



上海人工智能实验室
Shanghai Artificial Intelligence Laboratory



SCHOOL OF
COMPUTING &
DATA SCIENCE
The University of Hong Kong

From Digital Agents to AI Co-Scientists

Qiushi Sun

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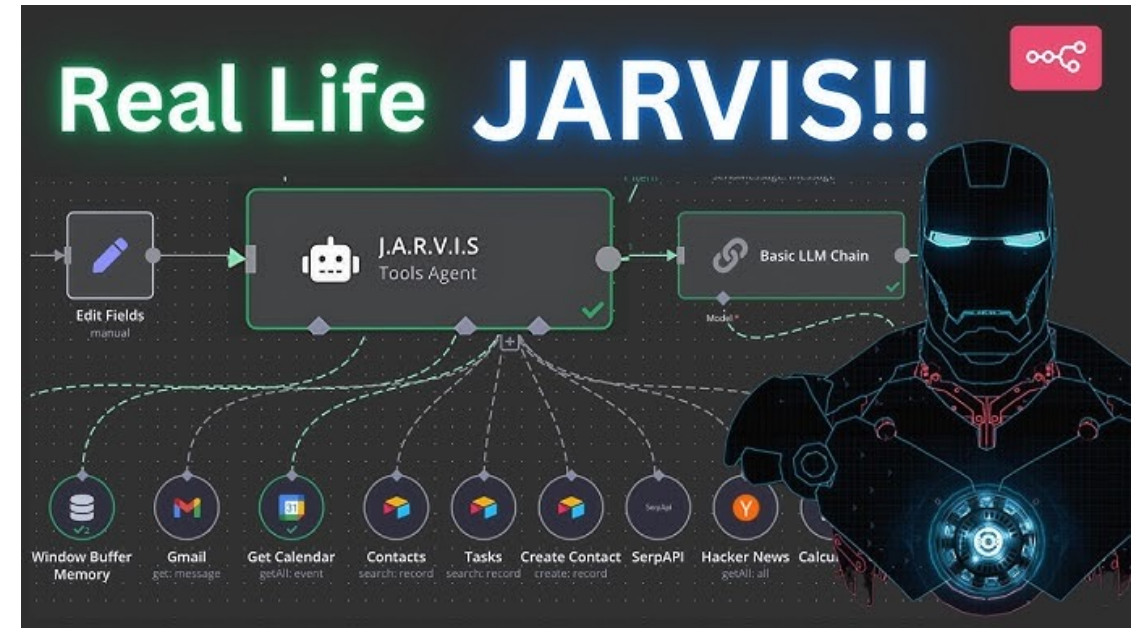
Part1 | Intro



Computer-Using Agents

A type of digital agent that operates within environments such as desktops, mobile devices, or web browsers.

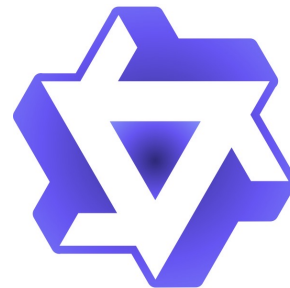
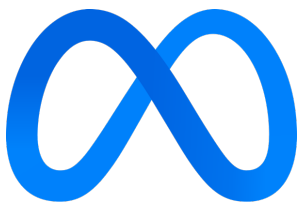
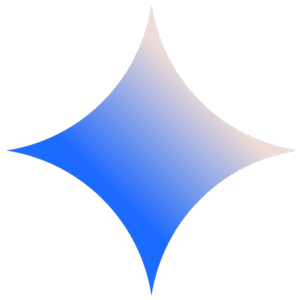
Computer-Using Agents



The Feasibility of Jarvis AI from Marvel in Real Life

Computer-Using Agents

Once out of reach, but we are turning it into reality.



Computer-Using Agents

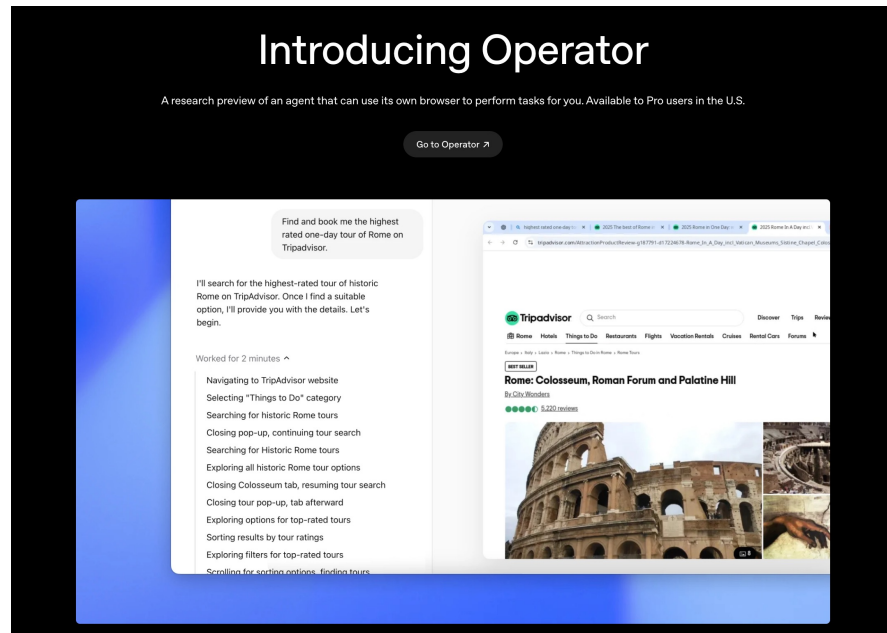
Both academia and industry are building **computer-using agents**



Claude Computer Use

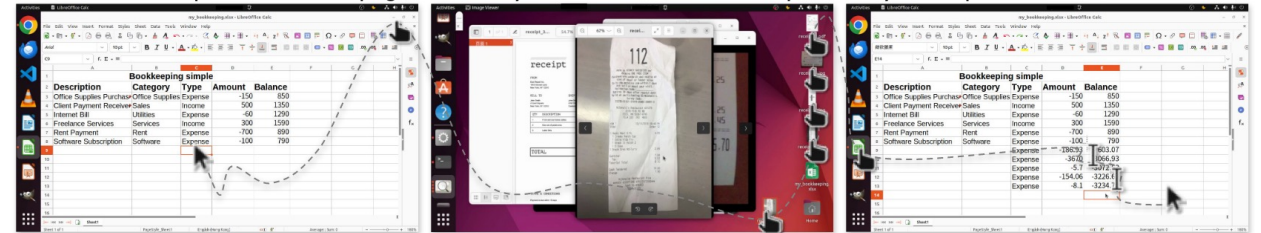
Computer-Using Agents

Automating daily computer tasks

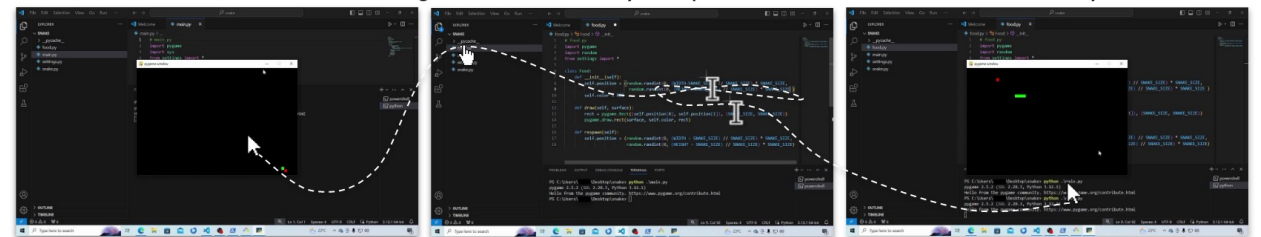


OpenAI Operator

Task instruction 1: Update the bookkeeping sheet with my recent transactions over the past few days in the provided folder.



Task instruction 2: ...some details about snake game omitted... Could you help me tweak the code so the snake can actually eat the food?



Daily Computer Use

[3] *Introducing Operator: A research preview of an agent that can use its own browser to perform tasks for you.*, Jan 23, 2025

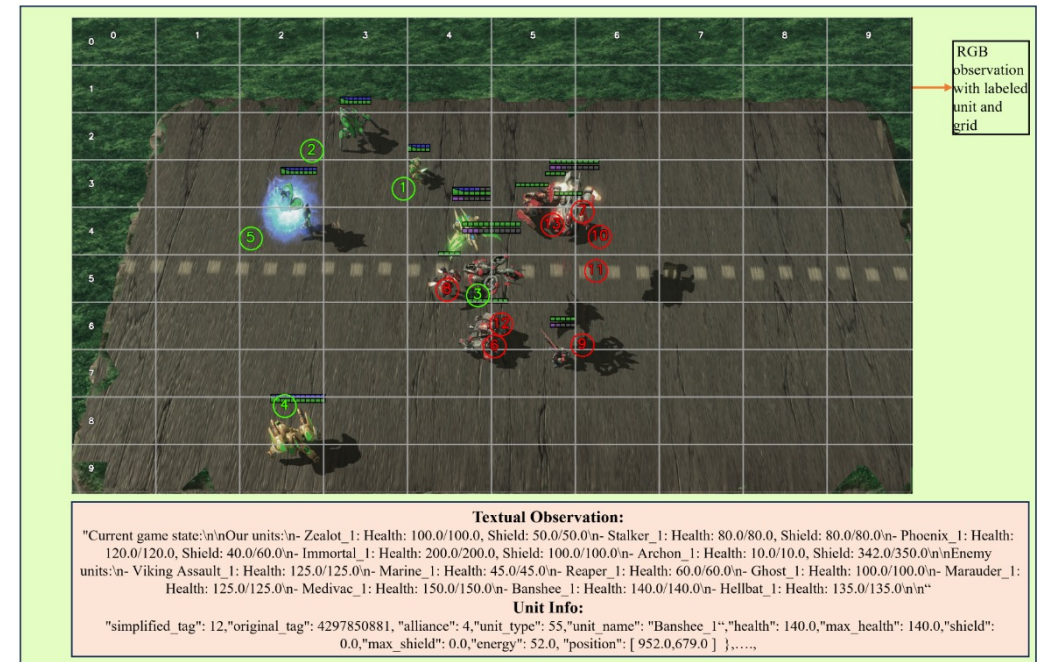
[4] *OSWorld: Benchmarking Multimodal Agents for Open-Ended Tasks in Real Computer Environments*

Computer-Using Agents

Playing Games



MineCraft



StarCraft II

Computer-using Agents

Startups



[Research](#) [Product](#) [About us](#) [Blog](#)

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[Github](#)

[macOS](#)

Private Public Run Similar locally on macOS

a web browser.

(Next Action) Now I need to open a web browser. I'll use the agent.open method to open Firefox, which is a common web browser on Linux systems.

(Grounded Action)

```
agent.open("firefox")
```

Agent S2 2 hours ago

Agent is executing task...

[Get in line to talk to Agent S](#)

Type your message here...

Shared Virtual Machine

40 viewers watching now

Seminal works on Computer-Using Agents



SeeClick: Harnessing GUI Grounding for Advanced Visual GUI Agents, [ACL 2024](#) Foundation Models



OS-ATLAS: A Foundation Action Model for Generalist GUI Agents , [ICLR 2025 Spotlight](#)



OS-Genesis: Automating GUI Agent Trajectory Construction via Reverse Task Synthesis , [ACL 2025](#)

Data



Breaking the Data Barrier -- Building GUI Agents Through Task Generalization, [COLM 2025](#)



AgentStore: Scalable Integration of Heterogeneous Agents As Specialized Generalist Computer Assistant , [ACL 2025](#)

Algorithm



OS-MAP: How Far Can Computer Use Agents Go in Breadth and Depth?

Evaluation



ScienceBoard: Evaluating Multimodal Autonomous Agents in Realistic Scientific Workflows

Frontier Application

Seminal works on Computer-Using Agents



SeeClick: Harnessing GUI Grounding for Advanced Visual GUI Agents, ACL 2024



OS-ATLAS: A Foundation Action Model for Generalist GUI Agents , ICLR 2025 Spotlight



OS-Genesis: Automating GUI Agent Trajectory Construction via Reverse Task Synthesis , ACL 2025



Breaking the Data Barrier -- Building GUI Agents Through Task Generalization, COLM 2025



AgentStore: Scalable Integration of Heterogeneous Agents As Specialized Generalist Computer Assistant , ACL 2025



OS-MAP: How Far Can Computer Use Agents Go in Breadth and Depth?



