

User-friendly workflows in quantum chemistry



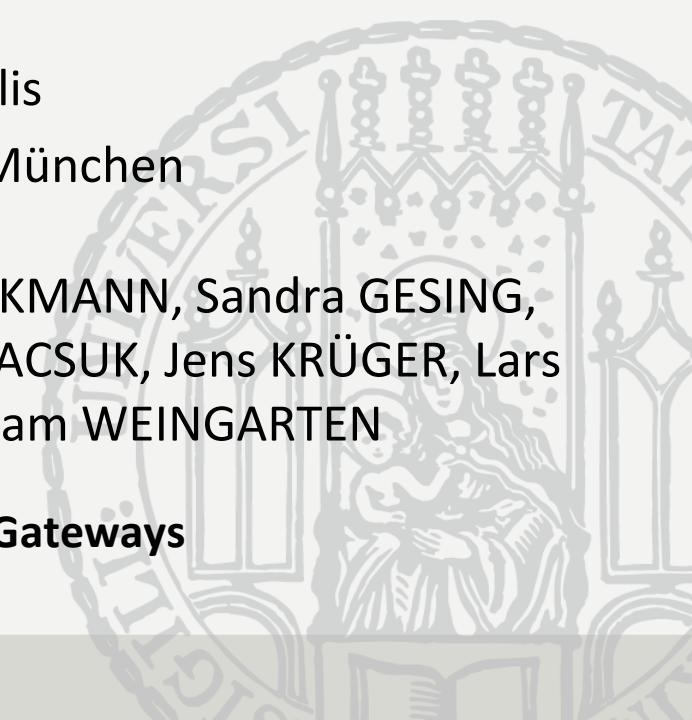
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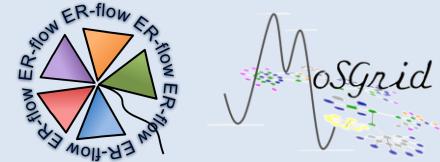
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PACKSCHIES, Gabor TERSTYANSKY, and Noam WEINGARTEN

International Workshop on Science Gateways

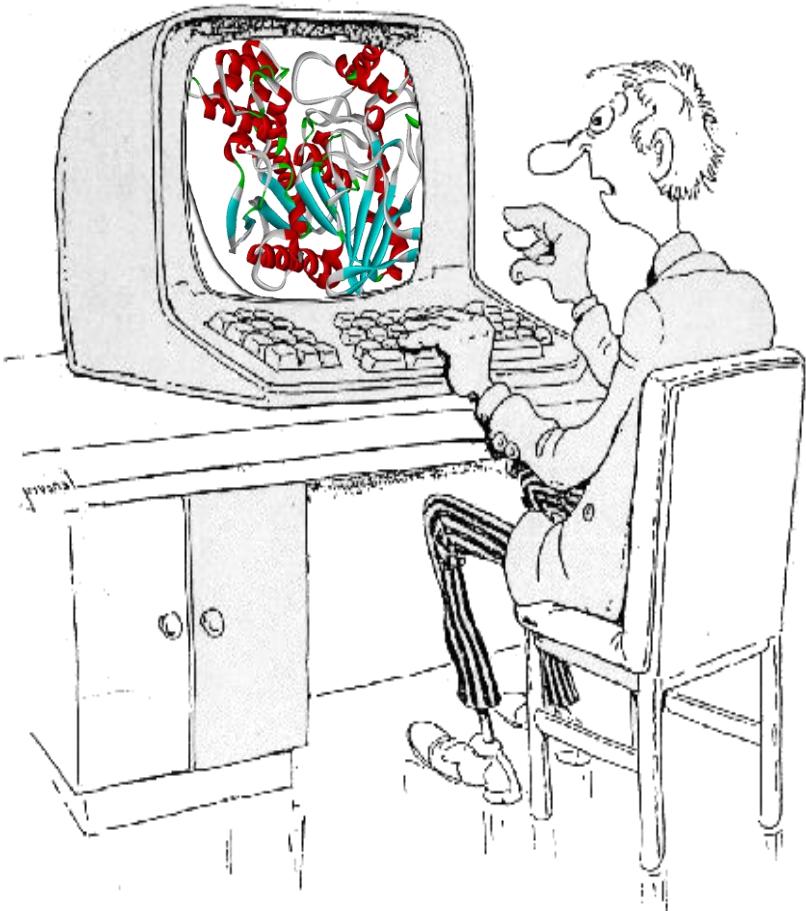
3.06.2013, Zürich



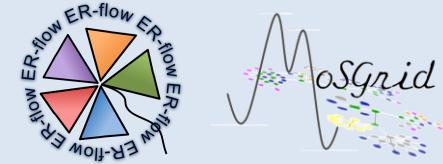
Motivation



- Excellent simulation codes available for all chemical domains
- Huge amount of computational resources
- Time consuming access even for experienced users
- Impassable obstacle for beginners

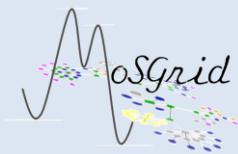


Aims of MoSGrid



- Enabling easier access to HPC (high-performance computing) facilities
- Allowing inexperienced scientists to run molecular simulations on DCIs
- Offering an intuitive user interface (gateway)
- Enabling easy access to recipes, workflows, data repositories

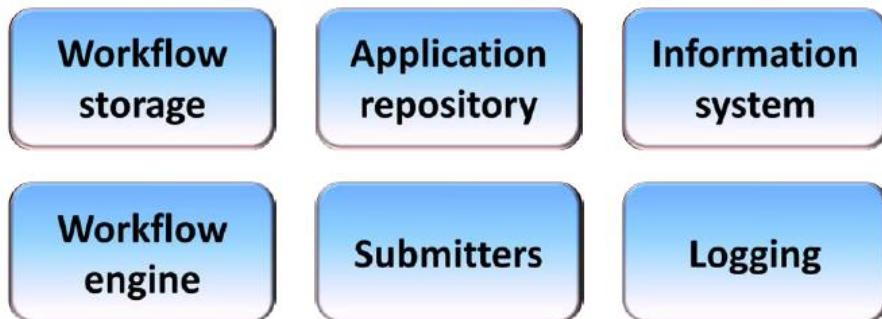
Molecular Simulation Grid



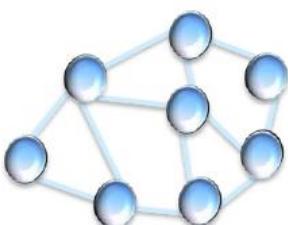
The MoSGrid portal

One goal of the interdisciplinary project [MoSGrid](#) (Molecular Simulation Grid) is to offer molecular simulation tools and docking tools via a port. MoSGrid portal is designed as workflow-enabled grid portal developed on top of the Lifcrai version of WS-PGRADE. WS-PGRADE is a usable, highly-flexible graphical user interface for the grid User Support Environment (gUSE) developed by [MTA SZTAKI](#).

