Quick Guide to Interpreting an EI Mass Spectrum

This guide is based on McLafferty and Turecek, *Interpretation of Mass Spectra*, which is available in the WCU library. Please consult it for more detail.

Before you begin, convert your spectrum from raw abundance (counts) into relative abundance (% of base peak).

1. Look for the M.+ (molecular ion) peak.

The molecular ion, sometimes called the *parent peak*, is the highest mass peak of significance in the spectrum (except for the $M^{+} + 1, \ldots$, etc. peaks). It represents the unfragmented molecule, and can therefore tell you the *monoisotopic* mass of the analyte. Not all compounds yield an M^{+} peak; some are too unstable. Some compounds yield very abundant M^{+} peaks; this stability can be a clue to the structure.

2. Use M.++1 peak to figure out C count.

Zoom in on the M^{·+} peak and look for a very small, but real, peak 1 m/z unit above your molecular ion peak. This peak is usually present for small, organic molecules and represents the fraction of the molecules containing one $^{1}3$ C atom. Take the ratio of these two abundances and divide by the natural abundance of $^{1}3$ C to find the approximate number (+/- 1) of carbon atoms in the molecule. (Or, remember, a ratio of $^{1}3$ C atom.)

The $M^{+}+1$ peak must be the highest mass in the spectrum, an odd-electron ion, and give logical, neutral losses (see 4 and 6 below). If you don't see an obvious $M_{-}+1$ peak or see multiple peaks in distinct patterns, you may have halogens or other non-C,H,O,N atoms in your molecule.

3. Apply the nitrogen rule.

Is M.+ odd or even? For small, organic molecules an even M.+ strongly suggests an even number of nitrogen atoms (0 is even), and an odd molecular weight (M.+) suggests an odd number of nitrogen atoms.

4. Calculate rings and double bonds (degree of unsaturation).

 $C_xH_yN_zO_n$ will have x-0.5y+0.5z+1 rings and/or double bonds. Integers represent odd-electron ions; results ending in 0.5 represent even-electron ions.

5. Look for the base peak.

The base peak is the largest peak in the spectrum (relative abundance = 100%). It represents the most stable fragment (keeping in mind that some "fragments" are actually rearrangements).

6. Calculate the identity of each peak using steps 2-4.

Attempt to identify any peaks possible. Look for logical losses (e.g. -OH or -CH₃) and series of logical losses (e.g. -CH₃, -CH₂). Look for isotopic abundances of fragments and assign compositions. The nitrogen rule still applies to fragments, but keep in mind any odd-mass element could cause the same result. When you've identified as many peaks as you can, attempt to deduce a structure.