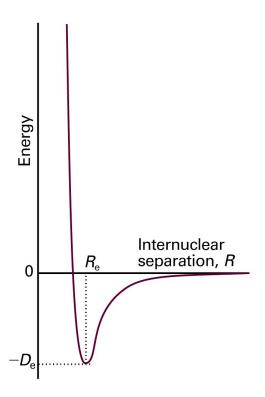
# CHAPTER 10 Molecular Structure

# I. Basic Concepts

- A. Born-Oppenheimer Approximation.
  - 1. All theories of molecular bonding make the same simplification to start the B.O. approximation
  - 2. Nuclei are sluggish relative to e<sup>-</sup> motion due to their masses. (Time scale separation)
  - 3. e cloud responds instantaneously to slowly changing nuclear positions.
  - 4. Separate problem into a) nuclear vibrations and b) electron wave function around fixed nuclear separation R.
  - 5. For  $H_2^+$ , 3 body problem  $\rightarrow$  1 body (1 e<sup>-</sup> moving around fixed nuclei)
  - 6. Potential energy curve for nuclear motion:
  - 7. Nuclei vibrate around  $R_e$ , the equilibrium bond Length
  - 8.  $D_o = bond dissociation energy$

$$D_e$$
 = well depth  
=  $D_o + \frac{1}{2} h_V$ 



# II. Valence Bond (VB) Theory

### A. Introduction.

- 1. First theory of the chemical bond, developed by Linus Pauling.
- 2. VB theory still largely affects the thinking of organic chemists.
- 3. Concepts such as spin pairing, orbital overlap, sigma and pi bonding, and hybridization carry over to all bonding theories, including Molecular Orbital Theory.
- 4. Bond forms when an electron in an atomic orbital on one atom pairs its spin with another electron in an atomic orbital on another atom.
- 5. 2 electrons = one bond
- B. Homonuclear Diatomics.
  - 1. Consider formation of the bond in  $H_2$  diatomic. Label the atoms A and B, each with one electron.
  - 2. At large distance  $\psi_{molecule} = A(1)B(2)$

where A() and B() are atomic orbitals, i.e. 1s.

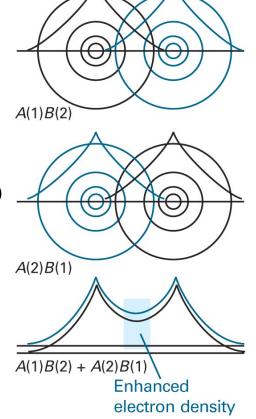
3. As atoms begin to interact at short distance, it is impossible to distinguish whether electron 1 is on A, or electron 2. An equally valid description would be:

$$\psi_{\text{molecule}} = A(2)B(1)$$

4. The true wavefunction is a superposition of these two possibilities, so in QM we construct two linear combinations:

$$\psi_{\scriptscriptstyle \pm} = A(1)B(2) \pm A(2)B(1)$$

5. The combination with the + sign is the lower energy one, so that becomes the VB wavefunction for H<sub>2</sub>.



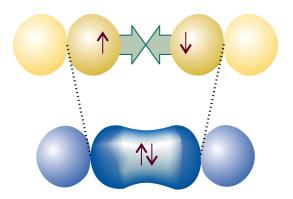
- 6. The electron distribution is called a sigma ( $\sigma$ ) **bond**, meaning it has cylindrical symmetry around the bond axis.
- 7. The VB wavefunction  $\psi_{\sigma} = A(1)B(2) + A(2)B(1)$  is only the <u>spatial</u> part of the wavefunction, and is <u>symmetric</u> with respect to interchange of electron labels 1 and 2. Pauli Principle requires a total wavefunction which is <u>antisymmetric</u>, so the spin part of the wavefunction must be antisymmetric.

Therefore the spin wavefunction we saw before must be used, and that means **spins paired**. (one up, one down)

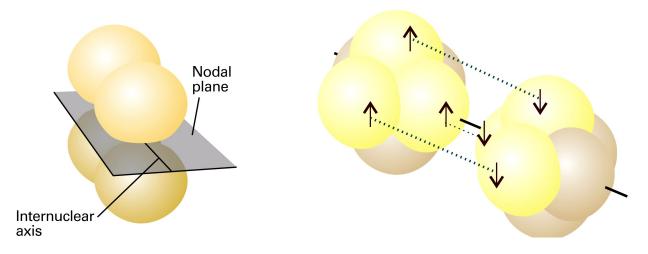
8. VB can easily treat other homonuclear diatomics, like  $N_2$ .

Each N atom has  $1s^22s^22p_x^{1}2p_y^{1}2p_z^{1}$ 

Take z-direction to be the internuclear axis. The two  $2p_z^1$  orbitals from separate atoms overlap to produce a  $\sigma$  bond.



The two neighboring  $2p_x^1$  orbitals overlap to give a  $\pi$  **bond**, and the two neighboring  $2p_y^1$  orbitals overlap to give another  $\pi$  **bond**.

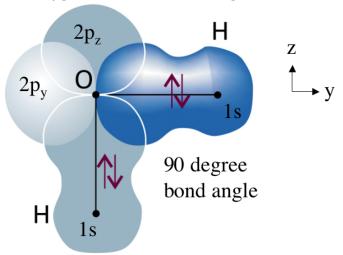


### C. VB treatment of polyatomic molecules.

1. Now let's try to use this idea to predict the bonding in water.

Each H atom brings one electron to the molecule as 1s1

The central oxygen atom has a configuration  $1s^22s^22p_x^22p_y^{-1}2p_z^{-1}$ 



Uh, oh! This is not a good prediction. It predicts two single bonds form, but the molecular geometry is not quite correct. Angle should be 104°, not 90°.

2. Let's try a different molecule, methane, CH<sub>4</sub>.

central atom Carbon =  $[He]2s^22p_x^12p_y^12p_z^0$ 

Only possibility of pairing up with a  $H=1s^1$  in two orbitals - prediction  $CH_2$ .

Carbon's known valency is 4, not 2.

3. VB Theory must introduce another two new ideas ("fixes") here:

# promotion and hybridization

1st, promote an electron out of 2s into the empty 2p

$$2s^{2}2p_{x}^{1}2p_{y}^{1}2p_{z}^{0} \rightarrow 2s^{1}2p_{x}^{1}2p_{y}^{1}2p_{z}^{1}$$

Now there are four locations for pairing up an electron from incoming atoms.

Problem: It would still not produce the known tetrahedral geometry of CH<sub>4</sub>.

2<sup>nd</sup>, form hybrid atomic orbitals by taking **linear combinations** of the 2s and the three 2p atomic orbitals to form four equivalent and **orthogonal** hybrid atomic orbitals. These hybrid orbitals are called **sp³ hybrid atomic orbitals**.

$$(sp^3)_1 = 2s + 2p_x + 2p_y + 2p_z$$

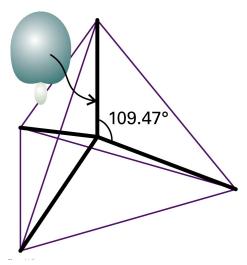
$$(sp^3)_1 = 2s - 2p_x - 2p_y + 2p_z$$

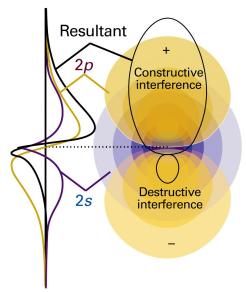
$$(sp^3)_1 = 2s - 2p_x + 2p_y - 2p_z$$

$$(sp^3)_1 = 2s + 2p_x - 2p_y - 2p_z$$

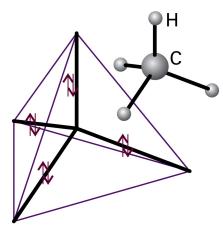
These have a tetrahedral geometry around carbon.

The shape of the hybrids result from constructive and destructive interference that results from adding 2s to a 2p.





Resulting molec. geometry of methane



4. Other hybridizations explain geometries of other molecular systems.

What about Carbon in Ethene? Makes use of sp<sup>2</sup> hybridization.

