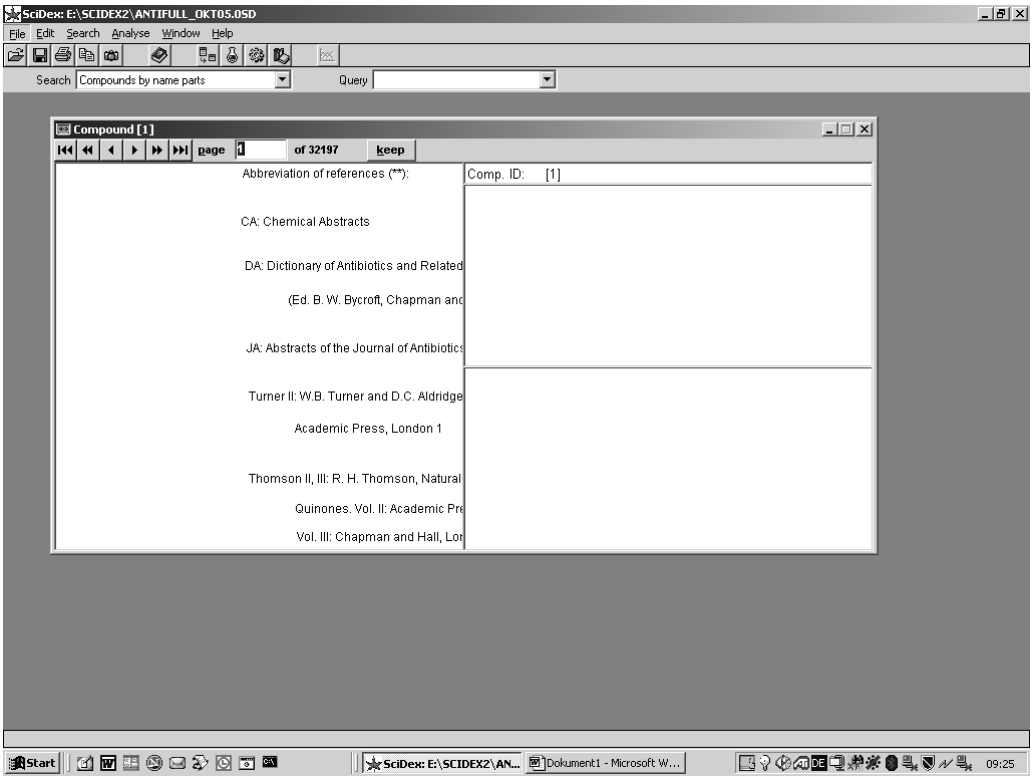
Installation of AntiBase on Scidex:

1) Start the self-extracting AntiBase.exe and install the program.
2) Start the newly installed AntiBase.exe in the respective folder.

Scidex is a database program of Prof. Vill, which distinguishes two types of AntiBase data: In the *list of compounds*, all structure-related data are saved (structures, molecular weight, formula and names). All other data are *properties* and stored there. This results in different search techniques, which will be described now very briefly to give a first introduction. A detailed manual will follow later.
As the program can be used intuitively and is self-explanatory, we strongly recommend just to try it. The data cannot be corrupted.

