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Erratum

Erratum to “Metabolite profiling of mycorrhizal roots of *Medicago truncatula*” [Phytochemistry 69 (2008) 112–146]

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The publishers regret that Fig. 19 (p. 135) was not printed in colour. The correct Fig. 19 is given herein.

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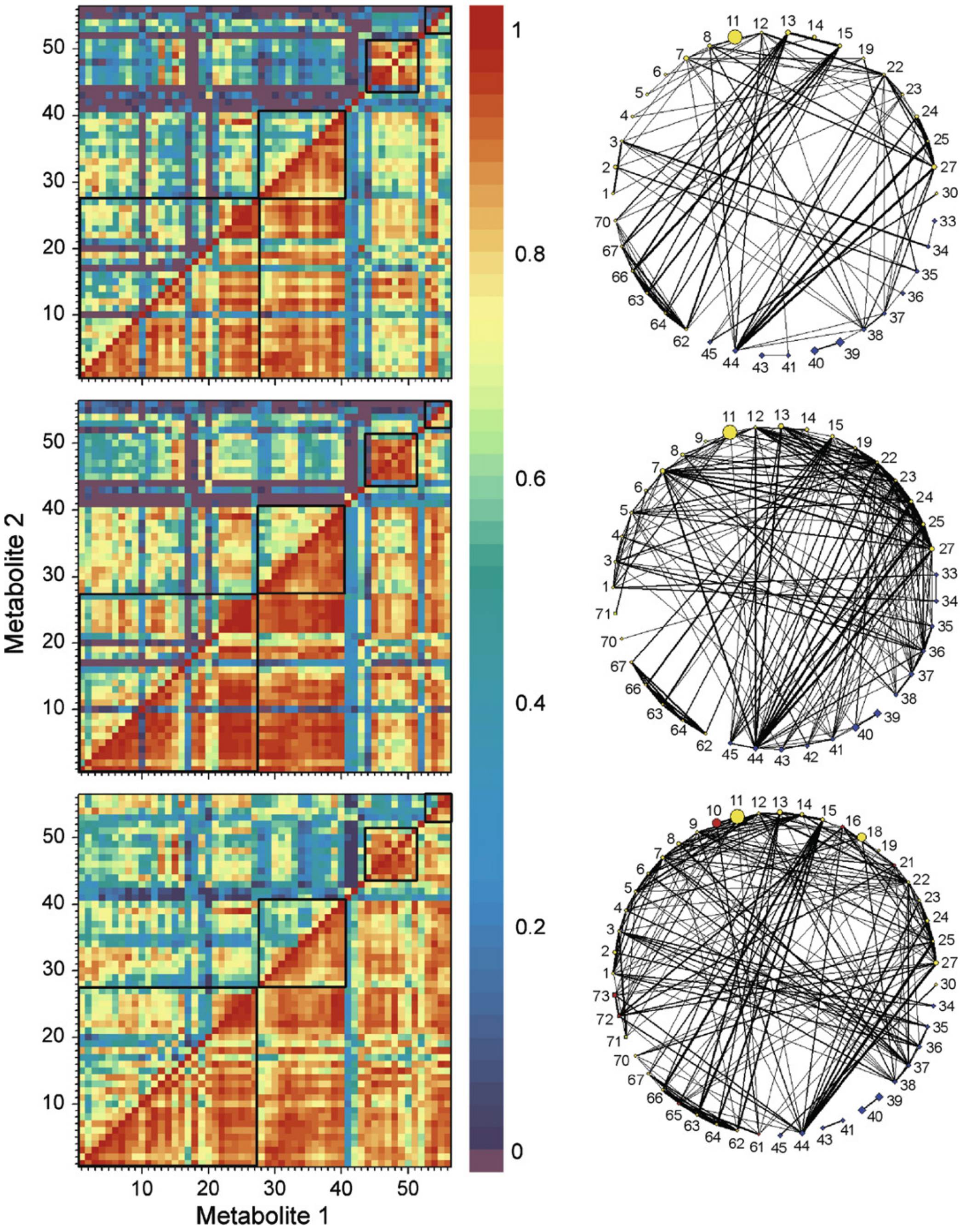

Fig. 19. Matrix plots of Pearson correlation coefficients of nonpolar metabolites (left) and circular network representations (right) for the plant sets NM20P (top), NM100P (middle) and M20P (bottom). The left triangle in the matrix plots are based on peak areas normalized to constant dry weight; the right triangle of peaks are normalized to individual root system. The boxes in the matrix plots group fatty acids, fatty alcohols, monoglycerides, and sterols (left to right). In the network representation, fatty acids (yellow circles), fatty alcohols (blue diamonds), monoglycerides (orange circles) and sterols (green boxes) are grouped in a circular arrangement. Fungus-related metabolites are shown in red color. The size of the symbols represent the relative amounts of the compounds deduced from characteristic fragment ions. The lines between the metabolites represent correlations higher than 0.8 with increasing thickness at increasing correlation. Numbers correspond to Table 2.