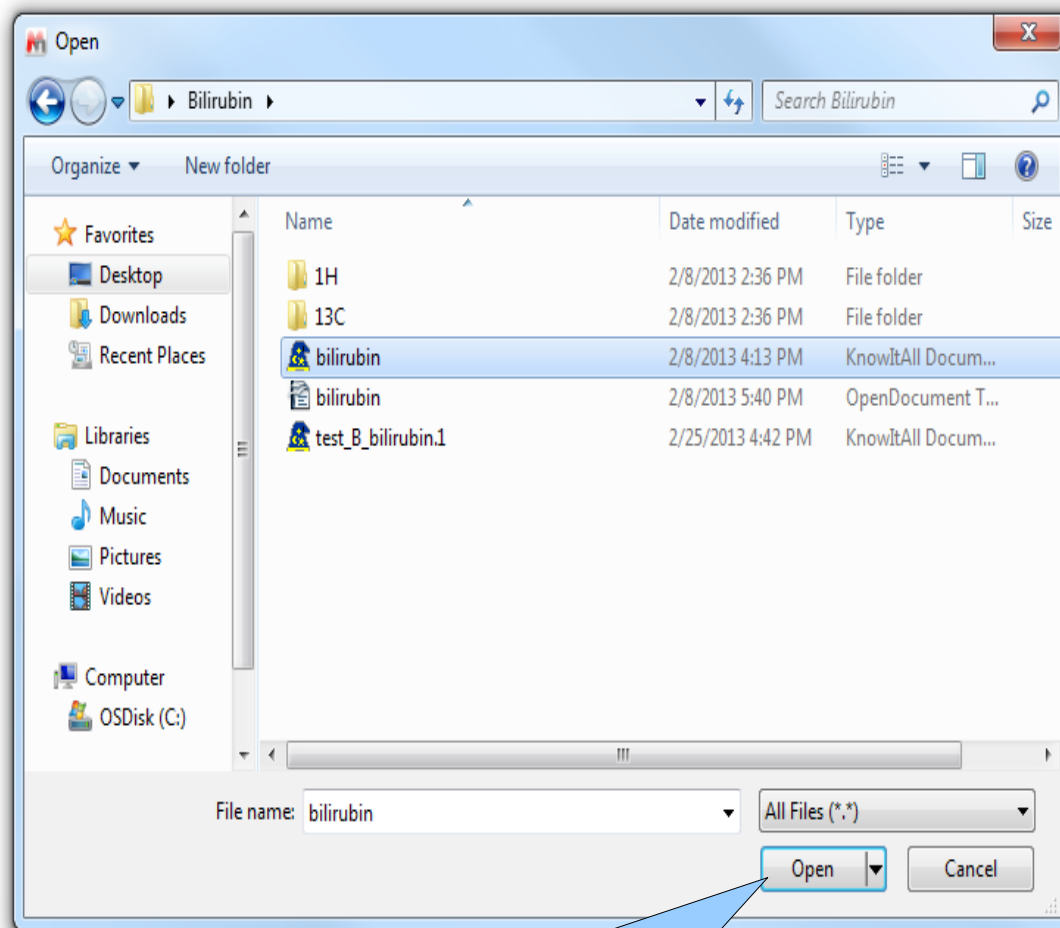


1

Click on 'File' and select 'Open...'

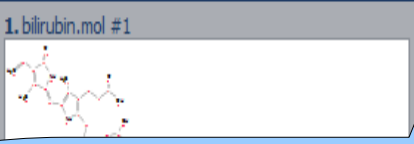


2

Select a structure file (.mol, .cdx, etc...) and click 'Open'

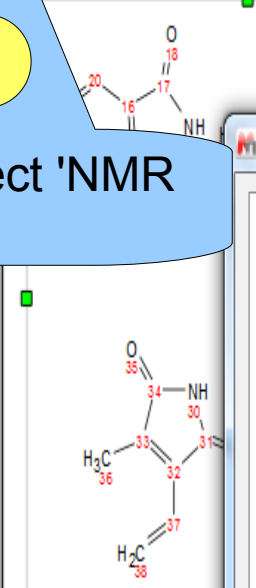
Pages

1. bilirubin.mol #1



1

Click on 'Predict' then select 'NMR Predictor Options'



NMR Predictor Options

1H 13C HSQC 31P 19F 15N 17O 29Si

From: -2.00 ppm

To: 10.00 ppm

Number of Points: 32K

Frequency: 500.13 MHz

Line Width: 0.75 Hz

Solvent: Chloroform-d

Exclude Labile Protons

Predictor: Modgraph NMRPredict Server

Predictor Properties...

