



OPEN **Publisher Correction:** Genetic co-expression networks contribute to creating predictive model and exploring novel biomarkers for the prognosis of breast cancer

Yuan-Kuei Li, Huan-Ming Hsu, Meng-Chiung Lin, Chi-Wen Chang, Chi-Ming Chu, Yu-Jia Chang, Jyh-Cherng Yu, Chien-Ting Chen, Chen-En Jian, Chien-An Sun, Kang-Hua Chen, Ming-Hao Kuo, Chia-Shiang Cheng, Ya-Ting Chang, Yi-Syuan Wu, Hao-Yi Wu, Ya-Ting Yang, Chen Lin, Hung-Che Lin, Je-Ming Hu & Yu-Tien Chang

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The original version of this Article contained errors in the author list as a previous version of the author list was erroneously published.

The order of author names was incorrect. Chien-An Sun, Chen Lin, and Hung-Che Lin were omitted from the author list. In addition, Yu-Jia Chang was omitted as an equally contributing author. The original author list and affiliations appear below.

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The original Article has been corrected.



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