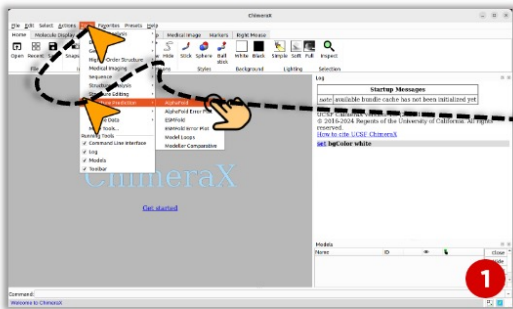
ScienceBoard: Evaluating Multimodal Autonomous Agents in Realistic Scientific Workflows

Frontier Application

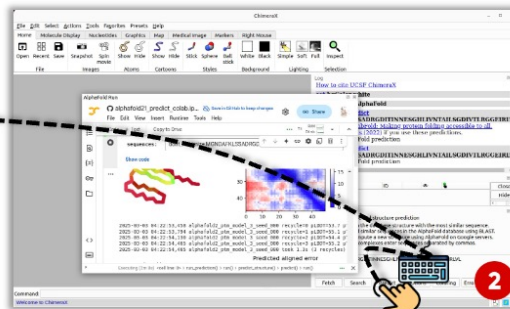
Computer-using Agents

Automate scientific workflows, be your co-scientist!

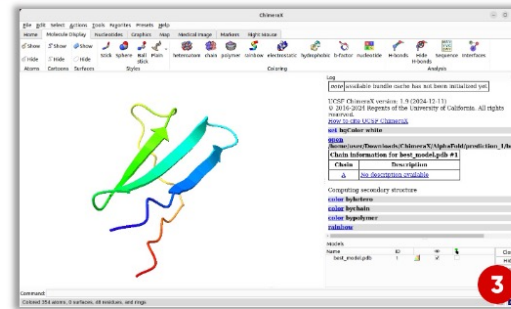
Instruction: Predict the protein structure for the amino acid sequence of 'MGND...' via AlphaFold in ChimeraX.



Step1: Toggle the widget of AlphaFold.

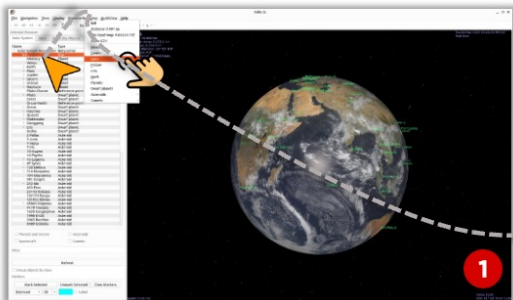


Step2: Input the given sequence and call out AlphaFold for structure prediction.

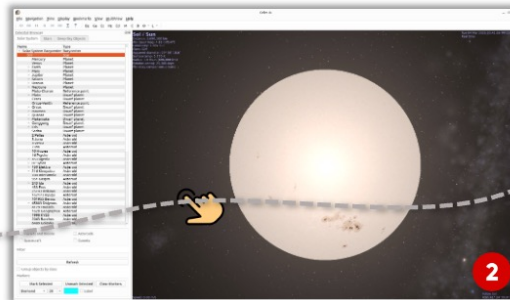


Step3: Wait until the prediction finished.

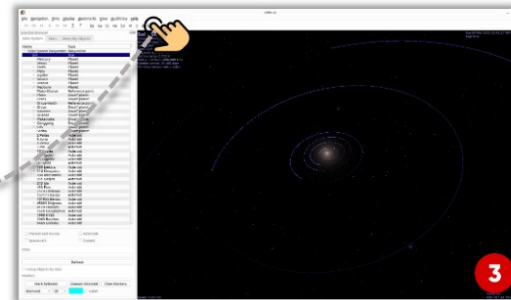
Instruction: Show planets' orbits of Solar System in Celestia.



Step1: Select the Sol and click 'Goto' in context menu.



Step2: Slide the mouse wheel to move the camera away from Sol.



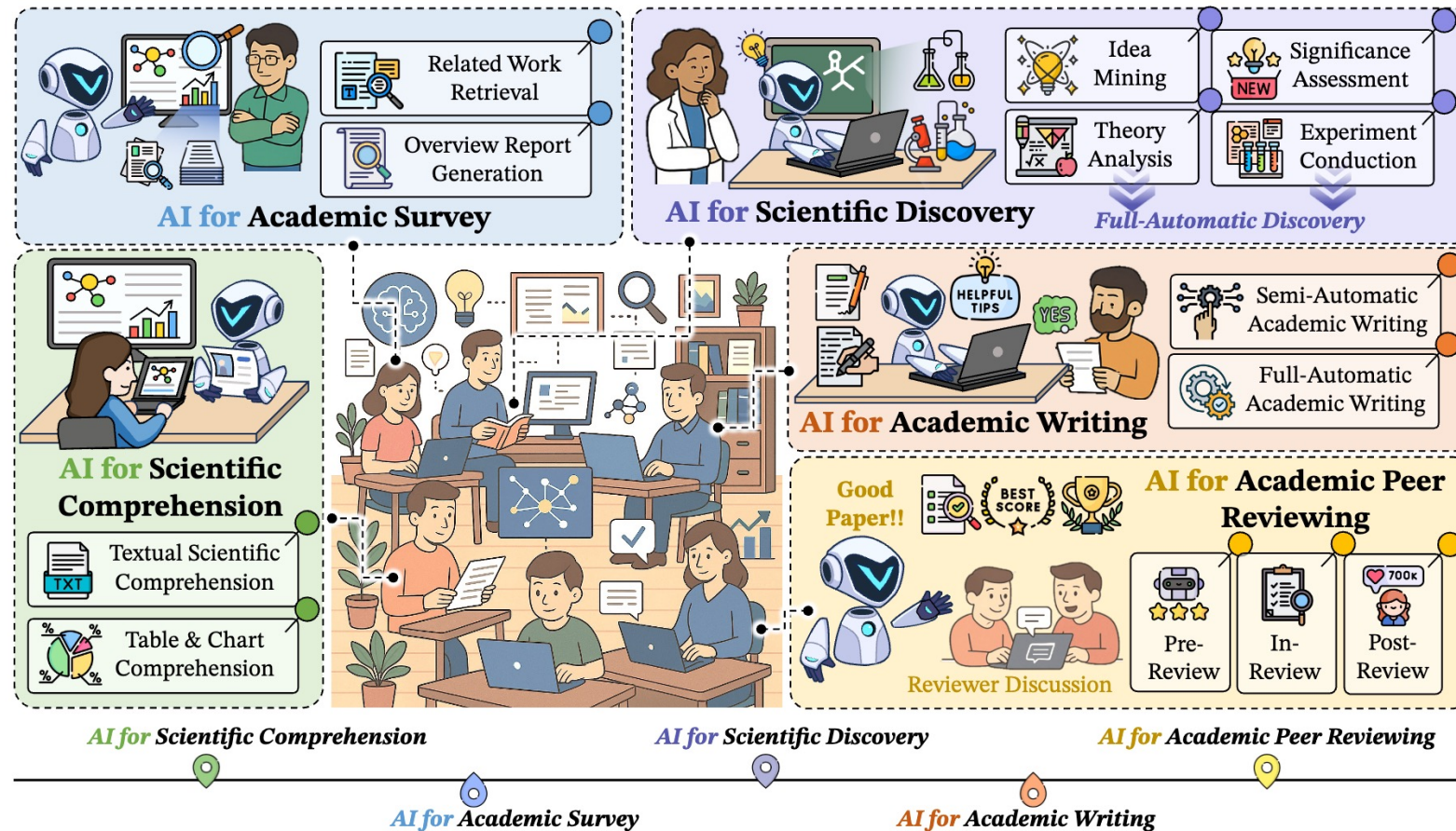
Step3: Click to show orbits of planets.

Part2 | AI4Research

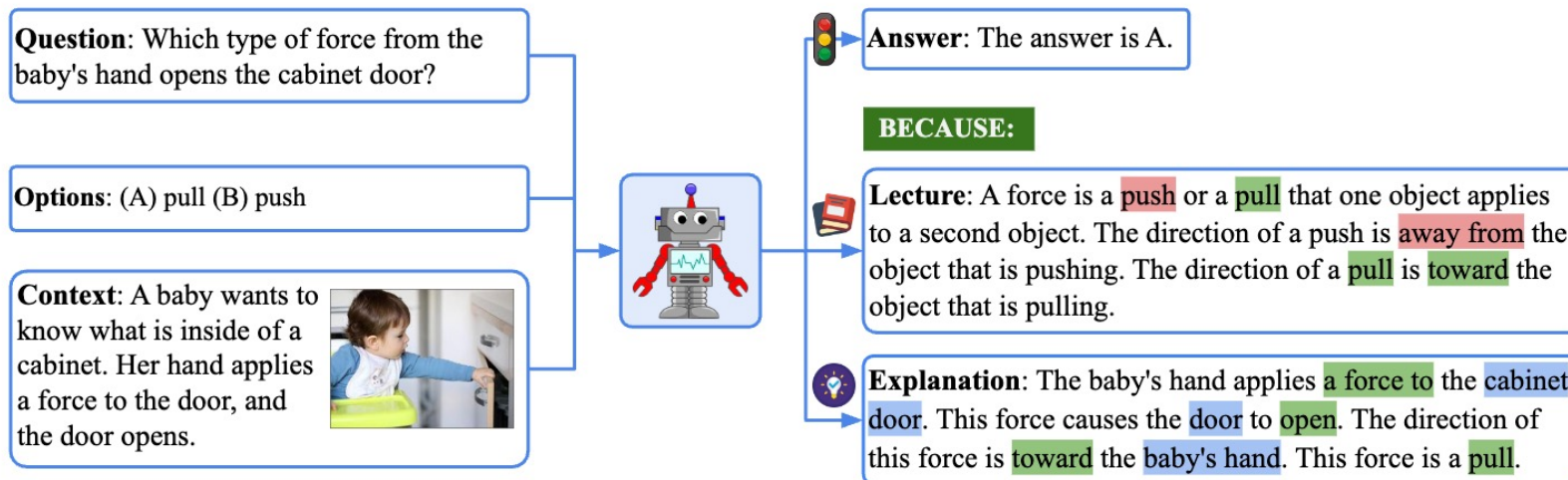


Backgrounds

AI4Research is a highly popular concept.



Backgrounds: Pastoral Age



ScienceQA (NIPS 2022)

- Multimodal Reasoning (Chain-of-Thought)
- Natural science, language science, and social science
- 12k Grade school-level MCQ

Backgrounds: Contemporary Era

SciCode (NIPS 2024)

- 16 subfields (e.g., math, physics, chem).
- 80 main problems, decomposed into 338 subproblems involving recall, reasoning, and code synthesis.
- Each question verified by 2 senior researchers to ensure scientific accuracy and relevance.