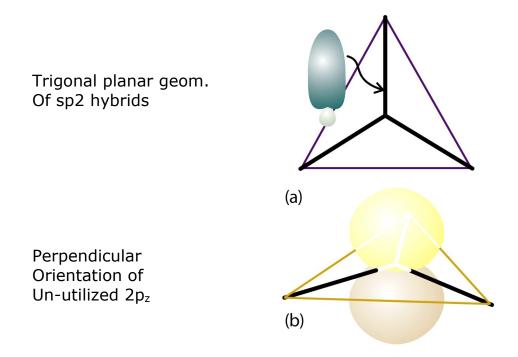
$$(sp^2)_1 = 2s + 2^{1/2} 2p_y$$

$$(sp^2)_2$$
 = 2s +  $(3/2)^{1/2}$  2p<sub>x</sub> -  $(1/2)^{1/2}$  2p<sub>y</sub>

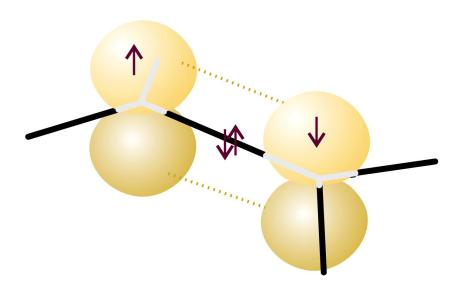
$$(sp^2)_3 = 2s - (3/2)^{1/2} 2p_x - (1/2)^{1/2} 2p_y$$

These need to be normalized.

Note that the 2pz orbital did not enter into the hybridization mix.



The un-utilized perpendicular  $2p_z$  on each carbon in ethene may overlap to create a  $\pi$  **bond**, non-cylindrically symmetric around the bond axis.



5. Hybrid atomic orbitals do not always form bonds. They may also contain a non-bonding pair of electrons, as in water or ammonia.

# 6. Other hybridization schemes:

**Table 10.1 Some hybridization schemes** 

Coordination number	Arrangement	Composition	
2	Linear	sp, pd, sd	
	Angular	sd	
3	Trigonal planar	$sp^2, p^2d$	
	Unsymmetrical planar	spd	
	Trigonal pyramidal	$pd^2$	
4	Tetrahedral	$sp^3$ , $sd^3$	
	Irregular tetrahedral	$spd^2, p^3d, dp^3$	
	Square planar	$p^2d^2$ , $sp^2d$	
5	Trigonal bipyramidal	$sp^3d$ , $spd^2$	
	Tetragonal pyramidal	$sp^2d^2$ , $sd^4$ , $pd^4$ , $p^3d^2$	
	Pentagonal planar	$p^2d^3$	
6	Octahedral	$sp^3d^2$	
	Trigonal prismatic	$spd^4, pd^5$	
	Trigonal antiprismatic	$p^3d^2$	

<sup>\*</sup> Source: H. Eyring, J. Walter, and G.E. Kimball, Quantum chemistry, Wiley (1944).

7. Can you show that a set of hybrid AO's are mutually orthogonal?

# III. Molecular Orbital Theory

### A. Basic Concepts.

- 1. Electrons are no longer regarded as belonging to particular bonds but are spread throughout the molecule to varying degrees.
- 2. Like VB Theory, the hydrogenic atomic orbitals are still the building blocks for the molecular orbitals.
- 3. MO Theory is less physically appealing to working chemists than VB in describing organic molecules, but it achieves much greater accuracy in predicting properties.
- B. The Hydrogen molecule-ion  $H_2^+$ .

Simplest molecular species known.  $H_2^+$  problem may be solved in terms of known functions, but they are too complicated to extend to polyatomics, so we adopt the LCAO-MO approximation.

# C. The LCAO - MO Approximation.

Is an orbital approximation= Electrons in a *molecule* reside in oneelectron orbitals, just like we saw for many-electron atoms. Thus the whole wavefunction is the product of one-electron wavefunctions.

Wave function  $\psi$  for one  $e^-$  in a molecule = "molecular orbital" = "M.O."

 $\psi^2$  = distribution of  $\text{e}^{\text{-}}$  in the molecule.

"LCAO-MO approximation." (linear combination of atomic orbitals [to give] molecular orbitals)

For  $H_2$ , take overlaps of 1s orbital from atom A and atom B

$$\psi_{1s\sigma} = N \left\{ \psi_{1s} (A) + \psi_{1s} (B) \right\}$$
 e occupies norm. cons. has cylindrical sym around internucl axis =  $\sigma$  molecular orbital, specific  $\underline{1s\sigma}$ 

The probability density for locating the  $e^-$  occupying  $1s\sigma$  orbital is:

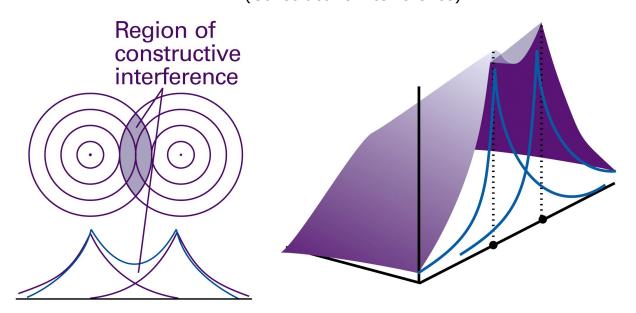
$$\psi_{1s\sigma}^2 = N^2 \Big\{ \psi_{1s} \big( A \big) + \psi_{1s} \big( B \big) \Big\}^2$$

$$= N^2 \Big\{ \psi_{1s}^2 \big( A \big) + 2 \psi_{1s} \big( A \big) \psi_{1s} \big( B \big) + \psi_{1s}^2 \big( B \big) \Big\}$$
prob dens
if e- were
confined to
1s orbital on
atom A
$$\text{extra contrib}$$

$$\text{called}$$

$$\text{"overlap density"}$$

$$\text{(Constructive interference)}$$



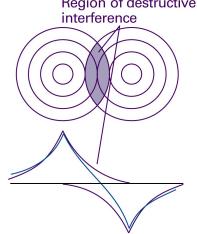
 $1s\sigma$  is a <u>bonding orbital</u>, since there is enhanced amplitude in  $\psi^2$  between the nuclei that hold molec together.

Region of destructive

Can also construct antibonding orbital 1s $\sigma^*$ .

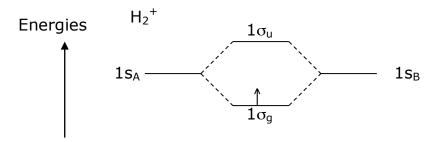
$$\psi_{1s\sigma^*} = N \left\{ \psi_{1s}(A) - \psi_{1s}(B) \right\}$$

Note: little electron probability in the internuclear region, so does not give rise to stability.



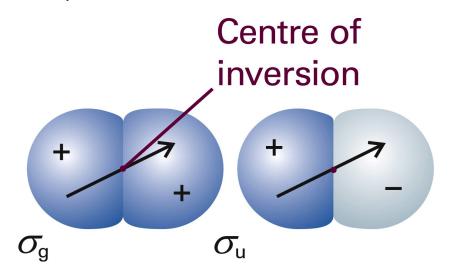
CHAPTER 10

# Conventional MO energy diagram

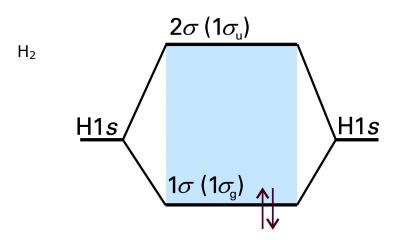


Modern notation focuses on the symmetry of the MO's. For homonuclear diatomics we classify MO's by looking at their **inversion symmetry**.

= the symmetry of MO through its center (a point midway between atoms A and B).





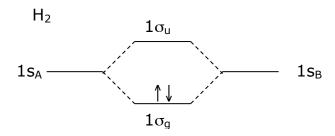


### IV. Homonuclear Diatomic Molecules

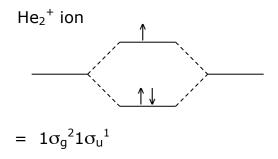
Form M.O.'s by LCAO, overlapping AO's.

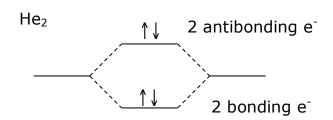
Then fill MO's with electrons just like Aufbau filling of AO's.

# A. Hydrogen and Helium.



 $H_2 = 1\sigma_q^2$  (electron configuration)





$$He_2 = 1\sigma_q^2 1\sigma_u^2$$
 not stable

Bond order of diatomics  $b = 1/2(n-n^*)$  n = # of  $e^-$  in bonding orbitals  $n^* = \#$  of  $e^-$  in antibonding orbitals

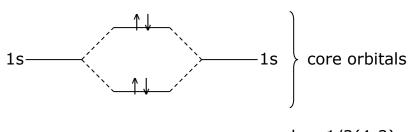
$$He_2^+$$
 b = 1/2(2-1) = 1/2 1/2 a bond  
 $H_2$  b = 1/2(2-0) = 1 "single bond"  
 $He_2$  b = 1/2(2-2) = 0 no bond

# B. Period-2 homonuclear diatomics.

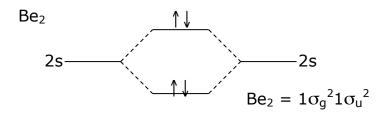
Homonuclear diatomics Hetero diatomics N<sub>2</sub>, etc. LiH, NO, etc.

