

PUBLISHER CORRECTION

Open Access



Publisher Correction: Analysis of transcriptomic features reveals molecular endotypes of SLE with clinical implications

Erika L. Hubbard^{1,2*} , Prathyusha Bachali^{1,2}, Kathryn M. Kingsmore^{1,2}, Yisha He³, Michelle D. Catalina⁴, Amrie C. Grammer^{1,2} and Peter E. Lipsky^{1,2}

Publisher Correction: *Genome Med* 15, 84 (2023)
<https://doi.org/10.1186/s13073-023-01237-9>

Figure 5 in the original publication of this article [1] was incorrectly cut-off during the publication process. The incorrect and correct version of Fig. 5 are shown in this correction article. Furthermore, in figure 7 the annotation bars above and below the heatmap were not properly represented, the correct version of figure 7 is shown in this correction article. The original article has been updated. The publisher apologizes for the inconvenience caused to authors & readers.

The original article can be found online at <https://doi.org/10.1186/s13073-023-01237-9>.

*Correspondence:

Erika L. Hubbard
erika.hubbard@ampelbiosolutions.com

¹ AMPEL BioSolutions, LLC, 250 W. Main St. #300, Charlottesville, VA 22902, USA

² RILITE Research Institute, Charlottesville, VA 22902, USA

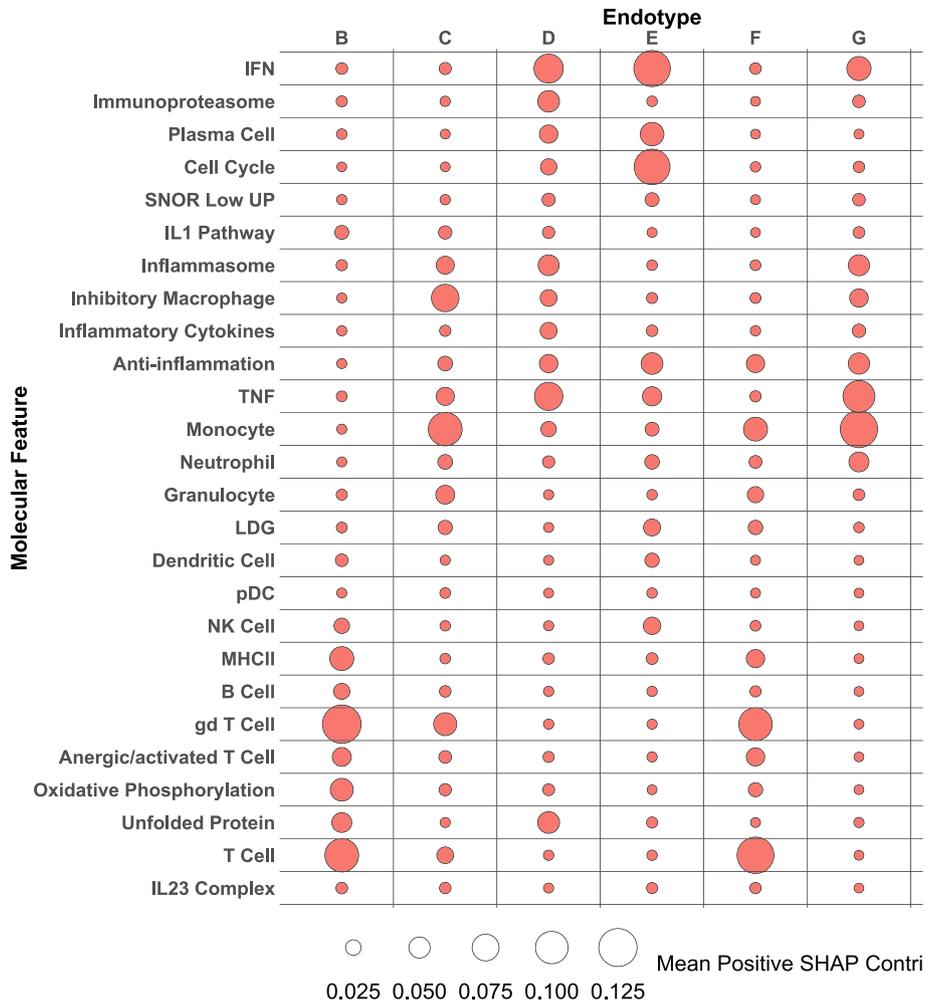
³ Altria, Richmond, VA 23230, USA

⁴ AbbVie, Worcester, MA 01605, USA



© The Author(s) 2023. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

Incorrect figure



Correct figure

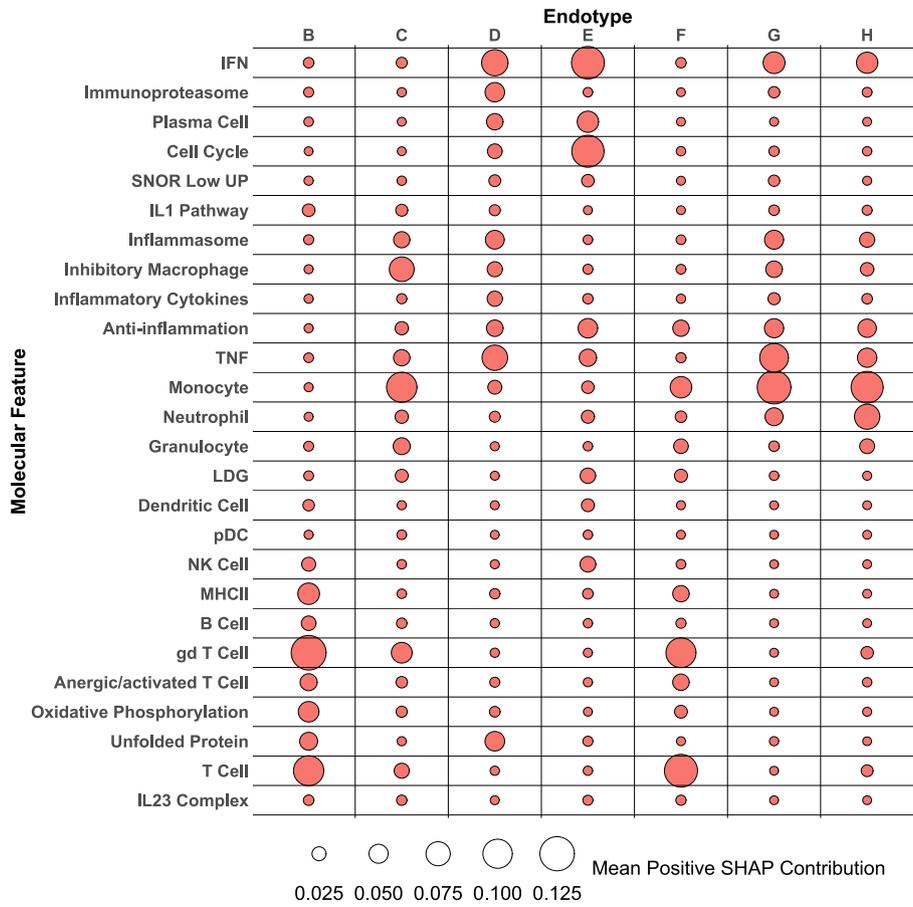
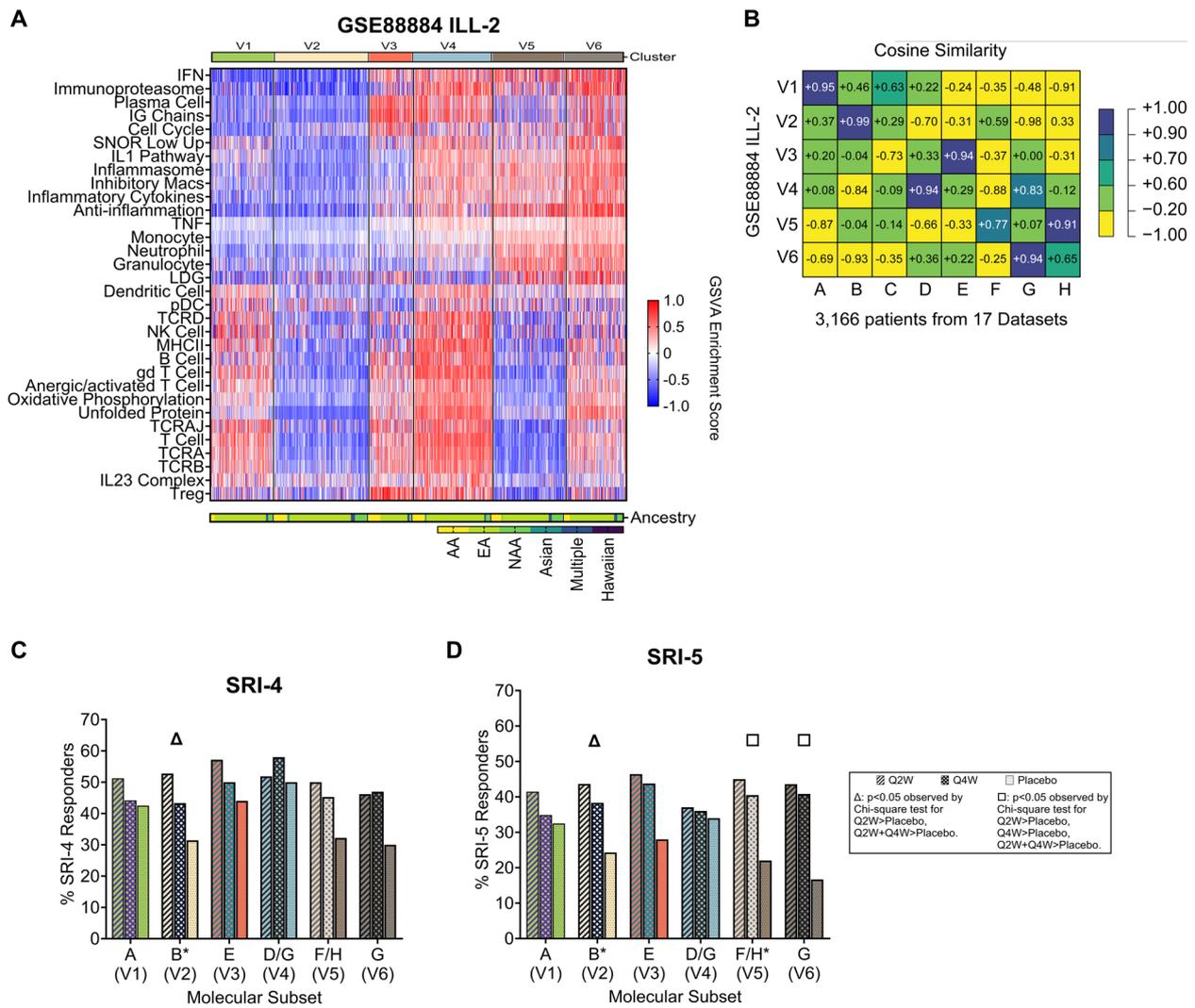


Fig. 7



Published online: 13 December 2023

Reference

- Hubbard EL, et al. Analysis of transcriptomic features reveals molecular endotypes of SLE with clinical implications. *Genome Med.* 2023;15:84. <https://doi.org/10.1186/s13073-023-01237-9>.