User interface WS-PGRADE

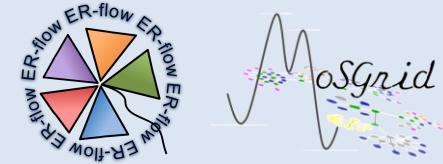


High-level
middleware
service layer
gUSE



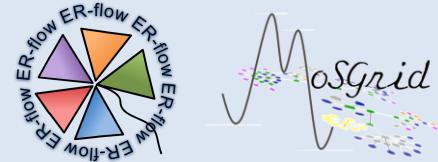
Grid resources
middleware layer
UNICORE 6

MoSGrid Science Gateway



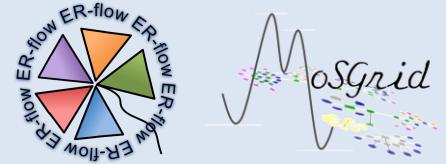
- Portal framework: Liferay
 - Versatile configuration possibilities
 - Allows to provide a user-friendly portal
- Certificate Portlet
 - Easy Management of access to DCIs
- Submission of chemical calculations *via* gUSE
- High-level middleware: WS-PGRADE
- Middleware: UNICORE
- Resources: Wide range of German D-Grid resources (in the future also European DCIs)

Molecular Simulation Grid



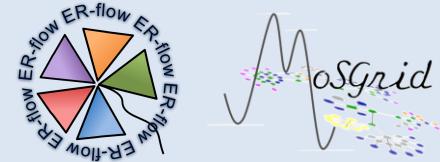
- Start in 2008: 50 associated chemical work groups
- October 2010: Mailing list with 100 addressees as MoSGrid Community
- December 2011: after the community meeting increase to 120 interested chemists from all three domains
- Community serves as pool for requirements analyses.
- Today: 20 real expert users
40 standard users

Use of Workflows in Domains



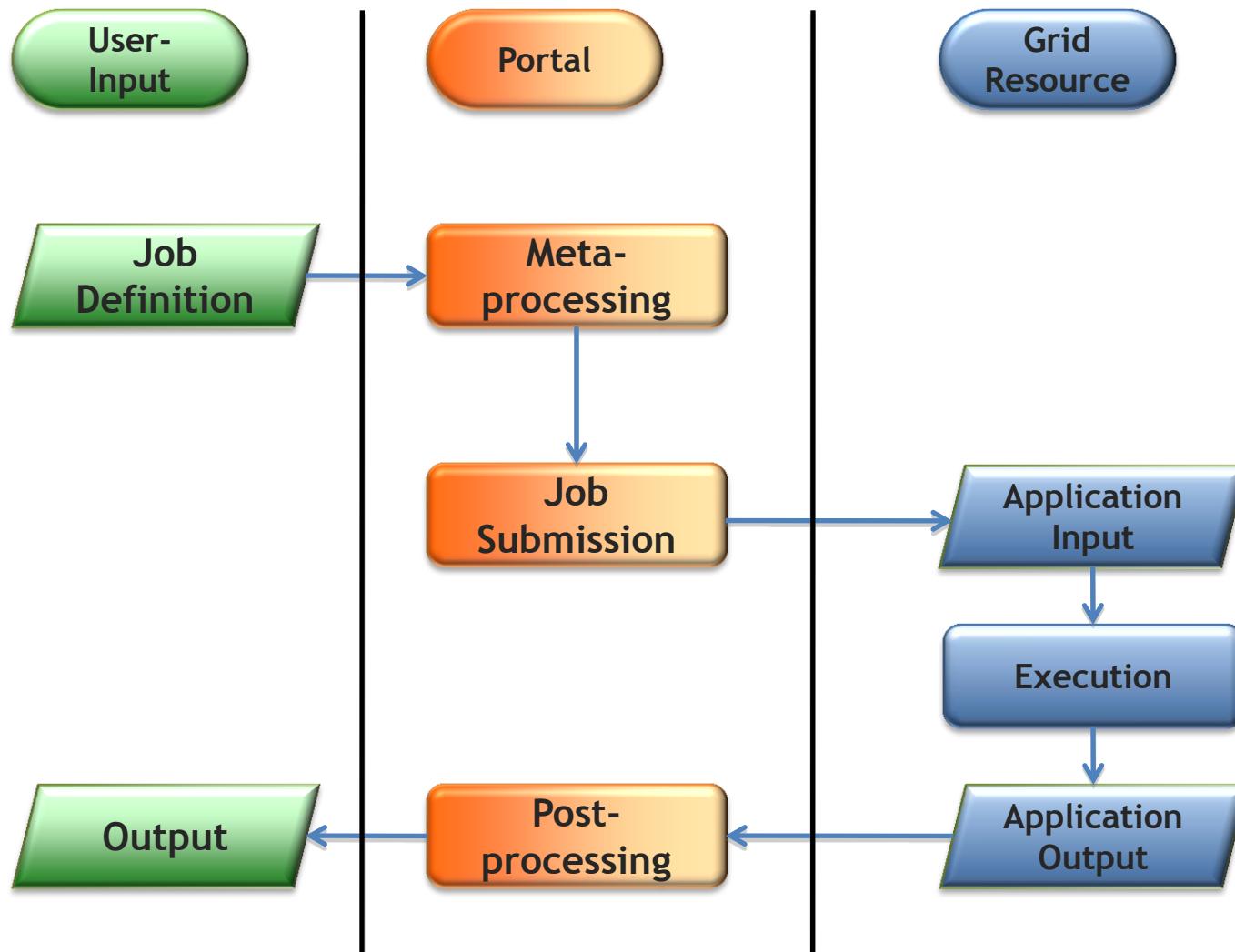
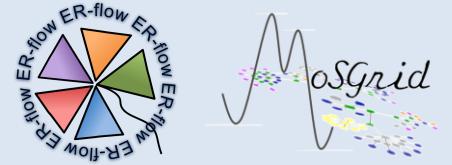
- Domains
 - Quantum chemistry
 - Molecular dynamics
 - Docking
- Identifying standard recipes
 - Modelling of suited workflows
- Goals:
 - For Power Users
 - Possibility to design own Workflows
 - For Novices
 - Offer standard workflows

