

# SCIENTIFIC REPORTS

**OPEN**

## **Corrigendum:** The genome sequence of the outbreeding globe artichoke constructed *de novo* incorporating a phase-aware low-pass sequencing strategy of F<sub>1</sub> progeny

Davide Scaglione, Sebastian Reyes-Chin-Wo, Alberto Acquadro, Lutz Froenicke, Ezio Portis, Christopher Beitel, Matteo Tirone, Rosario Mauro, Antonino Lo Monaco, Giovanni Mauromicale, Primetta Faccioli, Luigi Cattivelli, Loren Rieseberg, Richard Michelmore & Sergio Lanteri

*Scientific Reports* 6:19427; doi: 10.1038/srep19427; published online 20 January 2016; updated 23 May 2016

In this Article, there are typographical errors in the legend of Figure 5.

“(a) Overall distribution of the repetitive fraction. (b) Distributions of insertion ages in different families of LTR elements”.

should read:

“(b) Overall distribution of the repetitive fraction. (a) Distributions of insertion ages in different families of LTR elements”.



This work is licensed under a Creative Commons Attribution 4.0 International License. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in the credit line; if the material is not included under the Creative Commons license, users will need to obtain permission from the license holder to reproduce the material. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>