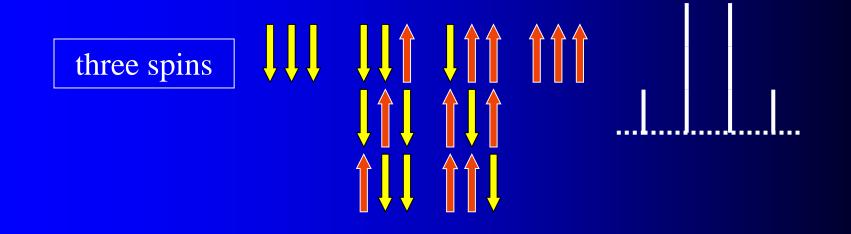
Lecture 5 Still More nmr

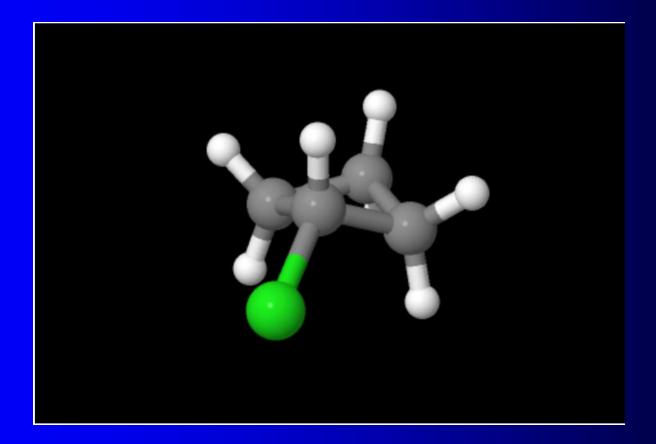






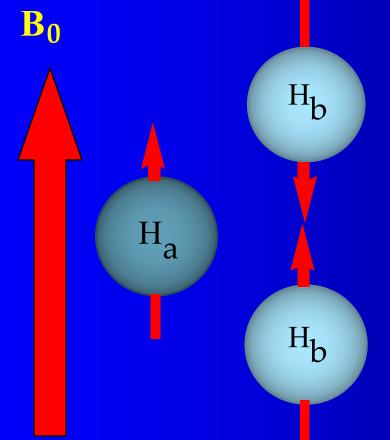
February 5, 2019

Supplemental Problem



Chlorocyclobutane

Origins of Signal Splitting

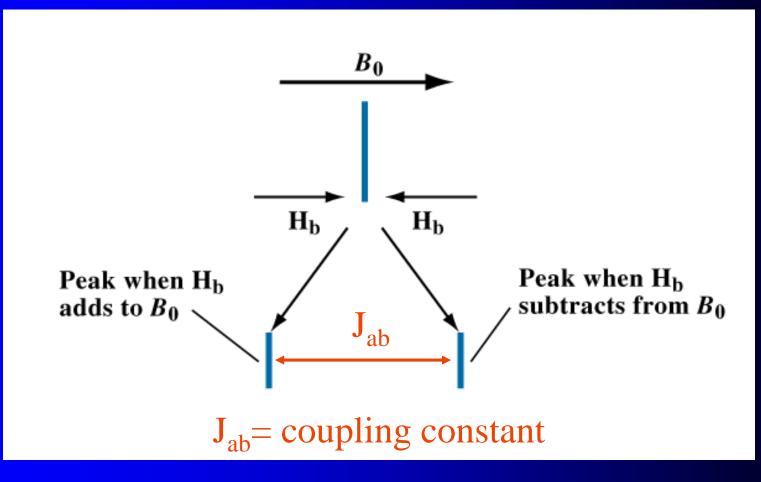


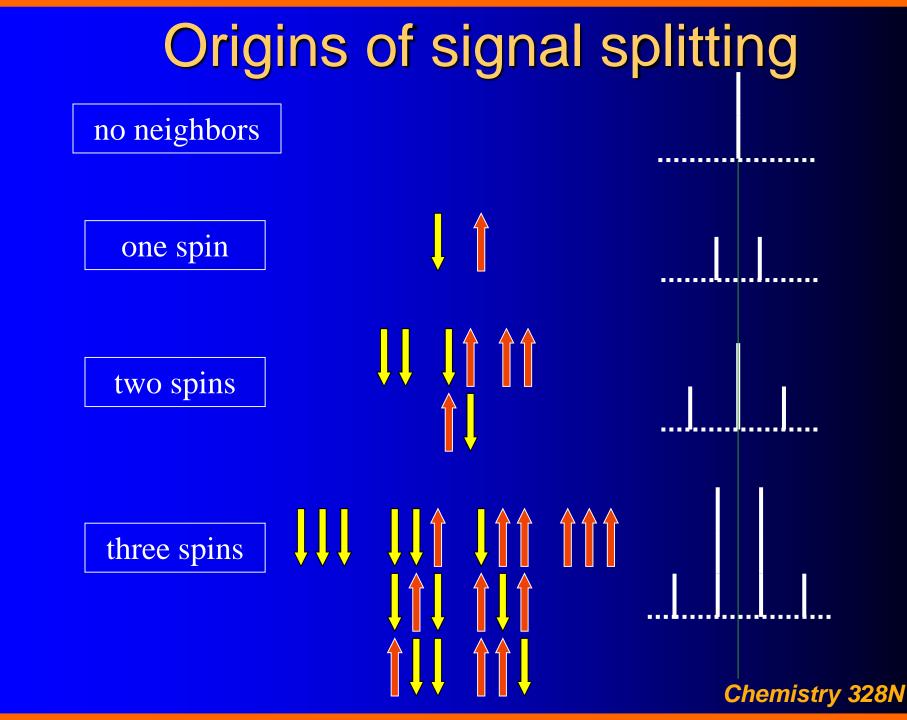
Magnetic field of H_b subtracts from the applied field; H_b signal appears at a higher applied field

Magnetic field of H adds to the applied field; H signal appears at a lower applied field

Remember...it is the **NET** field that counts

The signal of Ha is split into two peaks of equal area (a doublet)





Relative Intensity of Peaks

singlet	1
double	11
triplet	121
quartet	1331
quintete	14641
sextete	1 5 10 10 5

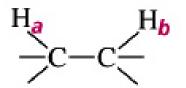
Pascal's triangle The binomial coefficients

The "N+1 Rule"

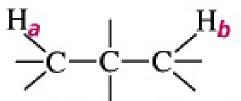
