

Smilgies: Molecular Axes and Planes as an Eigenvalue Problem (J Appl Cryst)

Supplementary Information

MathCAD code is replicated as it appears on the MathCAD graphical user interface

A copy of the MathCAD program can be obtained from the author on request (dms79@cornell.edu)

Pentacene, based on Campbell et al., Acta Cryst. (1961). 14, 705-711.

table of irreducible atom positions

$$R1 := \begin{pmatrix} 0.1484 & 0.0183 & 0.4071 \\ 0.1707 & 0.1361 & 0.3491 \\ 0.103 & 0.0604 & 0.2508 \\ 0.1274 & 0.165 & 0.1894 \\ 0.0572 & 0.0951 & 0.095 \\ 0.0925 & 0.2095 & 0.0335 \\ -0.0227 & -0.1227 & 0.0567 \\ -0.054 & -0.2325 & 0.1225 \\ 0.015 & -0.1516 & 0.2149 \\ -0.0152 & -0.2634 & 0.2786 \\ 0.0519 & -0.18 & 0.3706 \end{pmatrix} \quad R2 := \begin{pmatrix} 0.5627 & 0.5733 & 0.4014 \\ 0.5035 & 0.6805 & 0.3286 \\ 0.521 & 0.5876 & 0.2434 \\ 0.4582 & 0.701 & 0.1686 \\ 0.481 & 0.6062 & 0.0846 \\ 0.4231 & 0.7138 & 0.01 \\ 0.5622 & 0.3794 & 0.0754 \\ 0.6103 & 0.2743 & 0.1483 \\ 0.6061 & 0.3652 & 0.2342 \\ 0.6549 & 0.257 & 0.3094 \\ 0.6496 & 0.3487 & 0.3925 \end{pmatrix}$$

rows(R1) = 11

rows(R2) = 11

Complete molecule by adding symmetry-related positions (space group P-1)

$$\text{N} := \text{rows}(\text{R1}) \quad i := 0..N - 1$$

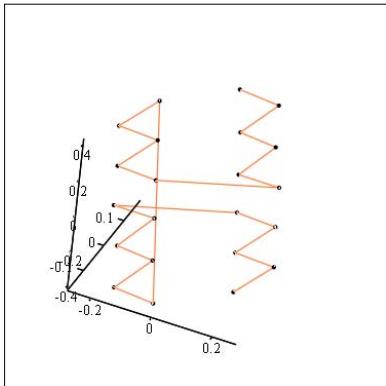
1st molecule

$$\begin{aligned} X1_i &:= (R1^{(0)})_i & Y1_i &:= (R1^{(1)})_i & Z1_i &:= (R1^{(2)})_i \\ X1_{N+i} &:= -X1_i & Y1_{N+i} &:= -Y1_i & Z1_{N+i} &:= -Z1_i \end{aligned}$$

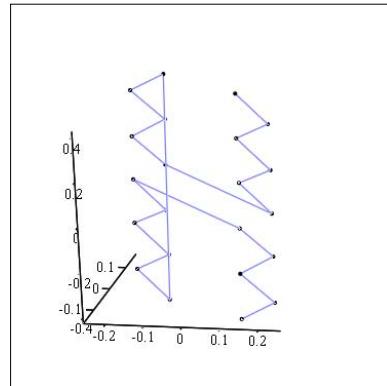
2nd molecule at Wyckoff position (0.5,0.5) was moved to the origin first

$$\begin{aligned} X2_i &:= (R2^{(0)})_i - 0.5 & Y2_i &:= (R2^{(1)})_i - 0.5 & Z2_i &:= (R2^{(2)})_i \\ X2_{N+i} &:= (-X2)_i & Y2_{N+i} &:= (-Y2)_i & Z2_{N+i} &:= -Z2_i \end{aligned}$$

graphical check



(X1, Y1, Z1)



(X2, Y2, Z2)

table of coordinates for complete molecule 1 and 2

$$\begin{aligned} \text{R1} &:= \text{augment}(\text{X1}, \text{augment}(\text{Y1}, \text{Z1})) & \text{rows}(\text{R1}) &= 22 \\ \text{R2} &:= \text{augment}(\text{X2}, \text{augment}(\text{Y2}, \text{Z2})) & \text{rows}(\text{R2}) &= 22 \end{aligned}$$

lattice parameters (from Campbell et al., erratum, Acta Cryst. (1962). 15, 289-290.)

