



OPEN ACCESS

APPROVED BY
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

*CORRESPONDENCE
Lu-Feng Cheng
✉ lfcheng@xjmu.edu.cn

SPECIALTY SECTION
This article was submitted to
Viral Immunology,
a section of the journal
Frontiers in Immunology

RECEIVED 18 February 2023
ACCEPTED 20 February 2023
PUBLISHED 28 February 2023

CITATION
Zhang Y-Z, Zeb A and Cheng L-F (2023)
Corrigendum: Exploring the molecular
mechanism of hepatitis virus inducing
hepatocellular carcinoma by microarray
data and immune infiltrates analysis.
Front. Immunol. 14:1168774.
doi: 10.3389/fimmu.2023.1168774

COPYRIGHT
© 2023 Zhang, Zeb and Cheng. This is an
open-access article distributed under the
terms of the [Creative Commons Attribution
License \(CC BY\)](#). The use, distribution or
reproduction in other forums is permitted,
provided the original author(s) and the
copyright owner(s) are credited and that
the original publication in this journal is
cited, in accordance with accepted
academic practice. No use, distribution or
reproduction is permitted which does not
comply with these terms.

Corrigendum: Exploring the molecular mechanism of hepatitis virus inducing hepatocellular carcinoma by microarray data and immune infiltrates analysis

Yong-Zheng Zhang, Amir Zeb and Lu-Feng Cheng*

Department of Pharmacology, School of Pharmacy, Xinjiang Medical University, Urumqi, China

KEYWORDS

hepatitis virus, hepatocellular carcinoma, immune infiltration, viral carcinogenicity, macrophage polarization, computer simulation

A Corrigendum on

Exploring the molecular mechanism of hepatitis virus inducing hepatocellular carcinoma by microarray data and immune infiltrates analysis

By Zhang Y-Z, Zeb A and Cheng L-F (2022) *Front. Immunol.* 13:1032819.
doi: 10.3389/fimmu.2022.1032819

In the published article, there was an error in **Figure 6** as published. The corrected **Figure 6** and its caption appear below.

Figure 6 caption (to be used in the figure only):

Expression correlation analysis of hub genes and Immune related genes (A) Correlation analysis of HBV-HCC immune pathway related genes and hub genes by using GSE55092. (B) Correlation analysis of HCV-HCC immune pathway related genes and hub genes by using GSE69715. Red was positively correlated and blue was negatively correlated. The size of the circle represented the magnitude of correlation, and the cross indicated no correlation

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.

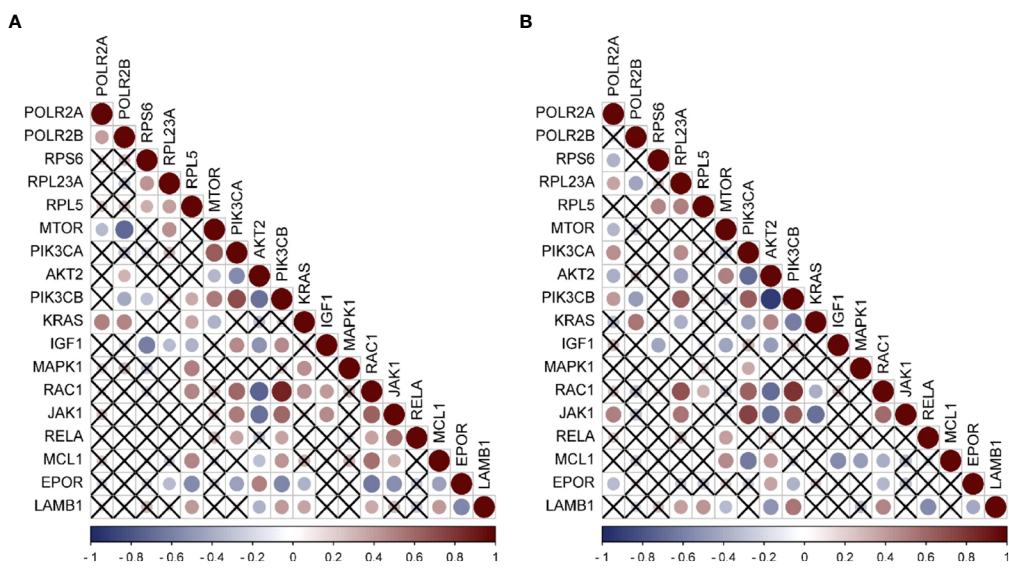


FIGURE 6
 Expression correlation analysis of hub genes and Immune related genes (A) Correlation analysis of HBV-HCC immune pathway related genes and hub genes by using GSE55092. (B) Correlation analysis of HCV-HCC immune pathway related genes and hub genes by using GSE69715. Red was positively correlated and blue was negatively correlated. The size of the circle represented the magnitude of correlation, and the cross indicated no correlation.

Publisher’s note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated

organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.