- The ¹H-NMR signal of a hydrogen or set of equivalent hydrogens is split into (N + 1) peaks by a set of N equivalent neighboring hydrogens
 - All neighboring hydrogens in the analysis must have the same chemical shift (magnetically equivalent)
 - If this condition is not met, a graphical tree or second order analysis must be used to predict the splitting pattern. We will explore this condition later

Splitting of Signals

 Nonequivalent protons split each other if they are on adjacent carbons, i.e. separated by 3 sigma bonds



H_a and H_b will split each other because they are separated by 3 σ bonds



 H_a and H_b will not split each other because they are separated by 4 σ bonds

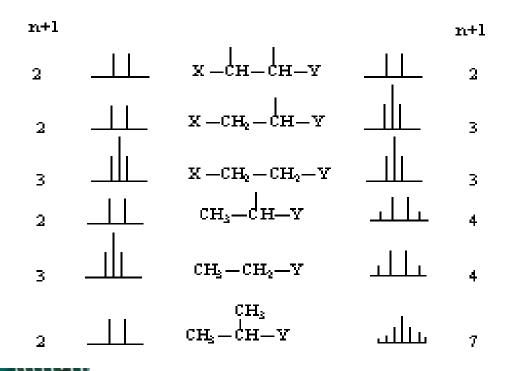
Splitting of Signals

Equivalent protons <u>never</u> split each other

CH₃Br bromomethane ClCH₂CH₂Cl 1,2-dichloroethane

both compounds have an NMR spectrum that shows one singlet because equivalent protons do not split each other's signals

