Journal of Cheminformatics

CORRECTION

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Correction: StreaMD: the toolkit for high-throughput molecular dynamics simulations

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Following publication of the original article [1], the authors identified that section Availability and requirements is missing.

Availability and requirements
Project name: StreaMD
GitHub: https://github.com/ci-lab-cz/streamd
Operating system(s): Linux
Programming language: Python 3
Other requirements: GROMACS, RDKit, ProLIF, Antechamber, MDAnalysis, Dask, Gaussian (optional, a license is required)
License: MIT
Any restrictions to use by non-academics: no

The original article has been corrected.

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Reference
1. Ivanova A, Mokshyna O, Polishchuk P (2024) StreaMD: the toolkit for high-throughput molecular dynamics simulations. J Cheminform 16:123. https://doi.org/10.1186/s13321-024-00918-w

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