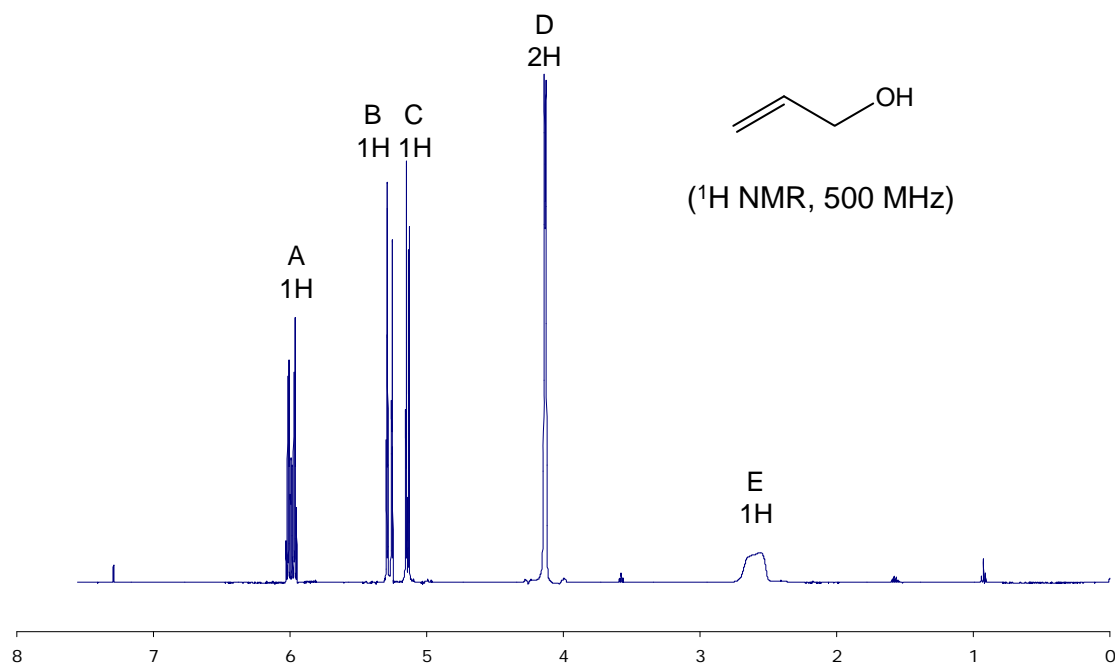
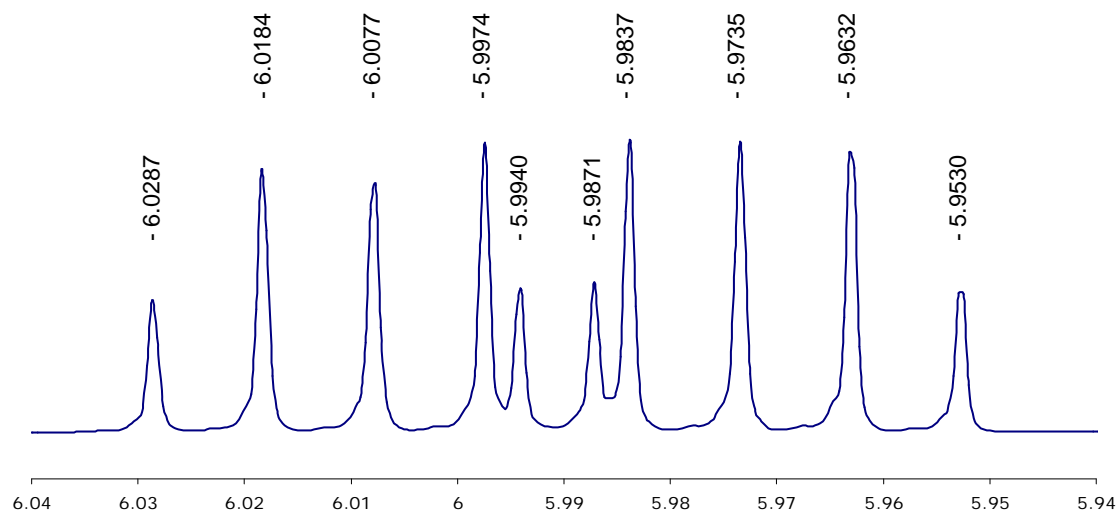