NMR Splitting Patterns

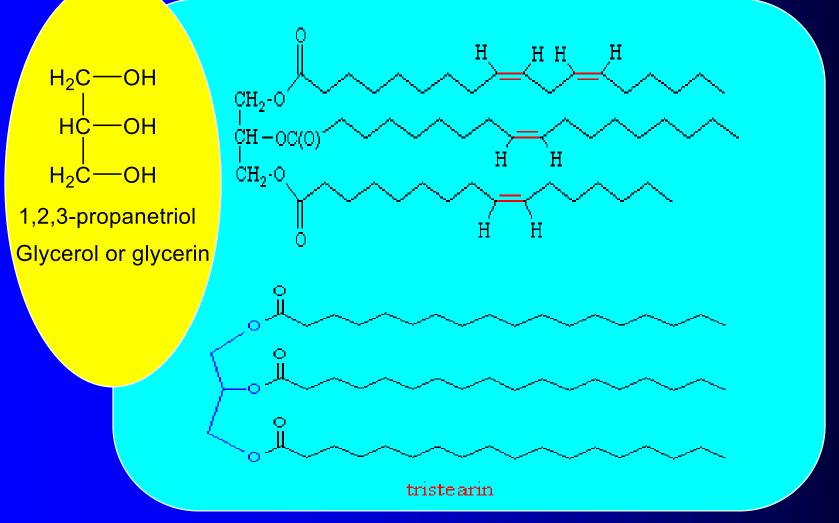




Who needs DNA??



Corn Oil and Tristearin



Coloring Margarine



Index of Hydrogen Deficiency

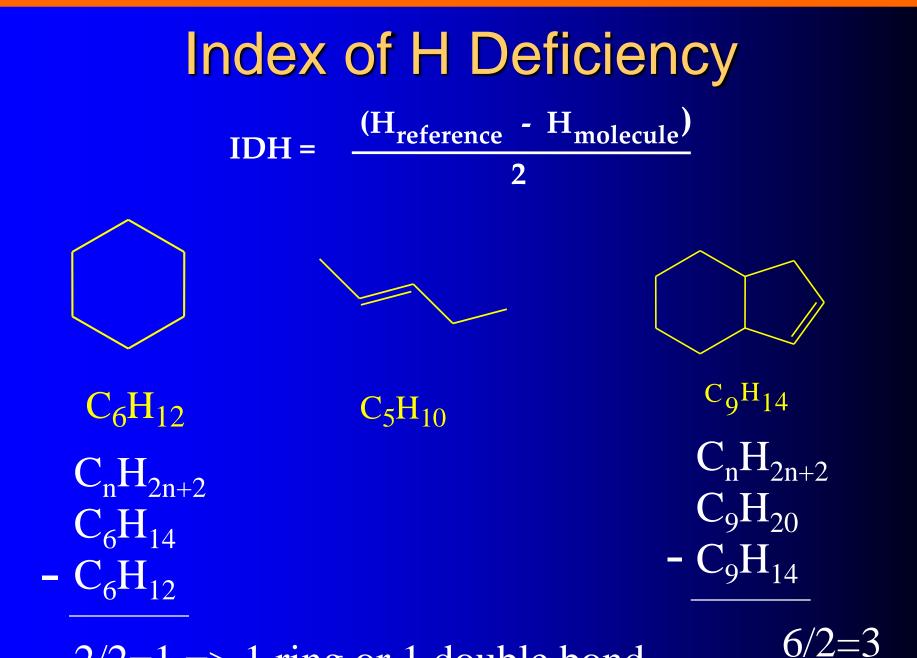
- I knew this as the unsaturation number
- Valuable characteristic of structure
 - Provides you with number of double bonds or rings in a compound
 - Easy with CHO formulas only...be careful with N
 - Simplest, "saturated" normal alkanes have:

 $\mathbf{H} - (\mathbf{CH}_2)_n - \mathbf{H} = \mathbf{C}_n \mathbf{H}_{2n+2}$

Index of H Deficiency

• IHD: the sum of the number of rings and pi bonds in a molecule....