$$a := 7.90 \quad b := 6.06 \quad c := 16.01 \quad \alpha := 101.9\text{-deg} \quad \beta := 112.6\text{-deg} \quad \gamma := 85.8\text{-deg}$$

reciprocal lattice (including factor of 2 PI)

$$V_{\text{rec}} := a \cdot b \cdot c \cdot \sqrt{1 + 2 \cdot \cos(\alpha) \cdot \cos(\beta) \cdot \cos(\gamma) - \cos(\alpha)^2 - \cos(\beta)^2 - \cos(\gamma)^2}$$

$$\text{as} := \frac{2 \cdot \pi \cdot b \cdot c \cdot \sin(\alpha)}{V} \quad \text{bs} := \frac{2 \cdot \pi \cdot c \cdot a \cdot \sin(\beta)}{V} \quad \text{cs} := \frac{2 \cdot \pi \cdot a \cdot b \cdot \sin(\gamma)}{V}$$

$$\gamma_s := \arccos\left(\frac{\cos(\alpha) \cdot \cos(\beta) - \cos(\gamma)}{|\sin(\alpha) \cdot \sin(\beta)|}\right)$$

$$\alpha_s := \arccos\left(\frac{\cos(\gamma) \cdot \cos(\beta) - \cos(\alpha)}{|\sin(\gamma) \cdot \sin(\beta)|}\right)$$

$$\beta_s := \arccos\left(\frac{\cos(\alpha) \cdot \cos(\gamma) - \cos(\beta)}{|\sin(\alpha) \cdot \sin(\gamma)|}\right)$$

numerical values

$$\begin{aligned} \text{as} &= 0.862 & \text{bs} &= 1.06 & \text{cs} &= 0.433 & d001 &:= \frac{2 \cdot \pi}{\text{cs}} & d001 &= 14.502 \\ \alpha_s &= 78.849 \text{ deg} & \beta_s &= 67.77 \text{ deg} & \gamma_s &= 89.619 \text{ deg} \end{aligned}$$

crystallographic B matrix (including 2 PI)

$$\text{BMat} := \begin{pmatrix} a & b \cdot \cos(\gamma) & c \cdot \cos(\beta) \\ 0 & b \cdot \sin(\gamma) & -c \cdot \sin(\beta) \cdot \cos(\alpha s) \\ 0 & 0 & \frac{2 \cdot \pi}{\text{cs}} \end{pmatrix} \quad \text{BMat} = \begin{pmatrix} 7.9 & 0.444 & -6.153 \\ 0 & 6.044 & -2.858 \\ 0 & 0 & 14.502 \end{pmatrix}$$

3D lattice vectors of pentacene unit cell

$$A_{\text{rec}} := \text{BMat}^{\langle 0 \rangle} \quad B_{\text{rec}} := \text{BMat}^{\langle 1 \rangle} \quad C_{\text{rec}} := \text{BMat}^{\langle 2 \rangle}$$

checks

$$|A| = 7.9 \quad a = 7.9 \quad |B| = 6.06 \quad b = 6.06 \quad |C| = 16.01 \quad c = 16.01$$

$$\arccos\left(\frac{B \cdot C}{b \cdot c}\right) = 101.9 \text{ deg} \quad \arccos\left(\frac{A \cdot C}{a \cdot c}\right) = 112.6 \text{ deg} \quad \arccos\left(\frac{A \cdot B}{a \cdot b}\right) = 85.8 \text{ deg}$$

$$\alpha = 101.9 \text{ deg} \quad \beta = 112.6 \text{ deg} \quad \gamma = 85.8 \text{ deg}$$

cartesian coordinates for molecule 1 and 2

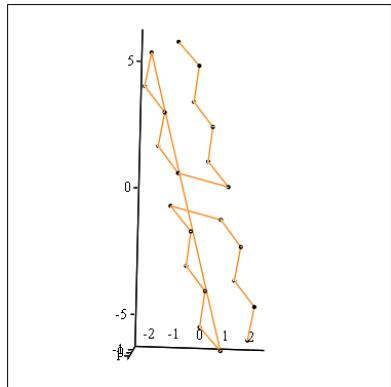
$$r1 := R1 \cdot BMat^T \quad N_{\text{rows}} := \text{rows}(R1) \quad N = 22$$

