

Chem 279 Final Presentation

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Overview

1. Introduction to Band Gap and Molecular Dipole
2. Refresher on CNDO/2
3. Extension to Energy Gap
4. Extension to Molecular Dipole
5. Results, Interpretations, and Next Steps

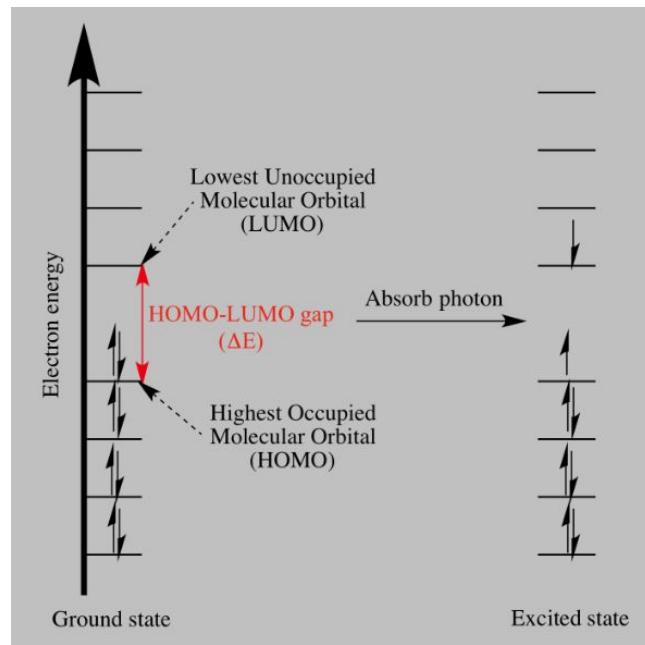


Band Gap and Molecular Dipole



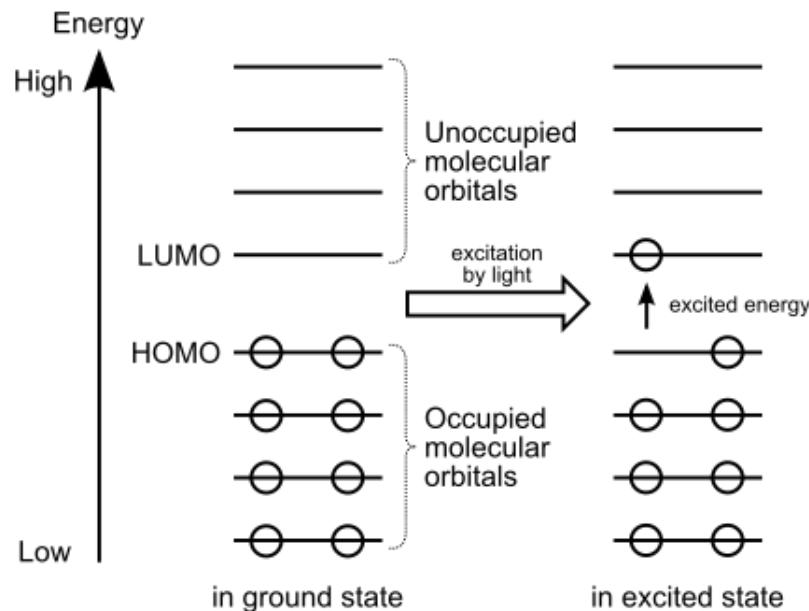
Energy Gap

- Energy difference between where electrons normally reside and where they are free to move
- HOMO (Highest Occupied Molecular Orbital)
 - Represents where the most weakly held electrons are.
 - Electrons in the HOMO are the first to participate in reactions (e.g. oxidation, nucleophilic).
 - A higher HOMO energy indicates electrons are easier to remove which means molecule is more reactive as an electron donor.



Energy Gap Continued

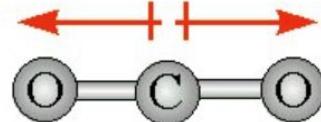
- LUMO (Lowest Unoccupied Molecular Orbital)
 - Represents the lowest available orbital electrons can jump into.
 - Determines how easily a molecule can accept electrons (reduction, electrophilic).
 - A lower LUMO energy indicates easier to add electrons which means better electron acceptor.
- Band Gap = LUMO – HOMO
 - Small Band Gap: Electrons can be excited easily → molecule is chemically reactive
 - Large Band Gap: Requires high energy to excite electrons → chemically inert



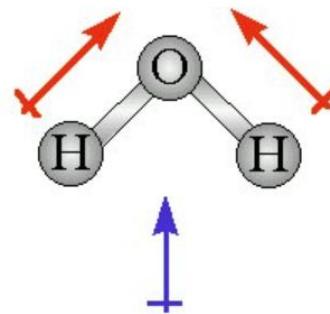
Molecular Dipole

- Separation of positive and negative charge in a molecule, creating a net polarity
 - Arises from polar bonds (unequal electron sharing) that don't completely cancel out due to molecule's asymmetrical shape
- Dipole-dipole interactions influence properties like solubility, boiling and melting point, and viscosity
 - Polar molecules are loosely attracted to each other because of slightly positive and negative regions

