ALCF Hands-on HPC Workshop

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Agentic Workflows at ALCF

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What are Agentic Workflows?

- Al-powered computational pipelines that can autonomously perform complex tasks, make decisions, and adapt in real-time based on intermediate results or user input
- Agentic workflows exhibit "agency/intelligence" can reason, plan, and execute tasks based on new results with minimal human intervention
- Key characteristics:
 - —Break down complex tasks into discrete steps, each step handled by an agent
 - Autonomous and real-time decision making
 - —Adaptive execution



Agentic Workflows for Science

• LLMs can provide immense value to scientific research, but struggle on complex topics (hallucinations)

Human

Provide the SMILES string corresponding to this molecule: 9-[(2,6-dichlorophenyl)methyl]-N-(furan-2-ylmethyl)purin-6-amine

** SMILES = Simplified Molecular Input Line Entry System

LLMs (with GPT-40)

The molecule **9-[(2,6-dichlorophenyl)methyl]-N-(furan-2-ylmethyl)purin-6-amine** has the following structure components:

- •A **purine** core
- •Substitution at position 9: a (2,6-dichlorophenyl)methyl group
- •Substitution at position 6: an amino group
- •That amino group is further substituted with a furan-2-ylmethyl group

Constructed SMILES:

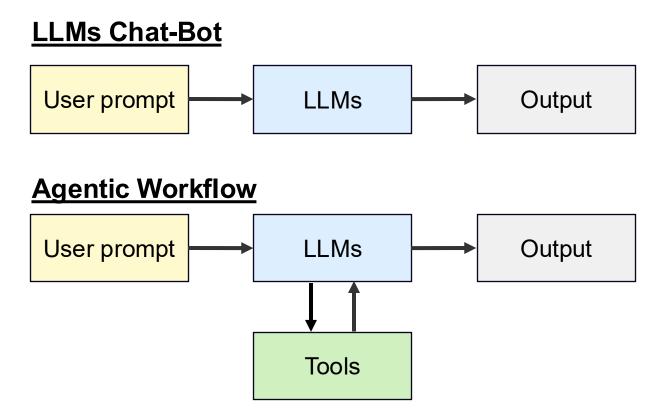
Clc1cccc(Cl)c1CN2c3nc(nc(NCC4=CC=CO4)c3nc2)N

Incorrect



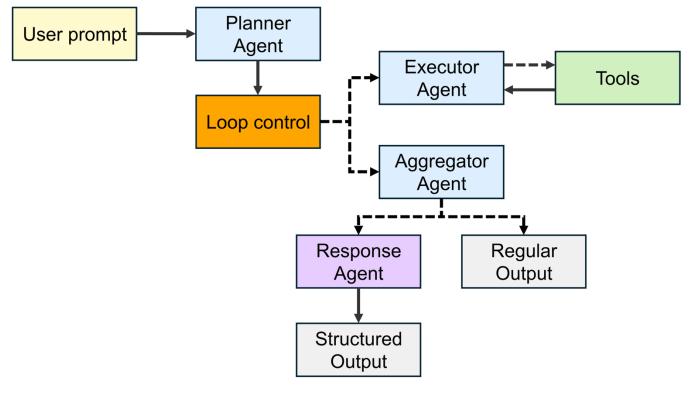
Agentic Workflows for Science

- Agentic workflows allow scientists to go beyond the "chat-bot" use of LLMs for question and answer
- They combine the power of gen-Al with computational science tools, research databases and HPC





ChemGraph: An Agentic Framework for Computational Chemistry Workflows



Software suite

















ChemGraph: An Agentic Framework for Computational Chemistry Workflows. (2025)Pham, T. D.; Tanikanti, A.; Keçeli, M. Under review.



ChemGraph: Providing Accurate SMILES strings

Human

Provide the SMILES string corresponding to this molecule: 9-[(2,6-dichlorophenyl)methyl]-N-(furan-2-ylmethyl)purin-6-amine

LLMs (with GPT-40)

The molecule **9-[(2,6-dichlorophenyl)methyl]-N-(furan-2-ylmethyl)purin-6-amine** has the following structure components:

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Constructed SMILES:

Clc1cccc(Cl)c1CN2c3nc(nc(NCC4=CC=CO4)c3nc2)N

[...]

