



Corrigendum: Novel Non-Histocompatibility Antigen Mismatched Variants Improve the Ability to Predict Antibody-Mediated Rejection Risk in Kidney Transplant

Silvia Pineda^{1,2}, Tara K. Sigdel¹, Jieming Chen², Annette M. Jackson³, Marina Sirota^{2,4*†} and Minnie M. Sarwal^{1*†}

¹ Division of Transplant Surgery, Department of Surgery, University of California, San Francisco (UCSF), San Francisco, CA, United States, ² Institute for Computational Health Sciences, University of California, San Francisco (UCSF), San Francisco, CA, United States, ³ Department of Medicine, Division of Immunogenetics and Transplantation Immunology, The Johns Hopkins University School of Medicine, Baltimore, MD, United States, ⁴ Department of Pediatrics, University of California, San Francisco (UCSF), San Francisco, CA, United States

OPEN ACCESS

Edited and Reviewed by:

Dominique Charron, Université Paris Diderot, France

*Correspondence:

Marina Sirota marina.sirota@ucsf.edu; Minnie M. Sarwal minnie.sarwal@ucsf.edu

[†]These authors have contributed equally to this work.

Specialty section:

This article was submitted to Alloimmunity and Transplantation, a section of the journal Frontiers in Immunology

Received: 21 December 2017 Accepted: 12 January 2018 Published: 31 January 2018

Citation:

Pineda S, Sigdel TK, Chen J, Jackson AM, Sirota M and Sarwal MM (2018) Corrigendum: Novel Non-Histocompatibility Antigen Mismatched Variants Improve the Ability to Predict Antibody-Mediated Rejection Risk in Kidney Transplant. Front. Immunol. 9:107. doi: 10.3389/fimmu.2018.00107 Keywords: kidney organ transplant, antibody-mediated rejection, exome sequencing, non-histocompatibility antigen, gene expression, machine learning

A corrigendum on

Novel Non-Histocompatibility Antigen Mismatched Variants Improve the Ability to Predict Antibody-Mediated Rejection Risk in Kidney Transplant

by Pineda S, Sigdel TK, Chen J, Jackson AM, Sirota M, Sarwal MM. Front Immunol (2017) 8:1687. doi: 10.3389/fimmu.2017.01687

Text Correction

In the original article, there was an error. The data were intended to be shared in dbGap, but because of some issues in the inform consent, the data cannot be uploaded to dbGap, so we are modifying the way the data will be shared.

A correction has been made to **Data and Material Availability** and **Paragraph 1**: "The data set for this study will be available at dbGap referenced by the PubMed ID and title of this article."

for: "The exome sequencing data set for this study is available upon request as a collaboration. Please contact Minnie Sarwal at minnie.sarwal@ucsf.edu."

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 was updated.

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Copyright © 2018 Pineda, Sigdel, Chen, Jackson, Sirota and Sarwal. 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 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.