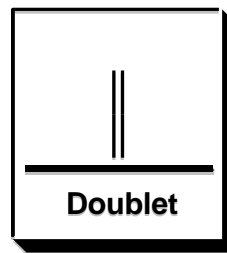
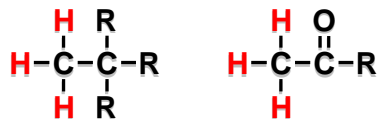
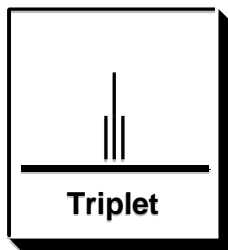
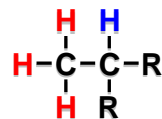


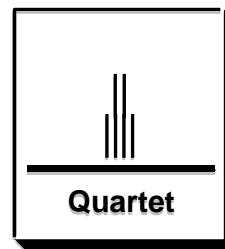
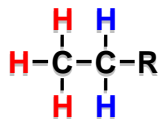
- No neighboring (adjacent) hydrogens



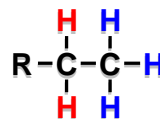
- One neighboring (adjacent) hydrogen



- Two neighboring (adjacent) hydrogens

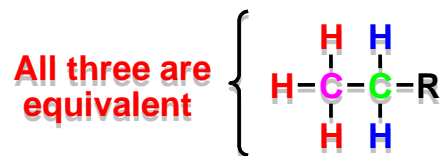
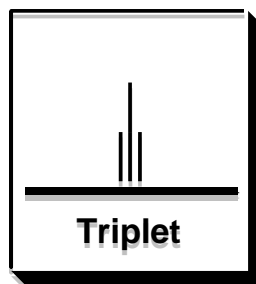


- Three neighboring (adjacent) hydrogens



**H** = Hydrogens that we are interested in  
**H** = Hydrogens that we use for the  $n+1$  rule

**Note:** the number of lines in the peak **does not** refer to the number of hydrogens; a singlet does not mean 1 hydrogen; a double does not mean 2 hydrogens

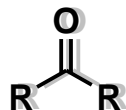


- Pick a specific carbon with hydrogens
- Look at adjacent carbon(s) and count the number of hydrogens...then add 1
- The  $n + 1$  value refers to the peak arising from the carbon with hydrogens you picked

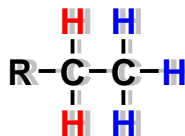
$$n + 1 = 2 + 1 = 3 \text{ (triplet)}$$

In this case, the triplet refers to signal (peak) from the three red hydrogens

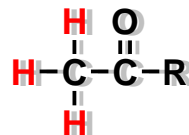
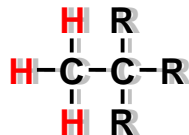
We have 1° of unsaturation and oxygens: carbonyl



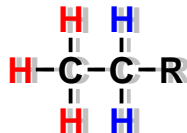
We have a quartet at ~ 4 ppm: these **hydrogens** must be next to **3 hydrogens**

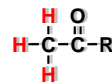
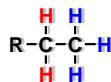
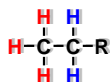


We have a singlet at ~ 2 ppm: these **hydrogens** are not next to any hydrogens

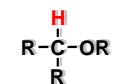


We have a triplet at ~ 1 ppm: these **hydrogens** must be next to 2 hydrogens



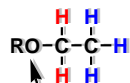


Where does the extra oxygen go? Let's look at our NMR charts:

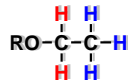


H = 3-4 ppm

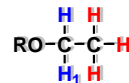
Therefore, we must have:



gives rise to 4 ppm shift

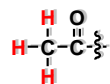


quartet (n+1)

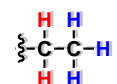


triplet (n+1)

Now, we have used 2 carbons, 5 hydrogens, and one oxygen. All we have left to figure out is the remaining 2 carbons, 3 hydrogens, and 1 oxygen. After our analysis, we know we have a carbonyl and a singlet peak at 2 ppm. Let's see how to put this together:



singlet (n+1)



(from above)

