ERRATUM

Erratum to: HS-SPME-GC×GC-qMS volatile metabolite profiling of *Chrysolina herbacea* frass and *Mentha* spp. leaves

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**Erratum to: Anal Bioanal Chem
                    DOI 10.1007/s00216-011-5600-4**

In Figure 3: Figure 3a corresponds to *Mentha spicata,*
Figure 3c corresponds to *Mentha longifolia* on the contrary
of what appeared in the published version. Please find the
correct Figure 3 with its legend below.

The online version of the original article can be found at http://dx.doi.
org/10.1007/s00216-011-5600-4.

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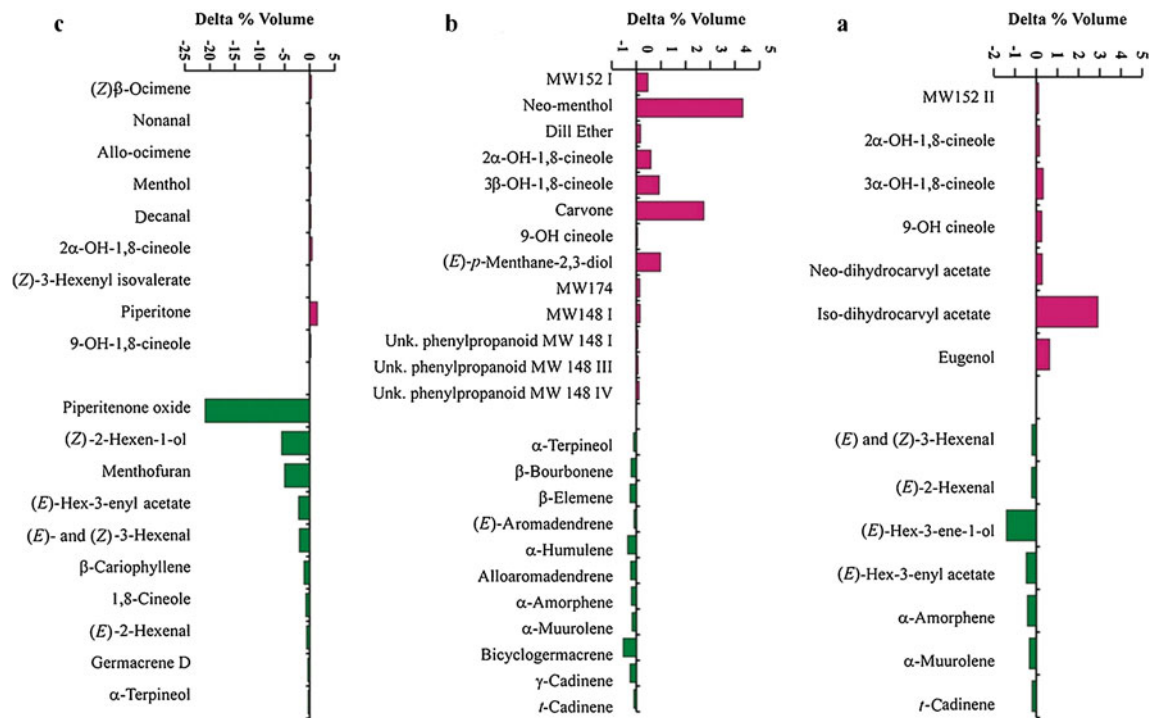Via Quarello 11/A,
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Fig. 3 Distribution of the main features revealed by the Comprehensive Template Matching fingerprinting for (**a**) *Mentha spicata*, (**b**) *Mentha piperita* and (**c**) *Mentha longifolia*. Variations are represented as the difference between 2D Peak Volumes % (PV%) from frass to leaf. The first group (upper) collects compounds revealed in frass samples and not in leaves, the second (bottom) those revealed in the leaf volatiles profile only