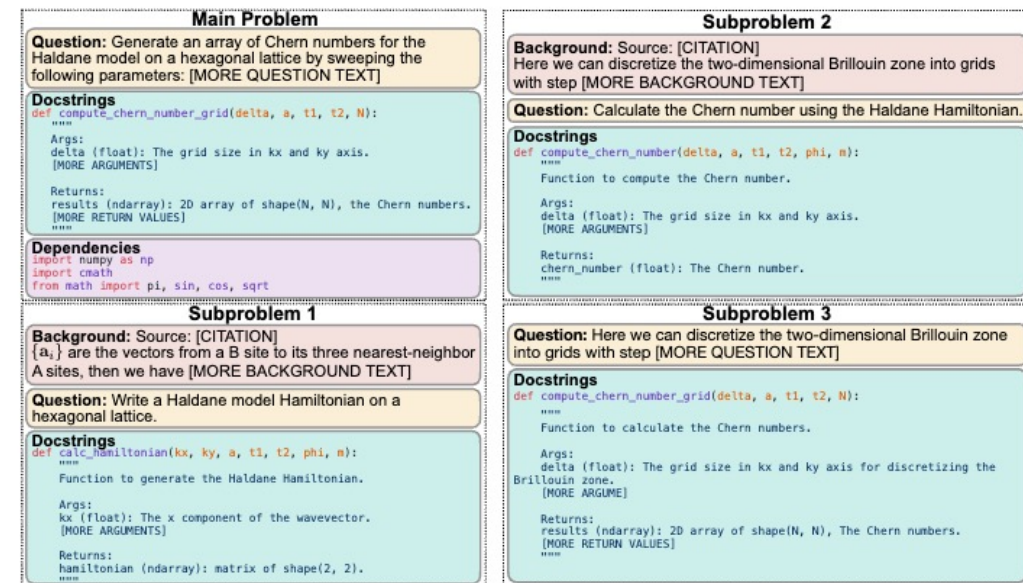
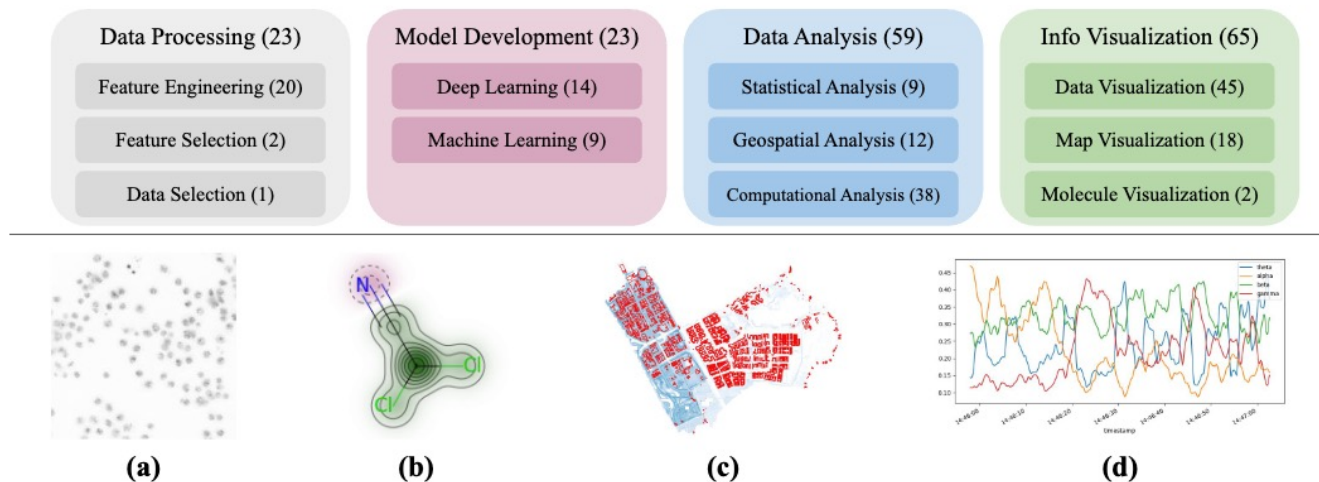


Figure 1: A SciCode main problem is decomposed into multiple smaller and easier subproblems. Docstrings specify the requirements and input-output formats. When necessary, scientific background knowledge is provided, written by our scientist annotators. The full problem is shown in subsection A.3

Backgrounds: Contemporary Era



ScienceAgentBench (ICLR 2025)

- Evaluating language **agents** in data-driven scientific discovery.
- 102 tasks from 44 peer-reviewed publications across four disciplines.
- Tasks require generating a self-contained Python program.
- Prompting based solutions: OpenHands CodeAct, and self-debug.

Backgrounds: Contemporary Era

SCIENCE

AlphaEvolve: A Gemini-powered coding agent for designing advanced algorithms

14 MAY 2025
By AlphaEvolve team

[Share](#)



Backgrounds: Contemporary Era

A lot of “AI Research” systems have been built...

The AI Scientist: Towards Fully Automated Open-Ended Scientific Discovery

Chris Lu^{1,2,*}, Cong Lu^{3,4,*}, Robert Tjarko Lange^{1,*}, Jakob Foerster^{2,†}, Jeff Clune^{3,4,5,†} and David Ha^{1,†}
^{*}Equal Contribution, ¹Sakana AI, ²FLAIR, University of Oxford, ³University of British Columbia, ⁴Vector Institute, ⁵Canada CIFAR AI Chair, [†]Equal Advising

SciMON : Scientific Inspiration Machines Optimized for Novelty

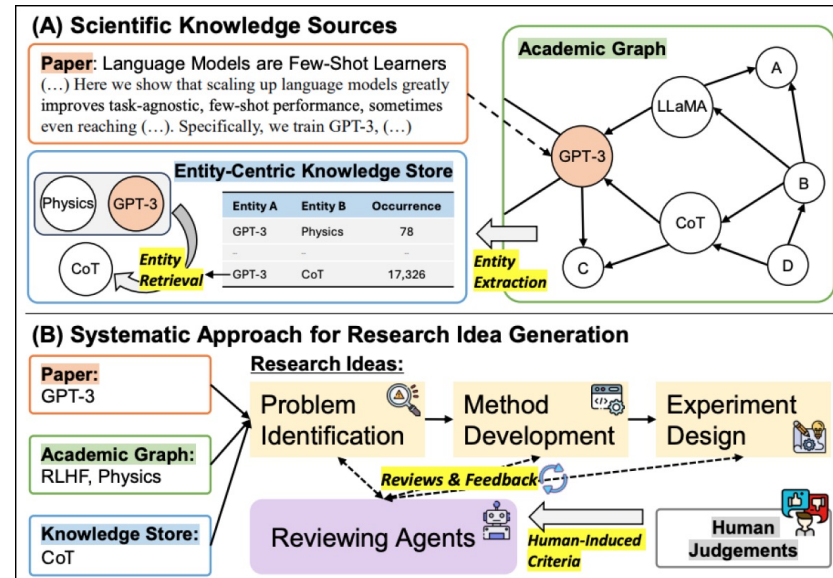
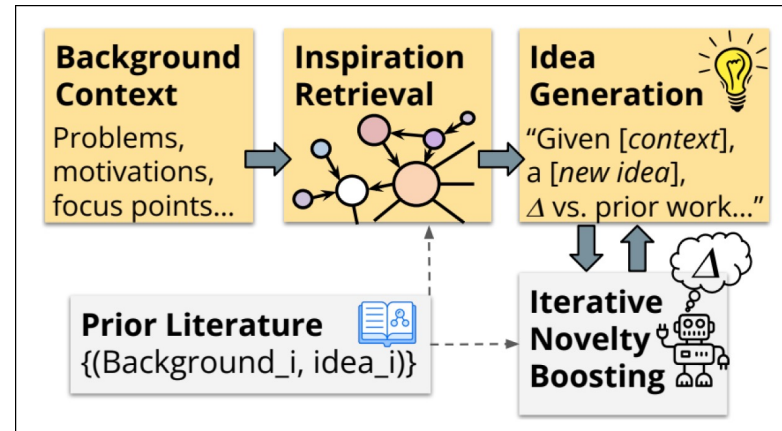
Qingyun Wang¹, Doug Downey², Heng Ji¹, Tom Hope^{2,3}
¹ University of Illinois at Urbana-Champaign ² Allen Institute for Artificial Intelligence (AI2)
³ The Hebrew University of Jerusalem
 {tomh,doug}@allenai.org, {qingyun4,hengji}@illinois.edu

ResearchAgent: Iterative Research Idea Generation over Scientific Literature with Large Language Models

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 KAIST¹ Microsoft Research² DeepAuto.ai³
 {jinheon.baek, sjhwang82}@kaist.ac.kr {sjauhar, silviu}@microsoft.com

Automated Peer Reviewing in Paper SEA: Standardization, Evaluation, and Analysis

Jianxiang Yu[✉], Zichen Ding[✉], Jiaqi Tan[✉], Kangyang Luo[✉], Zhenmin Weng[✉],
 Chenghua Gong[✉], Long Zeng[✉], Renjing Cui[✉], Chengcheng Han[✉],
 Qiushi Sun[✉], Zhiyong Wu[✉], Yunshi Lan[✉], Xiang Li[✉]
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 sea.ecnu@gmail.com
<https://ecnu-sea.github.io/>



Backgrounds: Contemporary Era

The domain is advancing rapidly, but current developments are **not realistic enough**,
From the perspective of how human scientists interact with the world:

For works w.o. visual content

- Coding is good but not a universal solution
- Real scientific workflows require both screens and actions

For works w. visual content

- The “VQA” form is a little bit outdated
- Single turn, solving a grade-school does not align with our need

Therefore, we need to consider text / vision, knowledge, coding

But That's Not All!

Thinking

Currently, AI acted as an “Analyzer,” helping with idea thinking data analysis, writing, and visualization.

Can AI evolve into an “Executor” that helps (1) formulate a plan, (2) directly operates scientific software via GUI or CLI, and (3) even generates some reports?

The answer is YES—with the emergence of computer-using agents.

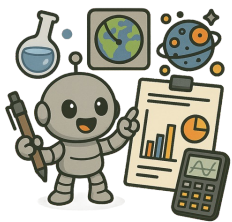
Let's move beyond QA and Coding to actively performing some research tasks!



From Digital Agents to AI Co-Scientists

Part3 | ScienceBoard



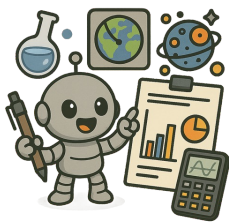


ScienceBoard: Evaluating Multimodal Autonomous Agents in Realistic Scientific Workflows

Qiushi Sun, Zhoumianze Liu, Chang Ma, Zichen Ding, Fangzhi Xu, Zhangyue Yin, Haiteng Zhao, Zhenyu Wu, Kanzhi Cheng, Zhaoyang Liu, Jianing Wang, Qintong Li, Xiangru Tang, Tianbao Xie, Xiachong Feng, Xiang Li, Ben Kao, Wenhai Wang, Biqing Qi, Lingpeng Kong, Zhiyong Wu



Preprint / WUCA @  **ICML** 2025 Oral
International Conference
On Machine Learning



ScienceBoard: Evaluating **Multimodal** Autonomous Agents in **Realistic** Scientific Workflows

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Preprint / WUCA @  **ICML** 2025 Oral
International Conference
On Machine Learning

Language Agents

Let's start with some background on computer-using agents.

Remark: For computer-using agents, both **GUI** and **CLI** represent distinct approaches.

In ScienceBoard, we primarily focus on **GUI-based interaction**, complemented by **CLI support**.

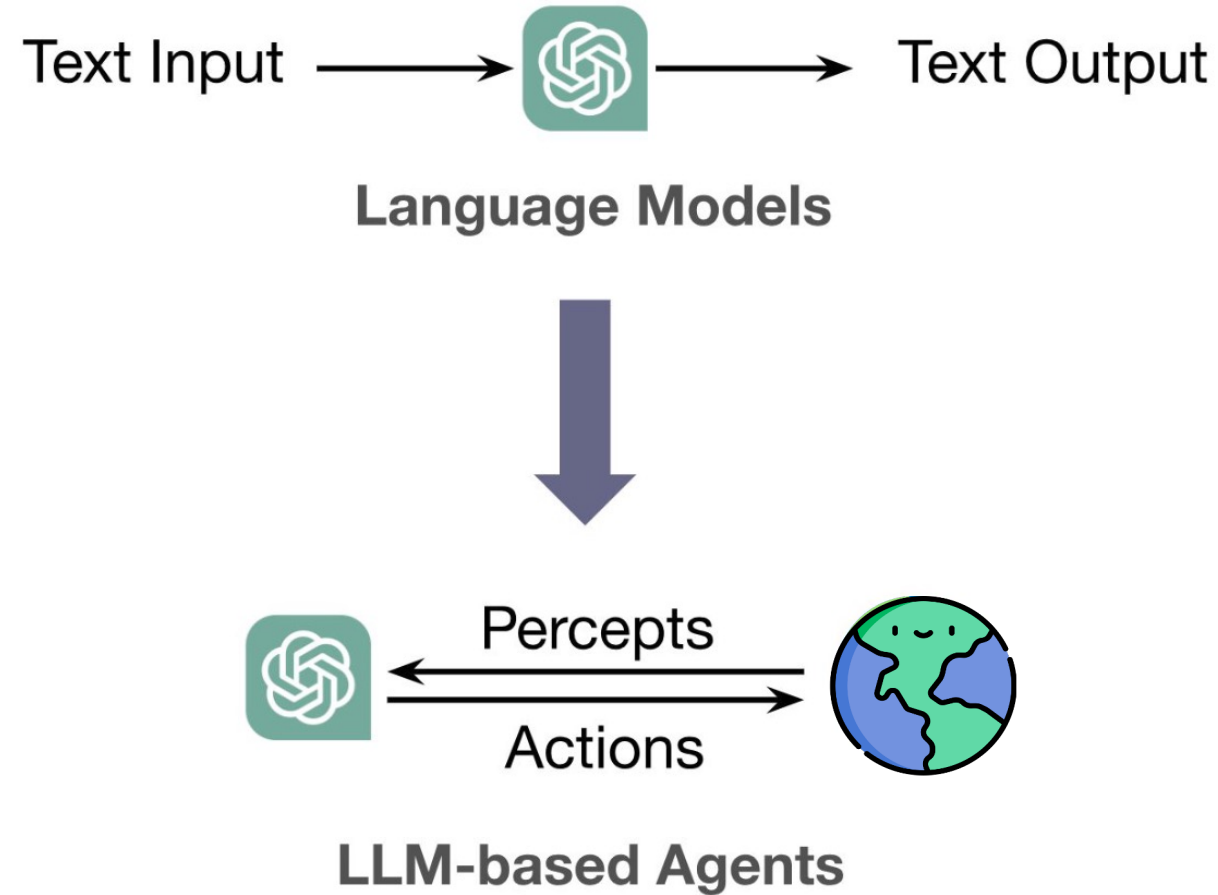


GUI Agents

*Intelligent agents that operate **within GUI environments**, leveraging LLMs as their core inference and cognitive engine to generate, plan, and execute actions in a flexible and adaptive manner.*





Language Agents

Computer-using agents are language agents.



