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Author Correction: MicroRNA-125b-5p regulates IL-1 β induced inflammatory genes via targeting TRAF6-mediated MAPKs and NF- κ B signaling in human osteoarthritic chondrocytes

Zafar Rasheed¹, Naila Rasheed¹, Waleed Al Abdulmonem² & Muhammad Ismail Khan³Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-019-42601-3>, published online 03 May 2019

This Article contains an error in the order of the Figure legends. The legends for Figures 1 and 2 were incorrectly given as those for 2 and 1 respectively.

The legend for Figure 1 should read:

“(A) MicroRNA database top predicted targets for microRNA hsa-miR-125b-5p. The number in the center of each circle represents target score of prediction. Full names of all genes are available at <https://www.ncbi.nlm.nih.gov/gene>. (B,C) Bioinformatics analysis of hsa-miR-125b-5p binding to 3’UTR of TRAF6 mRNA. (B) Seed-Matched sequences of hsa-miR-125b-5p in the 3’UTR of TRAF6 mRNA. (C) Target Scan prediction. (D) Cross-species conservation by Target Scan algorithm.”

The legend for Figure 2 should read:

“Expression of hsa-miR-125b-5p and TRAF6 in smooth and damaged OA cartilage tissues. (A) Smooth and damaged cartilage tissues from OA patients (B) Expression of hsa-mir-125b-5p in smooth and damaged cartilage tissues. *p < 0.01 versus smooth cartilage tissue. (C) Gene Expression of TRAF6 in smooth and damaged cartilage tissues. [®]p < 0.01 versus smooth cartilage tissue. (D) Protein expression of TRAF6 in smooth and damaged cartilage tissues (full length blot is provided in Supplementary Information File). *p < 0.01 versus SC. SC, smooth cartilage; DC, damaged cartilage. Band images were digitally captured by the Un-Scan-It software and the band intensities of TRAF6 proteins were divided by band intensities of β -actin and are expressed in average pixels.”

¹Department of Medical Biochemistry, College of Medicine, Buraidah, Qassim University, Buraidah, Saudi Arabia.

²Department of Pathology, College of Medicine, Buraidah, Qassim University, Buraidah, Saudi Arabia. ³Faculty of Medicine, School of Public Health, University of Queensland, Brisbane, Australia. Correspondence and requests for materials should be addressed to Z.R. (email: zafarrasheed@qumed.edu.sa)



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