Use of Workflows in Domains

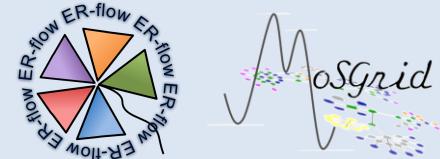


- Quantum Chemical Calculations
 - based on approximated solutions of the Schrödinger equation
 - Average scalability
 - Gaussian, NWChem, Turbomole, ...
- Molecular Dynamics
 - based on forcefields describing molecular interactions
 - Good scalability
 - Gromacs, NWChem, Amber, ...
- Docking
 - Based on simplified forcefields
 - Excellent scalability
 - CADDSuite, FlexX, ...

Basic Workflow



The QC Portlet @ MoSGrid.de



QCPortletVAPI

Import Submission Monitoring About

Welcome

Welcome to the Quantum Chemistry portlet.

Import a workflow

Please select a toolsuite
Gaussian 09

Please select a workflow
G09Minimal

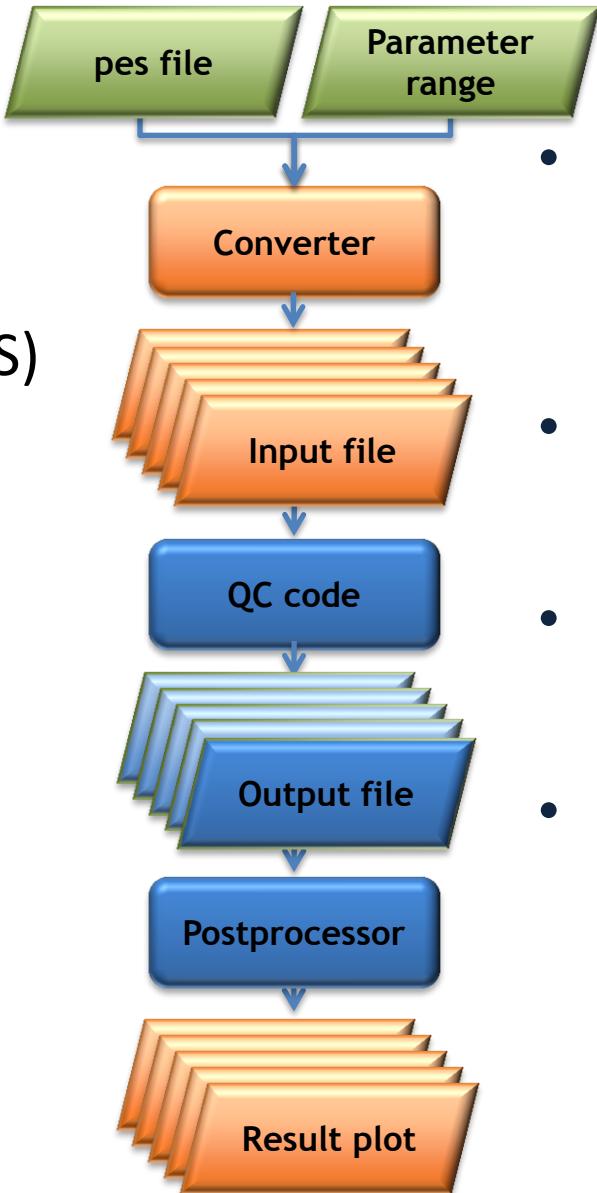
Please enter a name
TEGqu_26

- Specialised interface for quantum chemistry software (Gaussian, Turbomole, NWChem)
- Basic workflows
- Easy Generation or Uploading of Input Files
- Parsing of result files

Example: Parameter Sweep

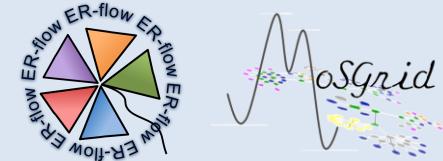


- Parameter Sweep
 - Scan of a potential energy surface (PES)
 - Change of one parameters crucial for theoretical analysis
 - Reaction path analysis
 - Pre-defined steps

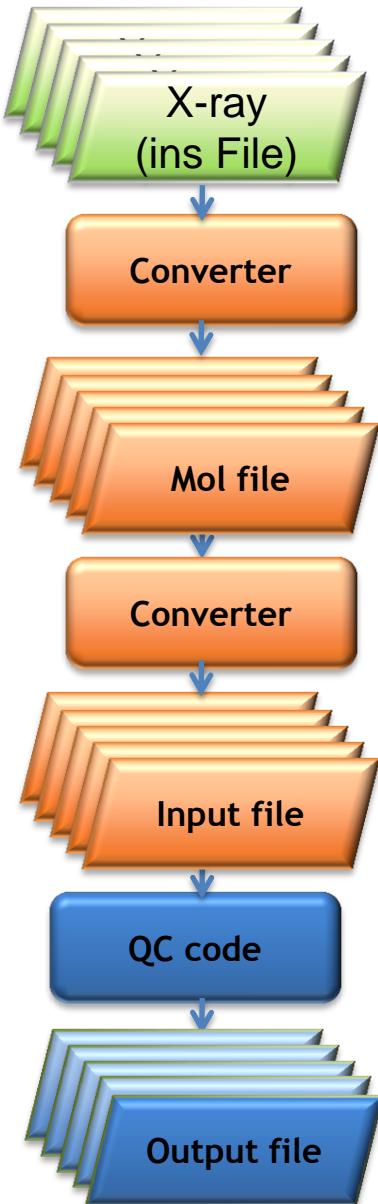


- Input
 - Molecular structure
 - Parameter
 - Parameter range
- First step
 - automatic generation of Inputfiles
- Second step
 - Submission into the Grid
- Third step
 - Plotting of obtained energies against the chosen parameter

