Whole-transcriptome analysis of aluminum-exposed rat hippocampus and identification of ceRNA networks to investigate neurotoxicity of Al

Chanting He, Xiaoyan Zhao, Yang Lei, Jisheng Nie, Xiaoting Lu, Jing Song, Linping Wang, Huan Li, Fangqu Liu, Yidan Zhang, and Qiao Niu

Correspondence: niuqiao55@163.com
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In the originally published version of this article, we, the authors, placed all the images in the PPT for analysis and organization and finally combined them into the images used in the article. Due to the large amount of data and numerous images in this article, errors occurred during the organization process, resulting in duplicated images. We sincerely apologize for this mistake.

After receiving feedback from readers, we carefully examined the original data and rechecked all the original images of each group (Western blotting bands, Nissl staining, immunohistochemistry, electron microscopy, etc.) and found that the statistical values were correct. However, during the combination process of electron microscopy images, errors occurred due to the copying and pasting process, resulting in duplicated images. To solve this problem, we have provided the corrected Figure 8 below. We greatly appreciate the help of readers in making our work more rigorous and scientific.
Figure 8. Blocking of miR-96-5p reduced the neurotoxicity of aluminum through the IRS1/PI3K/AKT pathway (corrected)