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*CORRESPONDENCE
Frontiers Production Office,
✉ production.office@frontiersin.org

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Erratum: Integrated weighted gene co-expression network analysis identified that TLR2 and CD14 are related to coronary artery disease

Frontiers Production Office*

Frontiers Media SA, Lausanne, Switzerland

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An Erratum on

[Integrated weighted gene co-expression network analysis identified that TLR2 and CD14 are related to coronary artery disease](#)

by Qi B, Chen J-H, Tao L, Zhu C-M, Wang Y, Deng G-X and Miao L (2021). *Front. Genet.* 11:613744.
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In the published article there was an error in the **article title**. Instead of “Integrated Weighted Gene Co-expression Network Analysis Identified That TLR2 and CD40 Are Related to Coronary Artery Disease”, it should be “Integrated Weighted Gene Co-expression Network Analysis Identified That TLR2 and CD14 Are Related to Coronary Artery Disease.”

The publisher apologizes for this mistake. The original version of this article has been updated.