Tip: Use the Molecule panel to see all predicted parameters.

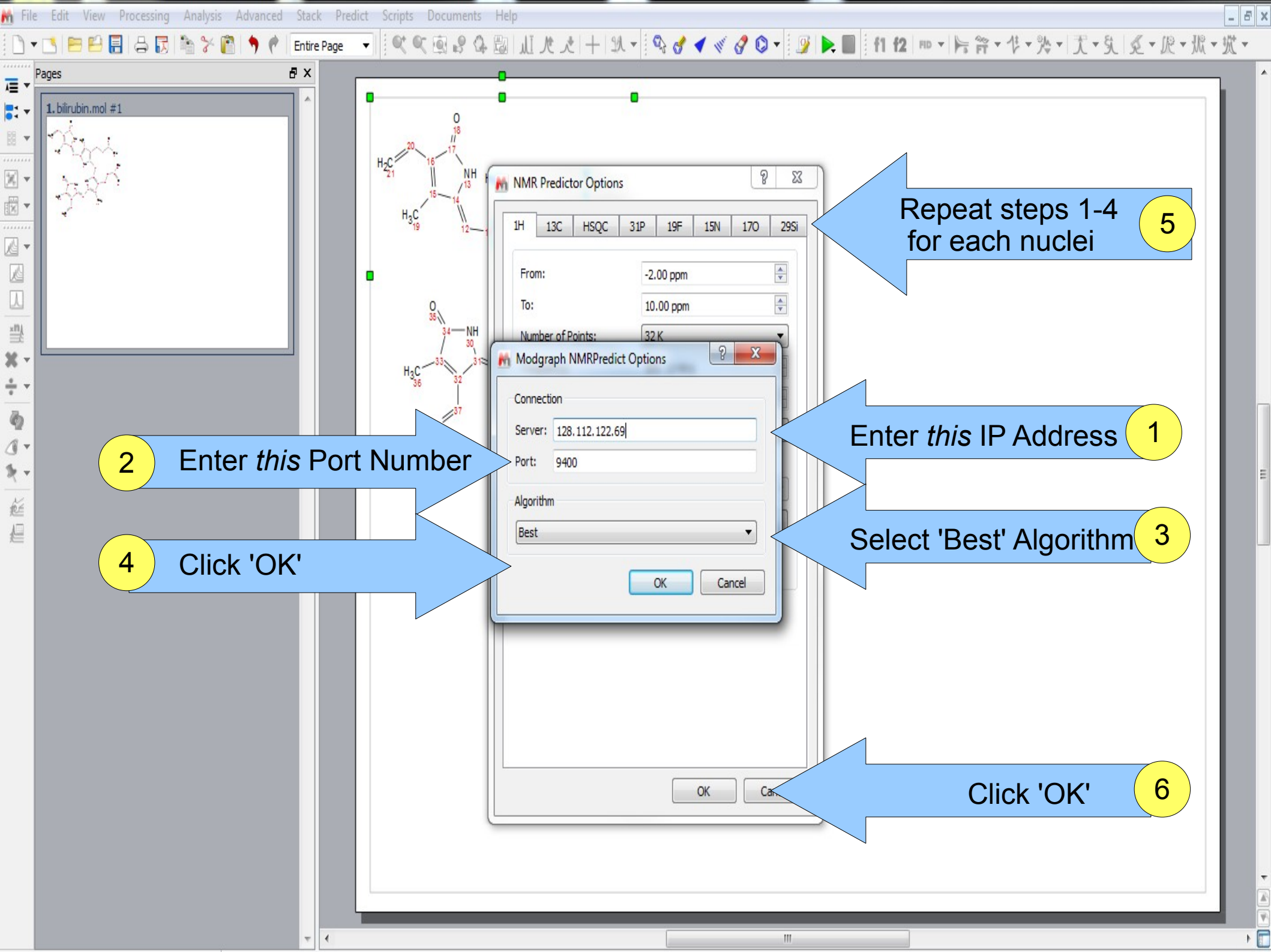
OK Cancel

2

Select 'Modgraph NMRPredict Server'

3

Click 'Predictor Properties...'



