

# Development of “Laser Ablation Direct Analysis in Real Time Imaging” Mass Spectrometry (LADI-MS): Application to Spatial Distribution Mapping of Metabolites Along the Biosynthetic Cascade Leading to Synthesis of Atropine and Scopolamine in Plant Tissue

## SUPPORTING INFORMATION

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This document contains Table S1 which lists mass measurement data for a typical DART-HRMS analysis of a *Datura leichhardtii* seed acquired in positive ion mode. The mass spectrum to which it corresponds is shown in Figure 4. Also within this document are the MALDI-MS/MS SpiralTOF mass spectra for confirmation of the presence of the ions depicted in Figure 4 (Figures S1-S6).

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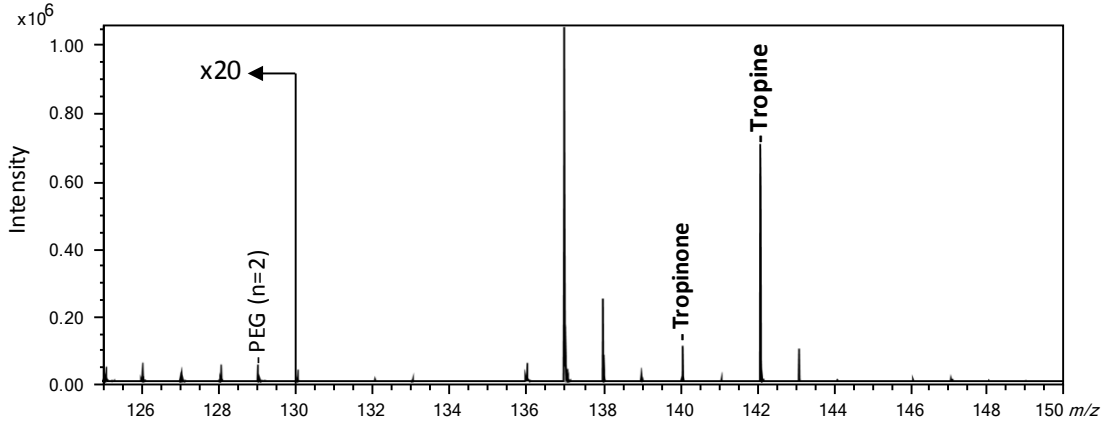
**Supporting Information Table S1.** DART-MS mass measurement data for the ions featured in this work.

Compound*	Composition	Observed Mass	Calculated Mass	Difference (mmu)
Unknown identity peak	-	124.1124	-	-
Unknown identity peak	-	131.0465	-	-
Tropinone	C <sub>8</sub> H <sub>13</sub> NO + H <sup>+</sup>	140.1079	140.1075	0.4
Tropine	C <sub>8</sub> H <sub>15</sub> NO + H <sup>+</sup>	142.1197	142.1232	3.5
Unknown identity peak	-	163.0742	-	-
Unknown identity peak	-	174.1124	-	-
Arginine	C <sub>6</sub> H <sub>14</sub> N <sub>4</sub> O <sub>2</sub> + H <sup>+</sup>	175.1217	175.1195	2.2
Unknown identity peak	-	179.0695	-	-
Unknown identity peak	-	197.0820	-	-
Atropine	C <sub>17</sub> H <sub>23</sub> NO <sub>3</sub> + H <sup>+</sup>	290.1744	290.1756	1.2
Littorine <sup>§</sup>	C <sub>17</sub> H <sub>23</sub> NO <sub>3</sub> + H <sup>+</sup>	290.1744	290.1756	1.2
Scopolamine	C <sub>17</sub> H <sub>21</sub> NO <sub>4</sub> + H <sup>+</sup>	304.1552	304.1549	0.3
Unknown identity peak <sup>‡</sup>	-	381.3340	-	-

