## **CORRIGENDUM**

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MicroRNA-1929-3p participates in murine cytomegalovirus-induced hypertensive vascular remodeling through Ednra/NLRP3 inflammasome activation

WEI ZHOU, DONGMEI XI, YUNZHONG SHI, LAMEI WANG, HUA ZHONG, ZHEN HUANG, YONGMIN LIU, YAN TANG, NING LU, YONGJIA WANG, ZHENGYU ZHANG, JIAXIN PEI, NA TANG and FANG HE

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Following the publication of this article, the authors have realized that the images selected for the 'MCMV+miR-NC' and 'MCMV+miR-1929-3p' data panels (15 months), as featured in the lower of the two rows of data shown in Fig. 4A on p. 726, were derived from the same experimental group. The authors re-examined their original data, and discovered that an error had inadvertently been made on account of the similar file names of the two images.

The corrected version of Fig. 4, featuring the correct data for the 'MCMV+miR-NC' experiment, is shown on the next page. The authors regret that this error was not picked up upon before the paper was sent to press, and thank the Editor of *International Journal of Molecular Medicine* for allowing them the opportunity to publish a corrigendum. Note that the error did not affect either the results or the conclusions reported in the study, and all the authors agree to this corrigendum. Furthermore, the authors regret any inconvenience caused to the readership.



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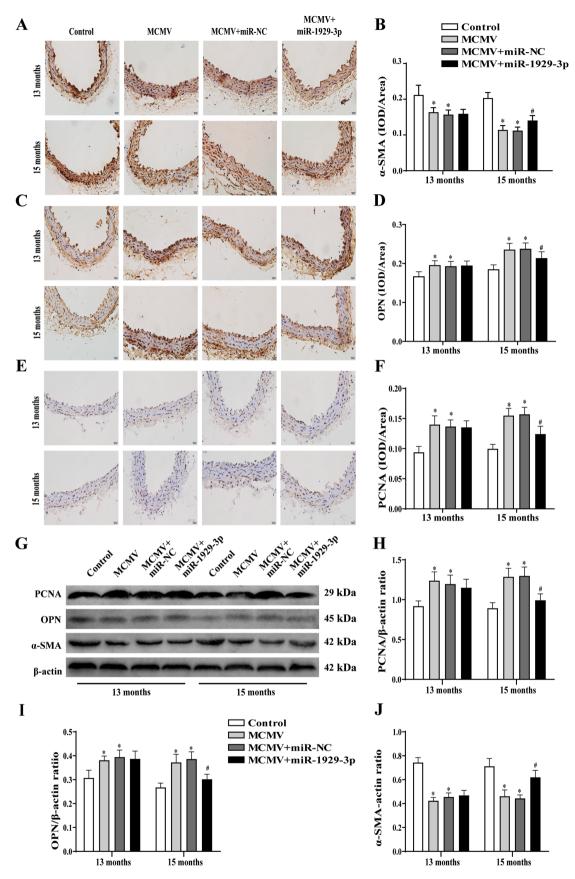


Figure 4. Improved MCMV-induced vascular phenotypic transformation by miR-1929-3p. (A-F) Expression of (A)  $\alpha$ -SMA, (C) OPN and (E) PCNA in mouse aorta as detected by immunohistochemical staining (magnification, x100), and statistical analysis of (B)  $\alpha$ -SMA, (D) OPN and (F) PCNA. (G) Western blotting and (H-J) statistical analysis of the expression of (H) PCNA, (I) OPN and (J)  $\alpha$ -SMA in aortic tissues. \*P<0.05 vs. control of the same age; \*P<0.05 vs. MCMV + miR-NC group of the same age. In panels A, D, F, H, I and J, data were analyzed using one-way ANOVA followed by Tukey's multiple comparisons post hoc. n=5 per group. MCMV, murine cytomegalovirus; SMA, smooth muscle.