Repeat steps 1-4 for each nuclei **5**

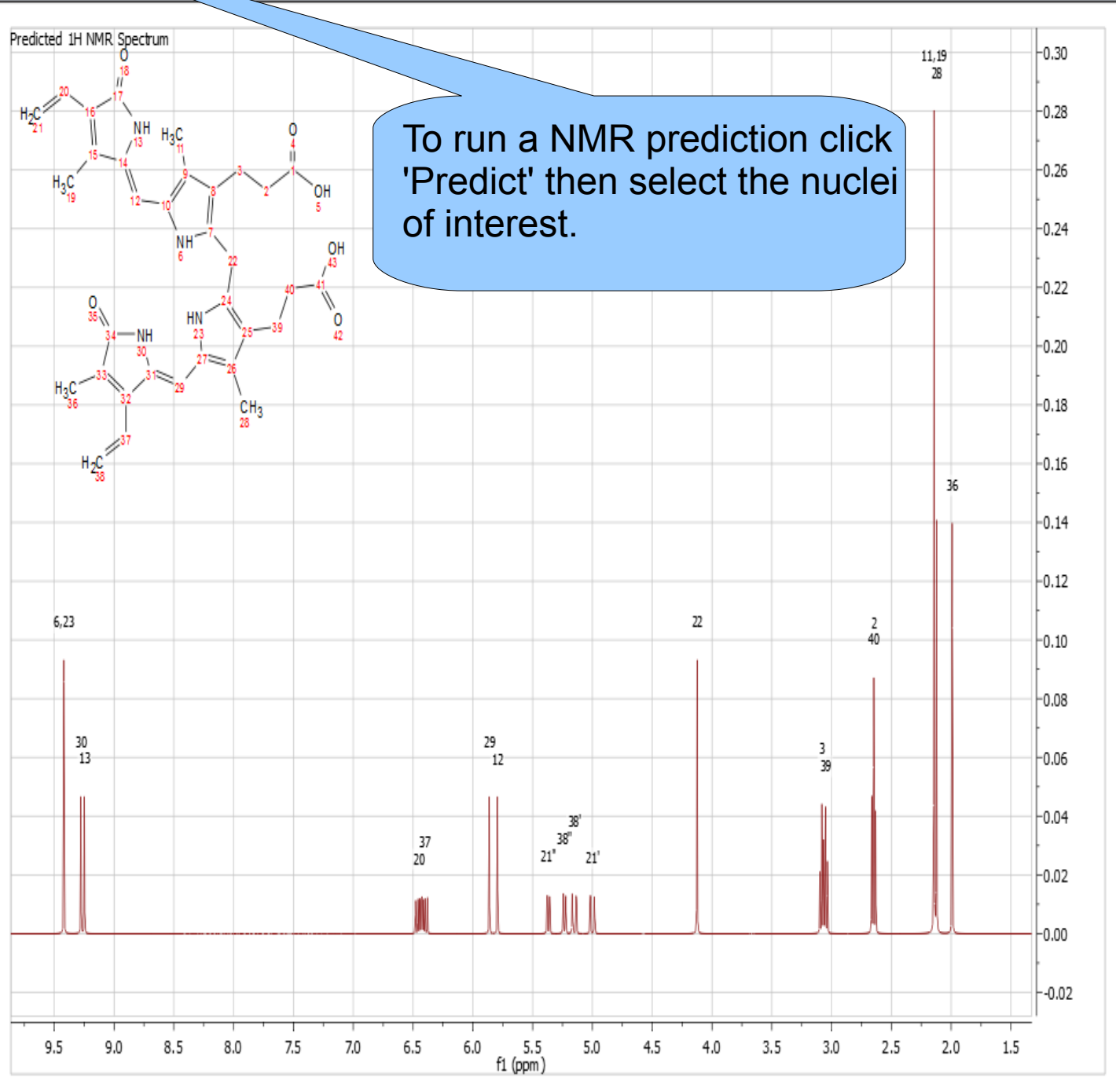
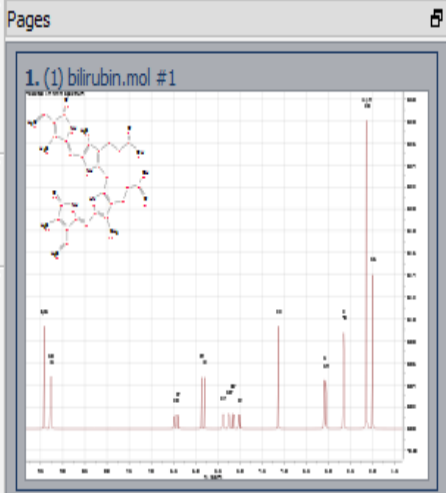
**2** Enter *this* Port Number

Enter *this* IP Address **1**

**4** Click 'OK'

Select 'Best' Algorithm **3**

Click 'OK' **6**



To run a NMR prediction click 'Predict' then select the nuclei of interest.