BMC Medical Genomics

CORRECTION Open Access

Correction to: A functional polymorphism in the promoter of miR-17-92 cluster is associated with decreased risk of ischemic stroke

Huatuo Huang[1,2†], Guijiang Wei[1†], Chunfang Wang[2], Yulan Lu[2], Chunhong Liu[2], Rong Wang[2], Xiang Shi[1], Jun Yang[3*] and Yesheng Wei[1,2*]

Correction to: BMC Med Genomics (2019) 12:159
https://doi.org/10.1186/s12920-019-0589-1

Following publication of the original article [1], it was reported that during the production process, Fig. 3b was omitted from the final article. The complete Fig. 3 is supplied in this correction. The original article [1] has been corrected.

Author details
[1]Department of Clinical Laboratory, The Affiliated Hospital of Guilin Medical University, Guilin 541001, Guangxi, China. [2]Department of Clinical Laboratory, The Affiliated Hospital of Youjiang Medical University for Nationalities, Baise 533000, Guangxi, China. [3]Southern Medical University, Guangzhou 510515, Guangdong, China.

Published online: 02 December 2019

Reference

1. Huang H, et al. A functional polymorphism in the promoter of miR-17-92 cluster is associated with decreased risk of ischemic stroke. BMC Med Genomics. 2019;12:159. https://doi.org/10.1186/s12920-019-0589-1.

The original article can be found online at https://doi.org/10.1186/s12920-019-0589-1

* Correspondence: junyang1232@163.com; yeshengwei22@163.com
†Huatuo Huang and Guijiang Wei contributed equally to this work.
[3]Southern Medical University, Guangzhou 510515, Guangdong, China
[1]Department of Clinical Laboratory, The Affiliated Hospital of Guilin Medical University, Guilin 541001, Guangxi, China
Full list of author information is available at the end of the article

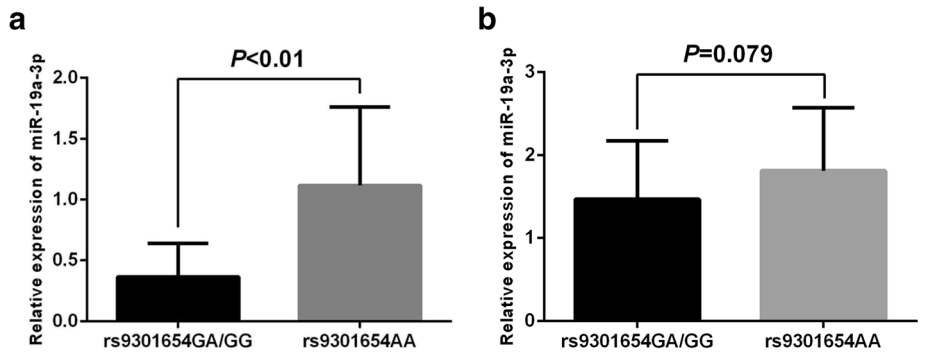

Fig. 3 (**a**) The association between rs9301654 polymorphism and the expression of miR-19a in ischemic stroke patients. Patients carrying rs9301654 GA or GG genotype (n=18) displayed a significant lower level of miR-19a as compared with those carrying rs9301654AA genotype (n=42); (**b**) The association between rs9301654 polymorphism and the expression of miR-19a in the control group (n=24 for GA/GG; n=36 for AA). The level of miR-19a showed no different in the control group between genotypes of the rs9301654 polymorphism.