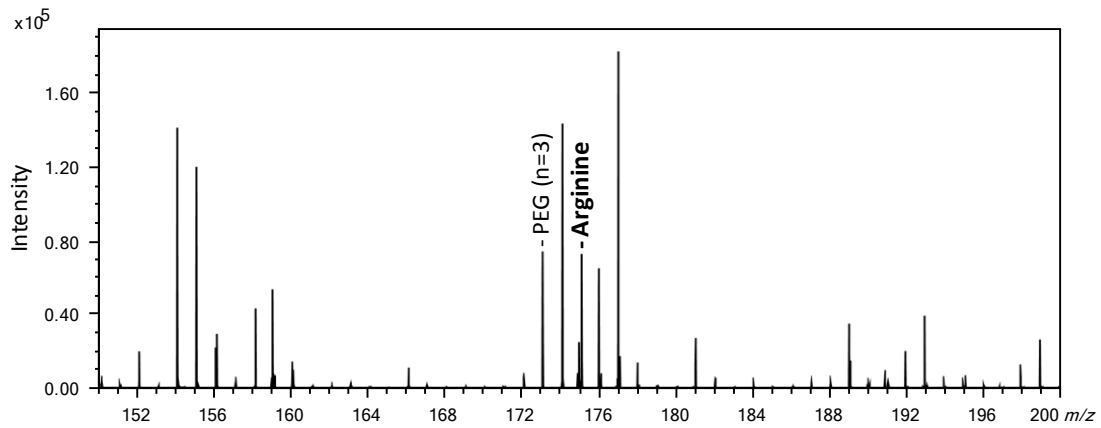
\*Unless otherwise stated, peaks of unknown identity that are listed are those of abundance  $\geq 9.2\%$  relative to the base peak. <sup>§</sup>Assignment is tentative. <sup>‡</sup>Spatial distribution of this mass was confined to the embryo and half of the seed coat.

**Supporting Information Figure S1.** MALDI mass spectra of an aqueous extract of the seed with PEG 200 as a calibrant. Peaks corresponding to tropinone, tropine, arginine, scopolamine and atropine are shown. Panels A, B, and C show the indicated mass ranges.