Language Agents

Agents are promising, but building powerful computer-using agents is challenging:

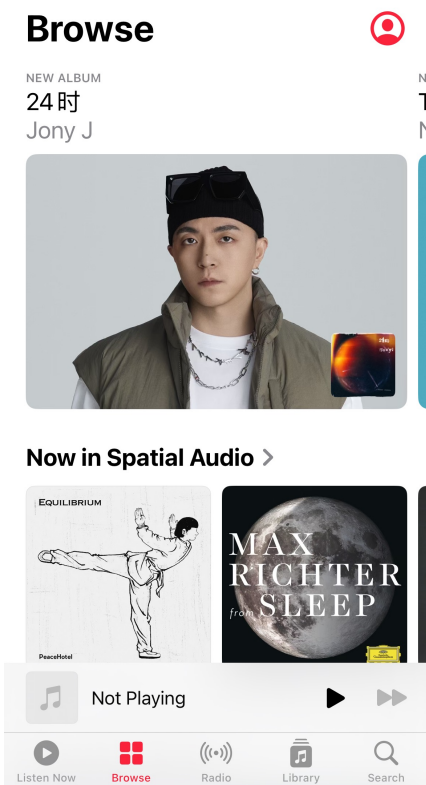
1. Agents need to follow human instructions. 
2. Agents need to perform planning and action. 
3. Agents need to perceive envs.  and the applications  they are interacting with.

Language Agents

What are “actions”

Language Agents

Typical **action**: GUI grounding – the capacity to accurately locate screen elements based on instructions, e.g., CLICK.



In order to view the new album of Jony J, where should I click?



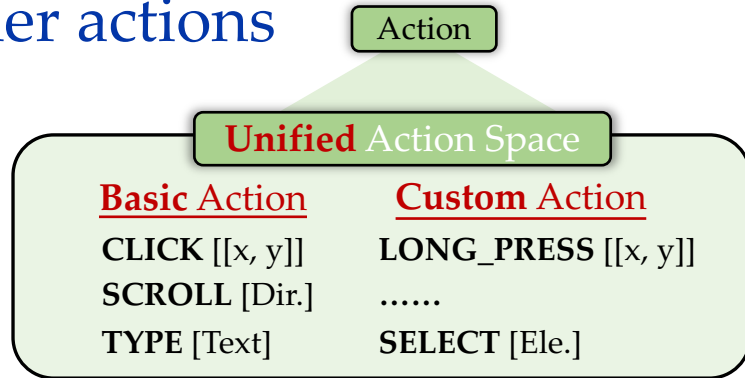
GPT-4o (an earlier version): hmmm... Sorry I don't know. ❌



SeeClick: (0.49, 0.40) ✅

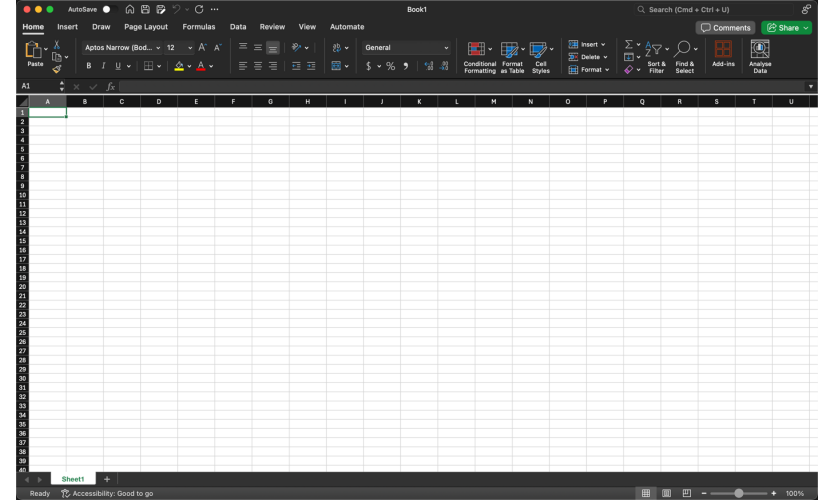
Language Agents

Other actions

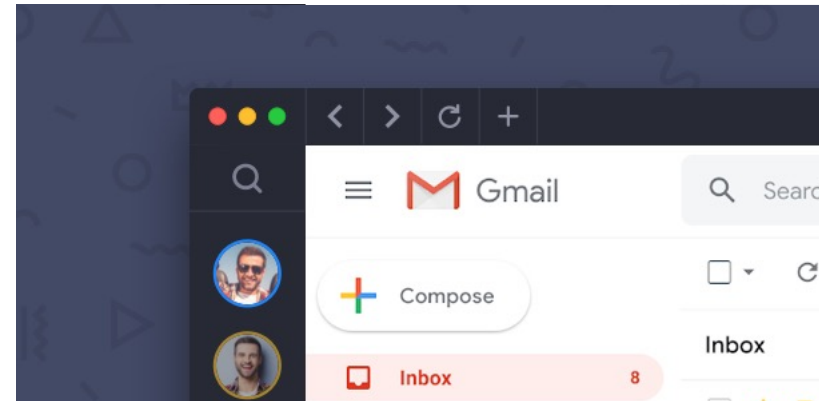


V.S. CLI

1. Screenshot is **information complete** for agentic tasks
2. GUI action space is much smaller and is shared across platforms/apps






"Close the app"



ScienceBoard Infra

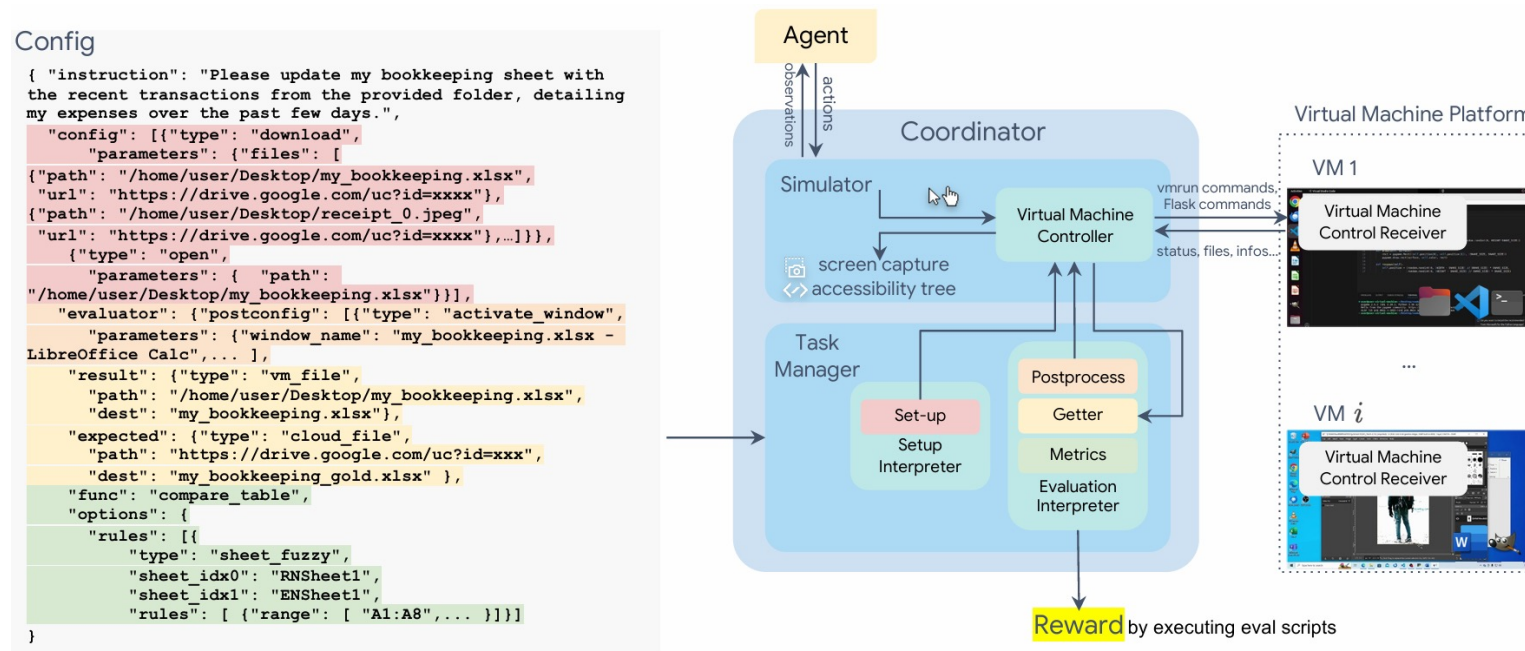
So to achieve our goal, we need an environment that allows agents to actively interact.

1. Supports native **multimodal** interaction 
2. Fully **compatible** with coding and conversational research assistance 
3. Enables rigorous **validation** 

A playground—a virtual machine pre-installed with well-adapted scientific software.

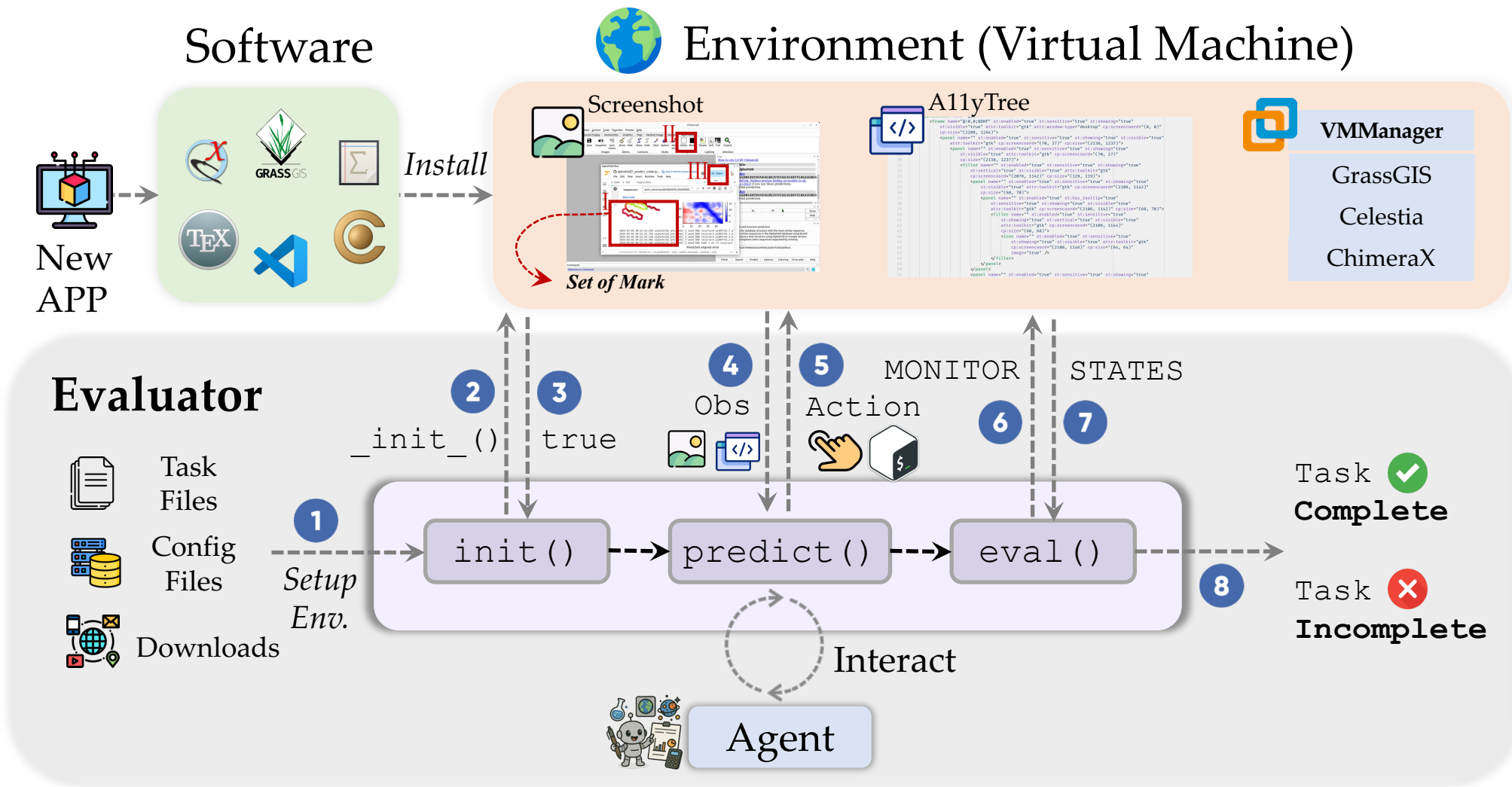
ScienceBoard Infra

We build upon the OSWorld infrastructure for GUI interaction.



