Erratum to: High-Resolution Melting approaches towards plant fungal molecular diagnostics

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In the initial version at the first page, the present address should be for A. Zambounis. The corrected affiliation is now shown below.

Also, there is a duplicate entry of a paragraph found in the section “Applications of HRM analysis in plant fungal diagnostics”. The following paragraph should have not appeared in the proof.

“Recently, we have also developed and applied a real-time PCR assay, using universal ITS primers,

coupled with HRM analysis to differentiate more accurately closely *F. oxysporum* formae speciales based on differences in melting curve characteristics (Ganopoulos et al. 2012). The melting curve analysis of these amplicons specifically classified all nine isolates into their unique seven formae speciales and generated respective HRM curve profiles. The smallest DNA sequence difference recognized in this study was one nucleotide. The results showed that HRM curve analysis is a simple, rapid, low cost and reproducible method that allows both the identification of seven *F. oxysporum* formae speciales and at the same time their screening for variants.

The online version of the original article can be found at <http://dx.doi.org/10.1007/s12600-014-0441-2>.

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