## <u>Equipment Overview</u>

## Mass Spectrometer Autoflex speed<sup>™</sup> MALDI TOF/TOF

<u>https://www.bruker.com/products/mass-spectrometry-and-separations/maldi-toftof/autoflex/overview.html</u>



Matrix-Assisted Laser Desorption/Ionization (MALDI) stands as one of the most powerful techniques invented for investigating:

- 1. Biopolymers (such as peptides, proteins, glycans and oligonucleotides)
- 2. Synthetic polymers
- 3. Organic molecules (>500 Da), especially in mixtures and crude samples

- 4. **MALDI Imaging. Distribution of proteins, lipid, small molecules and etc. in tissue sections.** Results are shown as color coded intensity plots and can be overlaid to an image of the investigated sample
- **5. TLC MALDI. Measurements directly from a thin layer chromatography (TLC)**
- 6. Microorganism and Fungi identification and classification
- 7. Detailed sample characterization by MS/MS. Top down proteomics.
- 8. Optimized software packages for arranging of protein, peptide, glycan and polymer analysis workflows. Microorganism and Fungi identification
  - a. ProteinScape<sup>™</sup> 4.0 Workstation data system with installed ProteinScape 4.0 software for Data Mining and Warehousing Database for Proteomics and Glycoanalysis(GlycoQuest)
  - b. Polymerix software provides sophisticated tools for analysis of homopolymer and copolymer composition and deconvolution of homopolymer and copolymer mixtures, including computation of the standard polymer metrics: Mn, Mw, Mz, PD, DPn, DPw, DPz values.
  - c. The MALDI Biotyper ( a bioinformatics package for based on MALDI-TOF MS analysis profile spectra of Microorganism and Fungi .)
- **9.** SCiLS Lab is the advanced software for statistical analysis of MALDI imaging data.