Design, Optimization, and Performance Evaluation of MALDI-TOF MS and MS-MS Instruments.

Ion Velocity

Poster Number ThP 624

Optimum Resolving Power and Mass Accuracy as Function of Effective Length



In earlier work, theoretical techniques for optimizing the resolving power of MALDI TOF MS systems were presented, and validated for a single reflector instrument. In the work presented here, all of the known contributions to peak width are considered and utimate limits with presently available technology are defined. **Objective**

Establish the practical limits on performance of MALDI-TOF instruments and design and build instruments that test these limits.

Space and Velocity Focusing in MALDI-TOF Ion Source



Contributions to Peak Width (dm/m): $R_{v1}=2K^{-1}(\delta x/D_v) = R_{v1}=[4d_1y/D_{eff}](\delta v_0/v_v)$ $R_{-}=R_{-1}[1-(m/m^2)^{1/2}]$ where $m^* =$ focused mass

 $\begin{array}{ll} & & & \\ R_{v2} = 2K^2(\delta v_d/v_n)^2 & & R_t = 2\delta t/t = 2\delta tv_d/D_e \\ R_{v3} = 2K^3(\delta v_d/v_n)^3 & & \delta t \text{ is width of single ion pulse} \end{array}$



Changes to correct mass independent errors

- Add voltage regulator at input to mirror HV supplies (removes low frequency noise)
- Correct trajectory error due to jon deflector
- · Use faster detector (0.5 ns)
- ETP DM167 in place of 5 μm dual channel plate
- Use faster digitizer
 - 0.5 ns bins>0.25 ns bins





Predicted Ultimate Resolving Power as Eurotions of Effective Length and



Optimum Resolving Power and Mass Accuracy as Function of Effective Length



RMS error for 10 peaks in spectrum of tryptic digest of BSA for all 675 spots on a 102x108 mm sample plate with automatic 2-point internal Calibration for 3.2 m analyzer



Physical length as function of effective length and number of mirrors.

Conclusions and future work

1. Experimental results on 3.2 m system in excellent agreement with theory 2. Trajectory error for single mirror analyzer <0.01 mm

- 3. Noise on mirror HV supplies <10 ppm in current analyzer
- 4. Noise on source HV pulser<500 ppm
- 5. Jitter of 10 ns between laser pulse and extraction pulse OK with reflector
- 6. Residual gas pressure<10-7 essential for high performance
- Resolving power of 1,000,000 and mass accuracy of 0.1 ppm is feasible, but requires effective length of ca. 60 m.
- Systems employing multiple reflections can provide practical configurations if trajectory error per mirror is sufficiently small.
- 9. Ultimate limits set by HV noise and residual gas pressure

Two analyzers with 10 m effective length are currently under construction to compare performance of single mirror vs. three mirrors.

Goal is >200,000 resolving power with RMS mass error <1 ppm

Reference: 1. M. L. Vetati, Modern MALDI Time of Fight Mass Spectrometry' J. Mass Spectrom. 44, 303-317(2009). Acknowledgements: This work was supported by the NIH, NCRR under grant RR 025705 and by the staff of Virgin Instruments.