$$x1 := r1^{(0)} \quad y1 := r1^{(1)} \quad z1 := r1^{(2)}$$

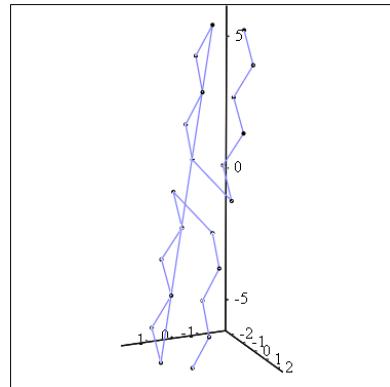
$$r2 := R2 \cdot BMat^T \quad N_{\text{rows}} := \text{rows}(R2) \quad N = 22$$

$$x2 := r2^{(0)} \quad y2 := r2^{(1)} \quad z2 := r2^{(2)}$$

graphical check



(x1, y1, z1)



(x2, y2, z2)

covariance matrices for molecule 1 and 2

$$\text{CoV1} := \frac{r1^T \cdot r1}{N}$$

$$\text{CoV2} := \frac{r2^T \cdot r2}{N}$$

$$\text{CoV1} = \begin{pmatrix} 1.309 & 1.454 & -3.569 \\ 1.454 & 1.772 & -3.27 \\ -3.569 & -3.27 & 12.921 \end{pmatrix}$$

$$\text{CoV2} = \begin{pmatrix} 1.293 & 0.467 & -3.862 \\ 0.467 & 1.71 & -3.051 \\ -3.862 & -3.051 & 13.312 \end{pmatrix}$$

Molecular main axes by diagonalization of covariance matrix

- > maximum variance: long molecular axis
- > minimum variance: normal vector of molecular plane

$$\text{var1} := \text{eigenvals}(\text{CoV1}) \quad \text{var1} = \begin{pmatrix} 1.148 \times 10^{-3} \\ 1.113 \\ 14.888 \end{pmatrix} \quad \min(\text{var1}) = 1.148 \times 10^{-3}$$

$$\text{var2} := \text{eigenvals}(\text{CoV2}) \quad \text{var2} = \begin{pmatrix} 1.172 \\ 6.21 \times 10^{-4} \\ 15.144 \end{pmatrix} \quad \min(\text{var2}) = 6.21 \times 10^{-4}$$

$$\max(\text{var1}) = 14.888$$

$$\max(\text{var2}) = 15.144$$

3D vectors of molecular axes as eigenvectors associated with the maximum variance

$$\text{MA1} := \text{eigenvec}(\text{CoV1}, \max(\text{var1})) \quad \text{MA1} = \begin{pmatrix} -0.271 \\ -0.261 \\ 0.926 \end{pmatrix} \quad |\text{MA1}| = 1$$

$$\text{MA2} := \text{eigenvec}(\text{CoV2}, \max(\text{var2})) \quad \text{MA2} = \begin{pmatrix} -0.269 \\ -0.222 \\ 0.937 \end{pmatrix} \quad |\text{MA2}| = 1$$

3D vectors of molecular planes as eigenvector associated with the minimum variance

$$\text{MP1} := -\text{eigenvec}(\text{CoV1}, \min(\text{var1})) \quad \text{MP1} = \begin{pmatrix} 0.858 \\ -0.502 \\ 0.11 \end{pmatrix} \quad |\text{MP1}| = 1$$

$$\text{MP2} := \text{eigenvec}(\text{CoV2}, \min(\text{var2})) \quad \text{MP2} = \begin{pmatrix} 0.871 \\ 0.36 \\ 0.335 \end{pmatrix} \quad |\text{MP2}| = 1$$

check of orthogonality

$$\text{MA1} \cdot \text{MP1} = 0 \quad \text{MA2} \cdot \text{MP2} = 0$$

Numerical results, part 1

Intermolecular angles

- angle between molecular axes

$$\delta_{12} := |\cos(\mathbf{MA1} \cdot \mathbf{MA2})| \quad \delta_{12} = 2.311 \text{ deg}$$

- angle between molecular planes (herring bone angle)

$$\varepsilon_{12} := |\cos(\mathbf{MP1} \cdot \mathbf{MP2})| \quad \varepsilon_{12} = 52.884 \text{ deg}$$