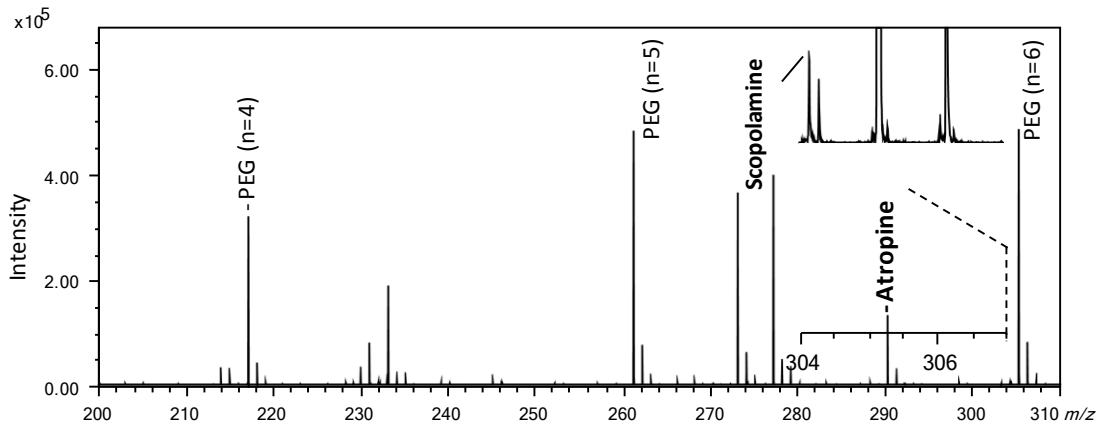
**Panel A**



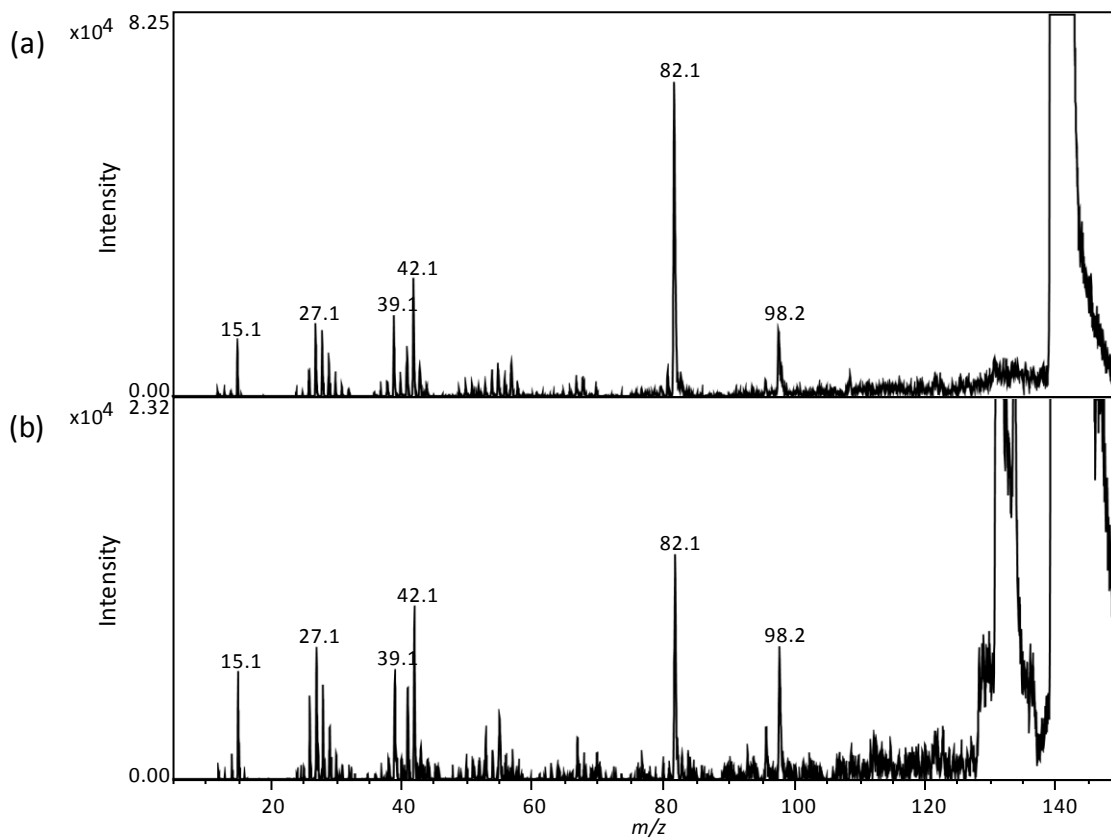
**Panel B**



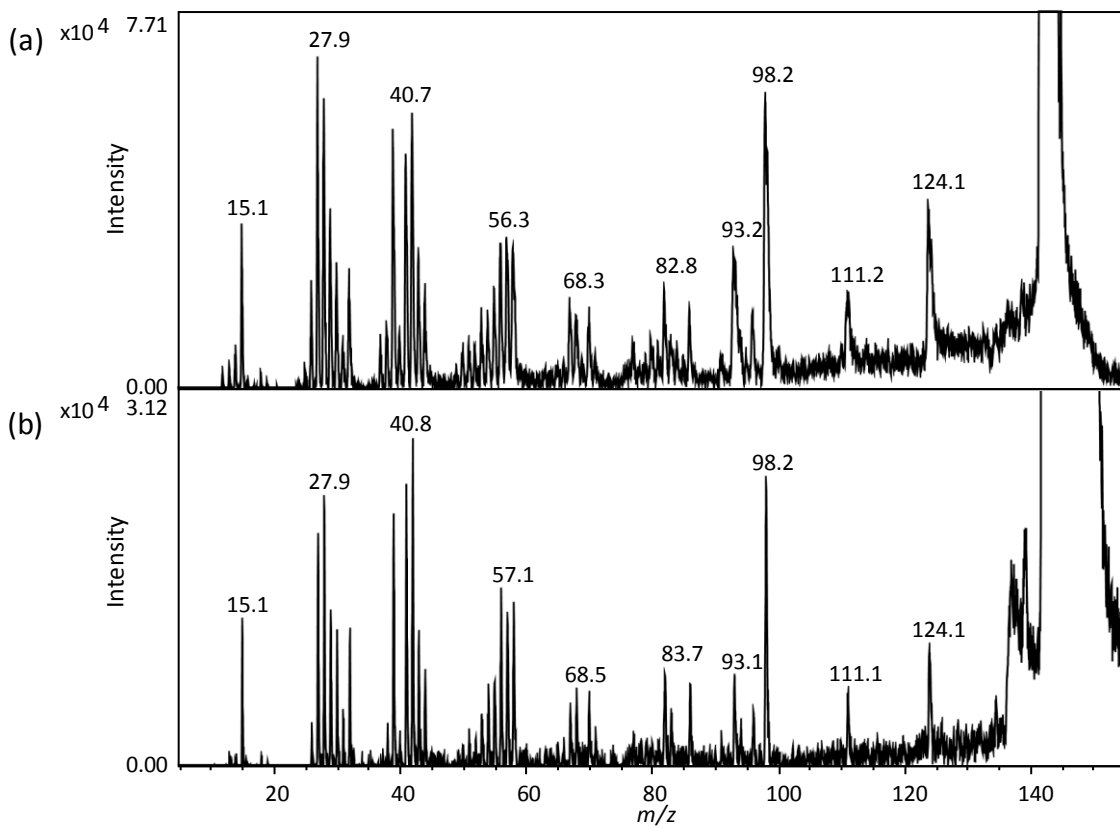
**Panel C**



