

POSTER PRESENTATION

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MoSGrid: efficient data management and a standardized data exchange format for molecular simulations in a grid environment

Georg Birkenheuer¹, Dirk Blunk², Sebastian Breuers², André Brinkmann¹, Ines dos Santos Vieira³, Gregor Fels¹, Sandra Gesing⁴, Richard Grunzke⁵, Sonja Herres-Pawlis³, Oliver Kohlbacher^{4*}, Jens Krüger¹, Ulrich Lang², Lars Packschies², Ralph Müller-Pfefferkorn⁵, Patrick Schäfer⁶, Thomas Steinke⁶, Klaus-Dieter Warzecha², Martin Wewior²

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The MoSGrid (Molecular Simulation Grid) project is currently establishing a platform that aims to be used by both experienced and inexperienced researchers to submit molecular simulation calculations, monitor their progress, and retrieve the results. It provides a web-based portal to easily set up, run, and evaluate molecular simulations carried out on D-Grid resources. The range of applications available encompasses quantum chemistry, molecular dynamics, and protein-ligand docking codes.

In addition, data repositories were developed, which contain the results of calculations as well as “recipes” or workflows. These can be used, improved, and distributed by the users. A distributed high-throughput file system allows efficient access to large amounts of data in the repositories. For storing both the input and output of the calculations, we have developed MSML (Molecular Simulation Markup Language), a CML derivative (Chemical Markup Language). MSML has been designed to store structural information on small as well as large molecules and results from various molecular simulation tools and docking tools. It ensures interoperability of different tools through a consistent data representation.

At <http://www.mosgrid.de> the new platform is already available to the scientific community in a beta test phase. Currently, portlets for generic workflows, Gaussian, and Gromacs applications are publicly accessible [1,2].

* Correspondence: oliver.kohlbacher@uni-tuebingen.de

⁴Eberhard-Karls-Universität Tübingen, Tübingen 72074, Germany
Full list of author information is available at the end of the article

Author details

¹Universität Paderborn, Paderborn 33098, Germany. ²Universität zu Köln, Köln 50923, Germany. ³Fakultät Chemie, Technische Universität Dortmund, Dortmund 44221, Germany. ⁴Eberhard-Karls-Universität Tübingen, Tübingen 72074, Germany. ⁵Technische Universität Dresden, Dresden 01187, Germany. ⁶Konrad-Zuse-Zentrum für Informationstechnik, Berlin 14195, Germany.

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