ERRATUM

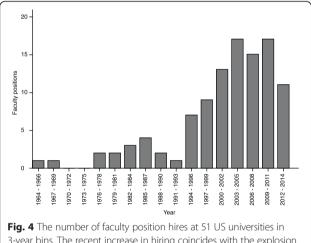
Open Access



Erratum to: The real cost of sequencing: scaling computation to keep pace with data generation

Paul Muir^{1,2,3}, Shantao Li⁴, Shaoke Lou^{4,5}, Daifeng Wang^{4,5}, Daniel J. Spakowicz^{4,5}, Leonidas Salichos^{4,5}, Jing Zhang^{4,5}, George M. Weinstock⁶, Farren Isaacs^{1,2}, Joel Rozowsky^{4,5} and Mark Gerstein^{4,5,7*}

After the publication of this work [1], the authors noticed an error in Fig. 4. The error occurred on the y-axis, instead of running from 0 to 20 it just has 20 at each point. The publisher apologises for the error and any inconvenience caused. The corrected figure is given below:



3-year bins. The recent increase in hiring coincides with the explosion in sequencing data. Data were obtained from http://jeffhuang.com/ computer_science_professors.html

Author details

¹Department of Molecular, Cellular and Developmental Biology, Yale University, New Haven, CT 06520, USA. ²Systems Biology Institute, Yale University, West Haven, CT 06516, USA. ³Integrated Graduate Program in Physical and Engineering Biology, Yale University, New Haven, CT 06520, USA. ⁴Program in Computational Biology and Bioinformatics, Yale University, New Haven, CT 06520, USA. ⁵Department of Molecular Biophysics and Biochemistry, Yale University, New Haven, CT 06520, USA. ⁶The Jackson Laboratory for Genomic Medicine, Farmington, CT 06032, USA. ⁷Department of Computer Science, Yale University, New Haven, CT 06520, USA.

Received: 1 April 2016 Accepted: 20 April 2016 Published online: 28 April 2016

Reference

 Muir P, Li S, Lou S, Wang D, Spakowicz DJ, Salichos L, Zhang J, Weinstock GM, Isaacs F, Rozowsky J and Gerstein M. The real cost of sequencing: scaling computation to keep pace with data generation. Genome Biol. 2016;17:53.

Submit your next manuscript to BioMed Central and we will help you at every step:

- We accept pre-submission inquiries
- Our selector tool helps you to find the most relevant journal
- We provide round the clock customer support
- Convenient online submission
- Thorough peer review
- Inclusion in PubMed and all major indexing services
- Maximum visibility for your research

Submit your manuscript at www.biomedcentral.com/submit



© 2016 Muir et al. **Open Access** This article is distributed under the terms of the Creative Commons Attribution 4.0 International License (http://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. 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.

* Correspondence: mark@gersteinlab.org
⁴Program in Computational Biology and Bioinformatics, Yale University, New

⁵Department of Molecular Biophysics and Biochemistry, Yale University, New Haven, CT 06520, USA



Haven, CT 06520, USA