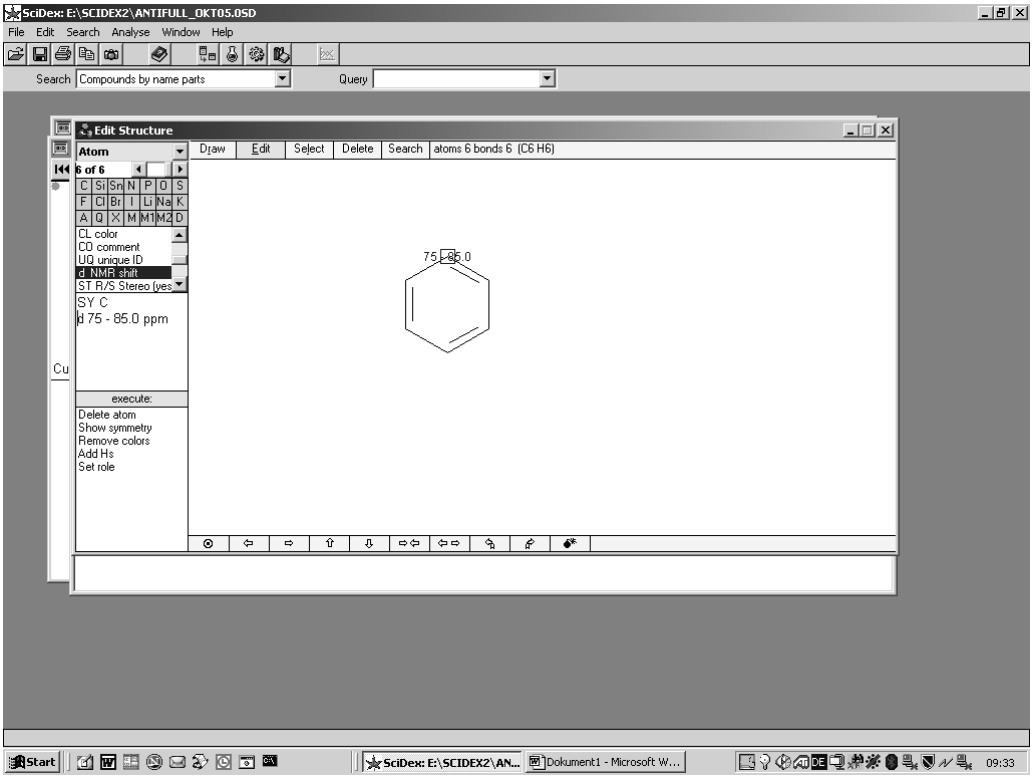
Entrance Screen after starting AntiBase



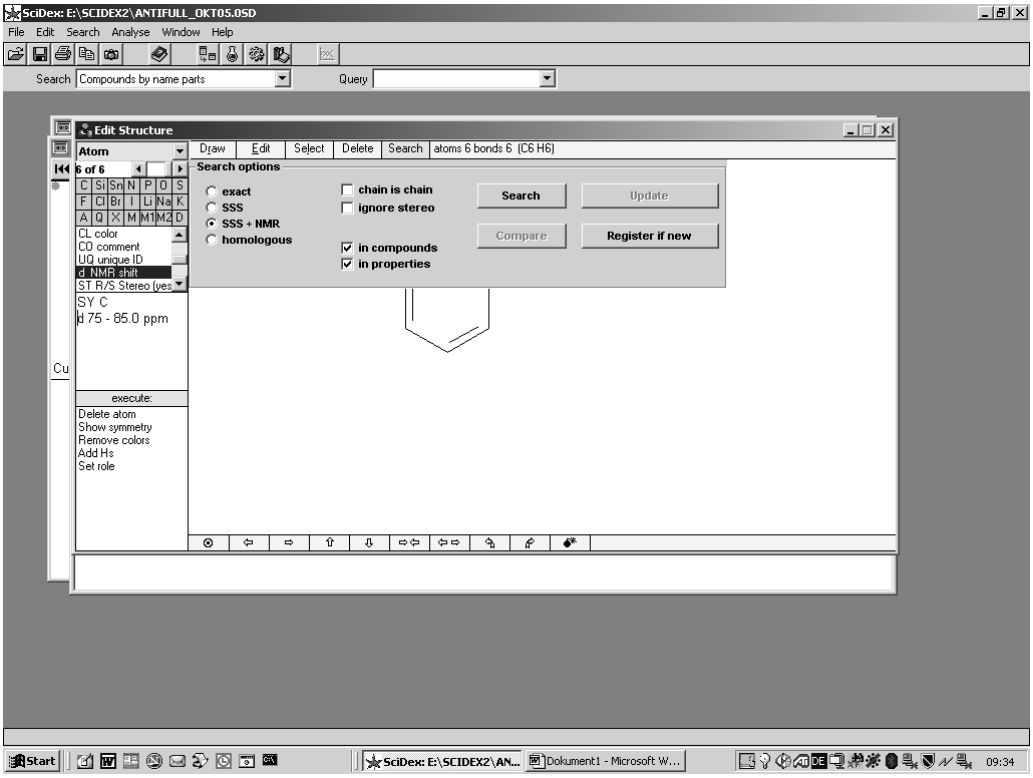
The appearance of the screen should be changed with WINDOW → CONFIGURATION