Example: High Throughput



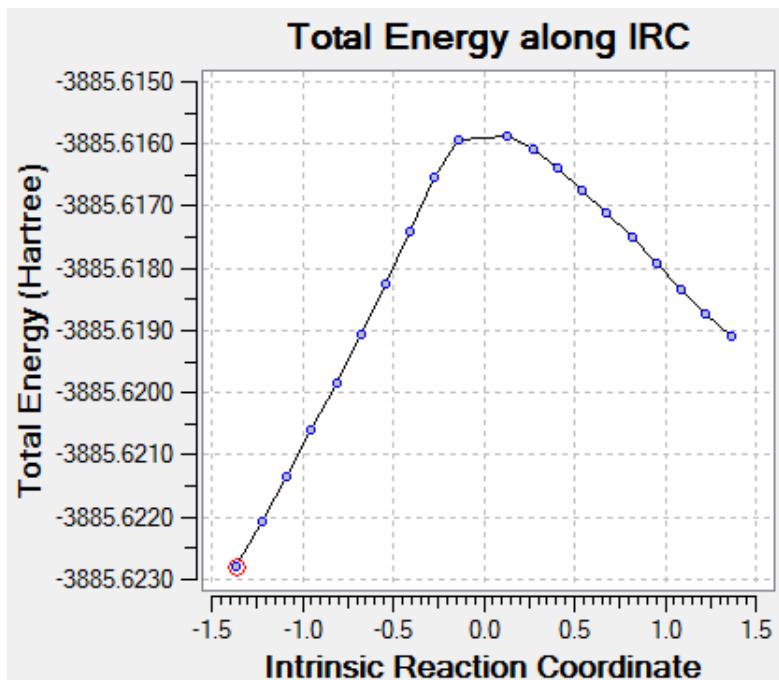
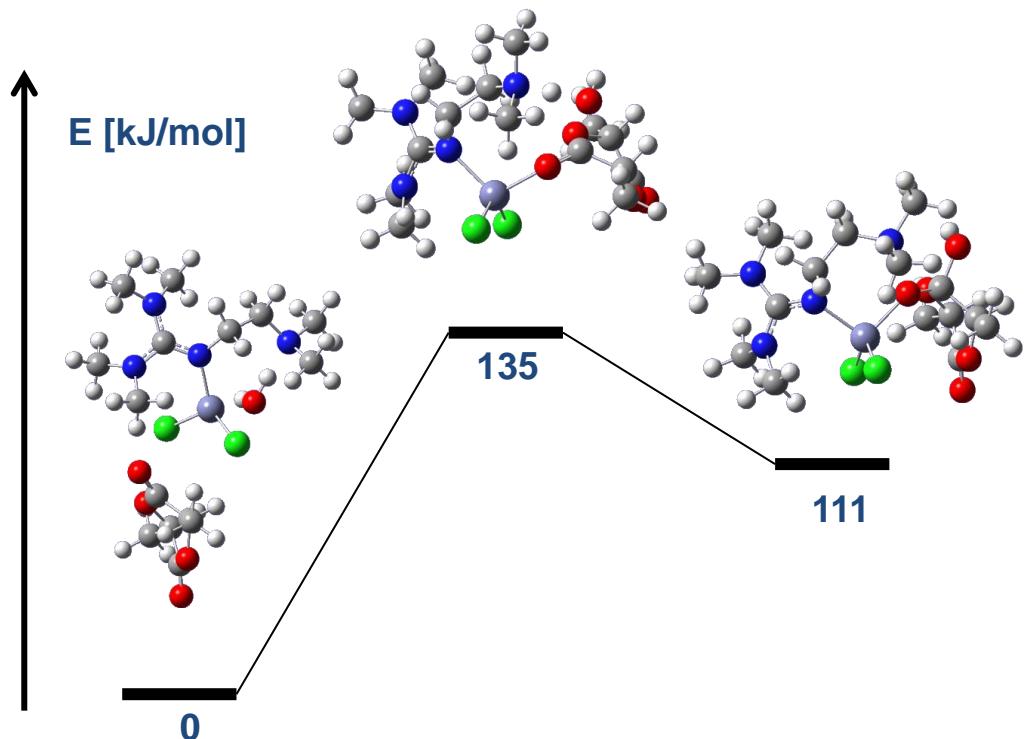
- Native X-ray format needs to be converted into a computational readable mol file.
- Time-consuming process of manual conversion can be transferred into the portlet.