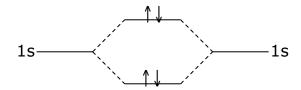
2s  $\longrightarrow$  2s  $\longrightarrow$  valence orb

$$\text{Li}_2 = 1\sigma_g^2$$
 (written config focuses on valence electrons only)



$$b = 1/2(4-2) = 1$$

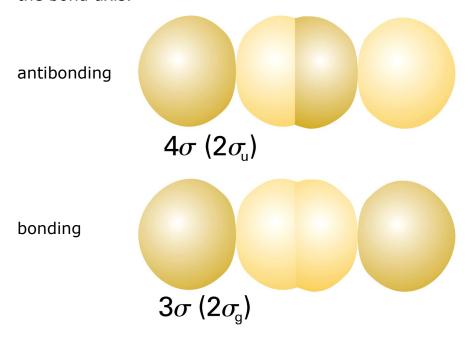


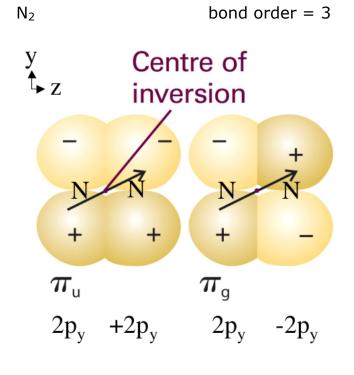


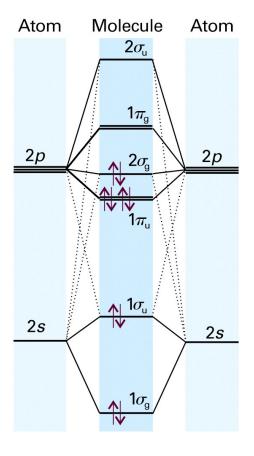
b = 0 unstable

Beyond Be<sub>2</sub>, the 2p orbitals begin to get involved.

Here are the two linear combinations of the  $2p_z$ , which are oriented along the bond axis.

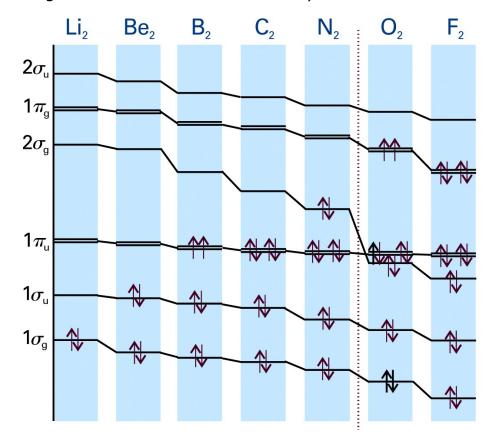






 $2p_x$ 's overlap the same way to give another  $\pi$  M.O. with same energy.

Relative energies of the  $2\sigma$  and  $1\pi$  M.O.'s vary across Row 2 diatomics



Why does  $2\sigma$  take dip down underneath  $1\pi$ ?

So far have constructed M.O.'s by adding together (1s + 1s), (2s + 2s)  $(2p_z + 2p_z)$ ,  $(2p_{x,y} + 2p_{x,y})$ 

In general however:

$$\psi_{MO} = \sum_{i=\text{orbitals}}^{\text{on A}} C_i \psi_i + \sum_{j=\text{orbitals}}^{\text{on B}} C_j \psi_j$$

in principle, all AO's contribute to all MO's

It just turns out that especially for diatomics, "like" orbitals seems to have the largest overlap with each other. e.g.

$$\psi_{2 \circ g} \approx 2p_z(A) + 2p_z(B) + \text{minor contribution from } 2s(A) + 2s(B)$$
dominant terms

Minor contributions do affect relative energies however.

 $O_2$ 

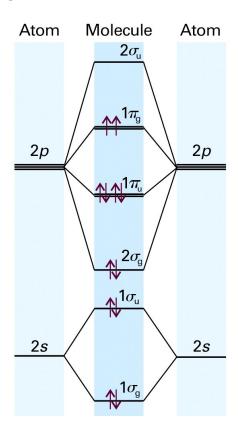
Bond order = 2

By Hund's Rule, the two electrons in the  $1\pi_g$  antibonding level go into separate MO's, giving rise to two unpaired electrons.

Molecular oxygen is paramagnetic.

 $F_2$  has two more electrons into  $1\pi_g$  antibonding level and so has bond order = 1 and no unpaired electrons.

Ne<sub>2</sub> has two more electrons going into antibonding MO, and has bond order = 0. Unstable.

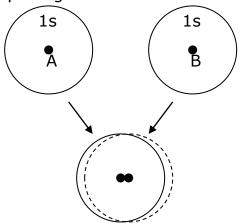


# Effectiveness of overlap:

The overlap integral S, determines the extent of overlap of two atomic orbitals, which determines whether they will form good M.O.

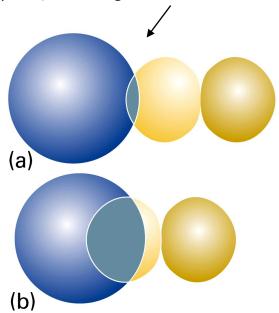
$$S = \int \psi(A) \psi(B) d\tau$$

E.g. If two 1s orbitals approach one another, one on A and one on B, their overlap integral increases to 1.

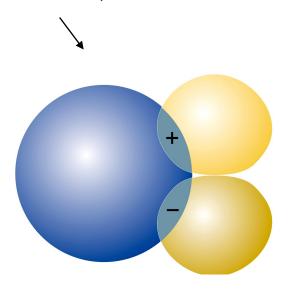


$$S = \int \psi_{1s}(A)\psi_{1s}(A) = 1$$

Overlap of s and p increases up to a point, then begins to decrease.



This s and p gives ZERO net overlap.



# Bond strength:

The greater the bond order, the greater the bond strength.

The greater the bond order, the shorter the bond.

**Table 10.2 Bond lengths** 

Bond	Order	R <sub>e</sub> /pm
НН	1	74.14
NN	3	109.76
HCl	1	127.45
CH	1	114
CC	1	154
CC	2	134
CC	3	120

\* More values will be found in the *Data section*. Numbers in italics are mean values for polyatomic molecules.

**CHAPT** 

Table 10.3 Bond dissociation energies

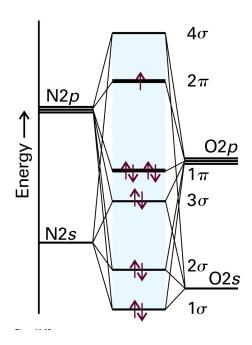
Bond	Order	$D_0/(\mathrm{kJ\ mol^{-1}})$
НН	1	432.1
NN	3	941.7
HCl	1	427.7
CH	1	435
CC	1	368
CC	2	720
CC	3	962

<sup>\*</sup> More values will be found in the *Data section*. Numbers in italics are mean values for polyatomic molecules.

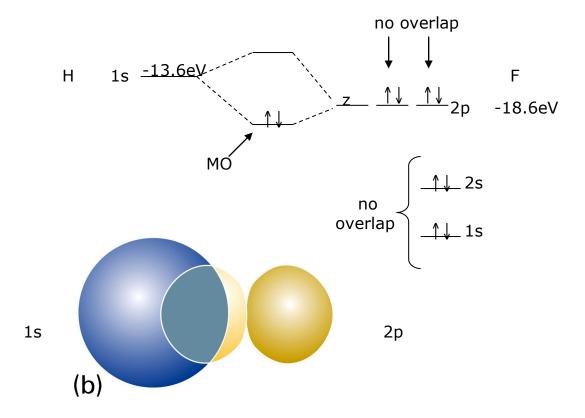
# C. Heteronuclear Diatomics.

M.O.'s are somewhat like homonuclear diatomics, but less so as EN difference between atoms increase.





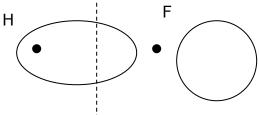
Now what about HF  $EN_F >> EN_H$ 



$$\psi_{MO} = 0.19 \psi_{1s}(H) + 0.98 \psi_{2p_z}(F)$$
 different weighting contribution

M.O. produced has much greater contrib. from  $2p_z$  on F (high Ea) than from 1s of H.

