Erratum to construction of a ceRNA immunoregulatory network related to the development of vascular dementia through a weighted gene coexpression network analysis

Editorial Office

Annals of Translational Medicine
Correspondence to: Editorial Office. Annals of Translational Medicine. Email: editor@atmjournal.org.

Erratum to: Ann Transl Med 2021;9:858

In the article entitled "Construction of a ceRNA immunoregulatory network related to the development of vascular dementia through a weighted gene coexpression network analysis" (1), there are errors.
In the flowchart in Figure 1, the authors marked the number of lncRNA, miRNA, and mRNA incorrectly. The correct number should be lncRNA: GSE122063; miRNA: GSE120584; mRNA: GSE122063.

In the "Data preprocessing and the differential analysis" section of the "Methods" module, "We transformed the probe into the corresponding gene symbol according to the annotation information." in lines 2–3 should be corrected to "We transformed the probe into the corresponding gene symbol according to the annotation information and the UniProt.".

The authors apologize for the negligent errors.

Click here to view the updated version of the article.

References

1. Shi H, Sun F, Yang T, et al. Construction of a ceRNA immunoregulatory network related to the development of vascular dementia through a weighted gene coexpression network analysis. Ann Transl Med 2021;9:858.