Starting the structure editor Edit → Edit structure, or CTRL F2

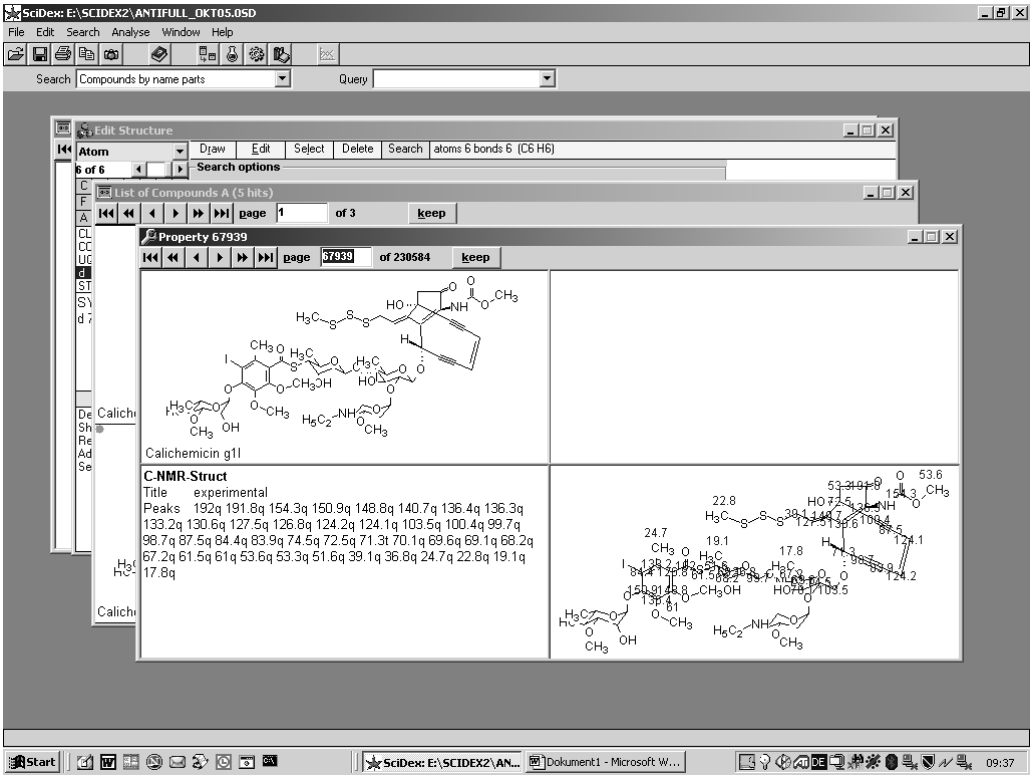
draw a structure, activate EDIT and click on an atom. Select in the tools menue *d NMR shift* and type value:



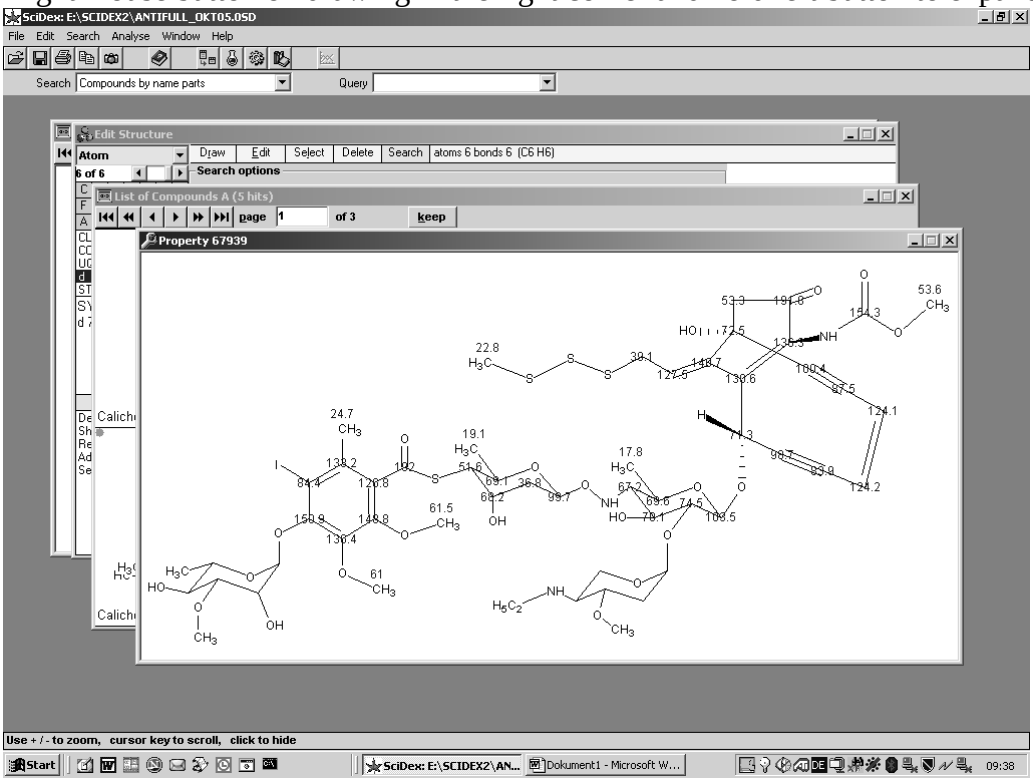
search with *SSS + NMR* and additionally in *compounds* and *in properties*. Search only in *SSS* and *in compounds* to ignore the NMR data.



in the list of results, double click on *C-NMR-Struct. experimenta* to see the data:

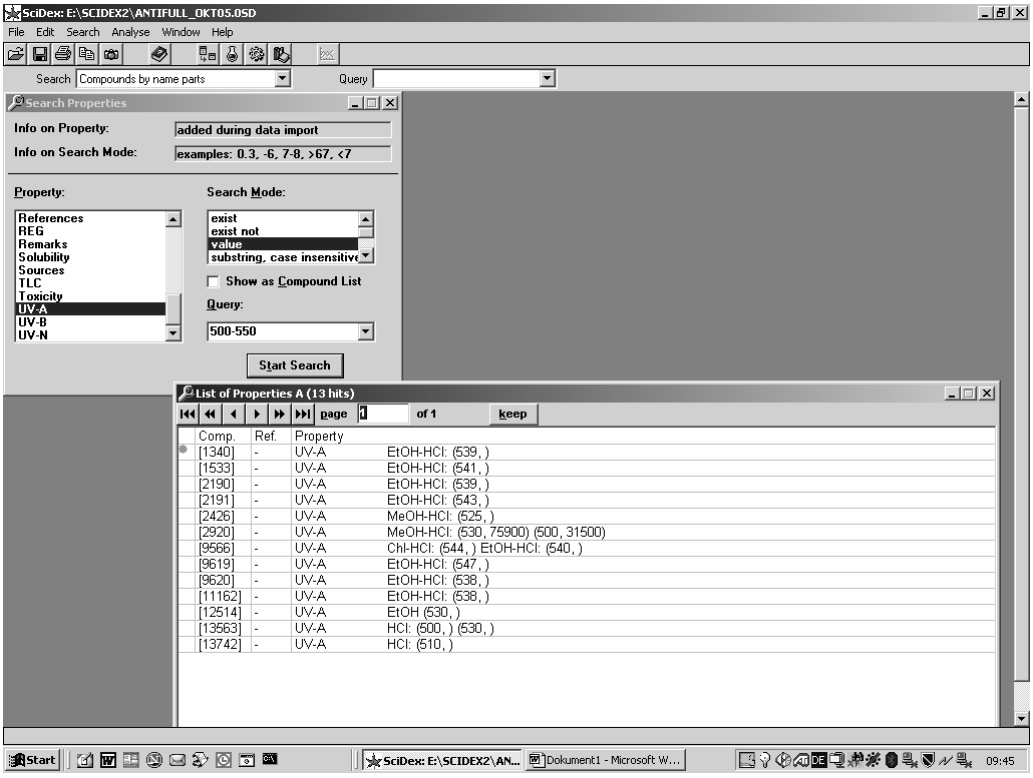


go with right mouse button on drawing in the right corner and hold left button to expand

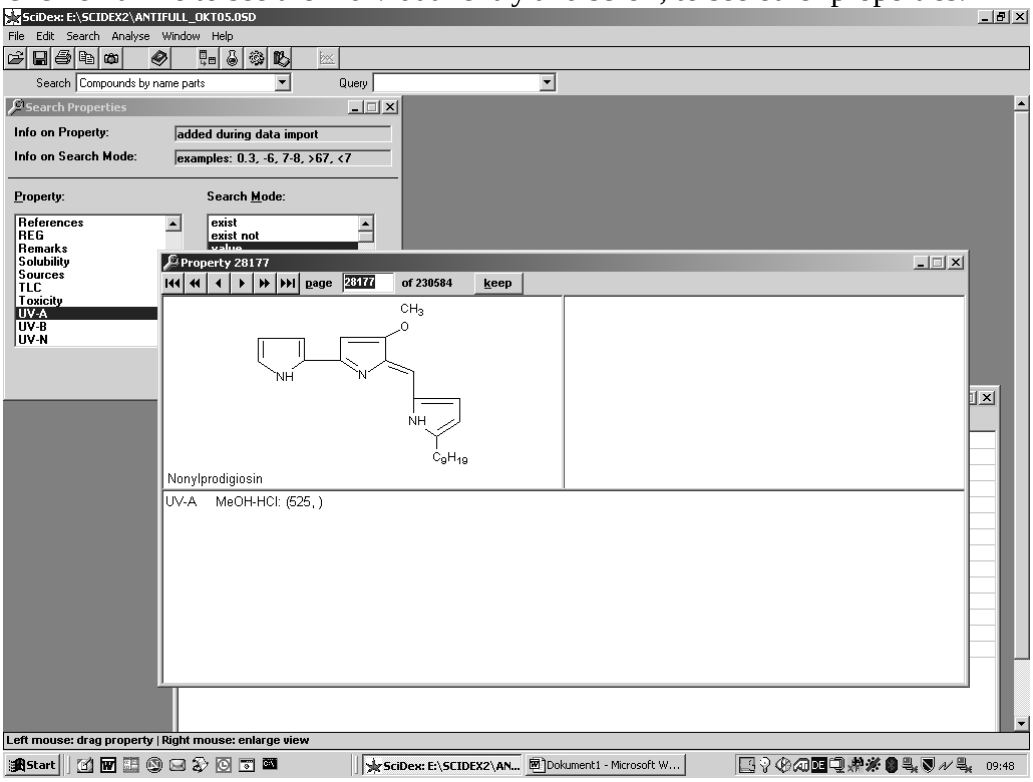


Search Data:

Goto SEARCH → SEARCH PROPERTIES, select the wanted property and the search mode and type values:



double click on a line to see the individual entry and scroll, to see other properties:



Click on *keep* for at least two result lists and then SEARCH COMBINE LISTS to perform logical operations with lists.