



Corrigendum: New Insights on Leucine-Rich Repeats Receptor-Like Kinase Orthologous Relationships in Angiosperms

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In the original article, the reference for the *Prunus persica* (PRUPE, peach) sequence genome was incorrectly written as (Ahmad et al., 2011). It should be (Verde et al., 2013). The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way.

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Verde, I., Abbott, A.G., Scalabrin, S., Jung, S., Shu, S., Marroni, F., et al. (2013). The high quality draft genome of peach (*Prunus persica*) identifies unique patterns of genetic diversity, domestication and genome evolution. *Nat. Genet.* 45, 487–494. doi: 10.1038/ng.2586

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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