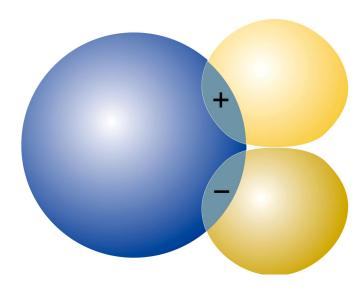
→a **polar covalent bond**. 2e- in the MO spend more time near F than near H. Shared unequally.



General Rule: atomic orbitals most effectively combine (overlap) to form M.O.'s when:

- 1) their energies are similar in the separated atoms.
- 2) their shapes are conducive to good overlap.

E.g. these give no overlap due to their shape and sign of wave function.



The construction of energy level diagrams for arbitrary heteronuclear diatomics is not straightforward, and requires <u>calculation</u>.

D. The Variation Principle – a systematic way of taking linear combinations of atomic orbitals in correct proportions to produce the most optimum approximation to the M.O.'s.

V.P. is based on the fact that approximate M.O.'s will always have a higher energy than the true exact M.O.'s energy.

E.g. in HF, if I had constructed the M.O. as:

$$\psi_{MO} = 1.0\psi_{1s}(H) + 1.0\psi_{2p_7}(F)$$
 with equal coefficients

the energy of this  $\psi_{mo} > E_{true}$ 

Consequence of V.P.  $\rightarrow$  vary the weighting coefficients until you get the lowest energy you can find. This will be the best approx you can make to the true wave function.

$$\psi_{trial} = C_A \psi(A) + C_B \psi(B)$$
 unknown coeff which are to be varied

$$\mathsf{E}_{trial} = \frac{\int \psi_{trial}^* \hat{H} \psi_{trial}}{\int \psi_{trial}^* \psi_{trial}}$$

Vary  $C_A$  and  $C_B$  until  $E_{trial}$  attains its lowest value.

Suppose  $\psi_{trial}$  is real:

$$\begin{aligned} \text{Denom} &= \int \psi_{trial}^2 d\tau = \int \left( C_A \psi(A) + C_B \psi(B) \right)^2 d\tau \\ &= C_A^2 \int \psi(A)^2 d\tau + C_B^2 \int \psi(B)^2 d\tau + 2C_A C_B \int \psi_A \psi_B d\tau \\ &= 1 &= 1 & \text{overlap integral S} \\ &= 1 & \text{wave functions from the separated atoms A + B} \end{aligned}$$

Denominator = 
$$C_A^2 + C_B^2 + 2C_AC_BS$$

$$\begin{aligned} \text{Numerator} &= \int \psi_{trial} \hat{H} \psi_{trial} \\ &= C_A^2 \int \psi(A) \hat{H} \psi(A) d\tau + C_B^2 \int \psi(B) \hat{H} \psi(B) d\tau + 2 C_A C_B \int \psi(A) \hat{H} \psi(B) d\tau \\ & \alpha_A \qquad \alpha_B \qquad \beta \end{aligned}$$

 $\alpha_{\text{A}}$ ,  $\alpha_{\text{B}}$  = "Coulomb integrals"  $\beta$  = "Resonance integral"

$$_{\text{Numerator}} = C_A^2 \alpha_A + C_B^2 \alpha_B + 2C_A C_B \beta$$

$$\mathsf{E}_{\mathsf{trial}} = \frac{\mathsf{C}_{\mathsf{A}}^2 \alpha_{\mathsf{A}} + \mathsf{C}_{\mathsf{B}}^2 \alpha_{\mathsf{B}} + 2 \mathsf{C}_{\mathsf{A}} \mathsf{C}_{\mathsf{B}} \beta}{\mathsf{C}_{\mathsf{A}}^2 + \mathsf{C}_{\mathsf{B}}^2 + 2 \mathsf{C}_{\mathsf{A}} \mathsf{C}_{\mathsf{B}} \mathsf{S}}$$

Must evaluate integrals  $\alpha_A$ ,  $\alpha_B$ ,  $\beta$  and S to obtain #'s. Then vary  $C_A$ ,  $C_B$  until  $E_{trial}$  achieves a minimum.

i.e. set 
$$\frac{\partial E_{trial}}{\partial C_A} = 0$$
 this gives equation: 
$$\left(\alpha_A - E_{trial}\right)C_A + \left(\beta - E_{trial} S\right)C_B = 0$$
 Then set  $\frac{\partial E_{trial}}{\partial C_B} = 0$  this gives equation: 
$$\left(\beta - E_{trial} S\right)C_A + \left(\alpha - E_{trial}\right)C_B = 0$$
 Two equations, two unknowns  $C_A$  and  $C_B$ .

This set of eqns has a solution when the determinant is zero:

$$\begin{vmatrix} \alpha_A - E_{trial} & \beta - E_{trial} S \\ \beta - E_{trial} S & \alpha_B - E_{trial} \end{vmatrix} = 0$$

this is called the "secular determinant"

Definition of "determinant"

$$\begin{vmatrix} A & B \\ C & D \end{vmatrix} = AD - BC$$

$$(\alpha_A - E)(\alpha_B - E) - (\beta - ES)^2 = 0$$

Solve for E (a quadratic equation)

Has 2 roots (corresponding to bonding and antibonding orbital energies)

Simple case: A = B (homonuclear diatomic)

$$\alpha_A = \alpha_B = \alpha$$

Two solutions: (1) bonding 
$$E_{+} = \frac{\alpha + \beta}{1 + S} \qquad C_{A} = \left(\frac{1}{2(1 + S)}\right)^{1/2} = C_{B}$$
 (2) anti 
$$E_{-} = \frac{\alpha - \beta}{1 - S} \qquad C_{A} = \left(\frac{1}{2(1 - S)}\right)^{1/2} = -C_{B}$$
 
$$\psi_{\pm} = \left(\frac{1}{2(1 \pm S)}\right)^{1/2} \left(\psi(A) \pm \psi(B)\right)$$

Even better approximate wave functions and energies for the molecule can be constructed by taking linear combinations of **lots** of atomic orbitals, not just two.

This is called expanding your basis set.

In principle, by using a "complete" basis set, you should obtain an exact solution upon using the variation principle.

In practice, a <u>complete</u> set would need to have <u>all</u> atomic orbitals (full and empty) on all the atoms in your molecule, and that is an infinite set.

# HF - MNDO CALCULATION RESULTS MOPAC: VERSION 6.00 CALC'D. 21-Oct-94 XYZ GEO-OK LARGE VECTORS FOCK MULLIK BONDS DENOUT 1ELECTRON

NO. 1 2	INPUT CARTESI ATOM H F FINAL HEAT O TOTAL ENERGY ELECTRONIC E CORE-CORE RE IONIZATION E NO. OF FILLE MOLECULAR WE SCF CALCULAT	X 0.0000 0.9200  F FORMATION INERGY PULSION OTENTIAL ID LEVELS GIGHT	Y 0.0000 0 0.0000 0 = -57 = -49 = -57 = 8 = 1	0.0000 69.73416 KCAL 64.25877 EV 76.87098 EV 62.61221 EV 64.89120	
ATOM	COMPUTATION  CHEMICAL E	TIME = 0.	180 SECONDS		
NUMBE		(ANGSTROMS)			
1 2	H F	0.95633 *			
	ET C'ENK	ECTORS			
ROOT	NO. 1	2	3	4	5
	-43.77134	-17.93881	-14.89120	-14.89120	5.80377
1S H	-0.37307	0.46609	0.00000	0.00000	0.80223
2S F 2P <sub>X</sub> F	-0.91567	-0.32434 -0.82314	0.00000	0.00000	
2P <sub>X</sub> F 2P <sub>Y</sub> F	0.14956 0.00000	0.00000	0.00000 0.92376	0.00000 -0.38298	0.54779 0.00000
2P <sub>Z</sub> F	0.00000	0.00000	0.38298	0.92376	0.00000
	NIETU NUVA	TO CUNDODO	AND DIDOLE C	YANTITO TEST WILL CALC	
		PE	CHARGE	CONTRIBUTIONS ATOM ELE	CTRON DENSITY
	1	Н	0.2872	0.7128	
	2		-0.2872	7.2872	
DIPOLE POINT-C	X HG1.319	Y 0.000	Z 0.000	TOTAL 1.319	
	-0.670				
SUM			0.000		
FOCK MAY	TRIX IS				
2001.121		2S F 2	2 2PX F 2	2PY F 2	2PZ F 2
1S H	1 -6.329743	 \			
2S F	2 -12.932297	-38.008471			
$2P_X$ F	2 11.390509	0.183063	-11.327472		
2P <sub>Y</sub> F	2 0.000000	0.000000	0.000000	-14.824289	
$2P_{\mathbf{Z}}$ F	2 0.000000			0.000000 -	14.824289
	BONDING CONT 1.1185 0.00 PU TIME:	0.0000			

#### MNDO QUANTUM MECHANICAL CALCULATION

HF 101 5 0 -5 -10 F ENERGY(eV) -15 2p -20 ΑO ΑO -25 11 -30 -35 -40 11

E.g. The  $2p\sigma$  M.O. of homonuclear diatomics can be improved by adding in 2s AO's.

