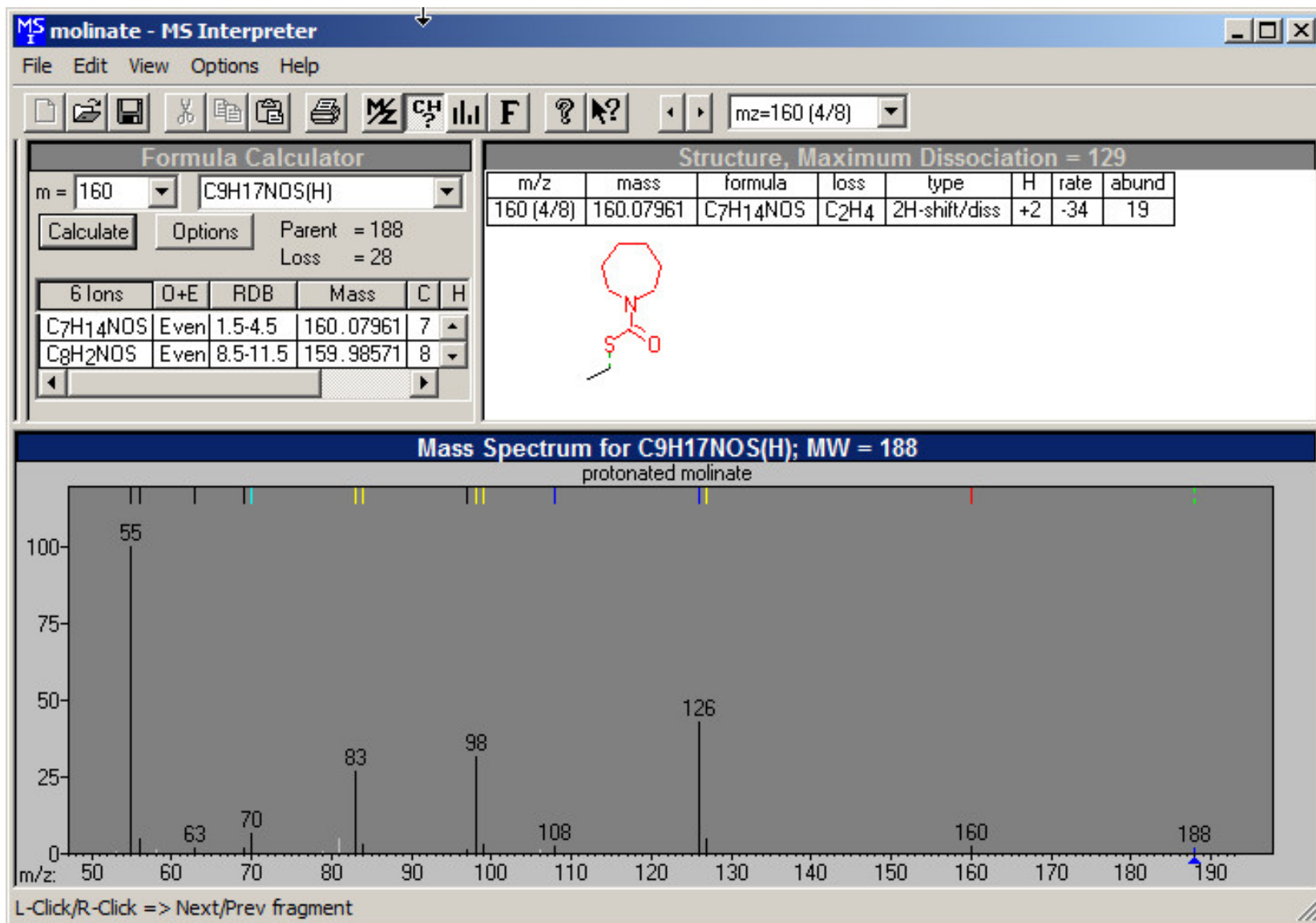


# NIST MS Interpreter Main Window



## NIST MS Interpreter Fragment Options Window

(right click in spectrum part of main window and select fragment options)

The image shows a dialog box titled "Fragment options" with a blue header bar. It contains several checked checkboxes: "Show fragments", "Show relative rate", "Show unspecified cleavage", "Mark Isotope Peaks", and "Protonate molecule". The "Show fragments with no peak" checkbox is unchecked. Below these are two input fields: "Display fragments for peaks not lower than:" with a dropdown menu set to "0" and a "Rate threshold" field set to "40". At the bottom left is a "Sort by" section with three radio buttons: "mz" (selected), "relative rate", and "abundance". At the bottom right is a "Tic coloring" section with five checked checkboxes: "Ring opening", "H-transfer", "Protonated", "Molecular ion", and "Unspecified". At the very bottom are "Cancel" and "OK" buttons.

**Fragment options**

- Show fragments
- Show relative rate
- Show fragments with no peak
- Show unspecified cleavage
- Mark Isotope Peaks
- Protonate molecule

Display fragments for peaks not lower than:  $\% \text{ max}$

Rate threshold

Sort by

- mz
- relative rate
- abundance

Tic coloring

- Ring opening
- H-transfer
- Protonated
- Molecular ion
- Unspecified