

CORRECTION

Correction: Functional analysis of the *GmESR1* gene associated with soybean regeneration

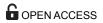
The PLOS ONE Staff

There are errors in the Funding section. The correct funding information is as follows: This research was supported through funding from The Funds for Creative Research Groups of Heilongjiang Province of China (JC2016004), Project of outstanding academic leaders in Harbin (2015RQXXJ018), and National Key Research Program of China (2016YFD0100201-21). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript. The publisher apologizes for the errors.

Reference

Zhang C, Wu X, Zhang B, Chen Q, Liu M, Xin D, et al. (2017) Functional analysis of the *GmESR1* gene associated with soybean regeneration. PLoS ONE 12(4): e0175656. doi:10.1371/journal.pone. 0175656 PMID: 28403182





Citation: The *PLOS ONE* Staff (2017) Correction: Functional analysis of the *GmESR1* gene associated with soybean regeneration. PLoS ONE 12(7): e0181576. https://doi.org/10.1371/journal.pone.0181576

Published: July 12, 2017

Copyright: © 2017 The PLOS ONE Staff. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.