

## Optimization of agroinfiltration in *Pisum sativum* provides a new tool for studying host plant adaptation mechanisms in the pea aphid complex

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*Submitted to Journal:*  
Frontiers in Plant Science

*Specialty Section:*  
Plant Biotic Interactions

*ISSN:*  
1664-462X

*Article type:*  
Correction Article

*Received on:*  
19 Dec 2016

*Accepted on:*  
21 Dec 2016

*Provisional PDF published on:*  
21 Dec 2016

*Frontiers website link:*  
[www.frontiersin.org](http://www.frontiersin.org)

*Citation:*  
Guy E, Boulain H, Aigu Y, Le\_pennec C, Chawki K, Morliere S, Schaedel K, Kunert G, Simon J and Sugio A(2016) Optimization of agroinfiltration in *Pisum sativum* provides a new tool for studying host plant adaptation mechanisms in the pea aphid complex. *Front. Plant Sci.* 7:2046.  
doi:10.3389/fpls.2016.02046

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Provisional

# Corrigendum: Optimization of agroinfiltration in *Pisum sativum* provides a new tool for studying host plant adaptation mechanisms in the pea aphid complex

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**Keywords:** Pea aphid, *Acyrtosiphon pisum*, Leguminosae, agroinfiltration, salivary proteins, biotypes, host specialization, effector.

Corrigendum on Supplemental Table 2.

In the original article, description of helper plasmids in C58C1 (pGV2260) and GV3101 (pMP90) were missing. The correct information of these two strains appears below. The authors apologize for the missing information. This error does not change the scientific conclusions of the article in any way.

Bacteria	Features	Reference or source
Bacteria		
<i>Agrobacterium tumefaciens</i> C58C1	Rif <sup>r</sup> , harbours pGV2260 (pTiB6S3ΔT-DNA)	(Deblaere <i>et al.</i> , 1985)
<i>Agrobacterium tumefaciens</i> GV3101	Rif <sup>r</sup> , harbours pMP90 (pTiC58ΔT-DNA)	(Koncz and Schell, 1986)

## References

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