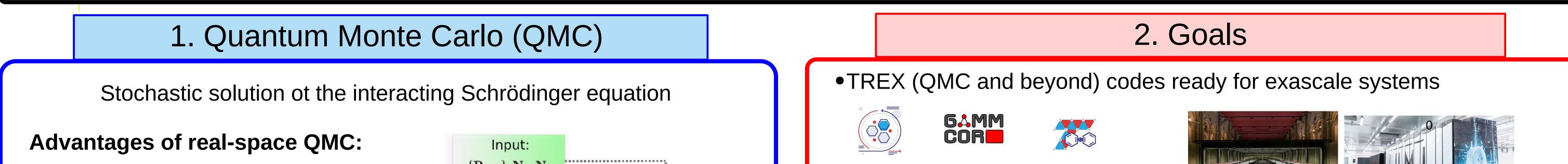
Leveraging stochastic electronic structure methods at the exascale

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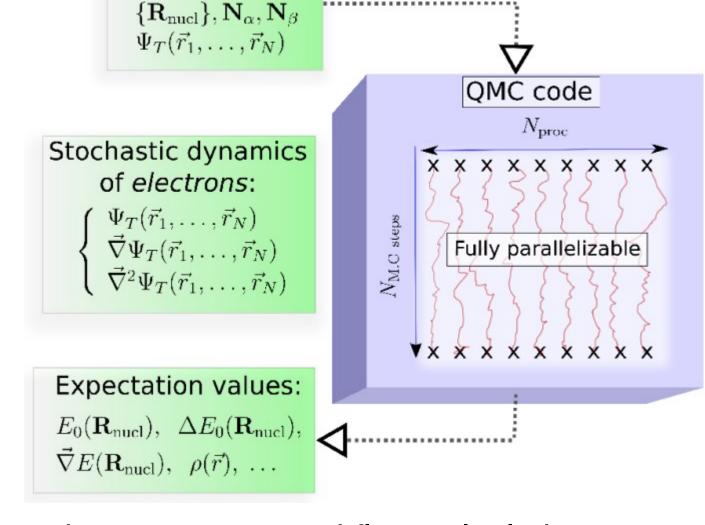
Abstract

We present here the software development strategy and the current achievements of the European Centre of Excellence "Targeting Real chemical accuracy at the EXascale" (TREX). The main objective of TREX is the development of a user-friendly and open-source software suite in the domain of stochastic electronic structure simulations, which integrates a set of flagship quantum Monte Carlo codes within an interoperable, high-performance platform. Core of our software efforts is the creation of two libraries, the TREXIO and the quantum Monte Carlo kernel library (QMCkl).



- Low memory requirements
- Distribute walkers on cores/nodes
- Often no blocking communication:
 - near-ideal scaling
- One QMC step is fast (~1ms)

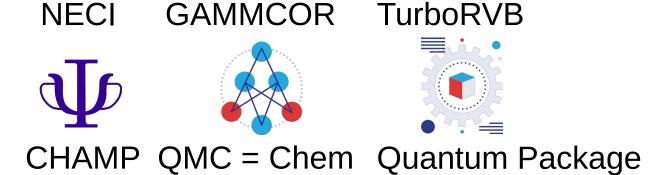
Challenges:

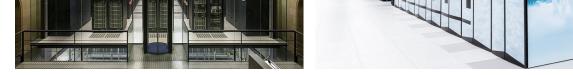


- Figure 1. QMC workflow calculation
- Linear algebra with small matrices
- Difficult to parallelize within a QMC trajectory

4. TREXIO library

- Source code in pure C (C99) for the best performance and portability
- Bindings in Fortran (using ISO_C_BINDING) and Python (using SWIG)
- Official releases are available via GitHub \mathbf{O} , Conda \mathbb{C} , and Spack





- Figure 2. EU exascale Leonardo and LUMI supercomputers
- Design ecosystem of efficient, scalable, and inter-operable software
- Robust management of complex workflows SAIIDA