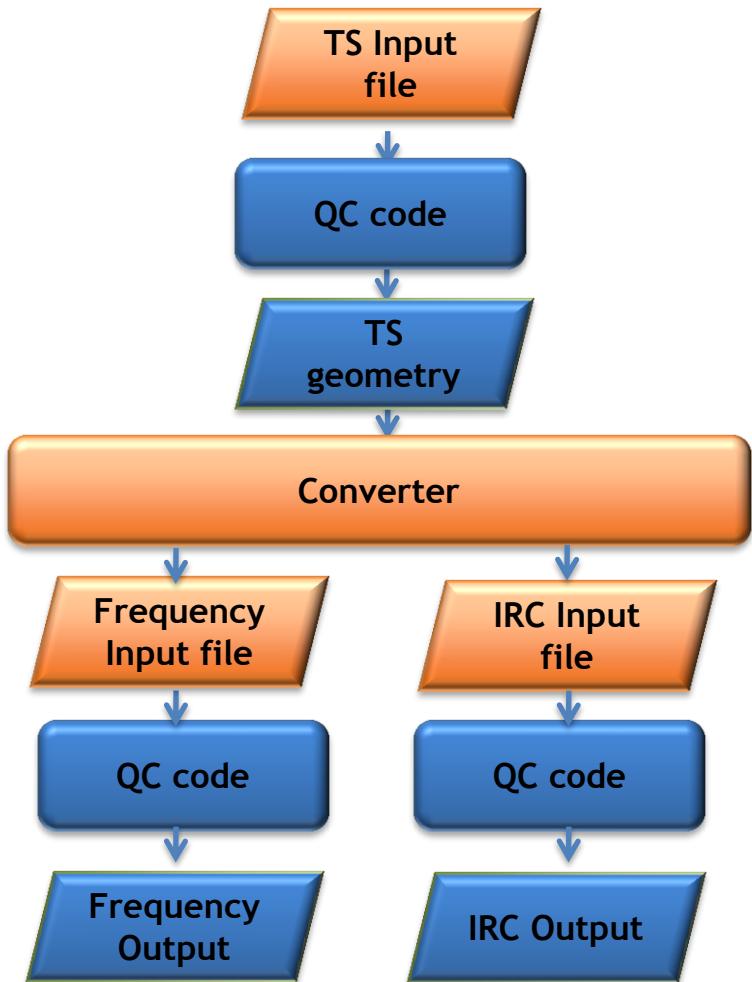
- Input
 - X-ray data
- First step
 - Conversion into mol file
- Second step
 - Conversion into job file with pre-defined job parameters (functional, basis set)
- Third step
 - Submission into the Grid

Example: Transition State Analysis

- Zinc complex reacting with lactide

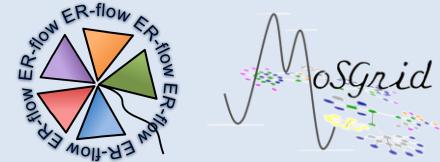


Example: Transition State Analysis

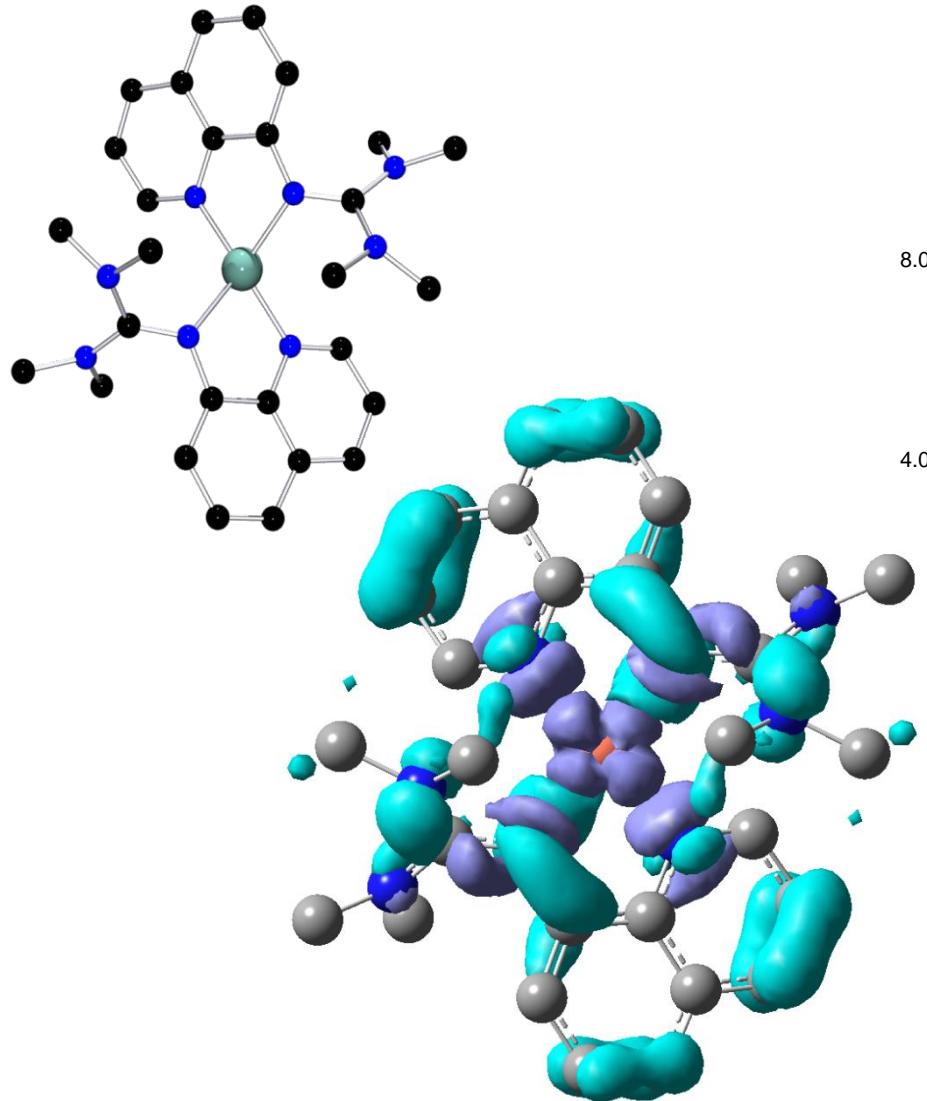


- **Input**
 - Transition state job file (e.g. QST3 file)
- **First step**
 - Calculation of TS geometry
- **Second step**
 - Conversion into job files for frequency and IRC calculations
- **Third step**
 - Calculation of frequency and reaction path