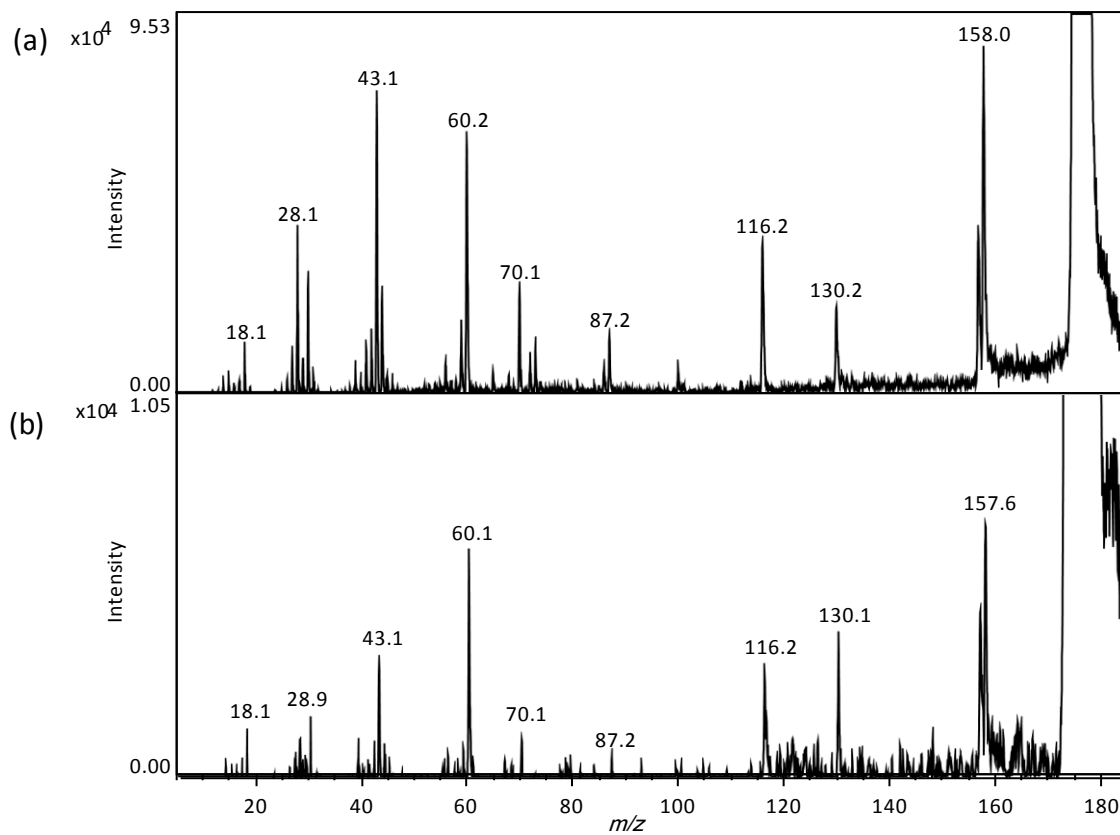
**Supporting Information Figure S2.** Product-ion spectra for tropinone ( $m/z$  140.1) authentic standard solution (a) and aqueous seed extract (b).