Angles between molecular axes and planes and the 3D lattice vectors

- angle of molecular axes with C lattice vector

$$\rho_1 := \cos\left(\frac{\mathbf{C}}{|\mathbf{C}|} \cdot \mathbf{MA1}\right) \quad \rho_2 := \cos\left(\frac{\mathbf{C}}{|\mathbf{C}|} \cdot \mathbf{MA2}\right)$$

$$\rho_1 = 8.103 \text{ deg} \quad \rho_2 = 7.305 \text{ deg}$$

- angle of molecular plane with A lattice vector

$$\sigma_1 := \sin\left(\frac{\mathbf{A}}{|\mathbf{A}|} \cdot \mathbf{MP1}\right) \quad \sigma_2 := \sin\left(\frac{\mathbf{A}}{|\mathbf{A}|} \cdot \mathbf{MP2}\right)$$

$$\sigma_1 = 59.096 \text{ deg} \quad \sigma_2 = 60.543 \text{ deg}$$

Transformation to Eulerian angles

- rotation of molecular axes from cartesian x-axis

$$\theta_1 := \text{atan2}(\text{MA1}_0, \text{MA1}_1) \quad \theta_1 = -136.12 \text{ deg}$$

$$\theta_2 := \text{atan2}(\text{MA2}_0, \text{MA2}_1) \quad \theta_2 = -140.417 \text{ deg}$$

- rotation of molecular axis into cartesian xz plane

$$Rz(\phi) := \begin{pmatrix} \cos(\phi) & -\sin(\phi) & 0 \\ \sin(\phi) & \cos(\phi) & 0 \\ 0 & 0 & 1 \end{pmatrix}$$

$$RMA1 := Rz(-\theta_1) \cdot MA1 \quad RMA1 = \begin{pmatrix} 0.377 \\ 0 \\ 0.926 \end{pmatrix}$$

$$RMA2 := Rz(-\theta_2) \cdot MA2 \quad RMA2 = \begin{pmatrix} 0.349 \\ 0 \\ 0.937 \end{pmatrix}$$

- tilt of molecular axis

$$\chi_1 := \text{atan}\left(\frac{RMA1_0}{RMA1_2}\right) \quad \chi_1 = 22.121 \text{ deg}$$

$$\chi_2 := \text{atan}\left(\frac{RMA2_0}{RMA2_2}\right) \quad \chi_2 = 20.413 \text{ deg}$$

- check: undo tilt of molecular axis

$$Ry(\chi) := \begin{pmatrix} \cos(\chi) & 0 & \sin(\chi) \\ 0 & 1 & 0 \\ -\sin(\chi) & 0 & \cos(\chi) \end{pmatrix}$$

$$RRMA1 := Ry(-\chi_1) \cdot RMA1 \quad RRMA1 = \begin{pmatrix} 0 \\ 0 \\ 1 \end{pmatrix}$$

$$RRMA2 := Ry(-\chi_2) \cdot RMA2 \quad RRMA2 = \begin{pmatrix} 0 \\ 0 \\ 1 \end{pmatrix}$$

-> all information from molecular axis is extracted

- rotate molecular planes with rotations determined from molecular axes

$$\text{RRMP1} := \text{Ry}(-\chi_1) \cdot \text{Rz}(-\theta_1) \cdot \text{MP1} \quad \text{RRMP1} = \begin{pmatrix} -0.292 \\ 0.956 \\ 0 \end{pmatrix} \quad \text{RRMP1} \cdot \text{RRMA1} = 0$$

$$\text{RRMP2} := \text{Ry}(-\chi_2) \cdot \text{Rz}(-\theta_2) \cdot \text{MP2} \quad \text{RRMP2} = \begin{pmatrix} -0.961 \\ 0.277 \\ 0 \end{pmatrix} \quad \text{RRMP2} \cdot \text{RRMA2} = 0$$

- azimuth angle of molecular planes with regard to x-axis

-> for the molecular plane to lie within the xz-plane,
the normal vector MP points along the y-direction

check

$$\phi_1 := \text{atan2}(\text{RRMP1}_0, \text{RRMP1}_1) - \frac{\pi}{2} \quad \phi_1 = 16.992 \text{ deg}$$

$$\text{Rz}(-\phi_1) \cdot \text{RRMP1} = \begin{pmatrix} 0 \\ 1 \\ 0 \end{pmatrix}$$

$$\phi_2 := \text{atan2}(\text{RRMP2}_0, \text{RRMP2}_1) - \frac{\pi}{2} \quad \phi_2 = 73.893 \text{ deg}$$

$$\text{Rz}(-\phi_2) \cdot \text{RRMP2} = \begin{pmatrix} 0 \\ 1 \\ 0 \end{pmatrix}$$