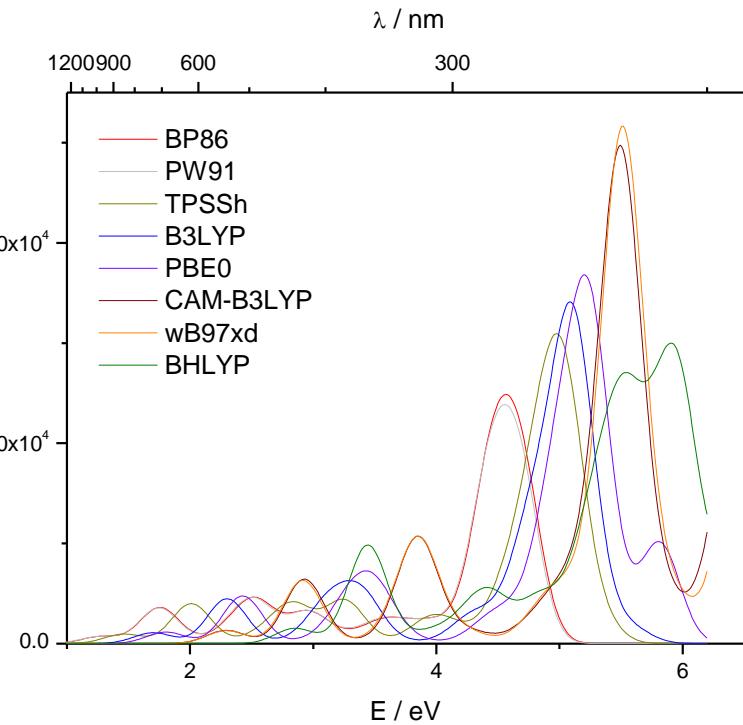
Example: Spectroscopic Analysis



Structure

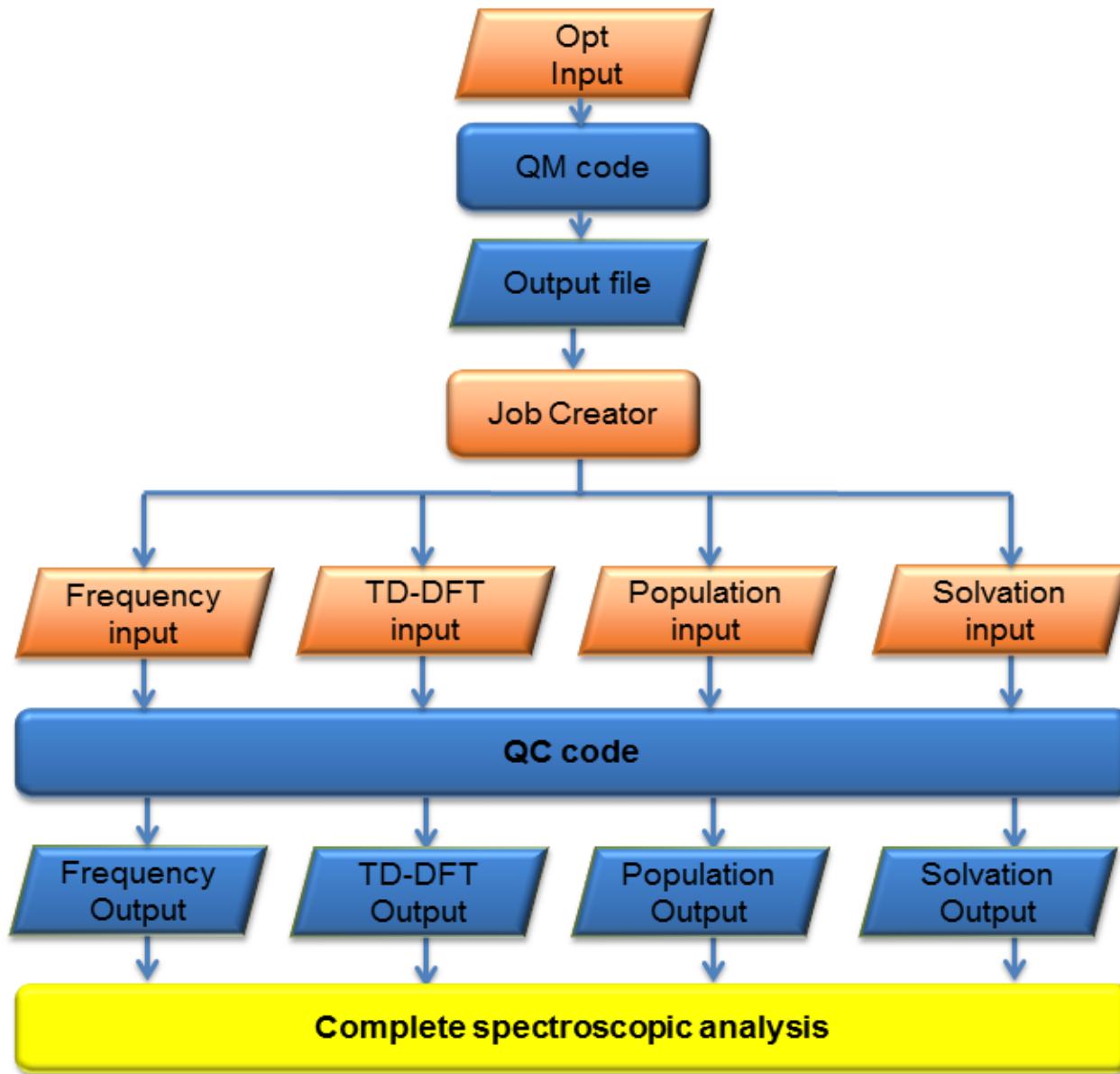
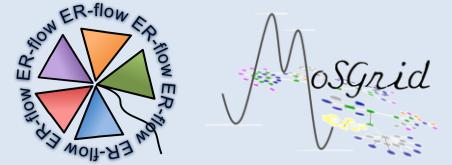


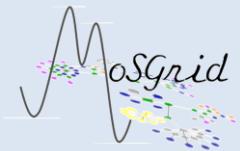
Time-dependent DFT for UV/Vis prediction



