CORRECTION



Correction: CKLF1 Aggravates Focal Cerebral Ischemia Injury at Early Stage Partly by Modulating Microglia/Macrophage Toward M1 Polarization Through CCR4

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The original version of this article unfortunately contained error in Fig. 10.

In Fig. 10B, the merge image of C021 100 nM group of microglia stained with CD206 is published incorrectly. Other images in the figure remains the same, and the interpretation of the results remains unchanged.

The corrected figure is presented here.

The authors would like to apologise for any inconvenience caused.

The original article has been corrected.

The original article can be found online at https://doi.org/10.1007/ \pm 10571-019-00669-5.

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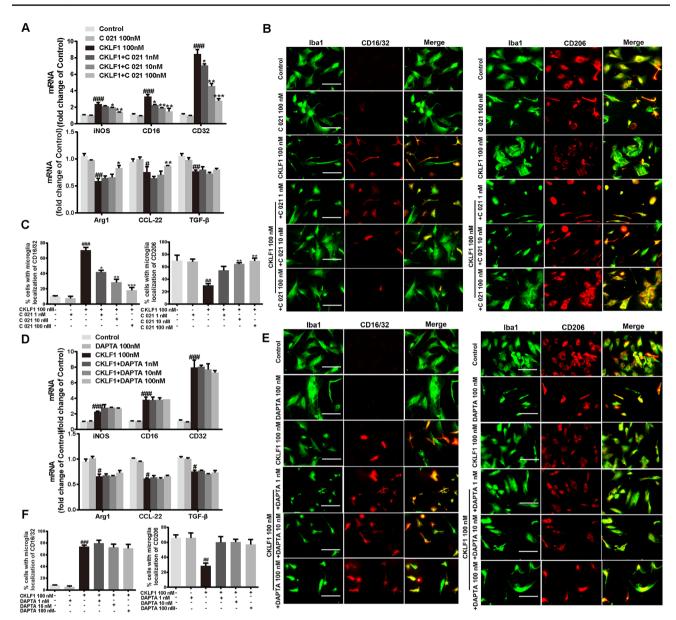


Fig. 10 CKLF1 modulates the microglia polarization through CCR4. **a** qPCR analysis of mRNA expression levels of M1 markers (iNOS, CD16, CD32) and M2 markers (Arg1, CCL-22, TGF-β) in primary microglia treated with CKLF1 and C 021 dihydrochloride for 24 h (n=6 cell samples). $^{*}P$ <0.05, $^{**}P$ <0.01, $^{***}P$ <0.001 versus control; $^{*}P$ <0.05, $^{**}P$ <0.01, $^{***}P$ <0.001 versus CKLF1 100 nM. **b** Representative photomicrographs of double-staining immunofluorescence of CD16/32 or CD206 with Iba1 in the primary microglia treated with CKLF1 and C 021 dihydrochloride for 24 h. Scale bars 100 μm. **c** Quantitative analysis of CD16/32-positive and CD206-positive microglia (n=6 cell samples). $^{**}P$ <0.01, $^{***}P$ <0.001 versus

control; *P<0.05, **P<0.01, ***P<0.001 versus CKLF1 100 nM. **d** qPCR analysis of mRNA expression levels of M1 markers (iNOS, CD16, CD32) and M2 markers (Arg1, CCL-22, TGF- β) in primary microglia after treatment with CKLF1 and DAPTA for 24 h (n=6 cell samples). *P<0.05, **P<0.001 versus control. **e** Representative photomicrographs of double-staining immunofluorescence of CD16/32 or CD206 with Iba1 in the primary microglia after treatment with CKLF1 and DAPTA for 24 h. Scale bars 100 μ m. **f** Quantitative analysis of CD16/32-positive and CD206-positive microglia (n=6 cell samples). **P<0.01, **P<0.001 versus control



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