MO's

$$\psi_{trial} = \underbrace{C_A \psi_{2p_Z}(A) + C_B \psi_{p_Z}(B)}_{usual\ combination} + \underbrace{C_A' \psi_{2s}(A) + C_B' \psi_{2s}(B)}_{extra}$$

The Variation procedure will generate 4 eqns and 4 unknowns  $C_A$ ,  $C_B$ ,  $C'_A$ ,  $C'_B$ .

Secular detm will be 4 x 4.

It will turn out that  $C_A$ ,  $C_B >> C'_A$ ,  $C'_B$ 

Therefore minor contrib. from 2s A.O.'s.

Using computers, basis sets can include dozens of A.O.'s to produce and solve huge matrices of numbers and extremely accurate solutions.

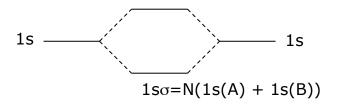
Still must evaluate  $\alpha$ ,  $\beta$  and S integrals. These are done two ways:

- 1) semi empirical estimate them from spectroscopic data.
- ab initio evaluate them numerically on the computer (Simpson's rule)

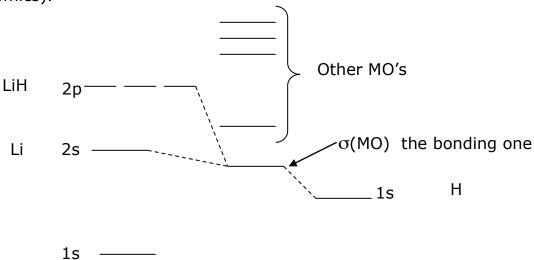
### E. Hybridization can be used in MO theory also.

Introduced to try to preserve the idea that every MO is constructed by linear combination of only 2 AO's.

E.g.



In reality, in advanced MO calcs, every MO is produced by a LC of <u>all</u> the atomic orbitals in the basis set. Turns out that usually only a single AO from each atom contributes significantly to a given MO (esp in homonucl diatomics).



$$\psi_{\sigma} = 0.414\psi_{2s}(Li) + 0.331\psi_{2p_{X}}(Li) + 0.848\psi_{1s}(H)$$

From Variation treatment, 3 AO's end up contributing to  $\sigma$  bond in Li-H.

To preserve concept of overlap of 2 AO's to get MO hybridization idea is introduced.

1s H is seen as overlapping with a hybrid orbital (call it  $\phi(Li)$ ),

 $\phi(Li)$  = mixture of Li atomic orbitals prior to interaction with H.

$$\phi_L(Li) = \phi(Li) + 0.715 \psi_{2p}(Li)$$

Now 
$$\psi_{\text{o}} = 0.323 \, \phi(\text{Li}) + 0.685 \, \psi_{1s} \, (\text{H})$$

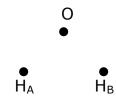
See that hybrid concept is cumbersome here.

### III. Polyatomic Molecules.

MO's are still constructed by LCAO, but using more atomic orbitals.

Ordinary procedure is to use all the AO's from the <u>valence</u> shell of every atom as our basis set.

- A. Treat H<sub>2</sub>O by LCAO-MO theory.
  - 1. Position nuclei at a trial starting position.



3. Use Variation Principle. Will solve 6 x 6 secular determinant.

Will find 6 roots which are the 6 energies of 6 MO's produced.

6 MO's produced 6 energies

4. Walsh diagram – plot of energies of the different MO's as we bend the molecule.

Procedure: vary angles until lowest energy is obtained, always putting the valence e<sup>-</sup> in the lowest energy orbitals possible.

Water,		MOPAC: V ear 180 de	ERSION 6.00 grees		CALC'D. 26	-0ct-94	
FINAL HEAT OF FORMATION = -16.01362 KCAL EIGENVECTORS							
ROOT	NO.		2	3	4	5	6
	•	-38.36620	-20.43460	-11.11085	-11.11085	3.66569	9.72546
S H	1	-0.29829	0.47730	0.00000	0.00000	-0.64111	0.52171
S 0		-0.90667		0.00000	0.00000	0.42185	0.00000
PX O		0.00000	-0.73782	0.00000	0.00000	0.00000	0.67500
PY 0	2		0.00000		-0.02782	0.00000	0.00000
PZ O	2	0.00000	0.00000	-0.02782	0.99961	0.00000	0.00000
S H	3	-0.29829	-0.47730	0.00000	0.00000	-0.64111	-0.52171
		NET ATO	MIC CHARGES	AND DIPOLE CO	NTRIBUTIONS		
	AT	OM NO.	TYPE	CHARGE		TRON DENSITY	
		1	Н	0.3664	0.6336		
		2		-0.7328	6.7328		
		3	Н	0.3664	0.6336		
DIPOLE		X	Y		TOTAL		
POINT-	CHG.	0.000			0.000		
HYBRID		0.000			0.000		
SUM		0.000	0.000	0.000	0.000		
Water,	opt	imum					
	F	INAL HEAT	OF FORMATION	l = -60	.92461 KCAL		
ATO		INAL HEAT CHEMICAL	OF FORMATION BOND LENGTH		.92461 KCAL TWIST A	NGLE	
• • • • •	4			BOND ANGLE	TWIST A	ES)	
• • • • •	4	CHEMICAL	BOND LENGTH	BOND ANGLE	TWIST A	ES)	NB NC
NUMBI	4	CHEMICAL	BOND LENGTH (ANGSTROMS) NA:I	BOND ANGLE (DEGREES)	TWIST A	ES)	NB NC
NUMBI (I) 1 2	4	CHEMICAL SYMBOL	BOND LENGTH (ANGSTROMS)	BOND ANGLE (DEGREES) NB:NA:I	TWIST A (DEGRE NC:NB:	ES) NA:I NA 1	
NUMBI (I) 1	4	CHEMICAL SYMBOL H O H	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000	BOND ANGLE (DEGREES)	TWIST A (DEGRE NC:NB:	ES) NA:I NA	NB NC
NUMBI (I) 1 2 3	ER	CHEMICAL SYMBOL H O H EIGE	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS	BOND ANGLE (DEGREES) NB:NA:I 106.9226	TWIST A (DEGRE NC:NB:	ES) NA:I NA 1 2	1
NUMBI (I) 1 2 3	4	CHEMICAL SYMBOL H O H EIGE	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000	BOND ANGLE (DEGREES) NB:NA:I	TWIST A (DEGRE NC:NB:	ES) NA:I NA 1	
NUMBI (I) 1 2 3	ER T NO	CHEMICAL SYMBOL H O H EIGE	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2	BOND ANGLE (DEGREES) NB:NA:I 106.9226	TWIST A (DEGRE NC:NB:	ES) NA:I NA 1 2	1
NUMBI (I) 1 2 3	ER T NO	CHEMICAL SYMBOL H O H EIGE	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354	TWIST A (DEGRE NC:NB:	ES) NA:I NA  1 2	1 6
NUMBI (I) 1 2 3 ROO	ER T NO	CHEMICAL SYMBOL H O H EIGE 1 -40.47670	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044	TWIST A (DEGRE NC:NB:	ES) NA:I NA  1 2  5  5.85841	1 6 5.97957 0.58139 -0.30659
NUMBI (I) 1 2 3 ROO	ER T NO	CHEMICAL SYMBOL H O H EIGE 1 -40.47670	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543	TWIST A (DEGRE NC:NB:	ES) NA:I NA  1 2 5 5.85841 -0.49342	1 6 5.97957 0.58139
NUMBI (I) 1 2 3 ROO	ER T NO	CHEMICAL SYMBOL H O H EIGE 1 -40.47670 -0.33152 -0.87433 0.10787	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118	TWIST A (DEGRE NC:NB:  8 * 4 -12.31455 0.00000 0.000000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308
NUMBI (I) 1 2 3 ROO	ER T NO 1 2 2	CHEMICAL SYMBOL H O H EIGE 1 -40.47670 -0.33152 -0.87433 0.10787	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997	TWIST A (DEGRE NC:NB:  8 *  4 -12.31455  0.00000  0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874	1 6 5.97957 0.58139 -0.30659 0.35296
NUMBI (I) 1 2 3 ROO S H S O PX O PY O	I NO	CHEMICAL SYMBOL H O H EIGE 1 -40.47670 -0.33152 -0.87433 0.10787 -0.12305 0.00000	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143 0.00000	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997 0.00000	TWIST A (DEGRE NC:NB:  8 *  4 -12.31455  0.00000  0.00000 0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308
NUMBI (I) 1 2 3 ROO S H S O PX O PY O PZ O	1 2 2 2 2 2	CHEMICAL SYMBOL H O H EIGE 1 -40.47670 -0.33152 -0.87433 0.10787 -0.12305 0.00000 -0.31444	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143 0.00000 -0.50373	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997 0.00000 -0.25673	TWIST A (DEGRE NC:NB:  4 -12.31455  0.00000 0.00000 0.00000 0.00000 0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241 0.00000	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308 0.00000
NUMBI (I) 1 2 3 ROO S H S O PX O PY O PZ O	1 2 2 2 2 3	CHEMICAL SYMBOL H O H EIGE 140.47670 -0.33152 -0.87433 0.10787 -0.12305 0.00000 -0.31444 NET AT	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143 0.00000 -0.50373	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997 0.00000 -0.25673 AND DIPOLE CO	TWIST A (DEGRE NC:NB:  8 *  4 -12.31455  0.00000 0.00000 0.00000 0.00000 0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241 0.00000 0.55562	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308 0.00000
NUMBI (I) 1 2 3 ROO S H S O PX O PY O PZ O	1 2 2 2 2 3	CHEMICAL SYMBOL H O H EIGE 1 -40.47670 -0.33152 -0.87433 0.10787 -0.12305 0.00000 -0.31444 NET AT	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143 0.00000 -0.50373 OMIC CHARGES TYPE	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997 0.00000 -0.25673 AND DIPOLE COCHARGE	TWIST A (DEGRE NC:NB:  8 *  4 -12.31455  0.00000 0.00000 0.00000 0.00000 0.00000 0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241 0.00000	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308 0.00000
NUMBI (I) 1 2 3 ROO S H S O PX O PY O PZ O	ER T NO	CHEMICAL SYMBOL H O H EIGE 1. 1 -40.47670 -0.33152 -0.87433 0.10787 -0.12305 0.00000 -0.31444 NET AT	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143 0.00000 -0.50373 OMIC CHARGES TYPE H	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997 0.00000 -0.25673 AND DIPOLE COCHARGE 0.1630	TWIST A (DEGRE NC:NB:  8 *  4 -12.31455  0.00000 0.00000 0.00000 0.00000 0.00000 0.00000 0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241 0.00000 0.55562	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308 0.00000
NUMBI (I) 1 2 3 ROO S H S O PX O PY O PZ O	ER T NO	CHEMICAL SYMBOL H O H EIGE 1 -40.47670 -0.33152 -0.87433 0.10787 -0.12305 0.00000 -0.31444 NET AT	BOND LENGTH (ANGSTROMS) NA:I 0.94000 0.94000 NVECTORS 2 -18.16014 0.43899 0.01750 -0.54937 -0.50143 0.00000 -0.50373 OMIC CHARGES TYPE	BOND ANGLE (DEGREES) NB:NA:I 106.9226 3 -15.45354 -0.34044 0.37543 0.54118 -0.61997 0.00000 -0.25673 AND DIPOLE COCHARGE	TWIST A (DEGRE NC:NB:  8 *  4 -12.31455  0.00000 0.00000 0.00000 0.00000 0.00000 0.00000 0.00000	ES) NA:I NA  1 2 5 5.85841 -0.49342 -0.01728 -0.51874 -0.42241 0.00000 0.55562	1 6 5.97957 0.58139 -0.30659 0.35296 -0.41308 0.00000

