



Correction to: The *CsGPA1-CsAQPs* module is essential for salt tolerance of cucumber seedlings

Yan Yan¹ · Mintao Sun¹ · Yansu Li¹ · Jun Wang¹ · Chaoxing He¹ · Xianchang Yu¹ 

Published online: 18 September 2021
© Springer-Verlag GmbH Germany, part of Springer Nature 2021

Correction to: Plant Cell Reports (2020) 39:1301–1316
<https://doi.org/10.1007/s00299-020-02565-5>

Figure 5 in the original publication contains a mistake. It shows the interaction between *CsGPA1* and *CsCOR413PM2* instead of *CsTIP1.1*. The correct Fig. 5, showing the interaction between *CsGPA1* and *CsTIP1.1*, is shown below.

The original article can be found online at <https://doi.org/10.1007/s00299-020-02565-5>.

✉ Chaoxing He
hechaoxing@caas.cn

✉ Xianchang Yu
yuxianchang@caas.cn

Yan Yan
yanyan@caas.cn

Mintao Sun
sunmintao@caas.cn

Yansu Li
liyansu@caas.cn

Jun Wang
wangjun01@caas.cn

¹ The Institute of Vegetables and Flowers, Chinese Academy of Agricultural Sciences, Haidian District, Zhongguancun South St, Beijing 100081, China

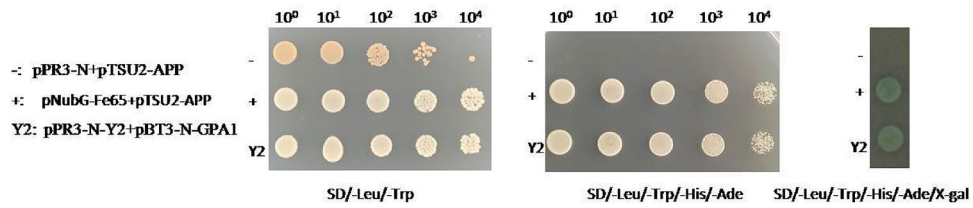


Fig. 5 Interaction between CsGPA1 and CsTIP1-1 using split-ubiquitin yeast two-hybrid System. **a** In *SD/-Leu/-Trp*, normal yeast colonies were grown proved that the vectors were successfully transferred into NMY51 yeast; **b** The negative control (–) did not produce colonies, indicating that there was no self-activation; **c** Color reaction.

‘–’ negative control, ‘+’ positive control, Y2 yeast two-hybrid of the target genes, *SD/-Leu/-Trp* two deficient media, *SD/-Trp-Leu-His-Ade* four deficient media, *X-gal* β -galactosidase, 10ⁿ is the dilution factor of yeast ($n=0, 1, 2, 3, 4$)

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.