

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

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

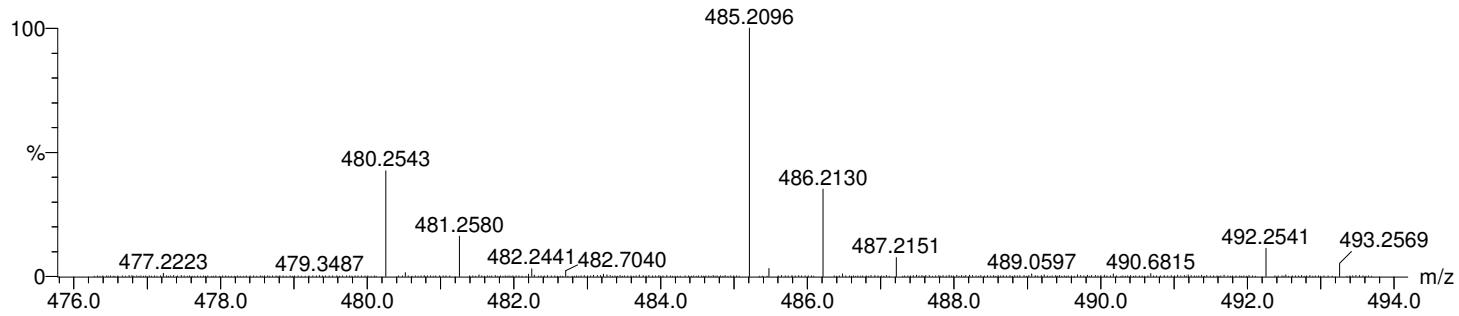
Elements Used:

C: 10-32 H: 5-30 O: 0-3 Na: 0-1

CyrBn2RphOPh

UNE BGreatrex CyrBn2RphOPh 19 (0.479) Cm (19:100)

1: TOF MS ES+
4.54e+004



Minimum: -1.5
Maximum: 55.0 5.0 120.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	i-FIT (Norm)	Formula
485.2096	485.2093	0.3	0.6	17.5	388.9	0.0	C32 H30 O3 Na