Erratum to: Study of Intra-Varietal Genetic Variability in Grapevine Cultivars by PCR-Derived Molecular Markers and Correlations with the Geographic Origins

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In the original manuscript, the legends for Figures 3 and 4 were reversed. The legend for Fig. 3 should read: “SAMPL digitalized electropherogram of 14 Malvasia nera di Brindisi/Lecce accessions using ALFexpress-II instrument. Genotype 1 (line 1, 1a, 1b, 1c) was used as internal reference to align the molecular profiles using the ALF-win software”.

The legend for Fig. 4 should read: “Dendrogram based on Dice’s Genetic Similarity (Coph. V. = 0.893) of the 26 accessions of Malvasia nera di Brindisi/Lecce (Malvasia n. BR/LE) using AFLP, SAMPL, M-AFLP, and ISSR molecular markers that discriminated all accessions according to the two geographic provinces, Brindisi (samples from 1 to 13) and Lecce (samples from 14 to 26)”.

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