Excellims MC3100 Compact High Performance Ion Mobility Mass Spectrometer

Integrated compact desktop chemical analysis / identification system based on both ion mobility and m/z measurements in seconds, providing capability of separating isobaric compounds and measuring molecular collision cross section

- Two dimensional chemical identification based on ion mobility library and m/z
- Mass analysis after high resolution ion mobility separation for high speed chemical ID in seconds
- Powerful tool for isomer analysis
- Total flexibility for user defined ion prefiltration prior to mass analyzer
- Flexible modes of operation allow freedom in experiment design
- Able to use to a variety of sample introduction / ionization sources, including Excellims' Infusion, Directspray[™], and Thermal Desorber



Excellims offers a compact ion mobility mass spectrometer system based on its proven high performance ion mobility spectrometry (HPIMSTM) in combination with a miniaturized ion trap mass spectrometer. The MC3100 is the first small-footprint analytical system that identifies chemicals based on both ion mobility and ion mass; it offers superior isomer separation and chemical identification capability while providing chemical structure information via direct collision cross-section measurements.

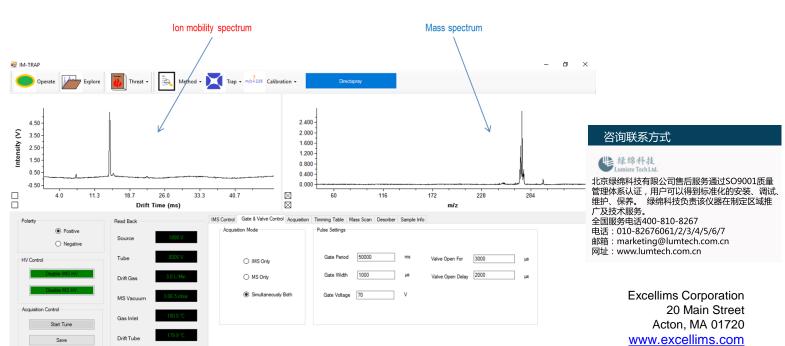
Several modes of operation are available, including IMS-only, MS-only, and simultaneous IMS-MS to produce a two dimensional mobility vs. m/z plot.

Alternatively, the IMS can be used as a prefilter for the mass spectrometer: particular mobility peaks can be selected for mass analysis, or particular mobility peaks or mobility ranges can be blocked from entering the mass spectrometer. By removing or including specific mobility-selected ions for subsequent MS and MS/MS analysis, spectral complexity and congestion can be reduced. By enriching desirable ion populations, the MC3100 delivers added confidence in compound identification.



Performance Characteristics Hardware Mobility resolving 60-120 **Excellims MC3100** power Drift tube length ~10 cm Drift time range 0-50 ms Drift tube voltage Up to 10 kV Drift time accuracy ±30 µs Ion gate pulse width From 30 µs up to the Drift gas N₂, Air, He, etc. (Bradbury-Nielson maximum drift range entrance & exit gates) setting 30-250°C Operating temperature Mass analyzer Ion trap Operating pressure Ambient conditions 10⁻⁵ torr Vacuum system Mass resolution Unit

Software Sample Introduction Options Excellims VisionTM MC3100 Software Infusion ESI source Continuous liquid sample introduction; compatible Trap package includes control and data acquisition with the use of an software for fully autosampler or HPLC integrated HPIMS and ion Directspray[™] ESI source Rapid liquid sample trap mass spectrometers screening; no additional Excellims VisionTM For post data processing pump needed **Analysis** of both ion mobility and Thermal Desorber with Solid phase sample m/z data, offering 2D Corona Discharge introduction data visualization and Ionization source chemical identification Direct gas phase sample Gas phase sample inlet introduction



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