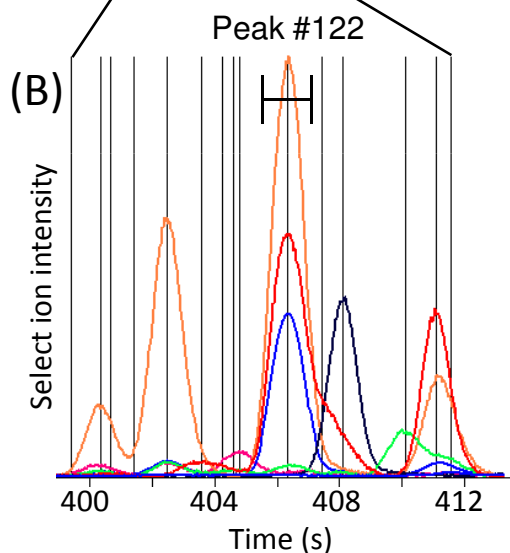
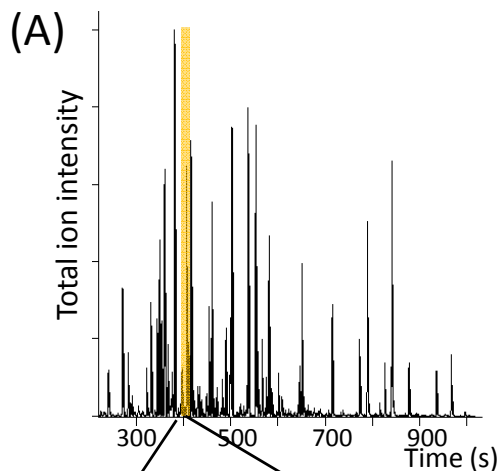
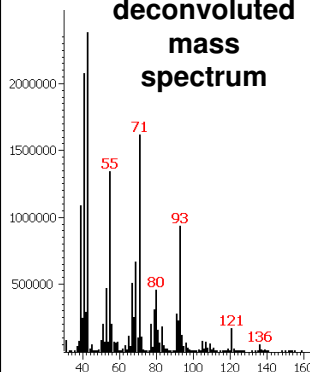




(C)

**Peak #122
deconvoluted
mass
spectrum**



Peak #122 metadata

retention time	406.343 s
unique ion	93
base mass	43
signal/noise	2013
purity	0.1137
apex masses	98+31+39+...

RI filter

8 possible Bin matches

**unique ion
filter**

2 possible Bin matches