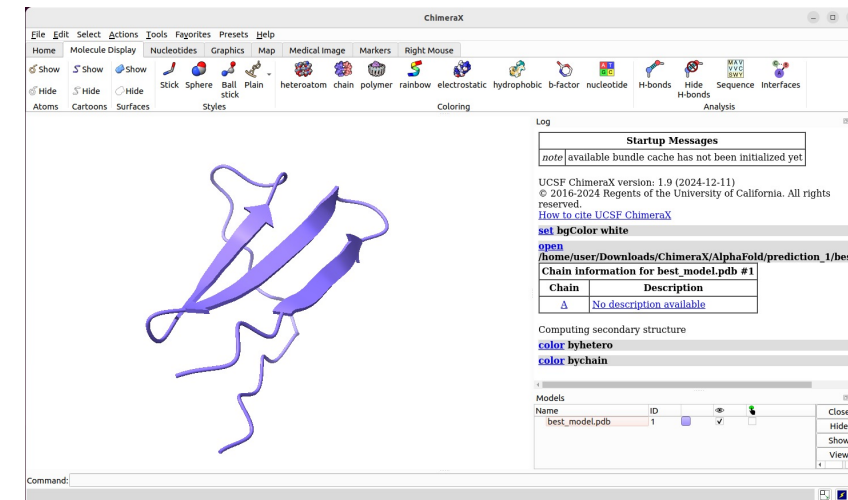
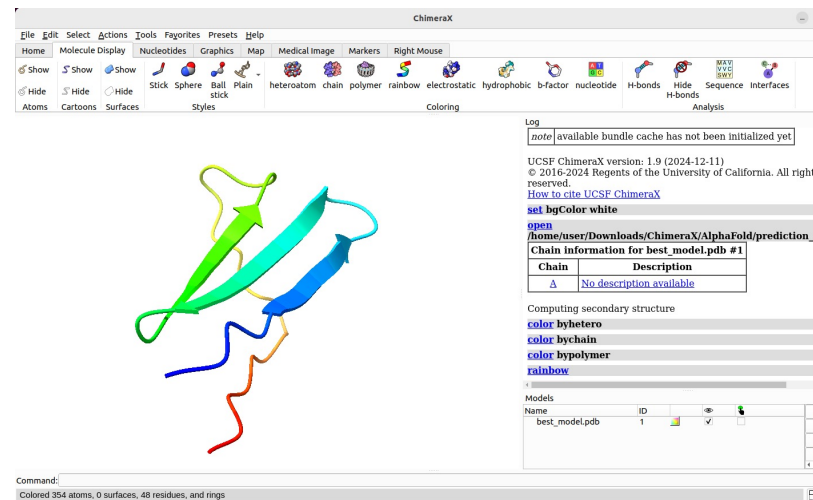
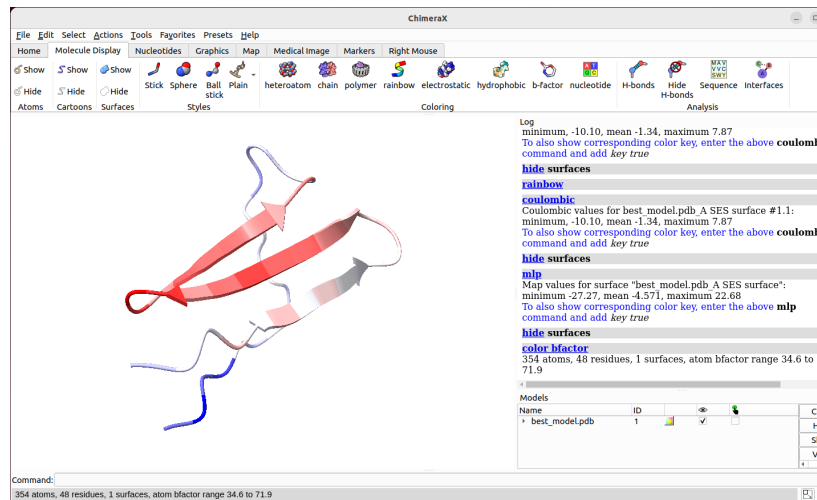
For CLI, we enable interaction by modifying the software itself and leveraging VSCode.

ScienceBoard Infra



The Dilemma of Evaluation

Evaluation is harder than we expect, let's take visualization as an example



It is impossible to “match”

How to Evaluate?

We rely on **internal states**.

By modifying the software, we access intermediate runtime states and enable precise state-based evaluation, e.g., UCSF ChimeraX 

Quick Links

[UCSF ChimeraX Home](#)
[Advantages](#)
[Documentation](#)
[User Guide](#)
[Tutorials and Videos](#)
[Presentations](#)
[Download](#)
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[Statistics](#)
[Citing ChimeraX](#)
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[Related Software](#)

Featured Citations

Effect of phosphorylation barcodes on arrestin binding to a chemokine receptor, Chen Q, Schafer CT et al. *Nature*. 2025 Jul 3;643(8070):280–287.

Zincore, an atypical coregulator binds zinc finger transcription factors to control gene expression, Bianchi D, Borza R et al. *Science*. 2025 Jul 3;389(6755):eadv2861.

Chromosome end protection by RAP1-mediated inhibition of DNA-PK, Eickhoff P, Sommer C et al. *Nature*. 2025 Jun 26;642(8069):1080–1086.

Structure of the ATP-driven methyl-coenzyme M reductase activation complex, Ramirez-Amador F, Paul S et al. *Nature*. 2025 Jun 19;642(8068):814–821.

Lewy-MSA hybrid fold drives distinct neuronal α -synuclein pathology, Enomoto M, Martinez-Vasbueno I et al. *Commun Biol*. 2025 Jun 16;8(1):929.

[More citations...](#)

UCSF ChimeraX

UCSF ChimeraX (or simply ChimeraX) is the next-generation molecular visualization program from the [Resource for Biocomputing, Visualization, and Informatics \(RBVI\)](#), following [UCSF Chimera](#). ChimeraX can be downloaded free of charge for academic, government, nonprofit, and personal use. Commercial users, please see [ChimeraX commercial licensing](#).

ChimeraX is developed with support from [National Institutes of Health R01-GM129325](#).

ChimeraX on Bluesky: [@chimerax.ucsf.edu](#)

Feature Highlight

AlphaFold Fetch

AlphaFold is an artificial intelligence method for predicting protein structures. With the [AlphaFold tool](#) or [command](#), ChimeraX can search for and load predicted structures from the freely available [AlphaFold Database](#), automatically coloring them by confidence value:

- 100 to 90 – high accuracy
- 90 to 70 – backbone accuracy
- 70 to 50 – low confidence, caution
- 50 to 0 – should not be interpreted, may be disordered

The figure shows the predicted structure of UniProt entry [TOM40_HUMAN](#), a channel protein needed to import other proteins into mitochondria. See the command file [tom40.cxc](#) for fetching data and other setup (background color, etc.).

Opening a sequence from UniProt also opens a [dialog](#) in which its annotations or “features” can be clicked to highlight those regions in both the sequence and the associated 3D structure. The low-confidence part of this structure (orange and red) maps to compositionally biased and likely disordered regions near the N-terminus of the sequence.

[More features...](#)

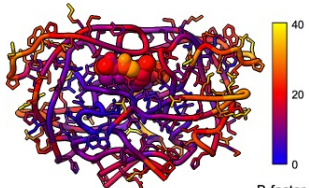
Example Image

B-factor Coloring

Atomic B-factor values are read from PDB and mmCIF input files and assigned as [attributes](#) that can be shown with [coloring](#) and used in [atom specification](#). This example shows B-factor variation within a structure of the HIV-1 protease bound to an inhibitor (PDB [4hvp](#)). For complete image setup, including positioning, [color key](#), and label, see the command file [bfactor.cxc](#).

Additional color key examples can be found in tutorials: [Coloring by Electrostatic Potential](#), [Coloring by Sequence Conservation](#)

[More images...](#)



B-factor

News


June 26, 2025

The ChimeraX 1.10 production release is [available!](#) See the [change log](#) for what's new.

May 7, 2025

The ChimeraX 1.10 release candidate is [available](#) – please try it and [report](#) any issues. See the [change log](#) for what's new.

March 19, 2025

 UCSF ChimeraX: Tools for structure building and analysis is one of the 10 most cited papers published in *Protein Science* in 2023!

[Previous news...](#)

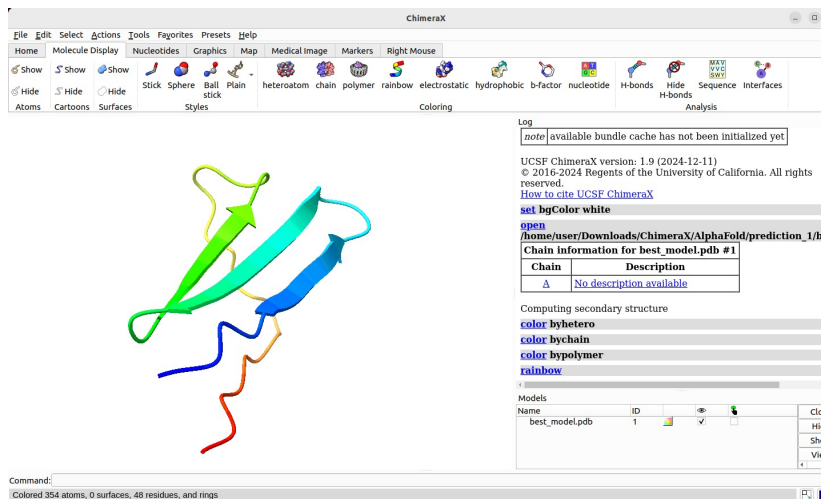
Upcoming Events

Page 36

How to Evaluate?