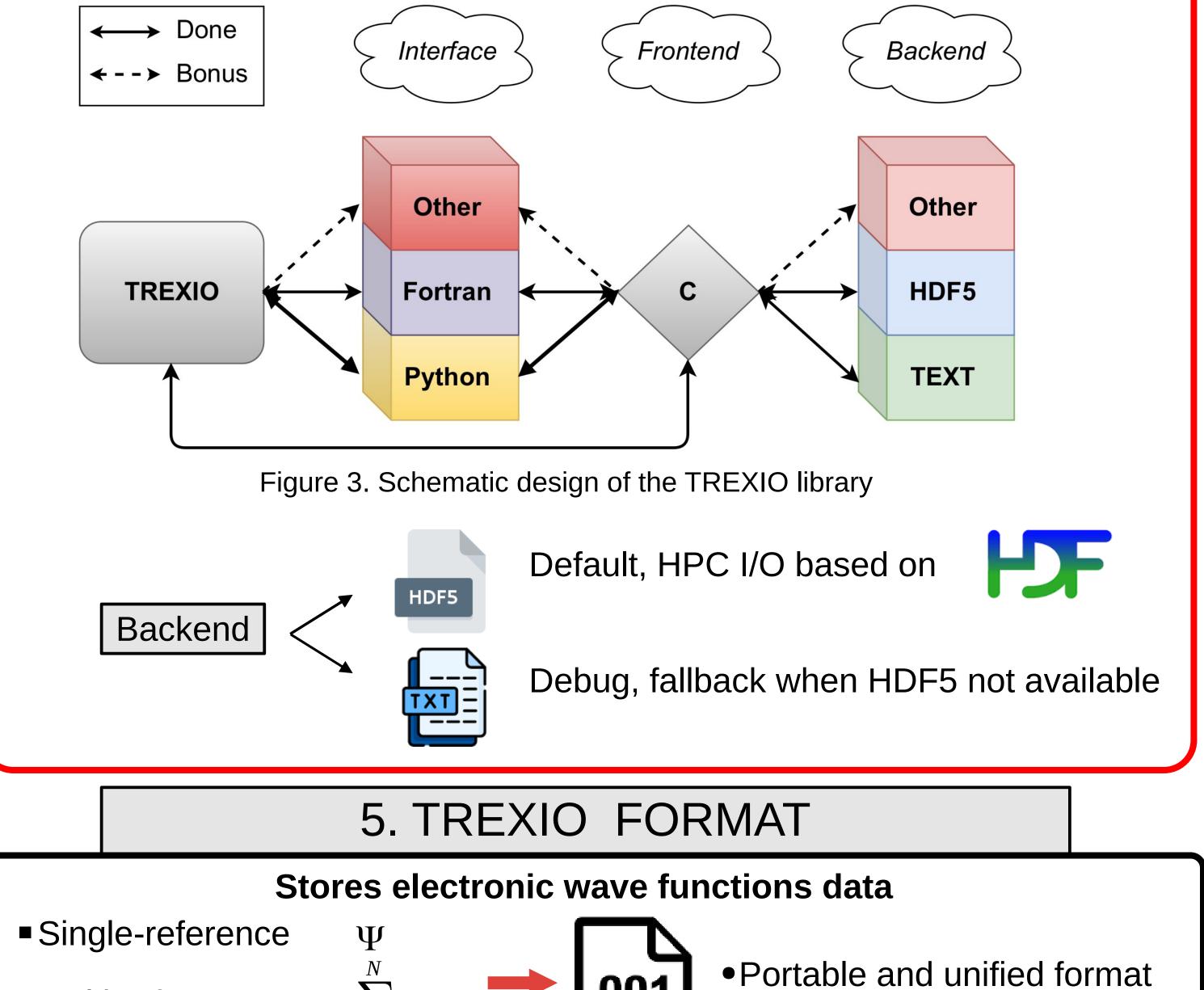
3. Strategy

Provide libraries with the common computational kernels for QMC instead of rewriting individual code

- Kernels developed by QMC experts in a human readable library
- Compile kernels into a HPC-library by HPC experts
- Scientist controls programming choices keeping the computational performance

6. QMCkl library

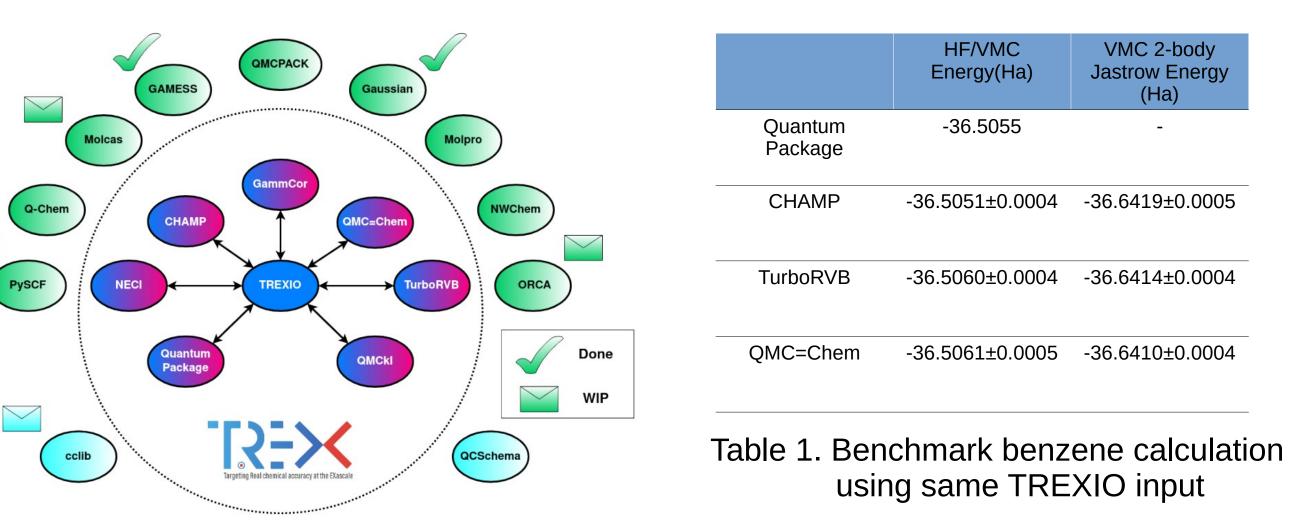
QMCkl is a C library with C-compatible API



