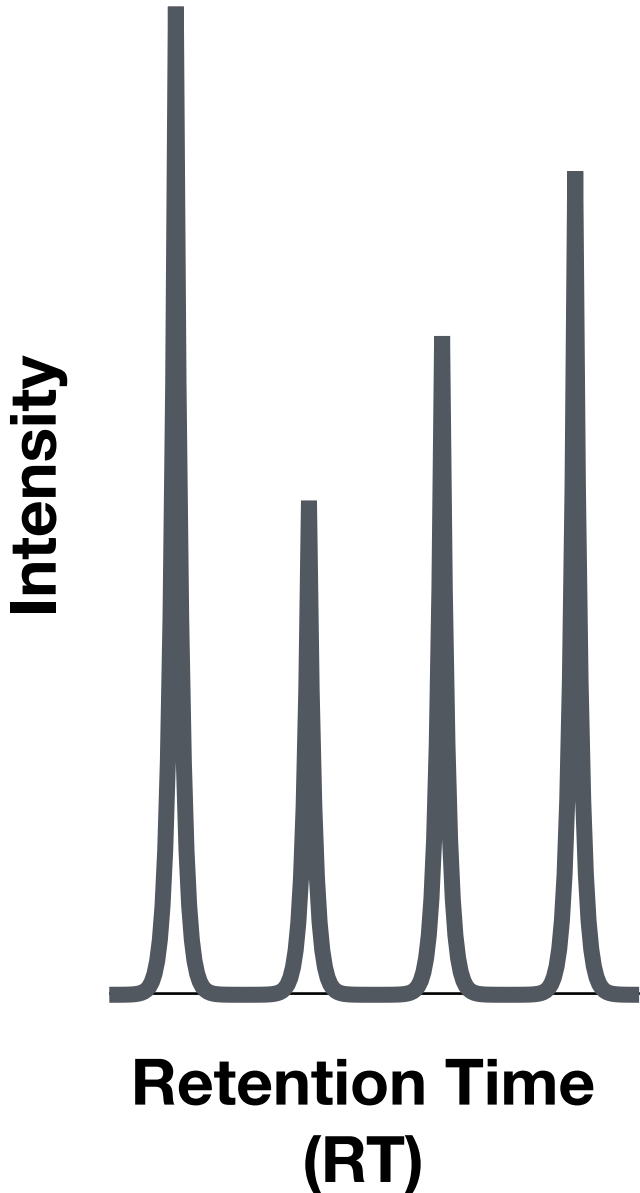


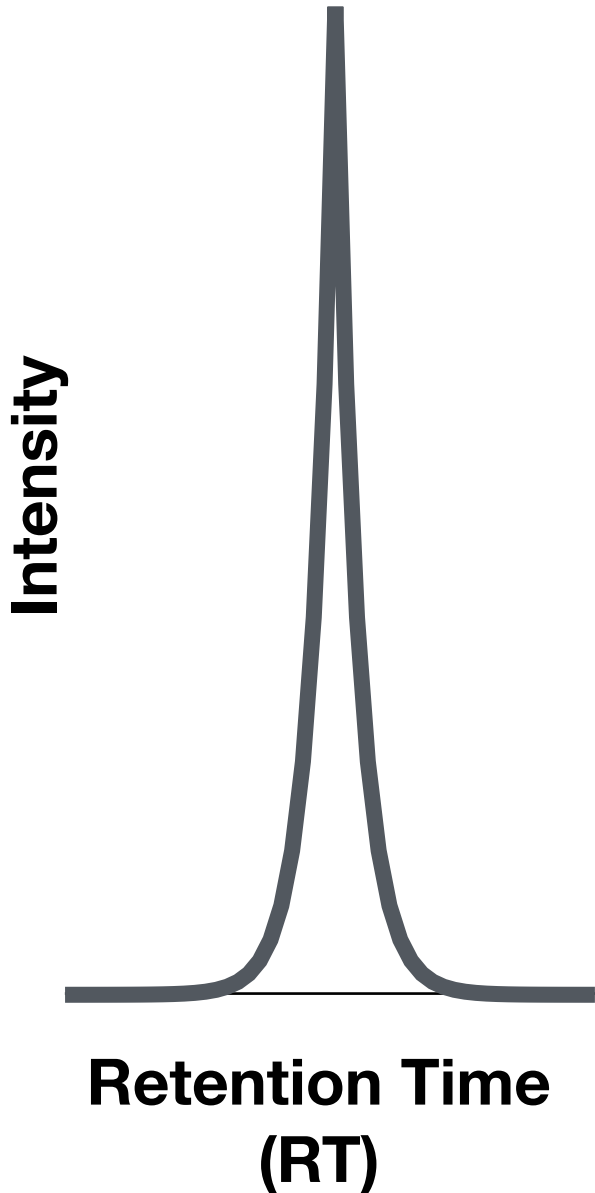
Chromatogram:

2 x 1D array
Intensity = f(RT)



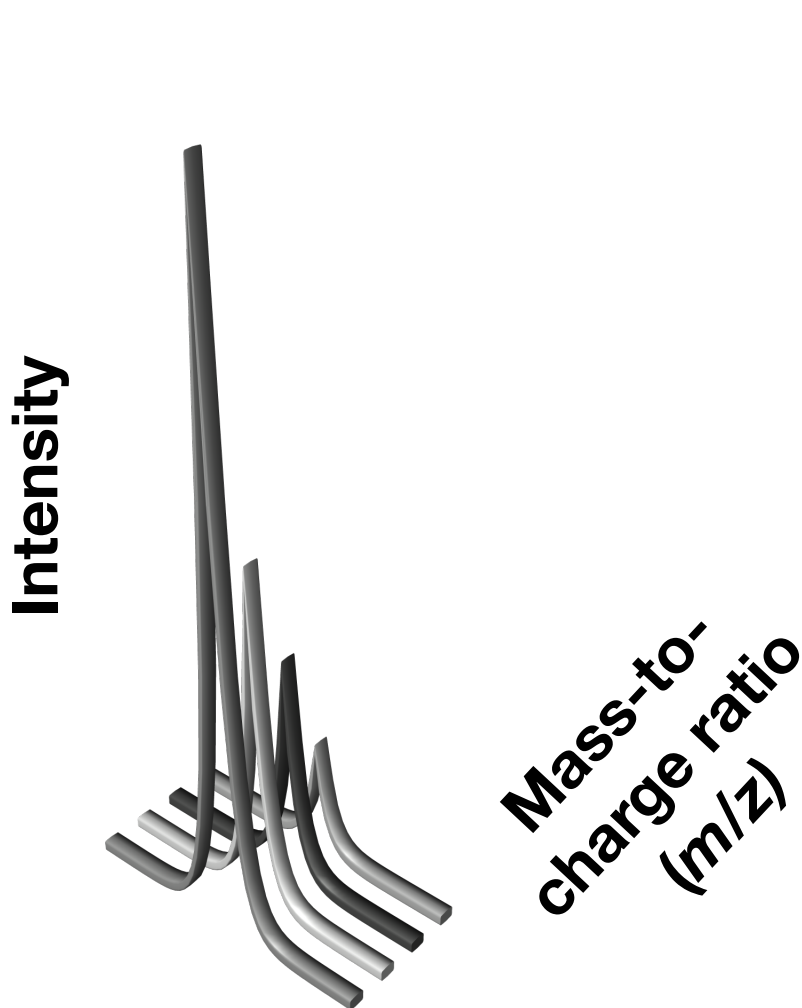
Peak:

2 x 1D array
Intensity = f(RT)



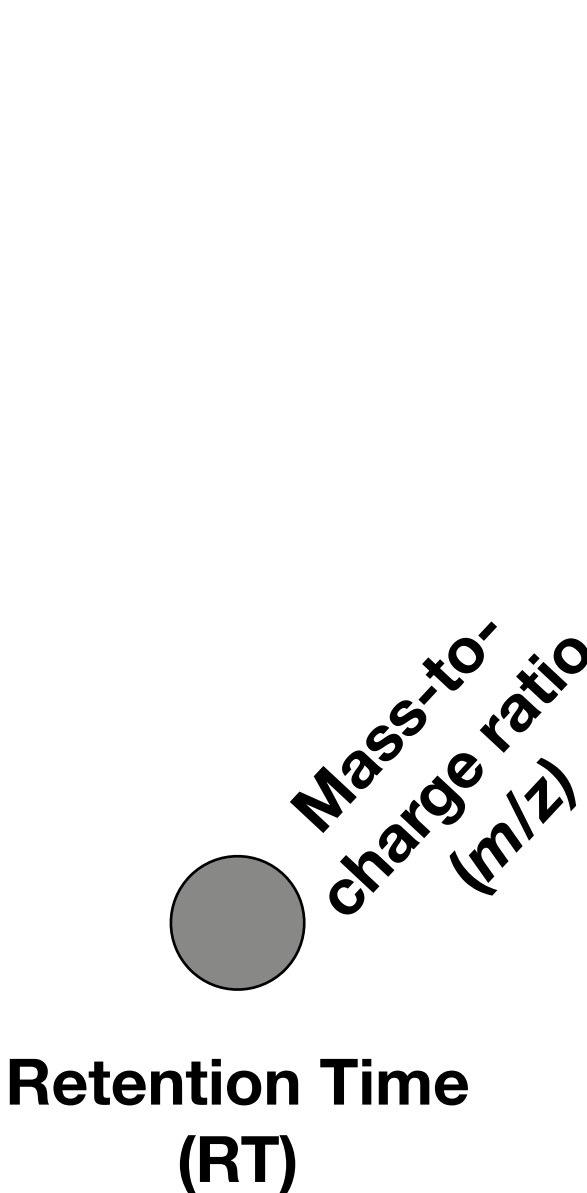
Spectrum:

2 x 1D array
Intensity = f(*m/z*)



Feature:

2 x pointer
RT @ *m/z*



Feature list (or table):

2D array

ID	RT	<i>m/z</i>	1D array of intensities	2D array of spectra
1	123.456	123.4567	(intensity@sample1, intensity@sample2, ...)	((spectrum1@sample1,...), (spectrum1@sample2), ...)
...