• Compare the number of hydrogens in an unknown compound with the number in the reference compound C_nH_{2n+2}



 $2/2=1 \Rightarrow 1$ ring or 1 double bond

Index of H Deficiency IDH = $\frac{(H_{reference} - H_{molecule})}{2}$

- No correction is necessary for the addition of atoms of O
- Subtract one H for each atom of halogen added (halogens like Cl replace H one for one)
- Add one hydrogen For each atom of N added to the reference hydrocarbon



"Unsaturation" Number or HDI

Η $C_8H_{14}ClN$

Add one H for N and subtract one for Cl Should be $C_n H_{2n+2+1-1} = C_8 H_{18}$ but it is... $C_{8}H_{14}$ H_4 Missing So index is.... 4/2=2

Calculate the Unsaturation Number

Η

 C_8H_9N

Add one H for N

 Should be $C_n H_{2n+2+1} = C_8 H_{19}$

 but it is...
 $C_8 H_9$

 Missing
 H_{10}

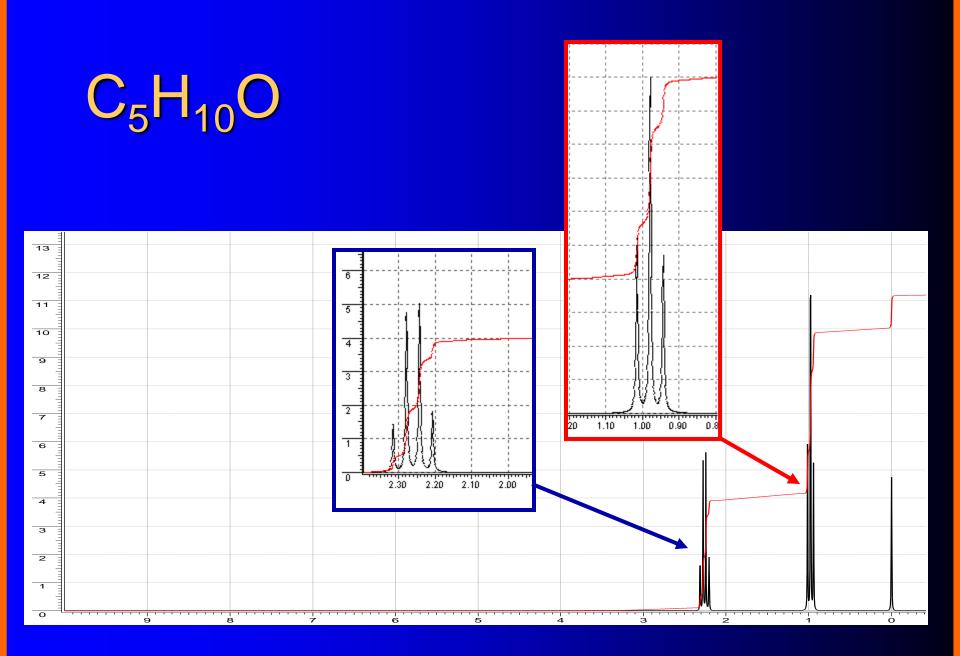
 So number is...
 10/2=5!

"Tricks" for solving unknowns

• Review....

