CORRECTION

Correction to: New traits in crops produced by genome editing techniques based on deletions

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Correction to: Plant Biotechnol Rep (2017) 11:1–8
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In the original publication of the article, Li et al. has been incorrectly cited in the following sentence “Interestingly, a complex trait par excellence, yield, was also shown to be amenable to an SDN-1 approach. Li et al. (2017) used CRISPR-Cas9 in rice to mutate the regulatory genes *Gna1*, *DEP1*, and *GS3* and obtained plants with increased grain numbers, dense erect panicles plus semi-dwarf phenotype, and larger grains, respectively.”

The sentence with the correct citation is provided.

Interestingly, a complex trait par excellence, yield, was also shown to be amenable to an SDN-1 approach. Li et al. (2016) used CRISPR-Cas9 in rice to mutate the regulatory genes *Gna1*, *DEP1*, and *GS3* and obtained plants with increased grain numbers, dense erect panicles plus semi-dwarf phenotype, and larger grains, respectively.

The reference is,

Li M, Li X, Zhou Z, Wu P, Fang M, Pan X, Lin Q, Luo W, Wu G, Li H (2016) Reassessment of the four yield-related genes *Gn1a*, *DEP1*, *GS3*, and *IPA1* in rice using a CRISPR/cas9 system. *Frontiers in Plant Science* 7:377. <https://doi.org/10.3389/fpls.2016.00377>.

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The original article can be found online at <https://doi.org/10.1007/s11816-017-0425-z>.

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