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MolSysMT: A computational laboratory bench to work with biomolecular systems in Python.

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Abstract

Summary: Bla, bla, bla.

Availability and implementation: The open source code is available at the uibcdf/MolSysMT GitHub repository to be used, modified and shared according to the MIT License. A conda package with the latest stable version is distributed through the "uibcdf" conda channel. And installation instructions, user's documentation and jupyter notebooks with illustrative examples are available at http://www.uibcdf.org/MolSysMT.

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3 Implementation

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