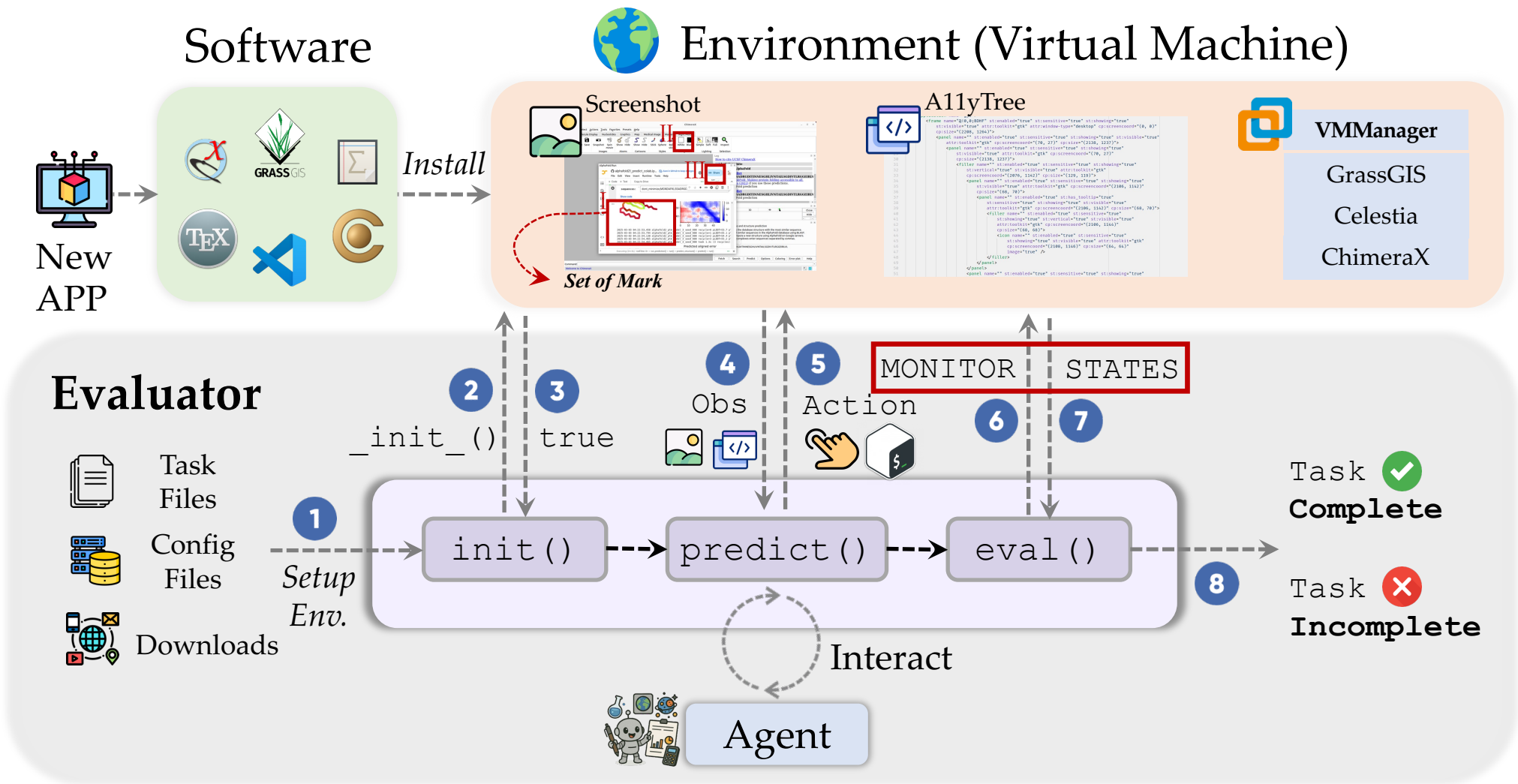
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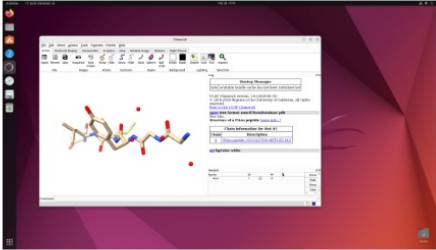
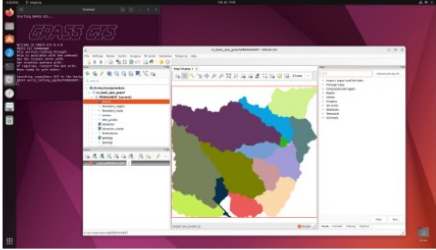
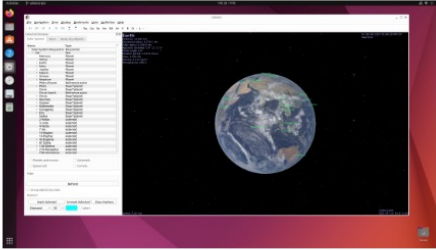


```
_state_containers.tools._tool_instances.#  
0.tool_window._ToolWindow__toolkit.main_w  
indow.tool_instance_to_windows.<chimerax.  
log.tool.Log object at  
0x0000020AA8023B50>.#0.tool_instance.page  
_source
```

ScienceBoard Infra



How to Evaluate?

Initial State	Instruction	Evaluation Script (Simplified)
	<i>Select all water molecules and draw their centroids with radius of 1Å in ChimeraX.</i>	<pre>{ "type":"info","key":"sell", "value":["atom id #!1/A:201@0 idatm_type 03" "...",] },{ "type":"states", "find":"lambda k,v:k.endswith('._name')", "key":"lambda k:'..._atoms_drawing'", "value":"[[13.0012 1.7766 21.3672 1.]]" }</pre>
	<i>Display and ONLY display the layer of 'boundary_region' in Grass GIS.</i>	<pre>{ "type":"info", "key":"lambda dump:len(dump['layers'])", "value":1 },{ "type":"info", "key":"lambda dump:dump['layers'][0]['name']", "value":"boundary_region@PERMANENT" }</pre>
	<i>Set the Julian date to 2400000 in Celestia.</i>	<pre>{ "type":"info", "key":"simTime", "value":2400000, "pred":"lambda left, right:abs(left-right) < 1", }</pre>

Evaluation

Approach to the Earth and display a solar eclipse in Celestia.

```
"evaluate": [  
  {  
    "type": "info",  
    "key": "lambda dump: dump['entity']['Earth']['distance']",  
    "value": 0,  
    "pred": "lambda key, value: abs(key - value) < 450000"  
  },  
  {  
    "type": "info",  
    "key": "lambda dump: dump['entity']['Sol']['visible']",  
    "value": false  
  },  
  {  
    "type": "info",  
    "key": "lambda dump: dump['entity']['Moon']['visible']",  
    "value": true  
  },  
  {  
    "type": "info",  
    "key": "lambda dump: (s := dump['entity']['Sol']['position'], e := dump['entity']['Earth']['position'], m  
:= dump['entity']['Moon']['position'], mv := [m[i] - e[i] for i in range(3)], sv := [s[i] - e[i] for i in  
range(3)], dp := mv[0] * sv[0] + mv[1] * sv[1] + mv[2] * sv[2], _mv := __import__('math').sqrt(mv[0]**2 + mv  
[1]**2 + mv[2]**2), _sv := __import__('math').sqrt(sv[0]**2 + sv[1]**2 + sv[2]**2), dp / (_mv * _sv))[-1]",  
    "value": 0.99,  
    "pred": "lambda key, value: key > value"  
  }  
]
```

Next

Now agents can freely **explore the environment** and **execute any actions** they choose.

But how well do these agents actually perform to automate science tasks?

What kind of benchmark is needed to truly evaluate their capabilities?

ScienceBoard Benchmark



We aim to build a benchmark with:

1. **Real-world** tasks that human actually perform
2. Coverage across **multiple disciplines**
3. Graded **difficulty** levels
4. Support for **cross-application** workflows
5. Cross “modality” GUI + CLI

And more

ScienceBoard Benchmark



Coverage:

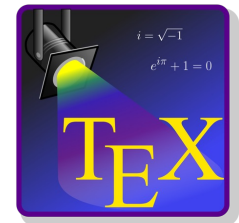
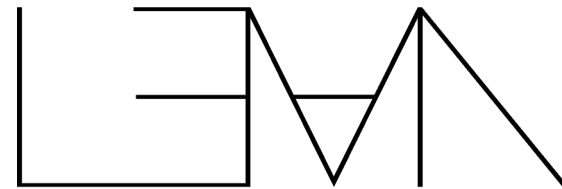
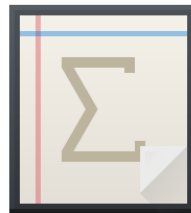
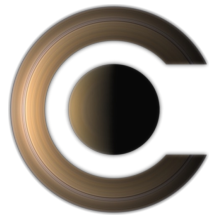
Biochem, GIS, Astronomy, Algebra, ATP, Documentation

ScienceBoard Benchmark



Our criteria for selecting software:

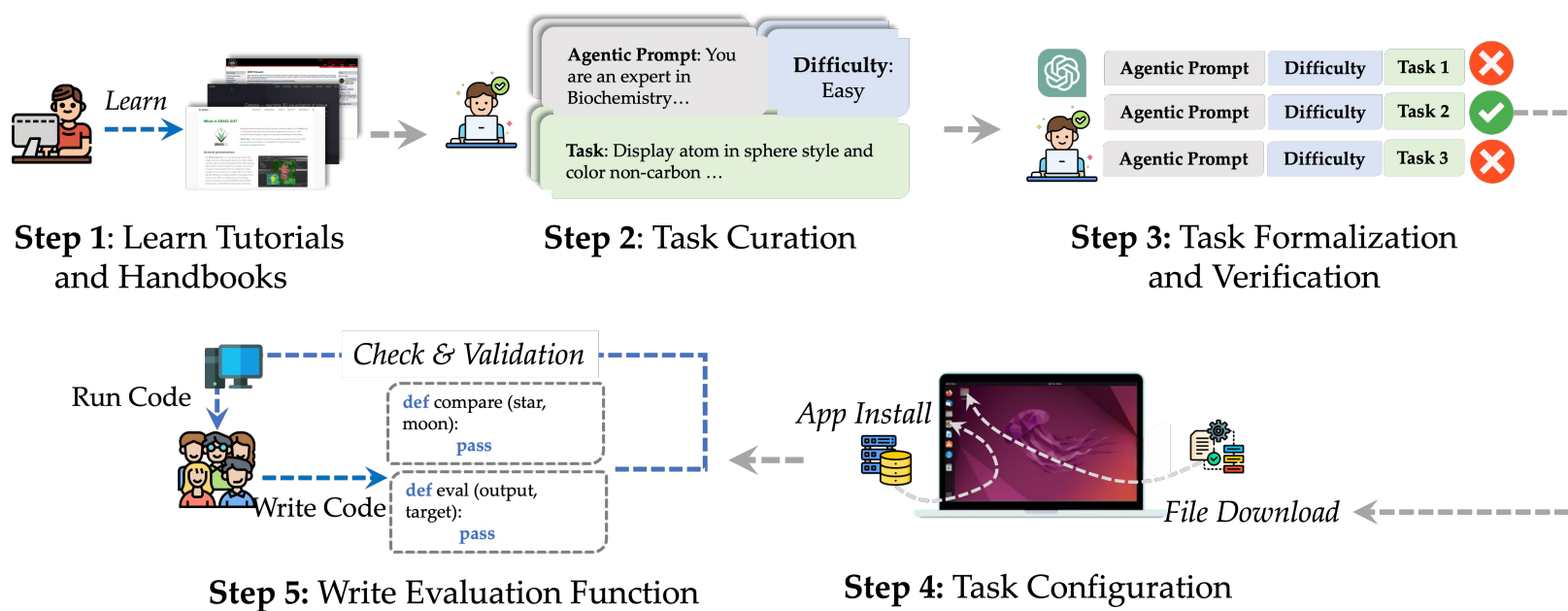
1. Stable operation on **Ubuntu** 
2. Open-source, allowing for **adaptation**
3. Accessible **a11ytree**, enabling text-based agents to participate (e.g., o1-mini)
4. License



ScienceBoard Benchmark




How the benchmark is built










ScienceBoard Benchmark






 Celestia Users Guide


by Frank Gregorio

-  MS Word document (zipped) or Read Online
-  Документ MS Word или PDF файл или Читать онлайн
-  Document PDF (pour Celestia 1.6.0)
-  PDF Datei (für Celestia 1.6.0-1)
-  Documento MS Word e OpenOffice (per Celestia 1.6.1, archivio zip)
-  PDF 文档 (Celestia 1.6.1)



 Celestia Key Chart


by The Learning Technologies Project Office of NASA


-  PNG Image
-  PNG изображение
-  PNG 图像

 CEL Scripting Guide

by Don Goyette

-  MS Word document or Read Online
-  Документ PDF (архив ZIP)

 SSC File Scripting Guide

-  PDF document (zipped)

Celestia tutorials

ChimeraX Quick Start Guide

[UCSF ChimeraX](#) is the next-generation visualization program from the [Resource for Biocomputing, Visualization, and Informatics](#) at UC San Francisco, following [Chimera](#). See also: [ChimeraX tutorials](#)

Many ChimeraX actions require typing commands. The help for a specific command can be shown with the [help](#) command (for example, [help style](#)). Other ways to interact with the program include:

- clicking [toolbar icons](#), optionally after making a [selection](#)
- [graphical tools](#)
- context menus shown by right-click (Ctrl-click on Mac, Alt-click on Windows trackpad)

Command-Execution Links

Clicking command links in the examples below will execute them in ChimeraX *if this page is shown in the ChimeraX internal browser*, such as with [Help... Quick Start Guide](#) in the ChimeraX menu.

Example Atomic-Structure Commands

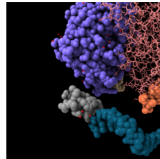
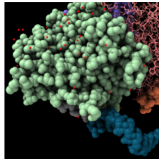
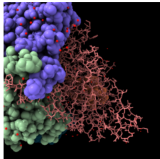
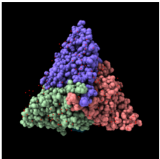
Example structure: Protein DataBank [2BBV](#), black beetle virus capsid

[open 2bbv](#)
[color bychain](#)

[style /b stick](#)

Mouse drag to move.
[color /n teal](#)

[hide /c](#)



File is fetched from the PDB in mmCIF format and cached locally. Lighting with shadows.

Change chain b to stick style.

Rotate by dragging, translate by dragging with middle mouse button or with option key pressed (Mac) ([more...](#)).

Hide chain c atoms.

