Specifications

Mass range: 6110 and 6120: m/z 10-1500

6130: *m/z* 2–3000 **6140**: *m/z* 10–1350

Mass accuracy: ± 0.13 u within the calibrated mass range in scan mode

Mass axis stability: Mass drift does not exceed the larger absolute value of \pm 0.1 u or \pm 100 ppm of measured mass

over a 12 hour period at constant temperature ± 3 degrees centigrade

Scan Speed: 6110: 2500 u/s 6120: 2500 u/s

6130: 2500 u/s in standard mode, 5250 u/s in fast-scan mode **6140:** 2500 u/s in standard mode, 10000 u/s in fast-scan mode

SIM sensitivity: • ESI at 400 μ L/min or APCI at 1000 μ L/min

• Selected ion monitoring of m/z 609.3

· Positive ionization

6110 and **6120**: 10 pg reserpine, 100:1 RMS (20:1 peak-to-peak) **6130** and **6140**: 1 pg reserpine, 100:1 RMS (20:1 peak-to-peak)

Scan sensitivity: • ESI at 400 μL/min or APCI at 1000 μL/min

Scan range m/z 100–650
Scan speed 2500 u/s
Extracted ion at m/z 609.3

· Positive ionization

6130 and 6140: 50 pg reserpine, 100:1 RMS (20:1 peak-to-peak)

Multiple signal acquisition: 6120, 6130 and 6140: Ability to cycle through four different acquisition modes

on a scan-by-scan basis within a single run.