ERRATUM



Open Access



Erratum: Computational identification of multi-omic correlates of anticancer therapeutic response

Lindsay C Stetson^{1,2}, Taylor Pearl³, Yanwen Chen¹ and Jill S Barnholtz-Sloan^{1,2*}

Erratum to: BMC Genomics doi:10.1186/1471-2164-15-S7-S2

Following the publication of our recent article in *BMC Genomics* [1] we wish to acknowledge the contribution to the concept for our study provided by the previous work of Papillon-Cavanagh et al. [2] and note that we should have included a reference to their paper in the legend to Figure 1. Furthermore when discussing our findings that NQO1 is a marker for sensitivity of TNBC to 17-AAG we should have cited Papillon-Cavanagh et al. [2] as a previous paper where similar results were reported. We apologize to all affected parties.

Author details

¹Case Comprehensive Cancer Center, Cleveland, OH, USA. ²Center for Proteomics and Bioinformatics, Case Western Reserve University, Cleveland, OH 44106, USA. ³Department of Biological Engineering, Massachusetts Institute of Technology, Cambridge, MA 02139, USA.

Received: 13 May 2015 Accepted: 13 May 2015 Published online: 30 June 2015

References

- Stetson LC, Pearl T, Chen Y, Sloan JS B. Computational identification of multi-omic correlates of anticancer therapeutic response. BMC Genomics. 2014;15 Suppl 7:S2.
- Papillon-Cavanagh S, De Jay N, Hachem N, Olsen C, Bontempi G, Aerts HJ, et al. Comparison and validation of genomic predictors for anticancer drug sensitivity. Journal of the American Medical Informatics Association: JAMIA. 2013;20(4):597–602.

* Correspondence: jsb42@case.edu

¹Case Comprehensive Cancer Center, Cleveland, OH, USA

 $^{2}\text{Center}$ for Proteomics and Bioinformatics, Case Western Reserve University, Cleveland, OH 44106, USA

Full list of author information is available at the end of the article



© 2015 Stetson et al. This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly credited. 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.