DIPOLE	X	Υ	Z	TOTAL
POINT-CHG.	-0.522	0.704	0.000	0.876
HYBRID	-0.539	0.728	0.000	0.906
SUM	-1.061	1.432	0.000	1.782

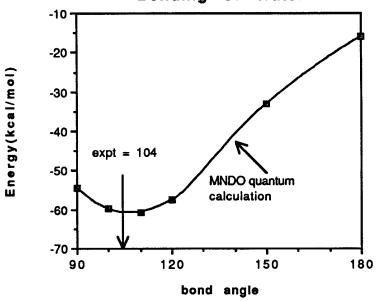
### Water, 90 degree

		F		OF FORMATION = -54.37055 KCAL NVECTORS				
	ROOT	NO		2	3	4	5	6
			-40.47670	-18.16014	-15.45354	-12.31455	5.85841	5.97957
S	H	1	-0.32425	0.46451	-0.32988	0.00000	-0.53313	0.53483
S	0	2	-0.87368	0.00000	0.39231	0.00000	0.00000	-0.28771
PΧ	0	2	0.11491	-0.53313	0.56056	0.00000	-0.46451	0.41541
PY	0	2	-0.11491	-0.53313	-0.56056	0.00000	-0.46451	-0.41541
PΖ	0	2	0.00000	0.00000	0.00000	1.00000	0.00000	0.00000
S	н	3	-0.32425	-0.46451	-0.32988	0.00000	0.53313	0.53483

#### NET ATOMIC CHARGES AND DIPOLE CONTRIBUTIONS ATOM NO. **TYPE** CHARGE ATOM ELECTRON DENSITY 0.8595 6.2811 1 2 0.1405 Н 0 -0.2811 Н 0.1405 0.8595 DIPOLE X TOTAL POINT-CHG. -0.635 0.897 0.635 0.000 HYBRID -0.650 0.919 0.650 0.000 SUM -1.284 0.000 1.816

### Walsh Diagram **Bending** of Water

1.284



CHAPTER 10 27 B. Huckel Molecular Orbital theory (HMO theory).

An approximate theory that gives us a very quick picture of the MO energy diagram and MO's of molecules without doing a lot of work.

Structure and bonding of the  $\pi$  electrons in highly conjugated systems nicely treated by HMO. Look at examples:

1. Ethene – look at the  $\pi$  bond in this simple molecule first.

Each carbon has a  $2p_z$  orbital perpendicular to plane of molecule. Use LCAO-MO to get two MO's, which gives rise to 2x2 secular determinant:

$$\begin{vmatrix} \alpha_{A} - E_{trial} & \beta - E_{trial} S \\ \beta - E_{trial} S & \alpha_{B} - E_{trial} \end{vmatrix} = 0$$

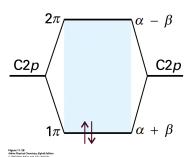
HMO approx:

Set all overlap integrals S = 0

Recognize the  $\alpha$  integrals are equal

$$\begin{vmatrix} \alpha - E & \beta \\ \beta & \alpha - E \end{vmatrix} = 0$$

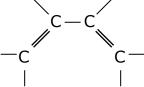
This gives quadratic  $(\alpha - E)^2 - \beta^2 = 0$ Roots  $E = \alpha + \beta$ ,  $\alpha - \beta$ 



Total  $\pi$  bond energy = 2 ( $\alpha$  +  $\beta$  ) Stabilization due to bond formation = 2 $\beta$ 

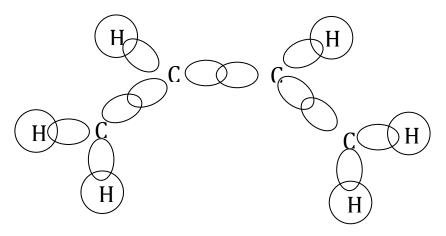
 $\beta$  is intrinsically negative and  $\sim$  -2.4 eV for C-C bond ( $\sim$  -230 kJ/mol)

2. Butadiene.



Treat sigma bonding framework using VB theory as follows:

 $sp^2$  around each C forms  $\sigma$ -bonding framework



Still have an unused p orbital available on each C, perpendicular to plane of molecule.

Edge on view:

These extensively overlap to form a delocalized  $\pi$  system.

Treat  $\boldsymbol{\pi}$  system by HMO as follow:

a. basis set is composed of four  $2p_z$  orbitals (perpendicular to plane).

Therefore will get 4 x 4 matrix or secular determinant

# b. HMO approx:

- 1) Set all overlap integrals S = 0
- 2) Set all resonance integrals between non-neighbors = 0  $\beta_{13}$ ,  $\beta_{14}$ ,  $\beta_{24}$  etc. = 0
- 3) Set all other resonance integrals ( $\beta$ ) equal to one another.

