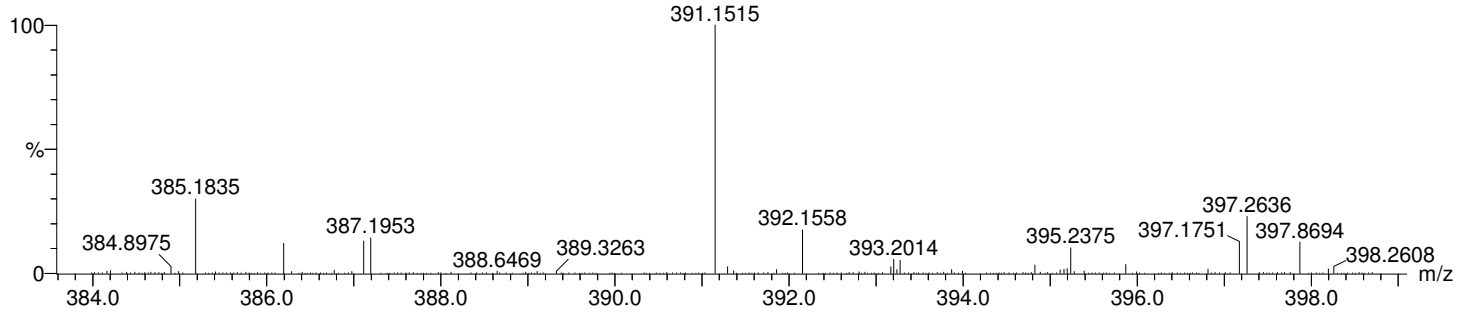


Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 120.0  
Element prediction: Off  
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions  
10 formula(e) evaluated with 1 results within limits (up to 20 closest results for each mass)  
Elements Used:  
C: 0-22 H: 0-26 O: 0-5 Na: 0-1  
HBr7  
UNE BGreatrex HBr7 40 (0.974) Cm (38:58)

1: TOF MS ES+  
2.43e+004



Minimum: -1.5  
Maximum: 55.0 5.0 120.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	i-FIT (Norm)	Formula
391.1515	391.1521	-0.6	-1.5	10.5	335.5	0.0	C22 H24 O5 Na