Incorrect

ChemGraph (with GPT-4o-mini)

Tool Call:

Tool name: molecule_name_to_smiles

Args: 9-[(2,6-dichlorophenyl)methyl]-N-(furan-2-ylmethyl)purin-6-amine

Output:

"C1=CC(=C(C(=C1)Cl)CN2C=NC3=C(N=CN=C32)NCC4=CC=CO4)Cl"

The SMILES string corresponding to the molecule "9-[(2,6-dichlorophenyl)methyl]-N-(furan-2-ylmethyl)purin-6-amine" is C1=CC(=C(C(=C1)Cl)CN2C=NC3=C(N=CN=C32)NCC4=CC=CO4)Cl

Correct



ChemGraph: Calculating Reaction Enthalpy

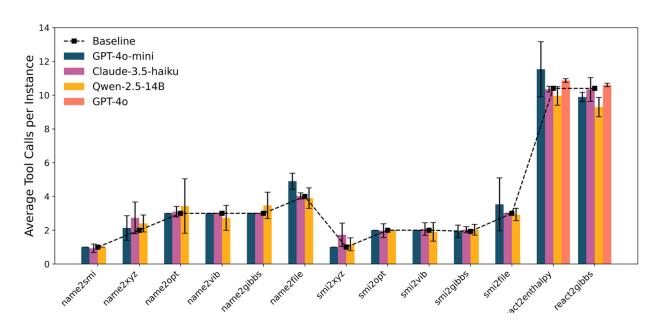
Human

You are given a chemical reaction: 1 (Methane) + 2 (Oxygen) -> 1 (Carbon dioxide) + 2 (Water). Calculate the enthalpy change for this reaction using GFN2-xTB at 400K.

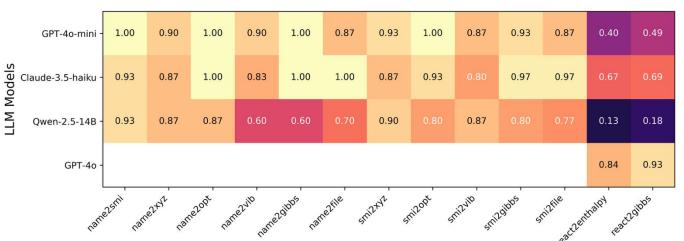
ChemGraph (multi-agent, GPT-4o-mini) Task 1: Calculate the Task 2: Calculate the Task 3: Calculate the Task 4: Calculate the Planner agent enthalpy of formation of enthalpy of formation of enthalpy of formation enthalpy of formation of methane (CH4) using carbon dioxide (CO2) of water (H2O) using oxygen (O2) using GFN2-xTB at 400K using GFN2-xTB at 400K GFN2-xTB at 400K GFN2-xTB at 400K **Executor agents** Executor 1 Executor 2 Executor 3 Executor 4 Tool call 1: molecule name to smiles(Methane) Tool call 2: smiles to atomsdata("C") Tool calls Tool call 3:run ase(Atomsdata(Methane), driver=thermo, temperature=400, "calculator"={"calculator type":"TBLite", "method": GFN2-xTB}) The thermochemical calculation for methane [...] [...] [...] (CH4) using the GFN2-xTB method at 400K has **Enthalpy of **Enthalpy of **Enthalpy of **Tool outputs** been successfully completed [...] Formation**: -214.93 eV Formation**: -280.0425 Formation**: -137.2903 **Enthalpy of Formation**: -112.2512 eV eV eV eV Aggregator agent [...] Thus, the enthalpy change (\u0394H) for the reaction is: Delta H = -12.5119 eV

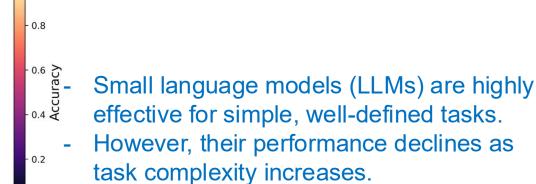


ChemGraph: Performance Across Task Complexity



Evaluation done on 13 different tasks, covering 360 instances and 200+ different molecules.







Agentic Workflows at ALCF

- Users can leverage ALCF systems to accelerate scientific agentic workflows with our HPC resources
- Public GitHub with examples: https://github.com/argonne-lcf/alcf-agentics-workflow

Multi-system (Remote) Workflows

- Agents run and launch components on various ALCF or local systems
- LLMs served remotely (e.g., ALCF Inference Service)
- Pros:
 - Lower implementation complexity
 - Can leverage ALCF services
 - Target specific system with each computational component (efficient use of allocation)
 - Better support for long-running workflows (agents can run on local system)
- Cons:
 - Higher latency between tasks
 - Subject to scheduler when launching tasks

Single-system (Local) Workflows

- All agents and components of the workflow run on the same system, even within same batch job
- LLMs served locally on system nodes
- Pros:
 - Lower latency between tasks
 - Not subject to scheduler when running tasks
 - Can serve custom LLMs if needed
- Cons:
 - Higher implementation complexity
 - Subject to queue limitations (e.g., max. run time)
 - May consume allocation inefficiently

