



Erratum: Proving sequence aligners can guarantee accuracy in almost $O(m \log n)$ time through an average-case analysis of the seed-chain-extend heuristic

Jim Shaw and Yun William Yu

Genome Res. 2024 34: 810

Access the most recent version at doi:[10.1101/gr.279618.124](https://doi.org/10.1101/gr.279618.124)

Open Access Freely available online through the *Genome Research* Open Access option.

License

Email Alerting Service Receive free email alerts when new articles cite this article - sign up in the box at the top right corner of the article or [click here](#).



To subscribe to *Genome Research* go to:
<https://genome.cshlp.org/subscriptions>

Published by Cold Spring Harbor Laboratory Press

Erratum

***Genome Research* 33: 1175–1187 (2023)**

Erratum: Proving sequence aligners can guarantee accuracy in almost $O(m \log n)$ time through an average-case analysis of the seed-chain-extend heuristic

Jim Shaw and Yun William Yu

In the initial publication of the article mentioned above, one of the corresponding authors' email addresses was inadvertently omitted. The correct email addresses are as follows:

Corresponding authors: ywyu@math.toronto.edu, jshaw@math.toronto.edu

This article has been corrected online.

The current affiliation of Yun William Yu is Carnegie Mellon University (ywyu@cmu.edu).

doi: 10.1101/gr.279618.124