



# Corrigendum: Assessing the Function of the ZFP90 Variant rs1170426 in SLE and the Association Between SLE Drug Target and Susceptibility Genes

Tingting Zhu<sup>1,2†</sup>, Yuandi Huang<sup>1†</sup>, Danfeng Qian<sup>3†</sup>, Yuming Sheng<sup>1</sup>, Chaowen Zhang<sup>1</sup>, Shirui Chen<sup>1</sup>, Hui Zhang<sup>1</sup>, Hui Wang<sup>1</sup>, Xuejun Zhang<sup>1</sup>, Junlin Liu<sup>4\*</sup>, Changhai Ding<sup>2,5,6\*</sup> and Lu Liu<sup>1,7\*</sup>

<sup>1</sup> Department of Dermatology, The First Affiliated Hospital, Anhui Medical University, Hefei, China, <sup>2</sup> Department of Rheumatology and Immunology, Arthritis Research Institute, The First Affiliated Hospital of Anhui Medical University, Hefei, China, <sup>3</sup> Department of Dermatology, Lu'an People's Hospital, Lu'an, China, <sup>4</sup> Department of Dermatology, The Second Affiliated Hospital, Hainan Medical University, Haikou, China, <sup>5</sup> Clinical Research Centre, Zhujiang Hospital, Southern Medical University, Zhuziang, China, <sup>6</sup> Menzies Institute for Medical Research, University of Tasmania, Hobart, TAS, Australia, <sup>7</sup> Department of Medical and Molecular Genetics, King's College London, London, United Kingdom

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Jeehee Youn,  
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### \*Correspondence:

Lu Liu  
liulu8887@163.com  
Changhai Ding  
changhai.ding@utas.edu.au  
Junlin Liu  
liujunlin0759@163.com;  
lu.2.liu@kcl.ac.uk

<sup>†</sup>These authors have contributed  
equally to this work

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## A Corrigendum on

### Assessing the Function of the ZFP90 Variant rs1170426 in SLE and the Association Between SLE Drug Target and Susceptibility Genes

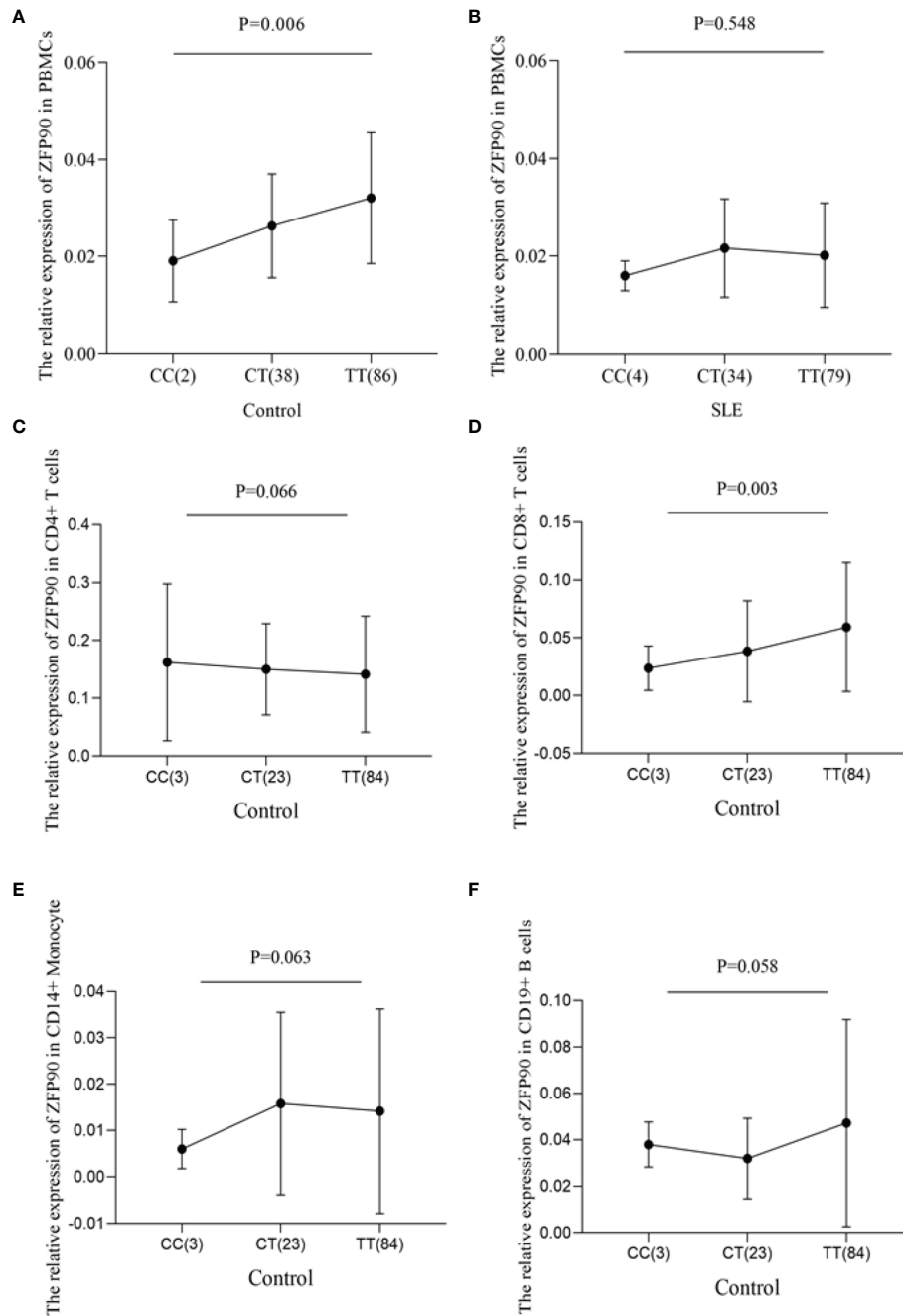
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In the original article, there were mistakes in **Figures 1, 3, 4** as published. The number of CC genotype is not 10 but 2 in the legend of **Figure 1A**. The SNP ID is not rs2297550 but rs1170426 and the number of CT genotype is not 33 but 34 in the legend of **Figure 1B**. The number of CT genotype is not 33 but 34 in the label of **Figure 1B**. In **Figure 3**, the legends of **Figures 3B, 3C** were inverted. In **Figure 4B**, the label of arthritis is not FDR  $p = 0.004$  but FDR  $p = 0.020$  as shown in **Table 3**. The corrected **Figures 1, 3, 4** appear below.

The authors apologize for these errors 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 | (A)** The effect of rs1170426 on *ZFP90* mRNA expression levels in PBMCs from healthy controls. Of the 126 controls, 2 individuals with CC, 38 with CT and 86 with TT were analyzed. The group with “CC” homozygous has the lowest expression levels ( $P = 0.006$ ). **(B)** The effect of rs1170426 on *ZFP90* mRNA expression levels in PBMCs from SLE cases. Of the 117 cases, 4 individuals with CC, 34 with CT and 79 with TT were analyzed. The expression did not significantly correlate with genotype of rs1170426 ( $P = 0.548$ ). **(C–F)** The effect of rs1170426 on *ZFP90* mRNA expression levels in CD4+ T cells, CD8+ T cells, CD19+ B cells, and CD14+ monocytes from other 110 healthy controls. Of the 110 controls, 3 individuals with CC, 23 with CT and 84 with TT were analyzed. *ZFP90* expression levels of samples with risk allele “C” of rs1170426 were significantly decreased in CD8+ T cells ( $P = 0.003$ ).