Resulting Sec detm

$$\begin{vmatrix} \alpha - E & \beta & 0 & 0 \\ \beta & \alpha - E & \beta & 0 \\ 0 & \beta & \alpha - E & \beta \\ 0 & 0 & \beta & \alpha - E \end{vmatrix} = 0$$

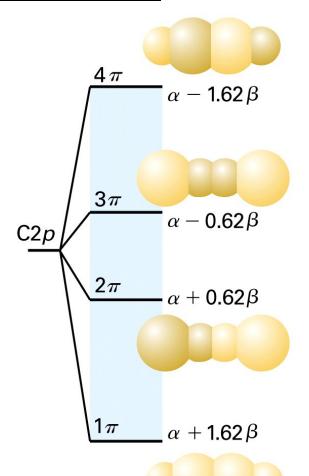
Generates a quartic polynomial in E which has four roots.

Therefore  $4 \pi$ -orbital MO

4 energies (roots of secular determinant)

They are:  $E = \alpha \pm 1.62 \beta$ ,  $\alpha \pm 0.62 \beta$ 

Energy diagram:



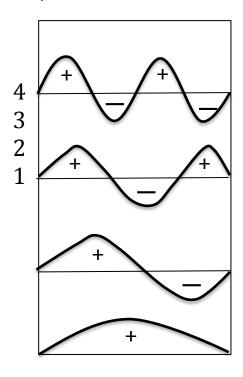
Spectroscopic transition HOMO → LUMO (called frontier orbitals)

HOMO= highest occupied molecular orbital (here MO  $2\pi$ )

LUMO = lowest unoccupied molecular orbital (here MO  $3\pi$ )

$$\Delta E_{trans} = LUMO - HOMO = (\alpha - 0.62 \beta) - (\alpha + 0.62 \beta) = -1.24\beta$$

Compare wave functions with particle in box:



Delocalized system have extra stability (=deloc energy  $\Delta E_{deloc}$ ) Calculate it:

$$\Delta E_{deloc} = \text{ Energy of c-c-c-c} - \text{ Energy of c=c-c=c}$$

$$Allow \ electrons \qquad 2 \ separate \\ \pi \ bonds \\ like \ in \ \underline{ethene}$$

$$= \left[2(\alpha + 1.62 \ \beta) + 2(\alpha + 0.62 \ \beta)\right] - 2(2\alpha + 2\beta)$$

$$= 4\alpha + 4.48\beta \quad - \quad (4\alpha + 4\beta)$$

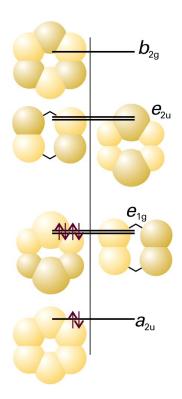
$$= 0.48\beta \quad \sim -110 \ kJ/mol \ of \ stabilization$$

3. Benzene – cyclic delocalized system.

6x6 secular determinant =

due to cyclic connection

=Energy eigenvalues are E =  $\alpha \pm 2\beta$ ,  $\alpha \pm \beta$ ,  $\alpha \pm \beta$ 



Calculate delocalization energy DE:

Total pi electron energy =  $2(\alpha + 2\beta) + 4(\alpha + \beta) = 6\alpha + 8\beta$ Energy of C=C in ethene =  $3(2\alpha + 2\beta)=6\alpha + 6\beta$ DE = difference =  $2\beta = -460$  kJ/mol

4. Solids are extreme case of delocalization.

Graphite is like benzene stretching to infinity.

- C. Computational Chemistry: Solving the Schrodinger Equation for Molecules.
  - 1. The molecular orbitals  $\phi_i$  are produced by a linear combination of atomic orbitals  $\psi_k$  that are the basis set. N atomic orbitals produce N molecular orbitals.

$$\phi_i = \sum_{k=1}^N C_{ik} \psi_k$$

Finding the weighting coefficients  $C_{ik}$  is the essence of the problem to be solved.

 $C_{ik}$  = contribution AO k makes to MO i.

2. The *variation principle* forms the basis for making sure we have the "best", most accurate wave function for our molecule.

Approximate MO energy = 
$$\frac{\int \phi_i^* H \phi_i d\vec{r}}{\int \phi_i^* \phi_i d\vec{r}}$$
 is always >  $E_i$  (true)

So, by varying the coefficients  $C_{ik}$  which make up the MO function  $\phi_i$  until we find the *minimum* energy possible, we are ensured of having the best  $\phi_i$  and energy  $E_i$  afforded by the *basis set* we have chosen. The more robust the basis set, the more accurate the  $\phi_i$  and  $E_i$  for the molecule.

3. The actual procedure for the above involves lots of matrix algebra with matrix elements derived from the following integral expressions:

$$S_{lk} = \int \psi_l \psi_k d\vec{r}$$
 overlap integrals   
  $H_{lk} = \int \psi_l H \psi_k d\vec{r}$  Coulomb integrals

- 4. The  $S_{lk}$  are a snap to solve. The harder problem is the  $H_{lk}$ . Different MO methods vary according to the way they handle this problem.
- 5. In the Coulomb integrals, *H* is an effective one-electron Hamiltonian which is complicated by the fact that it accounts for interactions with all the other electrons. *H* then has within it terms which require the electron density of *all the other electrons*.

Therefore one must "know" the solution  $\phi_i$  before one starts, just to have the effective one-electron Hamiltonians H.

6. This implies an iterative solution called the Hartree-Fock selfconsistent field (SCF) iteration. The Hartree-Fock one-electron

$$H^{SCF} = H^{N} + \sum_{j} J + \sum_{j} K$$

Hamiltonian is often written as, where sum j is over all the other e-.

$$J_j = \int \phi_j^2(2) \frac{1}{r_{12}} d\vec{r}_2$$
 Coulomb electronic interactions

$$K_j = \int \phi_j(1)\phi_i(2)\frac{1}{r_{12}}d\vec{r}_2$$
 exchange terms (due to Pauli)

 $H^N$  = attraction of electrons to fixed nuclei and simple kinetic energy terms.

- D. Ab Initio Hartree-Fock (HF) Methods:
  - 1. All integrals are actually directly solved for the basis set employed, then the matrix diagonalization is performed.

The number of integrals needing to be solved increases as  $N^4$ , where N = atoms.

- 2. ab initio HF methods then differ by the *basis set* employed.
- 3. Most intuitively obvious choice for basis set would be Slater-type atomic orbitals (STO). (Hydrogen-like with good  $\xi = Z_{eff}/n$  values):

$$\psi_k = Ce^{-\zeta r} Y_{lm}(\theta, \phi)$$

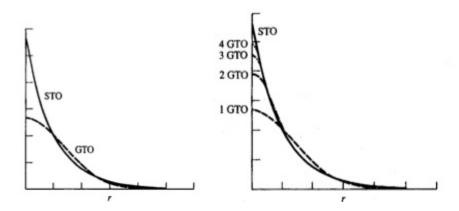
Then, e.g. for each Carbon atom we use a 1s, a 2s, and three 2p atomic orbitals.

4. Easier to do integrals if  $\psi_k$  are of a Gaussian form instead of a Slater exponential:

$$e^{-ar^2}$$
 instead of  $e^{-\zeta r}$ 

But the true wave function is more nearly like  $e^{-\zeta r}$  around nuclei, so we might use 3 Gaussian functions combined to approximate the Slater exponential. This is called:

meaning "Slater-type orbitals built up using 3 Gaussians" or STO - 4G is more accurate. (3 seems to be the absolute minimum number of Gaussians to treat each AO.



### 5. Gaussian basis sets:

- Each Gaussian shape is determined by an exponential parameter and an expansion coefficient.
- Most flexible basis allow these to be varied independently. (Uncontracted or primitive Gaussians)
- More commonly use *contracted* Gaussians = exponents are preset and remain constant during calculation.
- Further, it is common for efficiency's sake to use the same Gaussian exponents for the s and p orbitals in a given shell on a given atom. (Restricts flexibility but reduces number of integrals).
- 6. Minimal basis set = contains AO up through the valence shell on each atom.

e.g. all the atomic orbitals in the shell H and He, use 1s Li to Ne, use 1s, 2s, three 2p Third row, start adding 3d.

STO-nG are all minimal basis sets, using n Gaussians for each AO

### Deficiencies in STO-nG:

- atoms at the end of a period are described using same number of basis functions as atoms at the beginning. Oxygen and Fluorine versus Lithium. (Contracted Gaussian radial exponents are not allowed to vary – expand or contract according to molecular environment.)
- b. Can't treat non-spherical anisotropies in wave function because  $2p_x$ ,  $2p_y$  and  $2p_z$  are all the same size.
- 7. Short-hand notation schemes for the basis set used.

