

Table 1. Fragment ions found in positive ionisation mode for detected metabolites.

Compound	Retention time (min)	m/z	Mass error (ppm)	Elemental composition	Fragment ion	Formula	Mass error (ppm)
5-MeO-MiPT*	3.24	247.1810	0.0	C ₁₅ H ₂₃ N ₂ O ⁺	174.0916	C ₁₁ H ₁₂ NO ⁺	-1.7
					159.0683	C ₁₀ H ₉ NO ⁺	-0.6
					143.0733	C ₁₀ H ₉ N ⁺	-1.4
					131.0732	C ₉ H ₉ N ⁺	-2.3
					86.0972	C ₅ H ₁₂ N ⁺	2.3
Metabolite 1	2.94	233.1649	-0.4	C ₁₄ H ₂₁ N ₂ O ⁺	160.0751	C ₁₀ H ₁₀ NO ⁺	-3.7
					142.0648	C ₁₀ H ₈ N ⁺	-2.5
					132.0799	C ₉ H ₁₀ N ⁺	-6.7
					117.0573	C ₈ H ₇ N ⁺	-0.1
					115.0536	C ₉ H ₇ ⁺	-5.1
Metabolite 2	4.30	263.1753	-0.4	C ₁₅ H ₂₃ N ₂ O ₂ ⁺	86.0961	C ₅ H ₁₂ N ⁺	-4.3
					174.0913	C ₁₁ H ₁₂ NO ⁺	-0.3
					159.0685	C ₁₀ H ₉ NO ⁺	4.2
					143.0731	C ₁₀ H ₉ N ⁺	1.2
Metabolite 3	1.30	409.1965	-1.0	C ₂₀ H ₂₉ N ₂ O ₇ ⁺	131.0723	C ₉ H ₉ N ⁺	-4.8
					336.1061	C ₁₆ H ₁₈ NO ₇ ⁺	-4.9
					233.1637	C ₁₄ H ₂₁ N ₂ O ⁺	-4.8
					160.0750	C ₁₀ H ₁₀ NO ⁺	-4.1
Metabolite 4	1.72	439.2054	-4.9	C ₂₁ H ₃₁ N ₂ O ₈ ⁺	86.0963	C ₅ H ₁₂ N ⁺	-1.5
					366.1159	C ₁₇ H ₂₀ NO ₈ ⁺	-6.6
					190.0855	C ₁₁ H ₁₂ NO ₂ ⁺	-4.1
					175.0634	C ₁₀ H ₉ NO ₂ ⁺	-4.1
					158.0592	C ₁₀ H ₈ NO ⁺	-5.6
86.0961	C ₅ H ₁₂ N ⁺	-4.4					

*: The fragmentation of 5-MeO-MiPT corresponds to the observed in the analysis of *Estrella* sample.