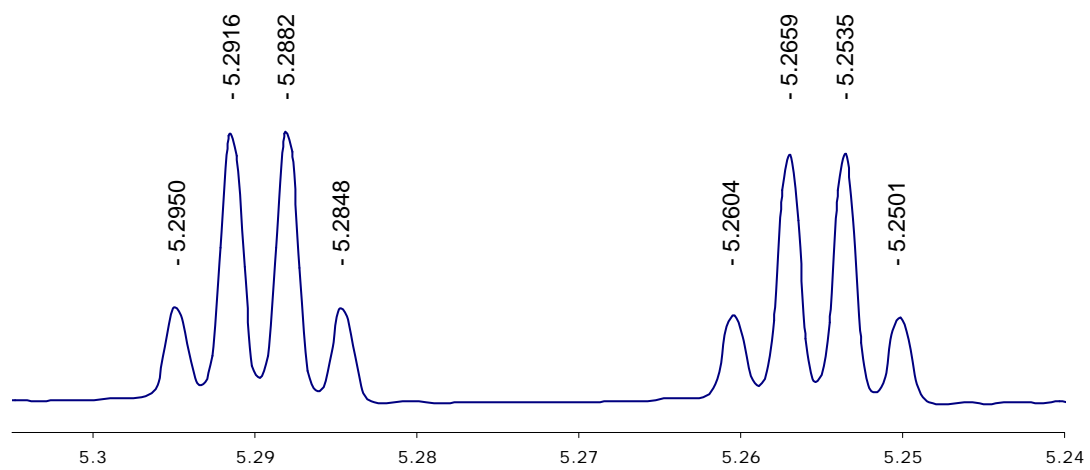
How Do We Deal with Many J 's?



