

## *Bruker MALDI-TOFMS Instructions*

### **Rules**

- 1) No sample preparation should occur in CB 103, prepare the sample plate in your lab
- 2) Do not exceed 200 laser shots/sample. The laser has a fixed number of shots it can do and once we hit it the laser must be replaced.
- 3) Any sample that gets on the instrument should be cleaned up immediately with isopropanol
- 4) When you leave the instrument make sure the sample carrier mechanism is in the **IN** position
- 5) Sign up for your instrumentation time

### **To collect data**

- 1) Launch flexControl from the desktop
- 2) After the software is fully loaded look in the lower right hand corner to determine whether the sample tray is **IN** or **OUT**.

It should be left in the in position when your done analyzing samples for the day.

- 3) click the eject/insert button (lower left side of window looks like a solid triangle over a line) to eject the sample carrier if it is **IN**
- 4) open the sample compartment and place your sample plate in the compartment
- 5) go back to the software and press the eject/insert button
- 6) wait for the insertion process to complete
- 7) load a data acquisition method by clicking the **Select...** button
  - LP : linear positive (enhanced sensitivity for positive ions)
  - LN: linear negative (enhanced sensitivity for negative ions)
  - RP: reflectron positive (enhanced resolution for positive ions)
  - RN: reflectron negative (enhance resolution for negative ions)
- 8) Do not modify or save over existing methods
- 9) using the mouse click on the gray circle where your sample is located, this will move the laser and camera to the sample
- 10) using the mouse in the camera window click on a spot that you want to shoot
- 11) Click the Start button to begin data collection
- 12) if you do not see any ions increase the laser power using the slider to the right of the camera window
- 13) if your baseline is not flat decrease the laser power
- 14) save your data **File → Save Spectrum to File As...**
- 15) repeat steps 9 through 14 until all your samples are analyzed
- 16) eject the sample plate by clicking the eject/insert button
- 17) open the sample compartment , remove the sample plate, and then **gently** close the sample compartment (the latch is made of plastic and will break if you let it free fall)
- 18) press the sample plate eject/insert button to move the sample carrier into the **IN** position and then close the flexControl software

### **To analyze data**

- 1) Launch flexAnalysis from the desktop
- 2) Open your data File → Open Single Analysis...
- 3) Navigate down your data folder until you get to fid file
- 4) Click on **Find Mass List** button to label peaks with their corresponding masses
- 5) Print data by going to **Report → Print...**
- 6) Close the software when you're done