Numerical results, part 2

- molecule 1: values of the Eulerian angles

$$\phi_1 = 16.992 \text{ deg} \quad \chi_1 = 22.121 \text{ deg} \quad \theta_1 = -136.12 \text{ deg}$$

- check

$$\text{Rz}(-\phi_1) \cdot \text{Ry}(-\chi_1) \cdot \text{Rz}(-\theta_1) \cdot \text{MA1} = \begin{pmatrix} 0 \\ 0 \\ 1 \end{pmatrix} \quad \text{Rz}(-\phi_1) \cdot \text{Ry}(-\chi_1) \cdot \text{Rz}(-\theta_1) \cdot \text{MP1} = \begin{pmatrix} 0 \\ 1 \\ 0 \end{pmatrix}$$

- molecule 2: values of the Eulerian angles

$$\phi_2 = 73.893 \text{ deg} \quad \chi_2 = 20.413 \text{ deg} \quad \theta_2 = -140.417 \text{ deg}$$

- check

$$\text{Rz}(-\phi_2) \cdot \text{Ry}(-\chi_2) \cdot \text{Rz}(-\theta_2) \cdot \text{MA2} = \begin{pmatrix} 0 \\ 0 \\ 1 \end{pmatrix} \quad \text{Rz}(-\phi_2) \cdot \text{Ry}(-\chi_2) \cdot \text{Rz}(-\theta_2) \cdot \text{MP2} = \begin{pmatrix} 0 \\ 1 \\ 0 \end{pmatrix}$$

molecules rotated to the main plane (cartesian xz plane)

$$Rr1 := \left(Rz(-\phi1) \cdot Ry(-\chi1) \cdot Rz(-\theta1) \cdot r1^T \right)^T \quad Rr2 := \left(Rz(-\phi2) \cdot Ry(-\chi2) \cdot Rz(-\theta2) \cdot r2^T \right)^T$$

	0	1	2
0	-0.68	0.042	6.103
1	-1.356	0.011	4.936
2	-0.713	-0.026	3.652
3	-1.328	0	2.449
4	-0.707	-0.078	1.222
5	-1.41	-0.004	-0.023
6	0.703	0.044	1.156
Rr1 =	7	1.368	-0.025
	8	0.694	0.02
	9	1.365	-0.031
	10	0.686	-0.005
	11	0.68	-0.042
	12	1.356	-0.011
	13	0.713	0.026
	14	1.328	-0
	15	0.707	0.078
			-1.222

	0	1	2
0	-0.725	0.006	6.134
1	-1.387	-0.015	4.947
2	-0.723	-0.003	3.692
3	-1.427	-0.03	2.472
Rr2 =	4	-0.738	0.012
	5	-1.395	0.003
	6	0.739	0.004
	7	1.354	-0.046
	8	0.743	0.027
	9	1.374	-0.029
	10	0.759	0.042
	11	0.725	-0.006
	12	1.387	0.015
	13	0.723	0.003
	14	1.427	0.03
	15	0.738	-0.012
			-1.228

graphical check

$$Rx1 := Rr1^{<0>}$$

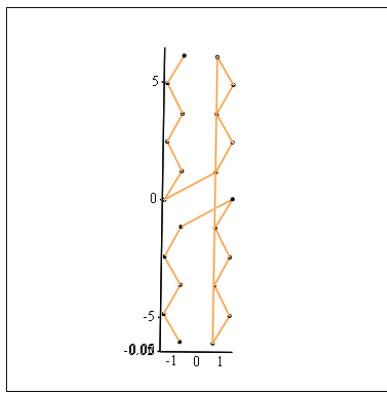
$$Ry1 := Rr1^{<1>}$$

$$Rz1 := Rr1^{<2>}$$

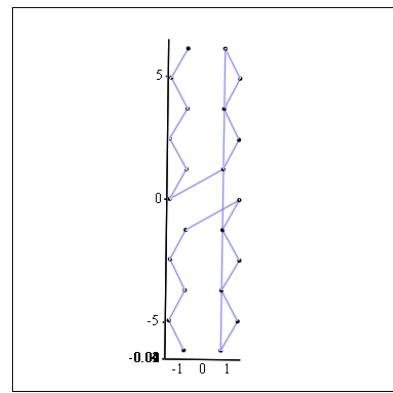
$$Rx2 := Rr2^{<0>}$$

$$Ry2 := Rr2^{<1>}$$

$$Rz2 := Rr2^{<2>}$$



(Rx1, Ry1, Rz1)



(Rx2, Ry2, Rz2)