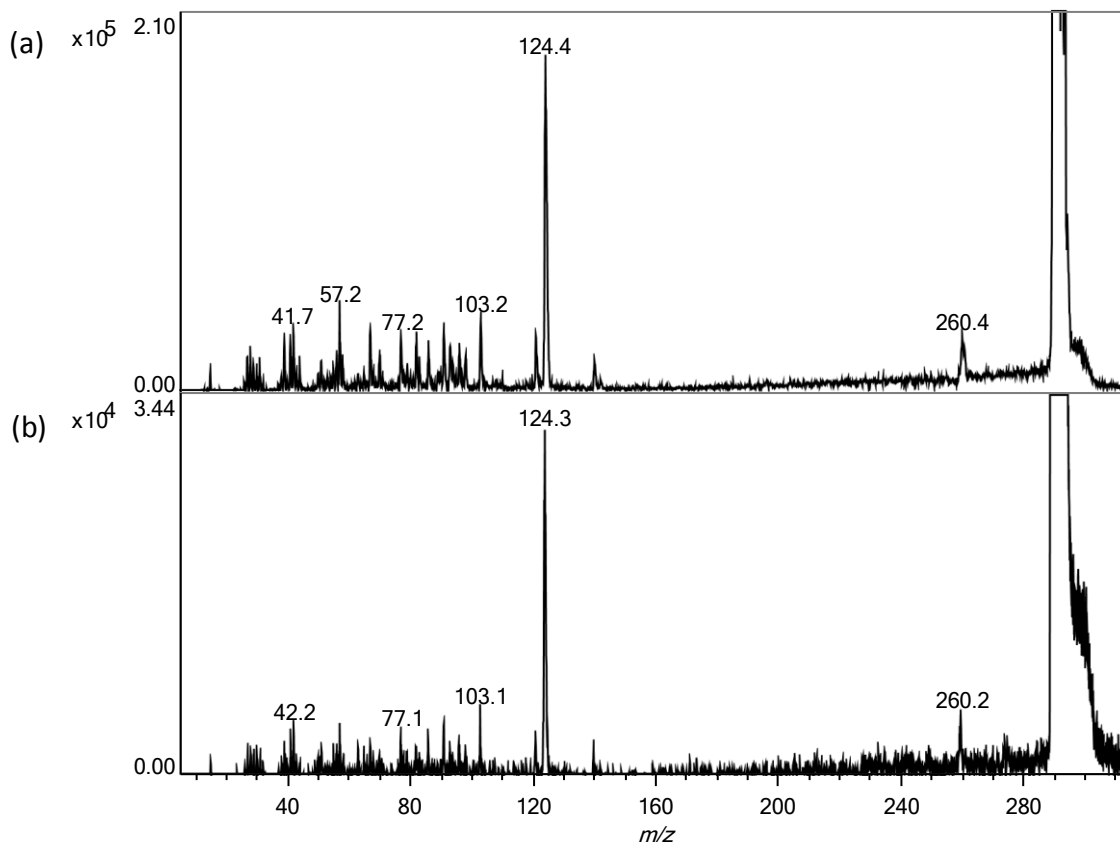
**Supporting Information Figure S3.** Product-ion spectra for tropine ( $m/z$  142.1) authentic standard solution (a) and aqueous seed extract (b).



**Supporting Information Figure S4.** Product-ion spectra for arginine ( $m/z$  175.1) authentic standard solution (a) and aqueous seed extract (b).



**Supporting Information Figure S5.** Product-ion spectra for atropine ( $m/z$  290.2) authentic standard solution (a) and aqueous seed extract (b).



**Supporting Information Figure S6.** Product-ion spectra for scopolamine ( $m/z$  304.2) authentic standard solution (a) and aqueous seed extract (b).

