



Corrigendum to "Genetics Variants and Serum Levels of MHC Class I Chain-related A in Predicting Hepatocellular Carcinoma Development in Chronic Hepatitis C Patients Post Antiviral Treatment" [EBioMedicine 15 (2017) 81–89]

Citation

Huang, C., C. Huang, M. Yeh, S. Wang, K. Chen, Y. Ko, C. Lin, et al. 2017. "Corrigendum to "Genetics Variants and Serum Levels of MHC Class I Chain-related A in Predicting Hepatocellular Carcinoma Development in Chronic Hepatitis C Patients Post Antiviral Treatment" [EBioMedicine 15 (2017) 81–89]." EBioMedicine 17 (1): 237. doi:10.1016/j.ebiom.2017.02.027.

Published Version

doi:10.1016/j.ebiom.2017.02.027

Permanent link

http://nrs.harvard.edu/urn-3:HUL.InstRepos:34493131

Terms of Use

This article was downloaded from Harvard University's DASH repository, and is made available under the terms and conditions applicable to Other Posted Material, as set forth at http://nrs.harvard.edu/urn-3:HUL.InstRepos:dash.current.terms-of-use#LAA

Share Your Story

The Harvard community has made this article openly available. Please share how this access benefits you. <u>Submit a story</u>.

Accessibility

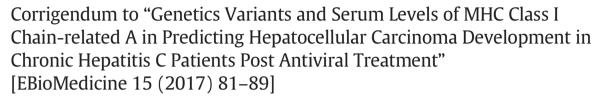
Contents lists available at ScienceDirect

EBioMedicine

journal homepage: www.ebiomedicine.com



Corrigendum





Chung-Feng Huang ^{a,b,c}, Ching-I Huang ^a, Ming-Lun Yeh ^{a,b}, Shu-Chi Wang ^a, Kuan-Yu Chen ^a, Yu-Min Ko ^a, Ching-Chih Lin ^a, Yi-Shan Tsai ^a, Pei-Chien Tsai ^a, Zu-Yau Lin ^{a,b}, Shinn-Cherng Chen ^{a,b}, Chia-Yen Dai ^{a,b,c,d}, Jee-Fu Huang ^{a,b}, Wan-Long Chuang ^{a,b}, Ming-Lung Yu ^{a,b,e,f,*}

- ^a Hepatobiliary Division, Department of Internal Medicine, Kaohsiung Medical University Hospital, Kaohsiung Medical University, Kaohsiung, Taiwan
- ^b Faculty of Internal Medicine, School of Medicine, College of Medicine, Kaohsiung Medical University, Kaohsiung, Taiwan
- ^c Department of Occupational Medicine, Kaohsiung Medical University Hospital, Kaohsiung Medical University, Kaohsiung, Taiwan
- d Department of Preventive Medicine, Kaohsiung Medical University Hospital, Kaohsiung Medical University, Kaohsiung, Taiwan
- ^e Institute of Biomedical Sciences, National Sun Yat-Sen University, Taiwan
- f Liver Center, Division of Gastroenterology, Massachusetts General Hospital, Harvard Medical School, Boston, MA, USA

The authors wish to point out that there is an error of abbreviation in the Abstract (Background paragraph) of this article. The correct Background paragraph of the Abstract should read as follows:

Background

The genome-wide association study has shown that MHC class I chain-related A (MICA) genetic variants were associated with hepatitis C virus (HCV) related hepatocellular carcinoma (HCC). The impact of the genetic variants and its serum levels on post-treatment cohort is elusive.

Also in the Abstract (Results paragraph), "HR/CI: $5\cdot93/1\cdot86-26.38\cdot61$, $P=0\cdot002$ " should read "HR/CI: $5\cdot93/1\cdot86-26\cdot38$, $P=0\cdot002$ ". The correct Results paragraph of the Abstract should read as follow:

Results

Fifty-eight (8·2%) patients developed HCC, with a median follow-up period of 48·2 months (range: 6–129 months). The MICA A allele was

associated with a significantly increased risk of HCC development in cirrhotic non-SVR patients but not in patients of non-cirrhotic and/or with SVR. For cirrhotic non-SVR patients, high sMICA levels (HR/CI: $5\cdot93/1\cdot86-26\cdot38$, $P=0\cdot002$) and the MICA rs2596542 A allele (HR/CI: $4\cdot37/1\cdot52-12\cdot07$, $P=0\cdot002$) were independently associated with HCC development. The risk A allele or GG genotype with sMICA >175 ng/mL provided the best accuracy (79%) and a negative predictive value of 100% in predicting HCC.

Finally, in the Results, Section 3.5. Impact of MICA SNP and sMICA on HCC Development in Non-SVR Patients, the last sentence should read as follow:

Cox regression analysis revealed that the factors independently associated with HCC development among cirrhotic patients without an SVR were high sMICA levels (HR/CI: 5.93/1.86-26.38, P=0.002) and the MICA rs2596542 A allele (HR/CI: 4.37/1.52-12.07, P=0.002).

E-mail address: fish6069@gmail.com (M.-L. Yu).

DOI of original article: http://dx.doi.org/10.1016/j.ebiom.2016.11.031.

^{*} Corresponding author at: Hepatobiliary Division, Department of Internal Medicine, Kaohsiung Medical University Hospital, Kaohsiung, Taiwan.