Dipoles



Overall
Dipole:
(none)



Refresher on CNDO/2

CNDO (Complete Neglect of Differential Overlap)

- What is CNDO/2?
 - Semi-empirical quantum chemistry method for approximating molecular electronic structure
 - Treats only valence electrons, reducing the basis size and computational effort
 - Uses STO 3G basis functions and parameterized atomic integrals
 - Produces a simplified Fock matrix that still captures essential bonding and charge distribution trends

CNDO/2 Fock Matrix Equations

1. Diagonal Fock Elements

$$F_{\mu\mu} = - \left(\frac{I_A + A_A}{2} \right) + (P_A - Z_A^*) \gamma_{AA} - \left(P_{\mu\mu}^\sigma - \frac{1}{2} \right) \gamma_{AA} + \sum_{C \neq A} (P_C - Z_C^*) \gamma_{AC}$$

2. Off-Diagonal Fock Elements

$$F_{\mu\nu} = \frac{1}{2} (\beta_A + \beta_B) S_{\mu\nu} - P_{\mu\nu}^\sigma \gamma_{AB} \quad (\mu \in A, \nu \in B, A \neq B)$$

3. Two-Center Coulomb Interaction

$$\gamma_{AB} = \iint \frac{\phi_A(\mathbf{r}_1)^2 \phi_B(\mathbf{r}_2)^2}{|\mathbf{r}_1 - \mathbf{r}_2|} d\mathbf{r}_1 d\mathbf{r}_2$$

Extension to Energy Gap

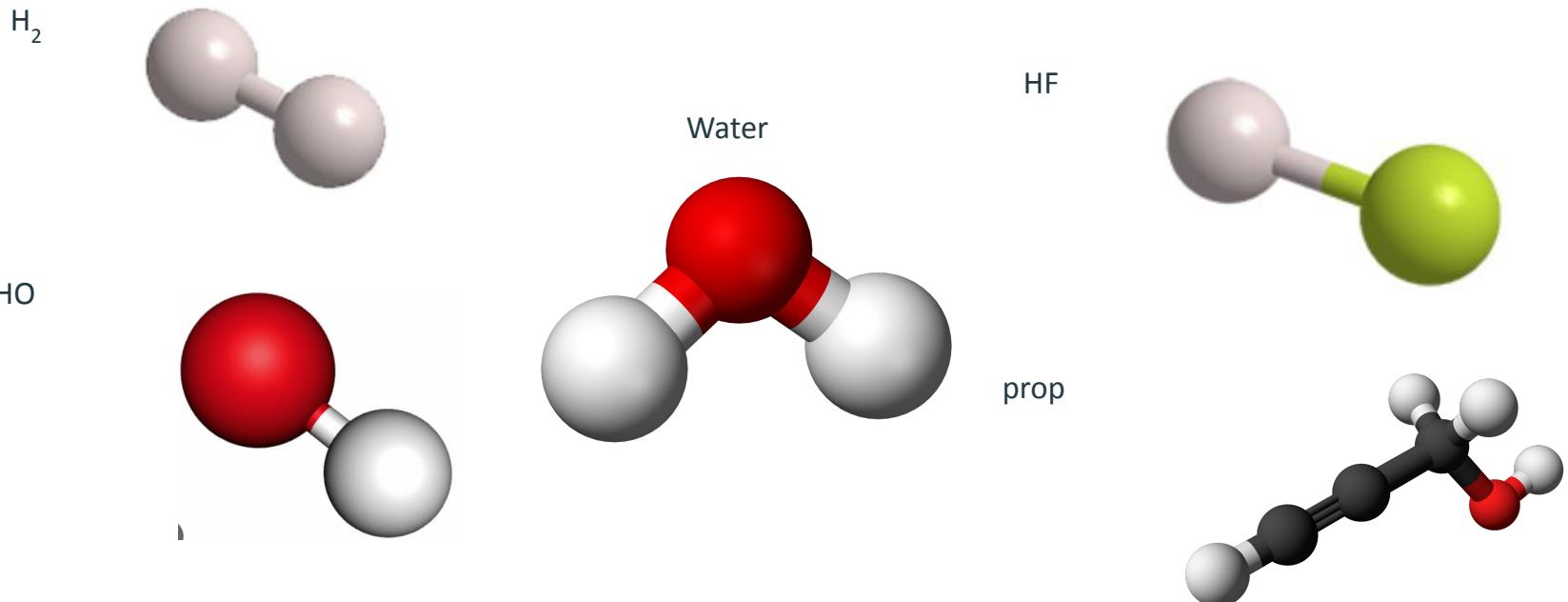
How We Computed Observables

- After SCF convergence, we diagonalize the final alpha and beta Fock matrices to get molecular orbital MO energies and coefficients.
- This produces two sets of MO energies because we use unrestricted Hartree Fock (alpha and beta spins treated separately).
- We sort the MO energies from lowest to highest.
- We determine which orbitals are occupied using the electron counts:
 - Fill the lowest num_alpha alpha-spin orbitals.
 - Fill the lowest num_beta beta-spin orbitals.
- The occupied orbitals are used to build the electron density matrix used in the SCF cycle.
- The eigenvectors give the shape of each MO; the eigenvalues are the MO energy levels.