[ribbon /c](#)

Mouse click with **ctrl** key pressed to select an atom, or command [select /N:4@C5'](#)

Press up-arrow key, or command [select up](#)

[color set gold](#)
[select clear](#)

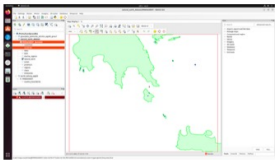
ChimeraX tutorials

ScienceBoard Benchmark



Develop a large set of evaluation scripts

Manually validate their correctness.

Initial State	Instruction	Evaluation Script (Simplified)
	Select all ligand(s) and color them into magenta in ChimeraX.	<pre>{ "type": "info", "key": "sel", "value": ["atom id /A:9@N1 idatm_type N3+", ...], }, { "type": "info", "key": "rescolor /A", "value": ["#1/A:1 color #d2b48c", ...], } }</pre>
	There is a point located in the Mediterranean Sea. Please find and delete it.	<pre>{ "type": "db", "cmd": "v.to.db", "kwargs": { "flags": "p", "map": "countries@PERMANENT", "type": "point", "option": "coor" }, "key": "lambda out: out.strip()", "value": "cat x y z n... 8.348947891274 0", "pred": "lambda key, value: key == value" } }</pre>
	Approach to the Earth and display a solar eclipse in Celestia.	<pre>{ "type": "info", "key": "lambda ...['Earth']['distance']", "value": 0, "pred": "lambda k, v: abs(k - v) < 450000" }, { "type": "info", "key": "lambda ...['Sol']['visible']", "value": false }, { "type": "info", "key": "lambda ...['Moon']['visible']", "value": true }, { "type": "info", "key": "lambda ...", "value": 0.99, "pred": "lambda key, value: key > value" } }</pre>
	<pre>theorem TP_3 [TopologicalSpace X] [TopologicalSpace Y] (f : X -> Y) (Z : Set X) (h1 : Continuous f) (h2 : IsConnected Z) : IsConnected {y : Y ∃ z ∈ Z, f z = y} := by sorry</pre>	<pre>{ "type": "placeholder" } }</pre>

ScienceBoard Benchmark



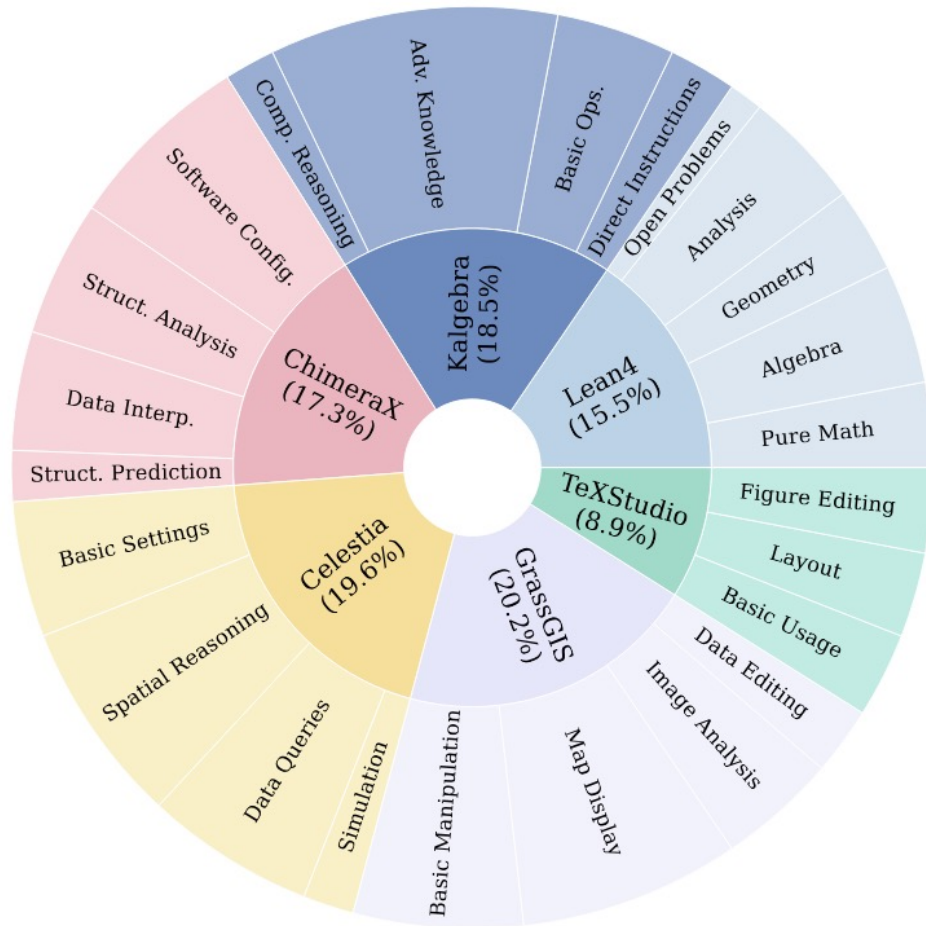
Task Type	Statistics
Total Tasks	169 (100%)
- GUI	38 (22.5%)
- CLI	33 (19.5%)
- GUI + CLI	98 (58.0%)
Difficulty	
- Easy	91 (53.8%)
- Medium	48 (28.4%)
- Hard	28 (16.6%)
- Open Problems	2 (1.2%)
Instructions	
Avg. Length of Task Instructions	20.0
Avg. Length of Agentic Prompt	374.9
Execution	
Avg. Steps	9.0
Avg. Time Consumption	124(s)



Evaluate autonomous computer-using agents in **realistic scientific workflows**.

Tasks require complex tool usage, scientific reasoning, and multi-step **GUI/CLI** operations

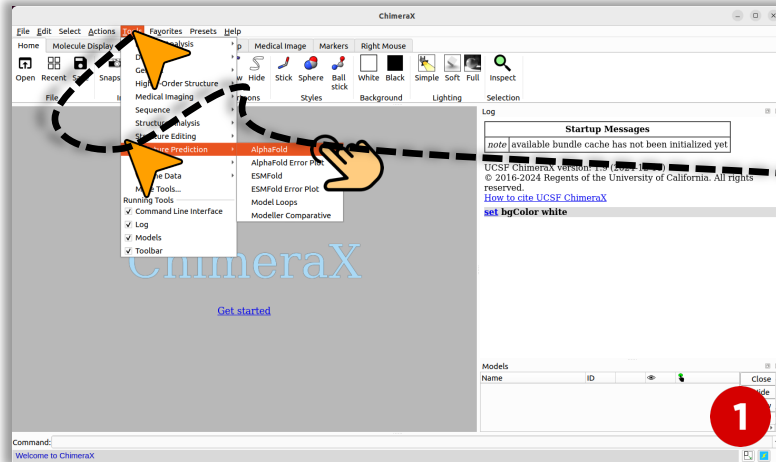
ScienceBoard Benchmark



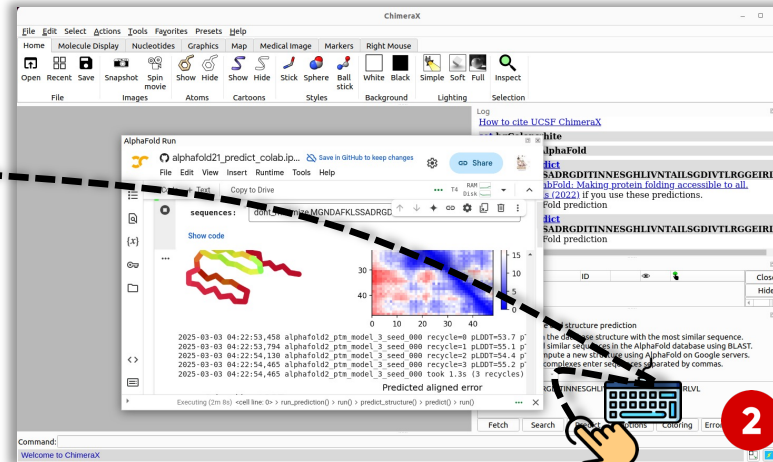
1. 169 high-quality tasks across 6 domains: Biochemistry, Algebra, Theorem Proving, GIS, Astronomy, Documentation.
2. Tasks require GUI operation, visual/textual reasoning, tool use, coding, spatial understanding.
3. Rich **mix**: CLI-only, GUI-only, hybrid workflows.

Use Cases

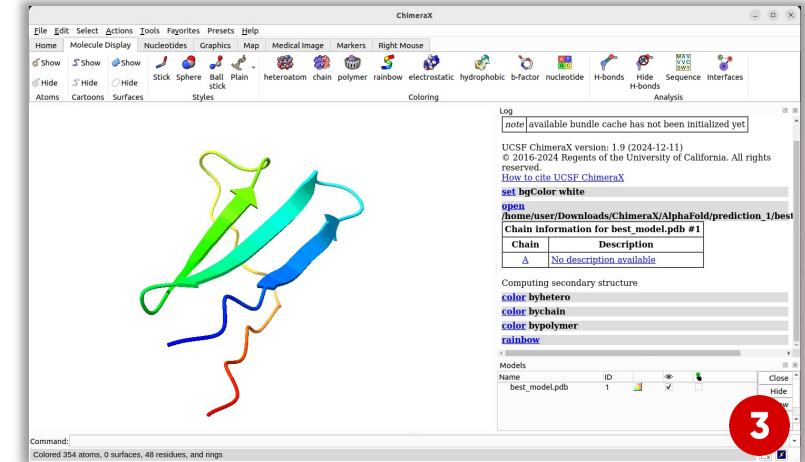
Instruction: Predict the protein structure for the amino acid sequence of 'MGND...' via AlphaFold in ChimeraX.



Step1: Toggle the widget of AlphaFold.



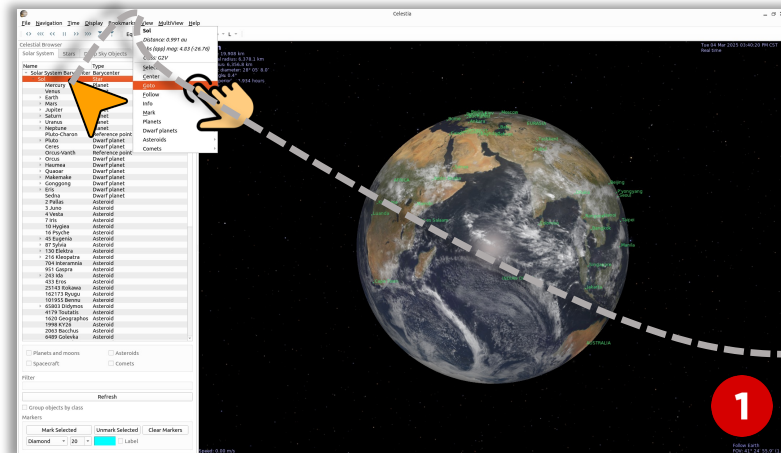
Step2: Input the given sequence and call out AlphaFold for structure prediction.



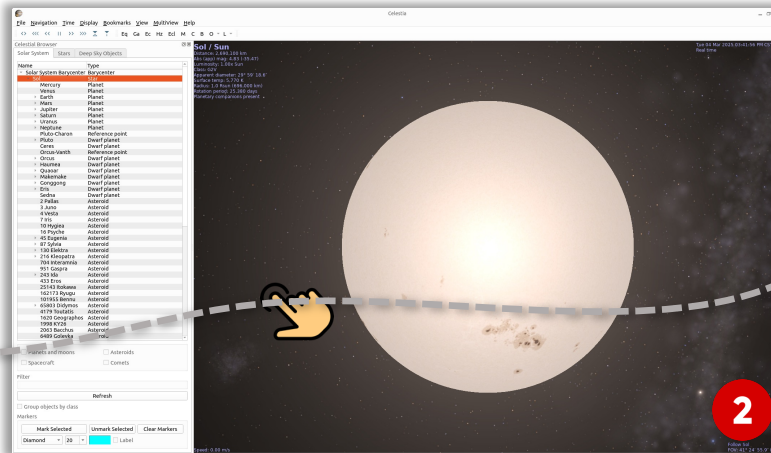
Step3: Wait until the prediction finished.