- System functions in C (memory allocation, thread safety, etc)
- Computational kernels in Fortran for readability
- Documentation library: Literate programming with org-mode

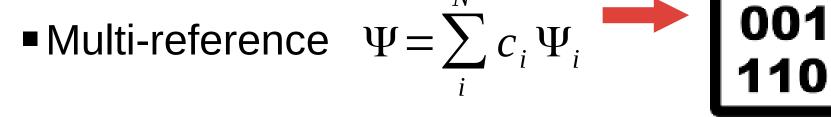
7. Demonstrations

QMC ecosystem Interoperability between codes through unified TREXIO format



Performance improvement

QMCkl Integration with codes and kernels customization



- Data is organized into groups
- Metadata • basis • cell • ao • mo • pbc •ao_1e_int •mo_1e_int Electron • ecp • Etc.. • qmc determinant
 ao_2e_int
 mo_2e_int Nucleus • rdm

•HPC I/O

Acknowledgments

Targeting Real chemical accuracy at the EXascale

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Spe.

customization for CHAMP 3-body

Jastrow factor and DGEMM in QMCkl

Get libraries here!

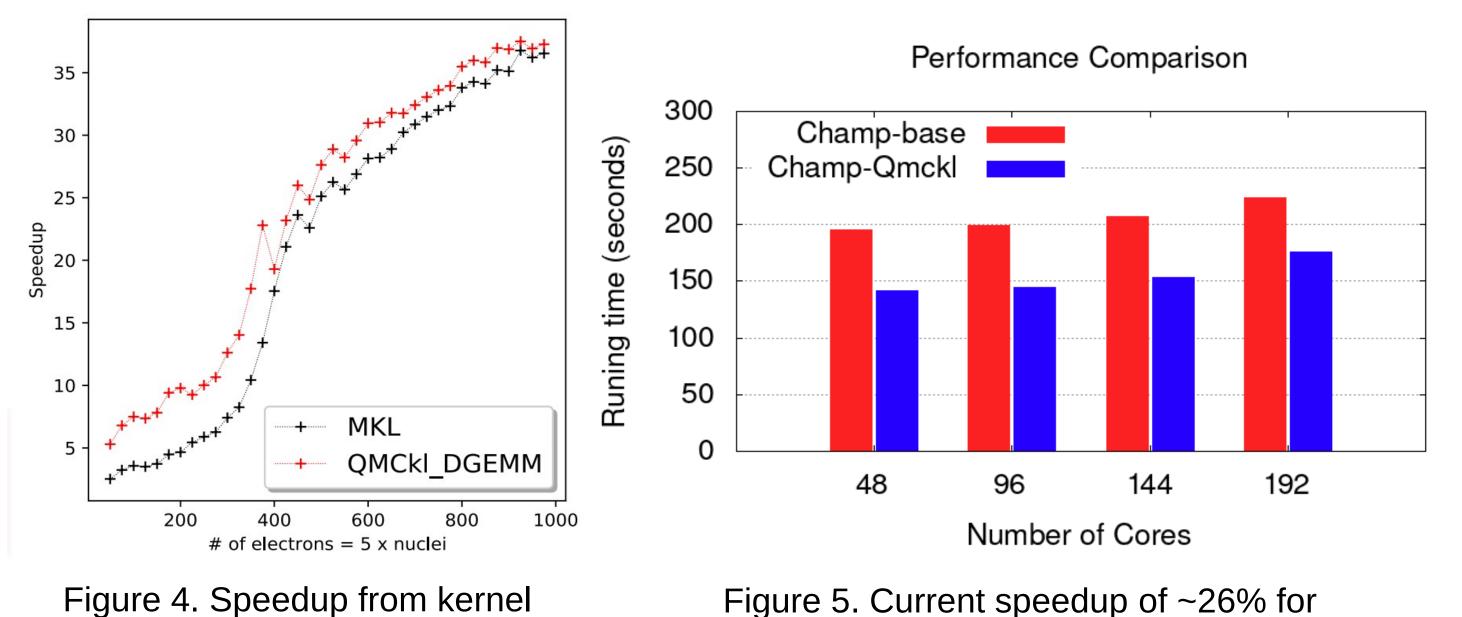


Figure 5. Current speedup of ~26% for energy calculation in CHAMP after integrating QMCkl in MO-subroutines