Check for updates

Publisher Correction: Rotamer-free protein sequence design based on deep learning and self-consistency

Yufeng Liu[®], Lu Zhang[®], Weilun Wang[®], Min Zhu, Chenchen Wang, Fudong Li, Jiahai Zhang, Houqiang Li[®], Quan Chen[®] and Haiyan Liu[®]

Correction to: Nature Computational Science https://doi.org/10.1038/s43588-022-00273-6, published online 21 July 2022.

In the version of this article initially published, reference 64 (Liu, Y. et al. ABACUS-R: Rotamer-free protein sequence design based on deep learning and self-consistency (Code Ocean, 2022); https://doi.org/10.24433/CO.3351944.v1) was omitted from the reference list and the Code availability section. It has been restored to the HTML and PDF versions of the article.

Published online: 2 August 2022

https://doi.org/10.1038/s43588-022-00305-1

© The Author(s), under exclusive licence to Springer Nature America, Inc. 2022