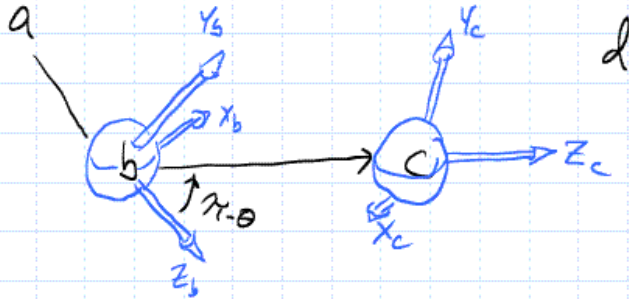
Let's write our coordinate frame  $(p, X, Y, Z)$  as four 4-element vectors:

$$\text{frame} = \begin{bmatrix} X_x & Y_x & Z_x & P_x \\ X_y & Y_y & Z_y & P_y \\ X_z & Y_z & Z_z & P_z \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

vectors
atom coordinates

Doing this lets us write both rotations & translations as  $4 \times 4$  matrices.  
 R. frame    T. frame

To get coordinates from parameters we can transform coordinate frames. Assume  $b$  is  $(0,0,0)$ , the origin



See also Kavraki's

<http://cnx.org/content/m11621/latest/>

$R_x$     first    Rotate around  $X$     by  $\theta - \pi$     to move  $Z_b$  to  $Z_c$   
 $R_z$     second    Rotate around  $Z$     by  $\phi$     to move  $X_b$  onto  $X_c$   
 $T_z$     finally    Translate along  $Z$     by  $-\text{length}(bc)$     to move  $c$  to origin.

$$R_x = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & \cos(\theta - \pi) & -\sin(\theta - \pi) & 0 \\ 0 & \sin(\theta - \pi) & \cos(\theta - \pi) & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$$R_z = \begin{bmatrix} \cos \phi & -\sin \phi & 0 & 0 \\ \sin \phi & \cos \phi & 0 & 0 \\ 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$$T_z = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & 1 & -\text{length}(bc) \\ 0 & 0 & 0 & 1 \end{bmatrix}$$

$\therefore$  new frame =  $T_z \cdot R_z \cdot R_x \cdot \text{frame}$

← note order  
 this is a single  $4 \times 4$  matrix,  
 since matrix multiplication  
 is associative.

Each bond gives us a transformation defined by its parameters.  
 12 residue loop  $\Rightarrow$  36 backbone bonds  $\Rightarrow$  36  $4 \times 4$  matrices.