



**Universiteit  
Leiden**  
The Netherlands

**Haplotype motif-based models for KIR-genotype informed selection of hematopoietic cell donors fail to predict outcome of patients with Myelodysplastic Syndromes or secondary Acute Myeloid Leukemia (vol 11, & nbsp;584520, 2021)**

Schetelig, J.; Baldauf, H.; Koster, L.; Kuxhausen, M.; Heidenreich, F.; Wreede, L.C. de; ... ; Robin, M.

**Citation**

Schetelig, J., Baldauf, H., Koster, L., Kuxhausen, M., Heidenreich, F., Wreede, L. C. de, ... Robin, M. (2021). Haplotype motif-based models for KIR-genotype informed selection of hematopoietic cell donors fail to predict outcome of patients with Myelodysplastic Syndromes or secondary Acute Myeloid Leukemia (vol 11, & nbsp;584520, 2021). *Frontiers In Immunology*, 12. doi:10.3389/fimmu.2021.813838

Version: Publisher's Version  
License: [Creative Commons CC BY 4.0 license](#)  
Downloaded from: <https://hdl.handle.net/1887/3276355>

**Note:** To cite this publication please use the final published version (if applicable).



# Corrigendum: Haplotype Motif-Based Models for KIR-Genotype Informed Selection of Hematopoietic Cell Donors Fail to Predict Outcome of Patients With Myelodysplastic Syndromes or Secondary Acute Myeloid Leukemia

Johannes Schetelig<sup>1,2\*</sup>, Henning Baldauf<sup>2</sup>, Linda Koster<sup>3</sup>, Michelle Kuxhausen<sup>4</sup>, Falk Heidenreich<sup>1,2</sup>, Liesbeth C. de Wreede<sup>2,5</sup>, Stephen Spellman<sup>4</sup>, Michel van Gelder<sup>6</sup>, Benedetto Bruno<sup>7</sup>, Francesco Onida<sup>8</sup>, Vinzenz Lange<sup>9</sup>, Carolin Massalski<sup>9</sup>, Victoria Potter<sup>10</sup>, Per Ljungman<sup>11</sup>, Nicolaas Schaap<sup>12</sup>, Patrick Hayden<sup>13</sup>, Stephanie J. Lee<sup>14</sup>, Nicolaus Kröger<sup>15</sup>, Kathy Hsu<sup>16</sup>, Alexander H. Schmidt<sup>2,9</sup>, Ibrahim Yakoub-Agha<sup>17</sup> and Marie Robin<sup>18</sup>

## OPEN ACCESS

### Edited and reviewed by:

Marina Cella,

Washington University School of Medicine in St. Louis, United States

### \*Correspondence:

Johannes Schetelig  
johannes.schetelig@ukdd.de

### Specialty section:

This article was submitted to NK and Innate Lymphoid Cell Biology, a section of the journal *Frontiers in Immunology*

**Received:** 12 November 2021

**Accepted:** 01 December 2021

**Published:** 21 December 2021

### Citation:

Schetelig J, Baldauf H, Koster L, Kuxhausen M, Heidenreich F, de Wreede LC, Spellman S, van Gelder M, Bruno B, Onida F, Lange V, Massalski C, Potter V, Ljungman P, Schaap N, Hayden P, Lee SJ, Kröger N, Hsu K, Schmidt AH, Yakoub-Agha I and Robin M (2021) Corrigendum: Haplotype Motif-Based Models for KIR-Genotype Informed Selection of Hematopoietic Cell Donors Fail to Predict Outcome of Patients With Myelodysplastic Syndromes or Secondary Acute Myeloid Leukemia. *Front. Immunol.* 12:813838. doi: 10.3389/fimmu.2021.813838

<sup>1</sup> Medizinische Klinik und Poliklinik I, University Hospital Dresden, Dresden, Germany, <sup>2</sup> DKMS Clinical Trials Unit, Dresden, Germany, <sup>3</sup> EBMT Data Office Leiden, Leiden, Netherlands, <sup>4</sup> Center for International Blood and Marrow Transplant Research, Minneapolis, MN, United States, <sup>5</sup> Leiden University Medical Center, Department of Biomedical Data Sciences, Leiden, Netherlands, <sup>6</sup> Maastricht University Medical Center, Department of Internal Medicine, Maastricht, Netherlands, <sup>7</sup> A.O.U. Citta della Salute e della Scienza di Torino, Turin, Italy, <sup>8</sup> Fondazione IRCCS Ca'Granda Ospedale Maggiore Policlinico, University of Milan, Milan, Italy, <sup>9</sup> DKMS Life Science Lab, Dresden, Germany, <sup>10</sup> GKT School of Medicine, London, United Kingdom, <sup>11</sup> Karolinska University Hospital and Karolinska Institutet, Stockholm, Sweden, <sup>12</sup> Radboud University Medical Centre, Nijmegen, Netherlands, <sup>13</sup> St. James's Hospital, Dublin, Ireland, <sup>14</sup> Fred Hutchinson Cancer Research Center, Seattle, WA, United States, <sup>15</sup> University Hospital Eppendorf, Hamburg, Germany, <sup>16</sup> Memorial Sloan Kettering Cancer Center, New York & Scientific Director, CIBMTR Immunobiology Working Committee, New York City, NY, United States, <sup>17</sup> CHU de Lille, LIRIC, INSERM U995, Université de Lille, Lille, France, <sup>18</sup> Hopital Saint-Louis, APHP, Université de Paris, Paris, France

**Keywords:** KIR, KIR2DS1, KIR3DL1, hematopoietic stem cell transplantation, donor selection, unrelated donor

## A Corrigendum on

### Haplotype Motif-Based Models for KIR-Genotype Informed Selection of Hematopoietic Cell Donors Fail to Predict Outcome of Patients With Myelodysplastic Syndromes or Secondary Acute Myeloid Leukemia

By Schetelig J, Baldauf H, Koster L, Kuxhausen M, Heidenreich F, de Wreede LC, Spellman S, van Gelder M, Bruno B, Onida F, Lange V, Massalski C, Potter V, Ljungman P, Schaap N, Hayden P, Lee SJ, Kröger N, Hsu K, Schmidt AH, Yakoub-Agha I, Robin M. (2021) *Front. Immunol.* 11:584520. doi: 10.3389/fimmu.2020.584520

There is an error in the Funding statement. The CIBMTR support erroneously included the grant number OT3HL147741 that is not associated with this work.

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.

**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.

Copyright © 2021 Schetelig, Baldauf, Koster, Kuxhausen, Heidenreich, de Wreede, Spellman, van Gelder, Bruno, Onida, Lange, Massalski, Potter, Ljungman, Schaap,

Hayden, Lee, Kröger, Hsu, Schmidt, Yakoub-Agha and Robin. 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.