 Empirical formula is lowest common denominator ratio of atomic composition

• From Homework: unknown has an empirical formula of C_4H_9a single high field peak in the ¹H nmr and a molecular ion at M/Z = 114.propose a structure..and predict the M/Z of the most intense fragment you expect to see in the mass spectrum



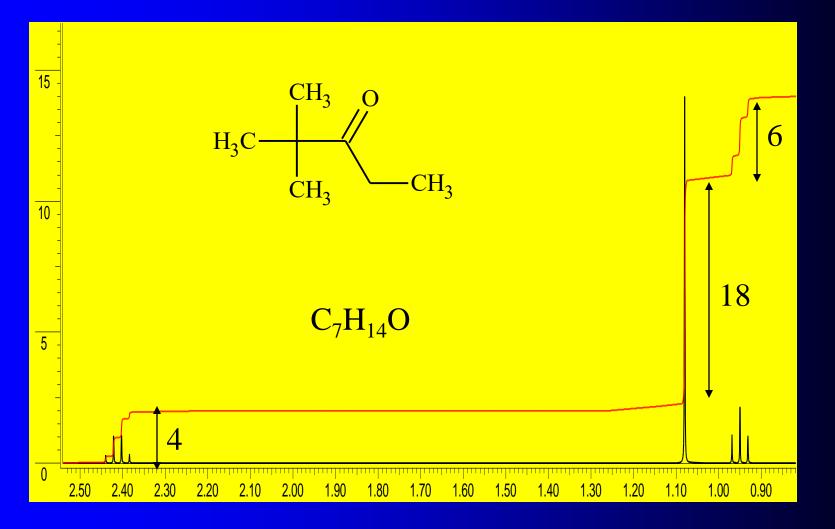
Chemical Shift - ¹H-NMR

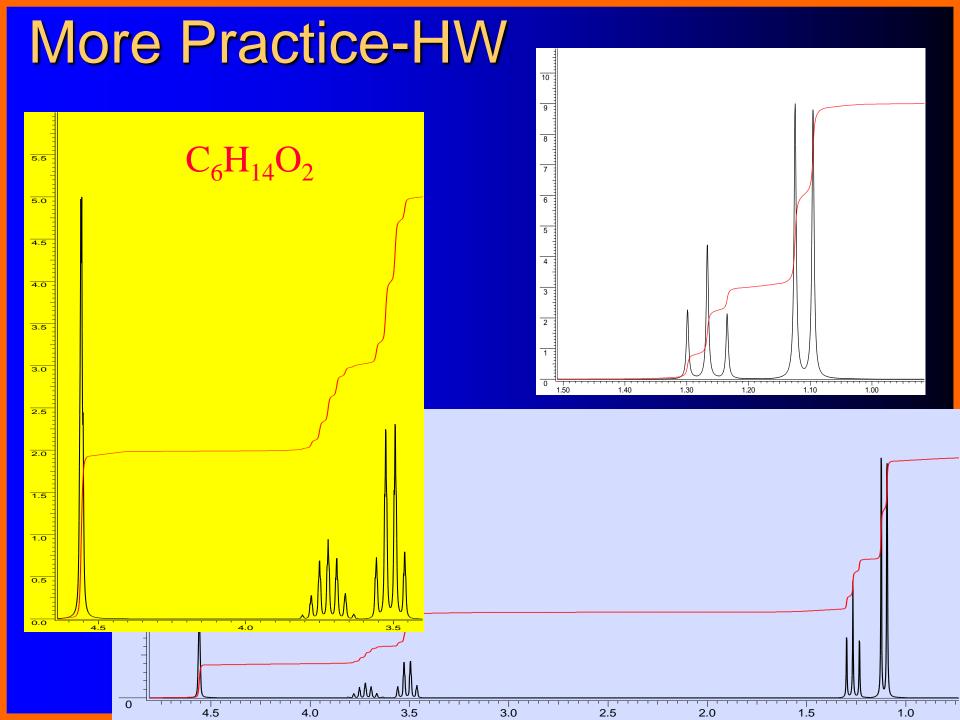
Type of H	δ	Type of H	δ
(C H ₃) ₄ Si	0	ROH	0.5-6.0
RCH ₃	0.9	RCH ₂ OR	3.3-4.0
RCH ₂ R	1.2-1.4	R ₂ NH	0.5-5.0
R ₃ CH	1.4-1.7	O	
R ₂ C=CRC HR ₂	1.6-2.6	RCCH3	2.1-2.3
RC≡CH	2.0-3.0		
ArC H ₃	2.2-2.5	RČCH ₂ R	2.2-2.6