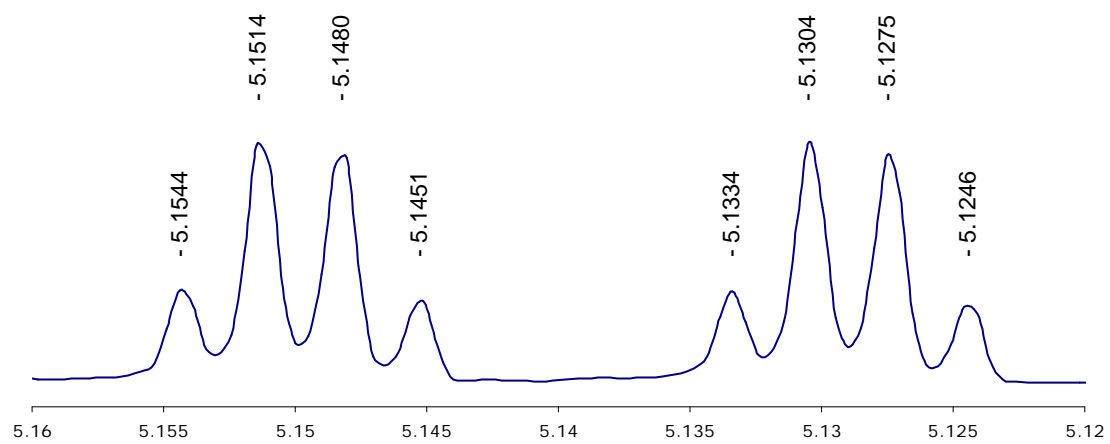
H_A : 1H



H_B: 1H



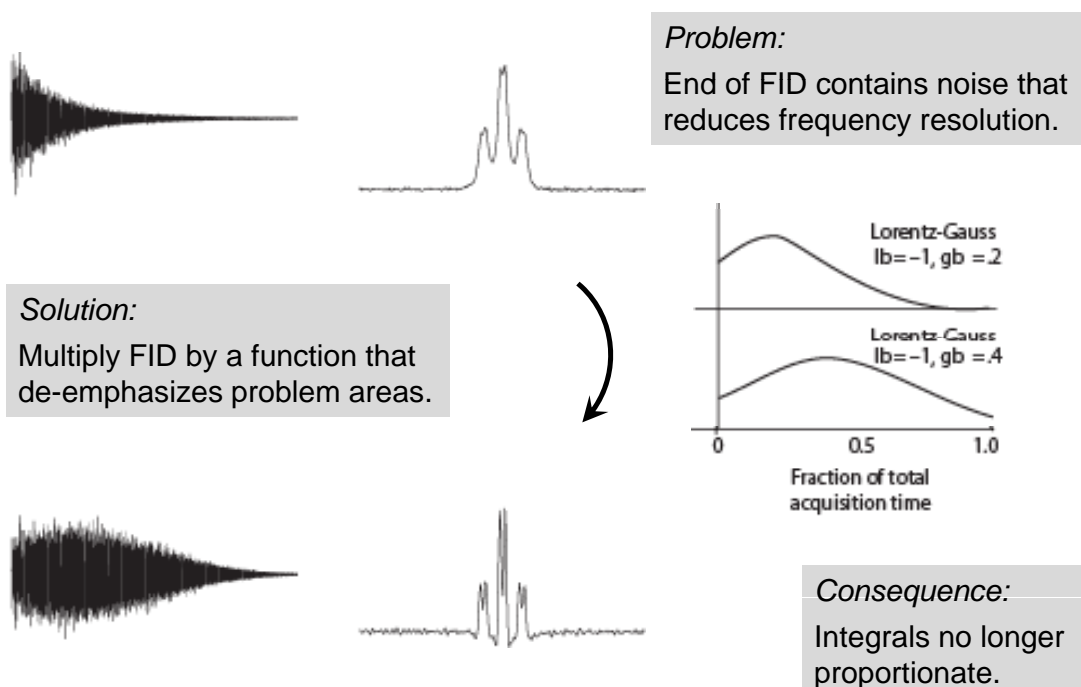
H_C: 1H



Shortcut for Determining J Values

1. Determine relative peak sizes
 - First peak is always intensity 1
 - Number all positions in order (a peak with intensity of 2 will have 2 numbers, etc.)
 - Sum of peak sizes must add up to 2^n
 - n = number of J values
2. Distance from 1 \rightarrow 2 = J_1
3. 1 \rightarrow 3 = J_2
4. Cross out number representing $J_1 + J_2$
5. 1 \rightarrow $n + 1$ = J_n
6. Cross out all additive combinations of $J_1, J_2 \dots J_n$ that haven't been crossed out yet (should be 2^{n-1} combinations)
7. Repeat 5, 6 ...

Resolution Enhancement with Window Functions



Resolution Enhancement with Window Functions

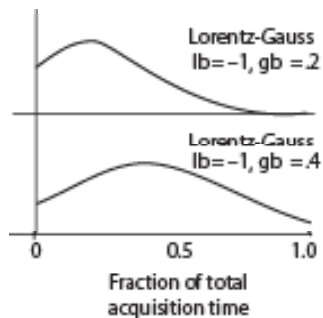


Problem:

End of FID contains noise that reduces frequency resolution.

Solution:

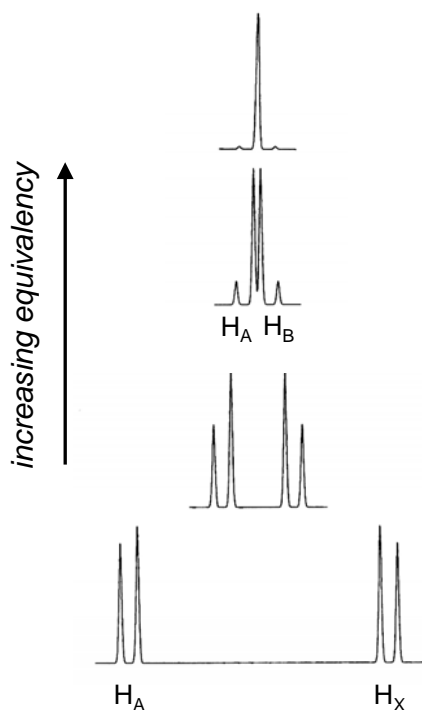
Multiply FID by a function that de-emphasizes problem areas.



Consequence:

Integrals no longer proportionate.

Spectral Distortions at Low $\Delta\nu/J$

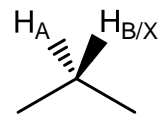


$\Delta\nu/J = 0.4$

$\Delta\nu/J = 1.0$

$\Delta\nu/J = 4.0$

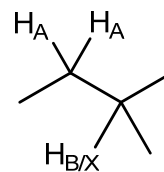
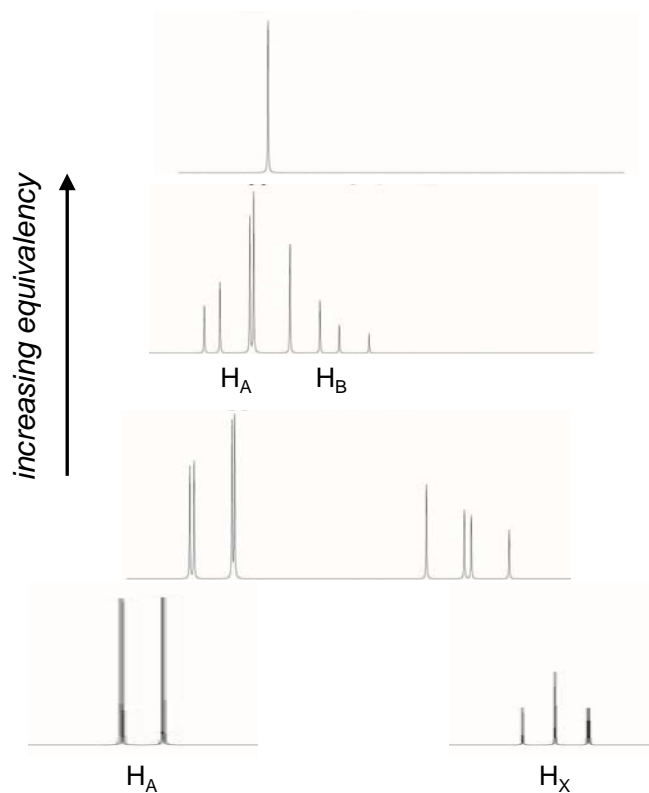
$\Delta\nu/J = 15.0$



- Say H's are coupled, inequivalent, but not very inequivalent.
- Multiplet intensities are distorted towards each other.
- As long as this is the only issue, J can sometimes still be determined by inspection.

Lambert has more discussion.

Spectral Distortions at Low $\Delta\nu/J$



- But sometimes multiplets get too complicated to evaluate J s.

Important takeaway:
Inequivalent protons with identical chemical shift do not split one another.