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

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