Determining HOMO, LUMO, and the Energy Gap

- HOMO = the last filled orbital in the **sorted** list.
- LUMO = the next unfilled orbital after the HOMO.
 - HOMO alpha/beta = orbital index (`num_alpha/beta - 1`)
 - LUMO alpha/beta = orbital index (`num_alpha/beta`)
- Overall HOMO = the larger of HOMO alpha and HOMO beta energies.
- Overall LUMO = the smaller of LUMO alpha and LUMO beta energies.
- Energy Gap = Change in Energy
 - This gap indicates how easily an electron can be promoted to an excited state.

Molecular Systems We Tested On



Results

Molecule	Our HOMO (eV)	Our LUMO (eV)	Our Gap (eV)	Expected Gap (eV)
H ₂	-20.87	6.52	27.39	~15
H ₂ O	-18.51	15.19	33.70	~12
HF	-21.26	8.99	30.25	~16
HO	-18.87	3.37	22.24	~11
Propynol	86.23	91.33	5.10	~5-10

Interpretation and Discussion

- HOMO–LUMO gap is still a good proxy for qualitative behavior
- Even when not numerically correct:
 - Small HF gap → molecule is more reactive, easier to excite
 - Large HF gap → more stable, harder to excite
- HF overestimates the gap, but the trend/motifs across molecules is preserved.
- HF gives qualitatively correct trends
 - Even though HF overestimates the numerical gap, it still gets the ordering right:
 - Conjugated molecules → smaller gaps
 - Radicals → intermediate gaps
 - Closed shell molecules → large gaps



Extension to Molecular Dipole



Basics of Dipole Calculation

- Molecular dipole is a vector that sums up all the individual bond dipoles
- Need to calculate positive and negative contribution to the dipole
- To perform this calculation, we need:
 - AO dipole integrals – x, y, z components of $\langle \mu | \hat{r} | v \rangle$ for every AO pair – product center of each basis-function pair and the contracted STO-3G normalization factors
 - Final MO coefficients in orthonormal representation – MOs must be S-orthonormalized before building a reliable density since CNDO/2 uses non-orthonormalized AO's
 - Final AO density matrix – distribution of electric charge — “negative” part of the dipole
 - CNDO-consistent nuclear charges – Z* for consistency with valence-only electronic model – “positive” part
 - A way to combine the electronic and nuclear terms
 - Conversion to Debye units

Translating Theory to Implementation

1. Build AO Dipole Integrals
 - a. Evaluated and assembled D_x , D_y , D_z into full AO dipole matrices
2. Prepare SCF Quantities
 - a. S-orthonormalized final MOs to ensure $C^T S C = I$
 - b. Constructed final AO density matrix from occupied orthonormalized MOs.
3. Electronic Dipole
 - a. Contracted density with AO dipole matrices: $\mu_{elec} = \text{Tr}(PD_k)$ for $k=x,y,z$
4. Nuclear Dipole (CNDO Effective Charges)
 - a. Used CNDO effective valence charges Z_A^*
 - b. Computed nuclear component: $\mu_{nuc} = \sum_A Z_A^* R_A$
5. Total Dipole + Units
 - a. Combined components: $\mu_{total} = \mu_{nuc} - \mu_{elec}$
 - b. Converted from a.u. to Debye

Calculation Results

Molecule	Our Dipole (Debye)	Literature Dipole (Debye)	Approximate Error	Comments
H ₂	0	0	0 %	exactly zero dipole by symmetry
H ₂ O	1.15	1.85	18 %	Trend is correct
HF	1.62	1.83	10.2 %	Semi empirical HF tends to give slightly smaller dipoles
HO	1.47	1.66	10.24 %	order of magnitude error as HF - makes sense
Propynol	141.16	~2	Way off	Breakdown of the model for large complex molecules

Interpretation and Discussion

- CNDO/2 captures correct polarity trends, but slightly underestimates magnitudes.
- Why results are reasonable for small systems:
 - Geometry is compact, minimizing origin effects.
 - Polarization is limited so CNDO/2's approximations remain stable.
 - Electron distribution is simple and dominated by a single bond dipole (O–H, H–F).
 - For water, our model is close showing that it can handle small bent polar molecules reasonably well despite approximation.
- Propynol dipole model breakdown.
 - Geometry is long and nearly linear, giving a very large "lever arm" for charge separation.
 - CNDO/2 significantly over-polarizes extended π systems and strongly electronegative atoms (O).
 - Minimal basis set cannot describe electron delocalization properly this forces electrons toward one end.
 - Dipole moment becomes highly origin dependent when the molecule spans a large coordinate range.
 - **Semi empirical parameters were built for small, compact molecules, not long chain alcohols with complex bonds.**

Thank you