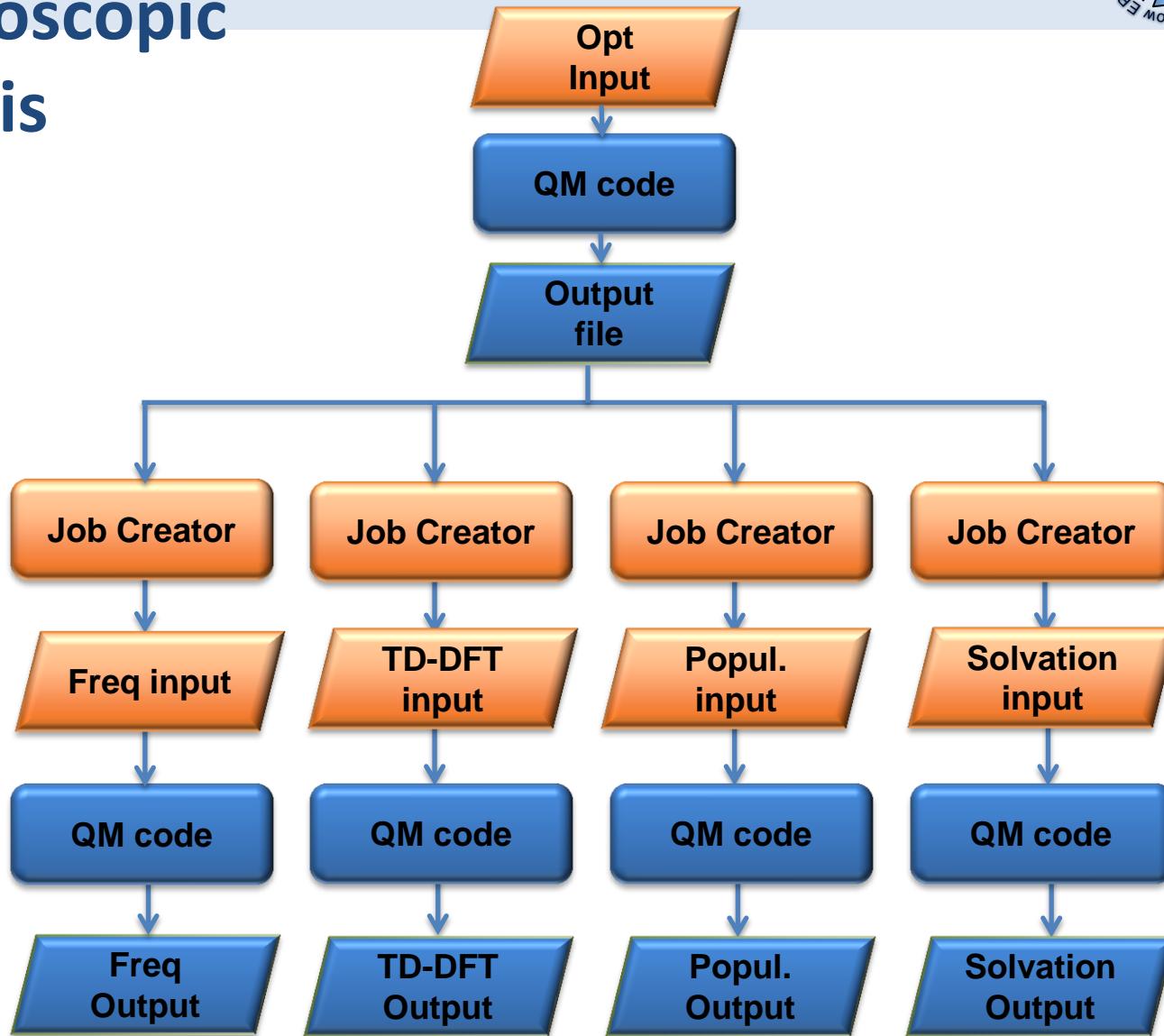
Orbital analysis (LMCT)