Double-zeta basis set uses a linear combination of two radial

exponents, one contracted and one diffuse, whose weightings are varied during the calc.

Split-valence double-zeta 3-21G basis set:

- 3 Gaussians to describe each core orbital.
- 3 Gaussians to describe each valence shell orbital, with 2 being contracted and one being diffuse.
- 6-31G is also common and works the same way.
- 8. Strong anisotropic charge distributions: all basis sets above use functions centered on atomic nuclei.

Treating non-isotropic charge distribution around an atom done by introducing *polarization* functions into the basis set.

e.g., electron cloud around a H atom bonded to another atom is distorted in direction of the other atom. Can mimic this by mixing with the 1s a p-type orbital to give a form of sp hybrid.

Similarly, p orbitals can be enhanced by adding in d type orbitals to give pd hybrids which are more flexible in treating the distortion.

Use of polarization basis functions is denoted with asterisk:

3-21G\* or 6-31G\* A single asterisk means polarization functions are used on only non-H atoms.

\*\* is used when polarization is also applied to H atoms (good for treating H-bonding better).

See reference books for other commonly used abbreviations.

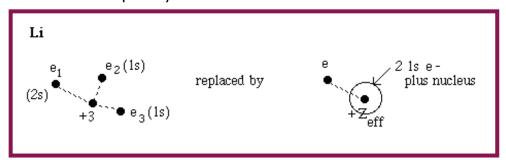
9. Another challenge is atoms with lone pairs, significant density away from nuclei. Highly diffuse functions are added to the basis set, denoted by using a "+" or "++"

$$6-311++G(3df,3pd)$$

- E. Semi-Empirical Methods:
  - 1. Less computationally intensive than ab initio.
  - 2. Many of the integrals are not actually directly solved, but are substituted for by parameters determined empirically, i.e., so as to

give good fit to specific experimental properties for a given class of molecules.

3. Particularly the  $H^N$  integrals, which treat electron interaction with nuclei, are replaced by  $H^{core}$ , which involves attractions to the inner core rather than a bare nucleus. Then <u>only valence shell</u> electrons are treated explicitly.



4. Some common semi-empirical quantum mechanics methods include:

CNDO - complete neglect of differential overlap

INDO - intermediate neglect of differential overlap

MNDO

AM1 - Austin Model 1

PM3 - Parameterized Model revision 3

Reference books abound which describe these various formulations.

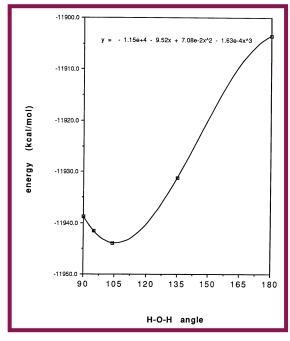
### F. Molecular Geometry Optimization:

- 1. All QM methods can solve the Schrodinger equation for MO's and energies (and everything else) for a **fixed** set of nuclear coordinates input by the user. This is called a *single point energy calculation*.
- 2. They are also capable of computing forces on the atoms and moving them in a direction of decreasing total energy until an optimum geometry is reached. This is called *geometry optimization*, and is done iteratively through many steps.
- 3. In the HF example, the user inputs starting positions of H and F such that the bond distance was 1.07Å. After 7 cycles of iteration, the energy decreased almost 30 kcal/mol and a final bond distance of 0.827Å was achieved. This is thus the predicted equilibrium bond length of H-F according to the AM1 Hamiltonian.
- 4. In a *manual* geometry optimization of  $H_2O$  using the CNDO method, I performed a single-point energy calculation for a series of H-O-H bond angles starting from bent configuration (90°) and going to

linear configuration (180°). The following potential energy curve was derived.

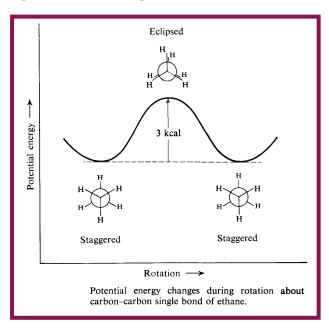
# e.g. CNDO Calculation Energy of Bending Water

Note that CNDO predicts an equilibrium bond angle of 105°, compared with the experimental value of 104°. Further, we see that it costs 40 kcal/mol energy to "straighten out" water to 180°.



### G. Molecular Conformation:

- 1. Most organic and biological molecules have freedom to rotate about bonds (A torsion angle rotation) and can exist in several conformational states. An example is butane, which rotates around the center C-C bond to produce g<sub>+</sub>, trans, and g<sub>-</sub> conformations.
- 2. Conformational analysis is process of the systematically exploring the conformational states of a molecule, at each point doing single-point calculation to generate potential energy surface.

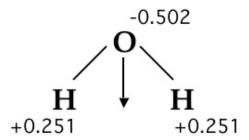


### H. Other Quantities Accessible to QM Calculations:

- 1. Vibrational spectra can be calculated by normal mode analysis. (useful for vibrational spectroscopy)
- 2. Ionization potentials and electron affinities.
- 3. Charge distribution and dipole moment.

Mulliken population analysis: Partitions the electron density between the nuclei so that each nucleus has a non-integer number of electrons associated with it.

e.g.  $H_2O$  calculation with AM1 (Dipole moment = 1.66 D)



Then these point charges can be used to estimate the electrostatic potential around a molecule in molecular mechanics methods.

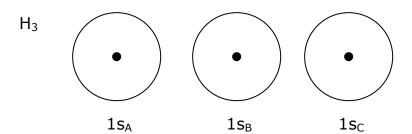
- 4. Electrostatic potential maps.
- 5. Frontier orbitals electron density:

LUMO = lowest unoccupied molecular orbital HOMO = highest occupied molecular orbital

Used to predict site of reactivity of molecules. Most highly reactive site on a molecule in an electrophilic reaction is the HOMO (the molecule is the e<sup>-</sup> donor). For a nucleophilic attack it is the LUMO (the molecule is the e<sup>-</sup> acceptor).

- 6. QSAR properties (Quantitative Structure-Activity Relationships) can be calculated:
  - a. hydrophobicity = log P , where P=partition coefficient between octanol and water.
  - b. molecular volume
  - c. molecular surface area.

I. Problem in Homework: Use HMO to treat a hypothetical molecule H<sub>3</sub>.

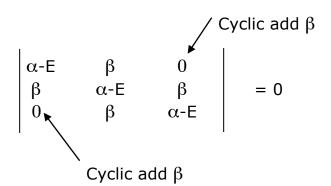


The secular determinant would be:

$$\begin{vmatrix} \alpha \text{-E} & \beta_{AB} \text{-E } S_{AB} & \beta_{AC} \text{-E } S_{AC} \\ \beta_{BA} \text{-E } S_{BA} & \alpha \text{-E} & \beta_{BC} \text{-E } S_{BC} \\ \beta_{CA} \text{-E } S_{CA} & \beta_{CB} \text{-E } S_{CB} & \alpha \text{-E} \end{vmatrix} = 0$$

Now, the HMO approximations are:

non-neighbor resonance integrals  $\beta$ 's = 0 all overlap integrals S=0 all resonance integrals between neighbors =  $\beta$ 



$$\left(\alpha - E\right) \begin{vmatrix} \alpha - E & \beta \\ \beta & \alpha - E \end{vmatrix} - \beta \begin{vmatrix} \beta & \beta \\ 0 & \alpha - E \end{vmatrix} + 0 = 0$$

$$(\alpha - E)\left\{\left(\alpha - E\right)^2 - \beta^2\right\} - \beta\left\{\beta\left(\alpha - E\right) - 0\right\} = 0$$
$$(\alpha - E)^3 - (\alpha - E)\beta^2 - \beta^2(\alpha - E) = 0$$

Cubic equation but one root is  $E = \alpha$ , so factor out  $(\alpha - E)$ .

$$(\alpha - E)^2 - \beta^2 - \beta^2 = 0$$

$$\left(\alpha - \mathsf{E}\right)^2 - 2\beta^2 = 0$$

$$\alpha^2 - 2\alpha E + E^2 - 2\beta^2 = 0$$

This is a quadratic equation that can be solved for E.

$$E=\alpha\pm\sqrt{2}\beta$$

 $E = \alpha$  is also a root

Energy diagram for linear H<sub>3</sub>:

$$\alpha - \sqrt{2}\beta$$

α

$$\alpha + \sqrt{2}\beta$$

# NOTES: