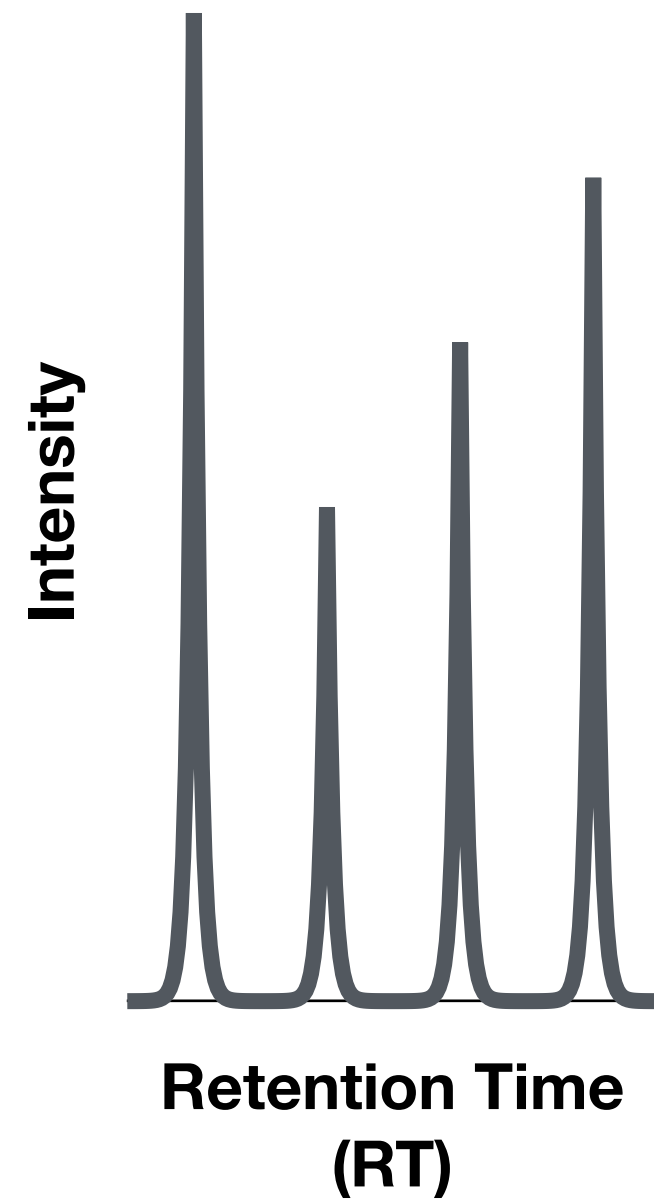


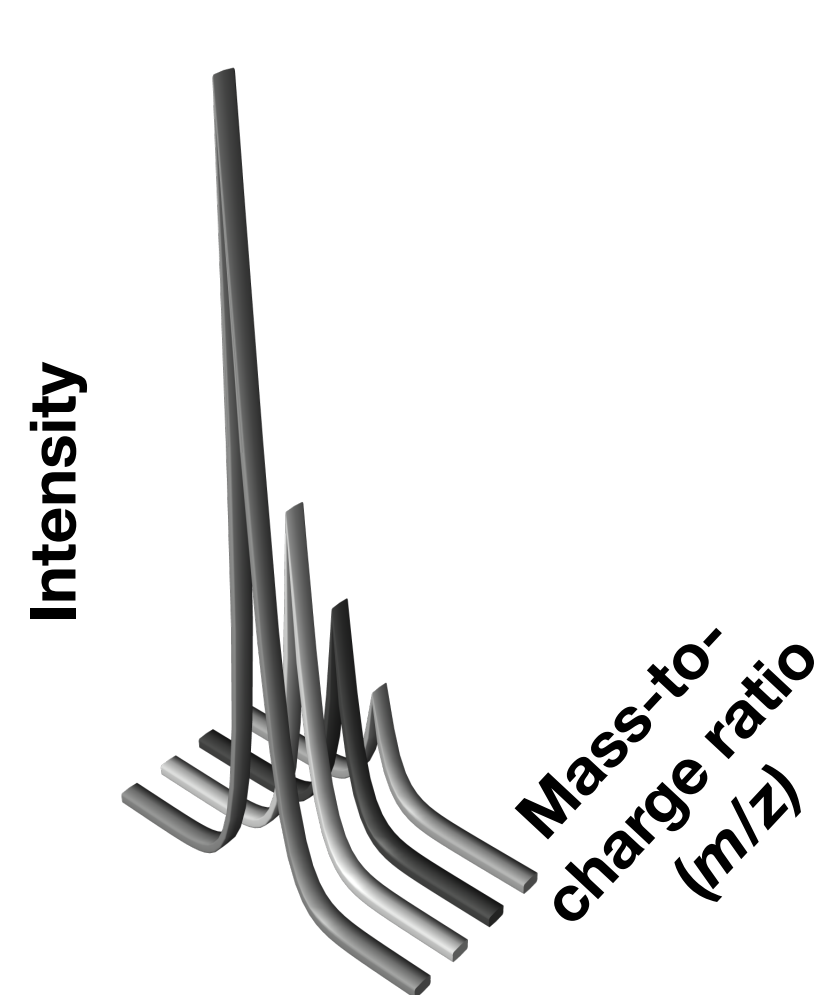
Chromatogram:

Intensity = f(RT)



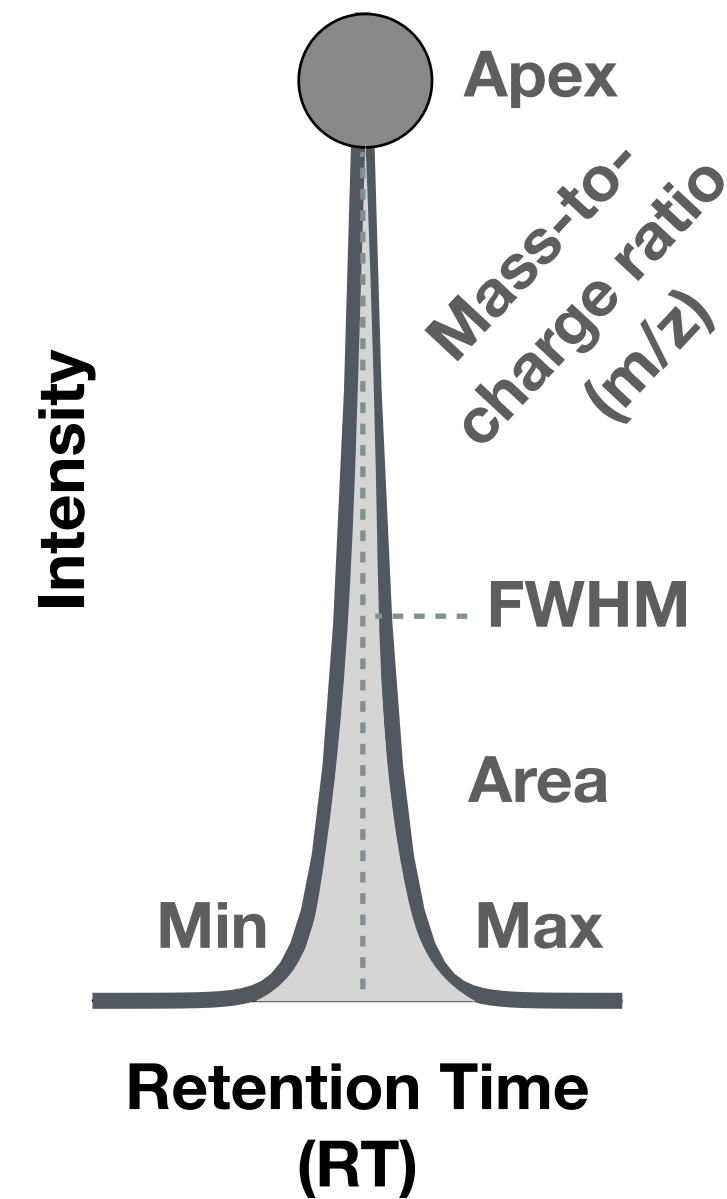
Spectrum:

Intensity = f(*m/z*)



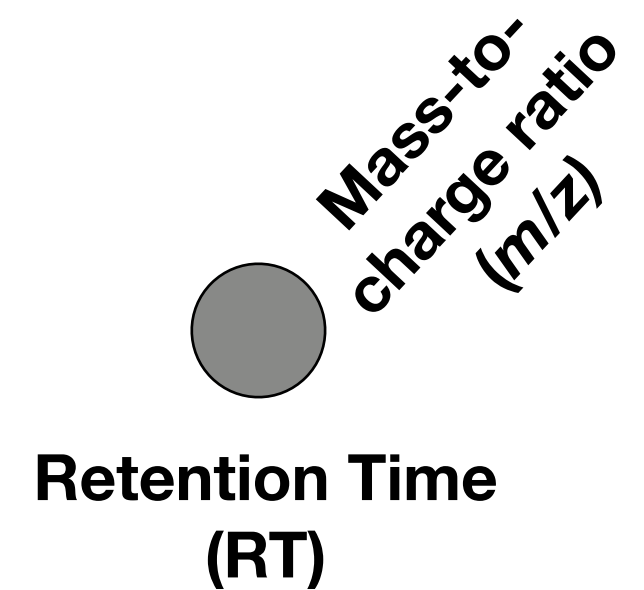
Peak:

Parameters with intensity, (*m/z*) and RT



Feature:

RT @ *m/z* pointer



Feature list (or table):

ID	RT	<i>m/z</i>	Peak area	Associated spectrum (or spectra)
1	123.45	123.4567	9876543.21	
...