



Corrigendum: Critical Role of Alternative M2 Skewing in miR-155 Deletion-Mediated Protection of Colitis

Jintao Li^{1,2*†}, Ji Zhang^{3†}, Hongxia Guo^{1,2†}, Shimin Yang⁴, Weiping Fan⁵, Nan Ye¹, Zhiqiang Tian³, Tiantian Yu¹, Guoping Ai¹, Zigang Shen³, Haiyang He³, Ping Yan⁶, Hui Lin¹, Xue Luo¹, Hongli Li^{7*} and Yuzhang Wu^{3*}

¹ Institute of Tropical Medicine, Army Medical University, Chongqing, China, ² Department of Microbiology, College of Basic

Medicine, Army Medical University, Chongqing, China, ³ Institute of Immunology, PLA, Army Medical University, Chongqing,

China, ⁴ Department of Gastroenterology, Xingiao Hospital, Army Medical University, Chongging, China, ⁵ Department of

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*Correspondence:

Jintao Li Ijtqms@qq.com Hongli Li wuyuzhang@tmmu.edu.cn Yuzhang Wu lihongli@tmmu.edu.cn

[†]These authors have contributed equally to this work

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A Corrigendum on

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In the original article, there were mistakes in **Figure 1J** and **Figure 6E** as published. **Figure 1J** (representative FACS plot of CD4⁺IFN- γ^+ cells) was mistakenly duplicated from **Figure 1I** (representative FACS plot of CD4⁺IL-17⁺ cells), and the tubulin band of **Figure 6E** was inadvertently covered by the band of that in **Figure 6C**. The corrected **Figure 1** and **Figure 6** appear below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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FIGURE 1 | Attenuated dextran sulfate sodium (DSS)-induced colitis in miR-155^{-/-} mice is dependent on commensal bacteria. WT or miR-155^{-/-} mice were given 3% DSS in drinking water for 5 days, followed by regular drinking water for 6 days. (A–E) Weight change (A), Kaplan–Meier plot of survival rate (B), stool blood (C), representative gross colon appearance [(D) left] and colon length [(D) right], and representative H&E-stained colon cross-sections [(E) left, original magnification, 200× or 400×] and semi-quantitative histopathology score [(E) right]. (F) WT and miR-155^{-/-} mice were treated with broad-spectrum antibiotics cocktail (ABX) for 4 weeks and then given 3% DSS, the body weight change (left) and DAI (right) were monitored daily. Ns vs WT control (G,H). The LPMCs were isolated from colon tissues of DSS-treated WT (*n* = 12) and miR-155^{-/-} (*n* = 15) mice, then the total number of LPMCs (CD45⁺) (G), T cells (CD4⁺ and CD8⁺), DCs (CD11c⁺CD11b⁻) and macrophages (CD11b⁺CD11c^{-//ow}) (H) were counted by flow cytometry. (I,J) Representative FACS showing CD4⁺IL-17⁺ cells (I) and CD4⁺IFN-γ⁺ cells (J) in isolated LPMCs of DSS-treated WT (*n* = 12) and miR-155^{-/-} (*n* = 15) mice. **P* < 0.05, ***P* < 0.01 vs WT control [Student's *t*-test in (A,D,E,G,H) and Kaplan–Meier analysis in (B,C)]. ns vs WT control. Data are representative of three independent experiments (mean and SD in A-D); n = 12–15 mice per group in (A–F) and n = 5–6 mice per group in (G). ns, not significant. WT, wild-type.

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FIGURE 6 | C/EBP β and SOCS1 are key functional targets in intestinal M2 polarization. (A) BMDMs isolated from WT and miR-155^{-/-} mice were treated with CBA (10 μ g/mL) and IFN- γ (20 ng/mL), and the relative expression of M1genes and M2 genes were determined by Q-PCR. (B) The absolute amounts of secreted cytokines IL-10 and IL-12 (as representative of M2 and M1 gene products, respectively) in the supernatants of WT or miR-155^{-/-} BMDMs that had been treated with M1 condition (CBA + IFN- γ) and M2 condition (IL-4) were measured by ELISA. (C) The protein expression level of C/EBP β in macrophages (CD11b⁺CD11c^{-/IOw}) isolated from LPMCs of dextran sulfate sodium colitis mice were determined by western blotting. (D) miR-155^{-/-} BMDMs were transferred with C/EBP β siRNA or control and then stimulated with CBA (10 μ g/mL) and IFN- γ (20 ng/mL), and the relative expression of M1genes and M2 genes were determined by Q-PCR. (E) The protein expression level of SOCS1 and SHIP1 in macrophages, as described in (C), was determined by western blotting. (F) miR-155^{-/-} BMDMs were transferred with SOCS1 and SHIP1 in macrophages, as described in (C), and the relative expressions of M1genes and M2 genes were determined by Q-PCR. **P* < 0.05, ***P* < 0.01 vs WT control or siRNA control [Student's *t*-test in (A,B,D)]. **P* < 0.05, ns > 0.05 vs. siRNA control (ANOVA with Bonferroni's posttest correction for multiple comparisons in (F). Data are representative of three independent experiments (mean and SD). Ns, not significant. BMDMs, bone marrow-derived macrophage. WT, wild-type. CBA, cecal bacterial antigen.