Chemical Shift - ¹H-NMR

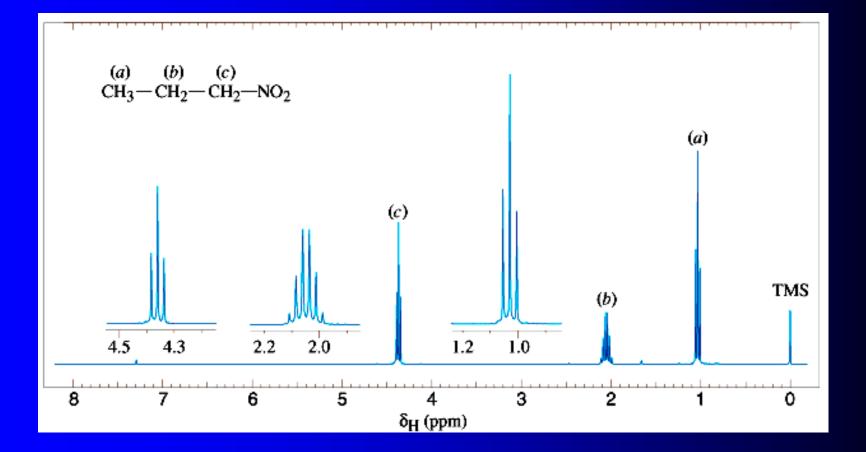
Type of H	δ	Type of H	δ
0			
RCOC H ₃	3.5-3.9	$R_2 C=C H_2$	4.6-5.0
O II		R ₂ C=C HR	5.0-5.7
RÖOC H ₂ R	4.1-4.7		
RCH ₂ I	3.1-3.3	ArH	6.5-8.5
RCH ₂ Br	3.4-3.6	RCH	9.5-10.1
RCH ₂ CI	3.6-3.8	Q	
RCH ₂ F	4.4-4.5	RCOH	10-13

Sample spectra

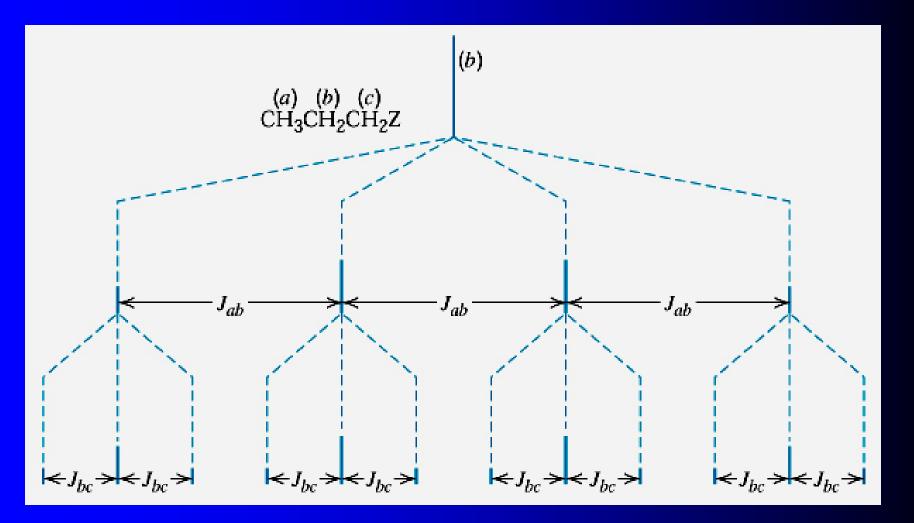




CH₃CH₂CH₂NO₂



Multiple interactions of *non-equivalent* neighbors here J_{ab} > J_{bc}



CICH₂CH₂CH₂CH₂CI