Example: Spectroscopic Analysis

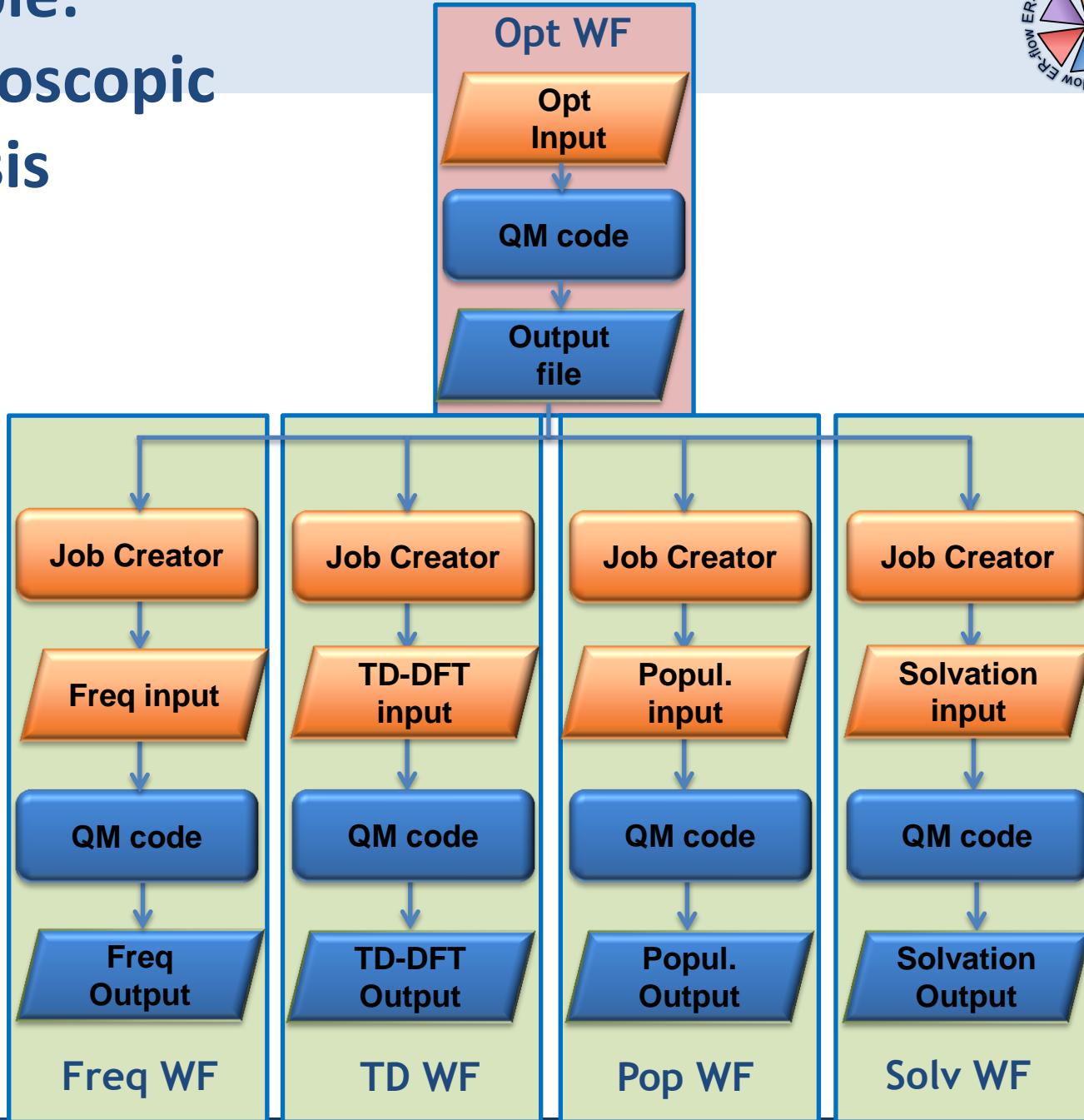




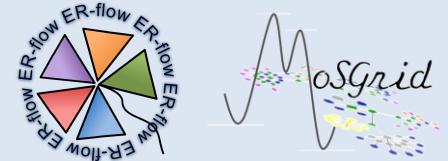
Example: Spectroscopic Analysis



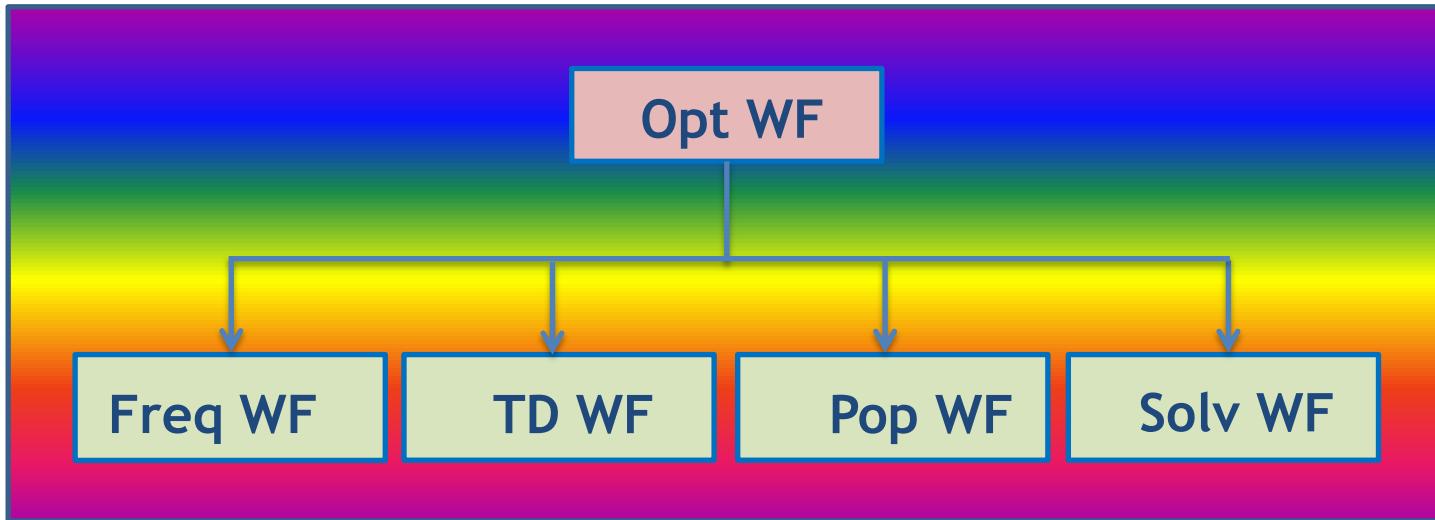
Example: Spectroscopic Analysis



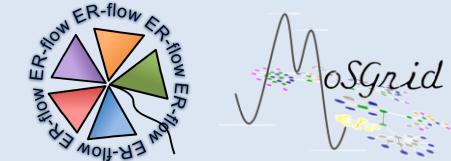
Example: Spectroscopic Analysis



Meta-Workflow



Workflows in Quantum chemistry



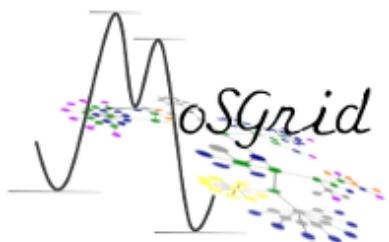
Application at the QC codes implemented in MoSGrid:

- ✓ Gaussian
- ✓ Turbomole
- ✓ NWChem
- Great Help for User (via standard recipes in the repository)
- Possibility to generate own complex workflows
- Facilitation of data extraction and postprocessing
- **Meta-workflows** allow re-use of basic workflows!

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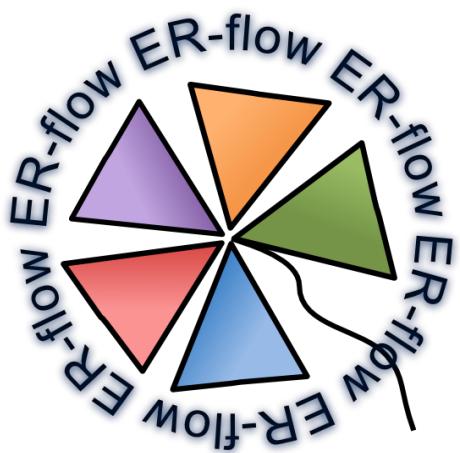
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