

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

5 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)

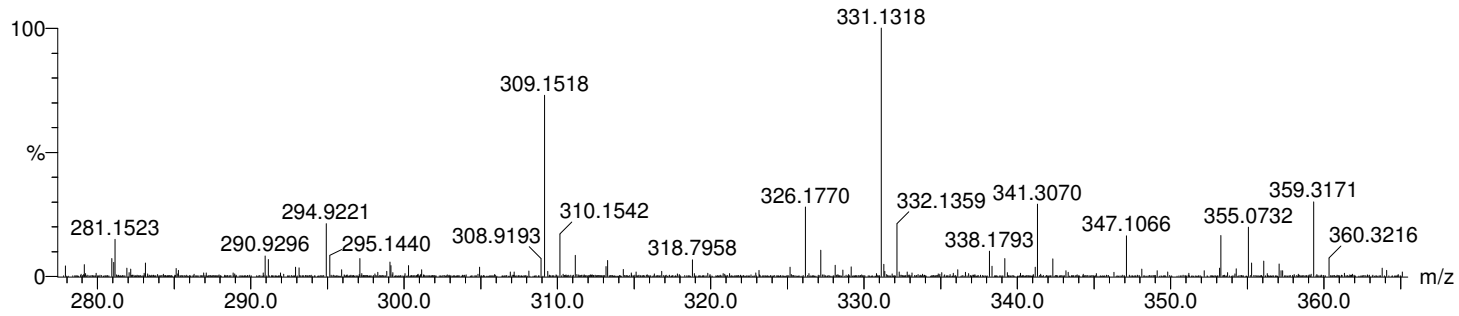
Elements Used:

C: 8-20 H: 5-20 O: 0-3 Na: 0-1

CyrBn2Rlactone

UNE BGreatrex CyrBn2Rlactone 81 (1.929) Cm (81:85)

1: TOF MS ES+
5.90e+003



Minimum: -1.5
Maximum: 55.0 5.0 120.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	i-FIT (Norm)	Formula
331.1318	331.1310	0.8	2.4	10.5	206.0	0.0	C20 H20 O3 Na