

## Erratum to: Association mapping of maturity and plant height using SNP markers with the sorghum mini core collection

Hari D. Upadhyaya · Yi-Hong Wang ·  
C. L. L. Gowda · Shivali Sharma

Published online: 4 May 2014  
© Springer-Verlag Berlin Heidelberg 2014

**Erratum to: Theor Appl Genet (2013) 126:2003–2015**  
**DOI 10.1007/s00122-013-2113-x**

Due to an error in the production process, legends of Fig. 1 and Fig. 2 are identical in the original publication. Given below is the correct legend for Fig. 1.

Fig. 1. Cumulative distribution of  $p$  values computed from 14,739 SNPs for naïve, Q, PCA, K, QK, and PK association mapping models with sorghum height and maturity. More uniform  $p$  value distribution indicates a better model. The lines of naïve and PCA overlap while those from K, QK and PK overlap in both traits.

---

The online version of the original article can be found under  
doi:[10.1007/s00122-013-2113-x](https://doi.org/10.1007/s00122-013-2113-x).

---

H. D. Upadhyaya · C. L. L. Gowda · S. Sharma  
Gene Bank, International Crops Research Institute for the Semi-  
Arid Tropics (ICRISAT), Patancheru 502 324, Andhra Pradesh,  
India

Y.-H. Wang (✉)  
Department of Biology, University of Louisiana at Lafayette,  
Lafayette, LA 70504, USA  
e-mail: [yxw9887@louisiana.edu](mailto:yxw9887@louisiana.edu)