

MATH 728A, BIOMOLECULAR GEOMETRY, HOMEWORK #1

Problem 1 (Methane). Consider an idealized molecule of methane CH_4 where the positions of the atoms are $C(1, 1, 1)$, $H_1(0, 0, 0)$, $H_2(2, 2, 0)$, $H_3(2, 0, 2)$, $H_4(0, 2, 2)$. Each of the hydrogen atoms is bonded to the carbon atom. Show that the four hydrogens are at the vertices of a regular tetrahedron, and the carbon is at its center. Compute the following.

- (1) The atom/bond tree $\Gamma^{(0)}$, where the edges are labelled with bond lengths.
- (2) A bond/angle tree $\Gamma^{(1)}$, where the vertices are the edges of $\Gamma^{(0)}$ and the edges (angles) are labelled with bond angles.
- (3) A triangle/wedge tree $\Gamma^{(2)}$, where the vertices are triangles corresponding to the edges of $\Gamma^{(1)}$ and the edges are (oriented) wedge angles.
- (4) For each edge in $\Gamma^{(2)}$, give the corresponding 3-simplex, and give the chosen orientation of that 3-simplex to match the assigned wedge angle.

Use this information to create this molecule in IMIMOL; export a MOPAC Z-matrix and a RASMOL label file, and visualize the molecule using RASMOL.

Problem 2 (Ethane). Consider an idealized molecule of ethane where the positions of the atoms are $C_1(1, 1, 1)$, $C_2(0, 0, 0)$, $H_1(2, 2, 0)$, $H_2(2, 0, 2)$, $H_3(0, 2, 2)$, $H_4(-1, -1, 1)$, $H_5(1, -1, -1)$, $H_6(-1, 1, -1)$. Compute the same four items as in Problem #1. Create this molecule in IMIMOL but change the CC bond length to 1.54 and all the CH bond lengths to 1.1. Draw the molecule using RASMOL.

Problem 3 (Stereo Viewing). Use cross-eyed stereo to look at the enclosed picture of cyclohexane. Rank order (approximately) the 18 atoms from closest to the viewer to farthest from the viewer.