Use Cases

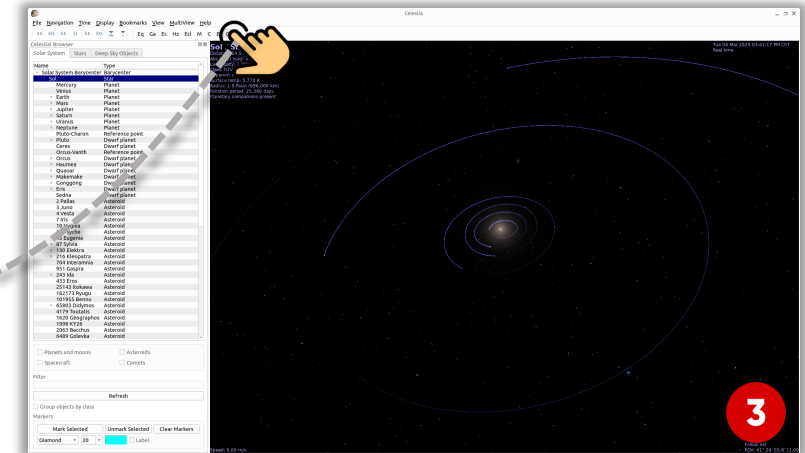
Instruction: Show planets' orbits of Solar System in Celestia.



Step1: Select the Sol and click 'Goto' in context menu.



Step2: Slide the mouse wheel to move the camera away from Sol.



Step3: Click to show orbits of planets.

Use Cases

Display constellations with their boundaries in Celestia.

I see the current state of Celestia, where the Earth is being displayed in the main view. The task is to display constellations with their boundaries in Celestia.

Looking at the accessibility tree, I can see there are several checkboxes including "C" which likely stands for Constellations and "B" which likely stands for Boundaries. These checkboxes are located in the main toolbar and need to be checked to display the constellations and their boundaries.

Currently, neither the constellations nor their boundaries are visible in the screenshot. I need to click on both the "C" checkbox and "B" checkbox to enable these features.

```
python
import pyautogui
import time
```

```
# Click on the "C" checkbox to enable constellations
```

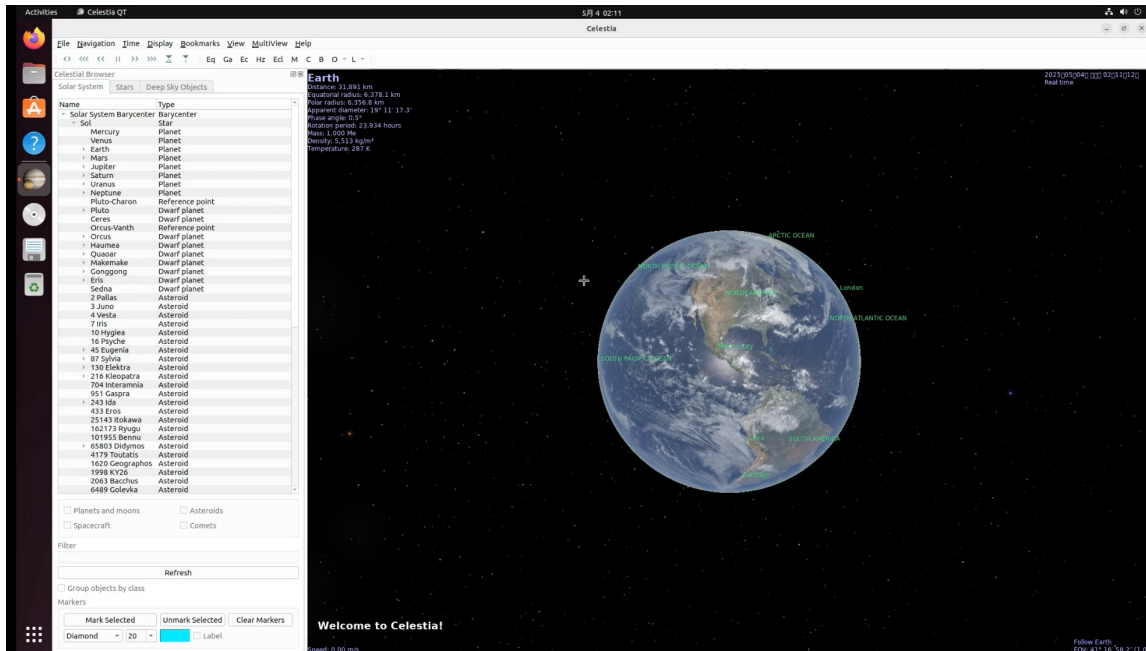
```
pyautogui.click(570, 93)
time.sleep(0.5)
```

```
# Click on the "B" checkbox to enable constellation boundaries
```

```
pyautogui.click(594, 93)
time.sleep(0.5)
```

```
# Zoom out a bit to get a better view of the constellations
```

```
pyautogui.press('minus')
time.sleep(0.5)
pyautogui.press('minus')
python
```



Use Cases

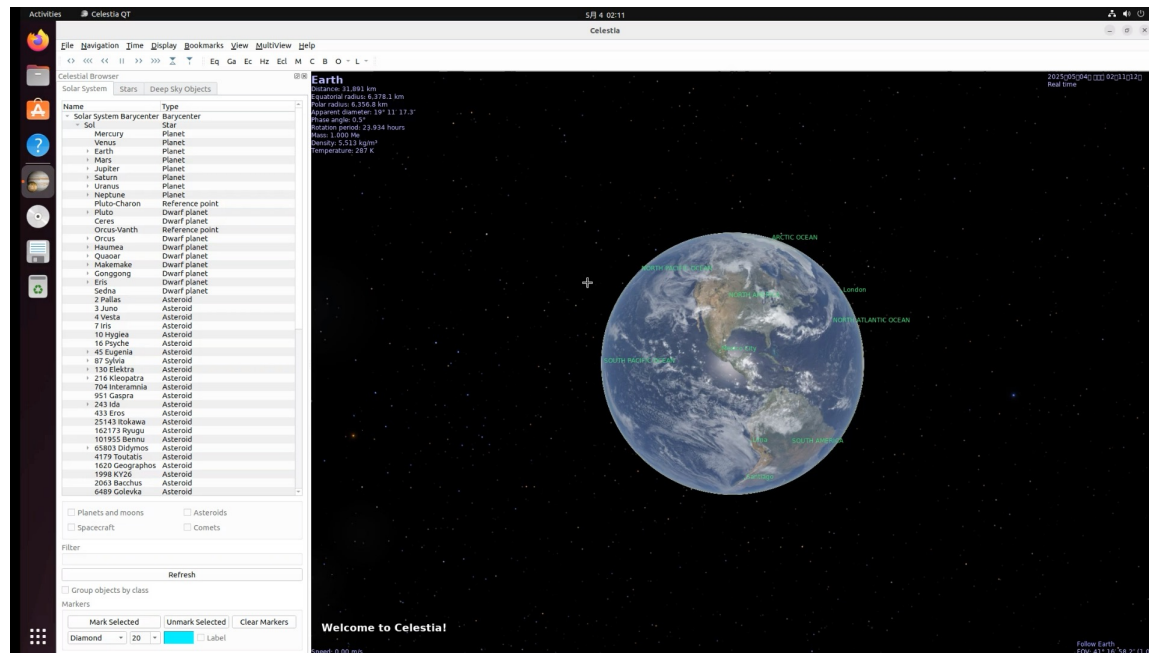
Looking at the screenshot, I can see that my previous action to click on "C" (Constellation) and "B" (Boundary) checkboxes worked correctly. I can see constellation lines displayed around the Earth (the blue lines showing the constellation patterns), and I can also see the boundary lines (the red/pink lines showing constellation boundaries).

To get a better view of the constellations and their boundaries, I should zoom out more and move away from Earth to see the full night sky. This will give a clearer view of all the constellations.

```
python import pyautogui import time
# Zoom out more to get a better view of the sky
pyautogui.press('minus')
time.sleep(0.5)
pyautogui.press('minus')
time.sleep(0.5)
pyautogui.press('minus')
time.sleep(0.5)
```

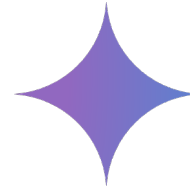
```
# Use the right arrow key to rotate away from Earth
pyautogui.keyDown('right')
time.sleep(2)
pyautogui.keyUp('right')
```

```
# Move slightly upward to get a better view of the sky
pyautogui.keyDown('up')
time.sleep(1)
pyautogui.keyUp('up')
```



Evaluation: Models

Proprietary Models



Opensource LLM / VLMs



InternVL

GUI Action Models



[16] *Navigating the Digital World as Humans Do: Universal Visual Grounding for GUI Agents*

[17] *UI-TARS: Pioneering Automated GUI Interaction with Native Agents*

[18] *GUI-Actor: Coordinate-Free Visual Grounding for GUI Agents*

Evaluation: Observations

Screenshot: Vision-only 

A11ytree: Text-only 

Screenshot + A11ytree: Hybrid  

Set-of-Mark 

Evaluation: Observations



Set-of-Mark ? “segmentation”

Partition an image into regions at different levels of granularity + label

Input: Image



Input: Image + SoM



Conversation

User What is on the left side of the right laptop?

GPT-4V On the left side of the right laptop, there is a **cup or mug**. ❌

User I want to find a seat close to windows, where can I sit?

GPT-4V You can sit on **either of the two black chairs** in front of the white desks, as they are closest to the windowed wall. ❌

Conversation + SoM

User What is on the left side of the right laptop?

GPT-4V On the left side of the right laptop (numbered as 9), there is **a lamp, which is numbered as 12**. ✔️

User I want to find a seat close to windows, where can I sit?

GPT-4V You can sit on the **chair numbered as 7**. It's the closest seat to the windows (represented by the area numbered as 1). ✔️

Evaluation: Inference

Our agents will generate **ReACT-Style** output

Examples:

Step 1: To create a new folder in Markor, I need to first open the Markor app.

```
action: {"action_type": "open_app", "app_name": "Markor"}
```

Step 2: To create a new folder, I need to click on the "Create a new file or folder" button, which is indicated by the plus icon.

```
action: {"action_type": "click", "x": 964.5, "y": 2074.5}
```

Step 3: I need to change the folder name to folder_20241224. The current text field for the folder name is visible and editable.

```
action: {"action_type": "type", "text": "folder_20241224", "x": 373.5, "y": 552.0}
```

Evaluation: General Setting

Overall success rate **remains low** (avg. ~15%)

Performance varies among domains

Best results achieved with **combined Screenshot + a11ytree** setting

Table 3: Success rates on SCIENCEBOARD. We present the performance of each agent backbone across different scientific domains under various observation settings. **Proprietary Models**, **Open-Source VLMs / LLMs**, and **GUI Action Model** are distinguished by color.

Observations	Model	Success Rate (↑)						
		Algebra	Biochem	GIS	ATP	Astron	Doc	Overall
Screenshot	GPT-4o	3.23%	0.00%	0.00%	0.00%	0.00%	6.25%	1.58%
	Claude-3.7-Sonnet	9.67%	37.93%	2.94%	0.00%	6.06%	6.25%	10.48%
	Gemini-2.0-Flash	6.45%	3.45%	2.94%	0.00%	0.00%	6.06%	3.15%
	Qwen2.5-VL-72B	22.58%	27.59%	5.88%	0.00%	9.09%	12.50%	12.94%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	0.00%	6.25%	2.69%
	UI-TARS-1.5-7B	12.90%	13.79%	0.00%	0.00%	6.06%	0.00%	2.69%
a11ytree	GPT-4o	12.90%	20.69%	2.94%	0.00%	6.06%	0.00%	7.10%
	Claude-3.7-Sonnet	19.35%	34.48%	2.94%	3.85%	12.12%	0.00%	12.12%
	Gemini-2.0-Flash	9.68%	17.24%	0.00%	0.00%	0.00%	0.00%	4.49%
	o3-mini	16.13%	20.69%	2.94%	3.85%	15.15%	6.25%	10.84%
	Qwen2.5-VL-72B	9.68%	10.34%	2.94%	0.00%	3.03%	0.00%	4.33%
	InternVL3-78B	3.23%	3.45%	0.00%	0.00%	0.00%	0.00%	1.11%
Screenshot + a11ytree	GPT-4o	22.58%	37.93%	2.94%	7.69%	3.03%	12.50%	14.45%
	Claude-3.7-Sonnet	12.90%	41.37%	8.82%	3.85%	9.09%	18.75%	15.79%
	Gemini-2.0-Flash	16.13%	24.14%	2.94%	0.00%	18.18%	12.50%	12.32%
	Qwen2.5-VL-72B	16.13%	20.69%	2.94%	0.00%	18.18%	12.50%	11.74%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	3.03%	6.25%	3.20%
Set-of-Mark	GPT-4o	6.45%	3.45%	0.00%	0.00%	3.03%	12.50%	4.24%
	Claude-3.7-Sonnet	16.13%	31.03%	5.88%	0.00%	6.06%	12.50%	11.93%
	Gemini-2.0-Flash	3.23%	0.00%	0.00%	0.00%	3.03%	6.25%	2.09%
	Qwen2.5-VL-72B	6.45%	6.90%	2.94%	0.00%	3.03%	12.50%	6.36%
	QvQ-72B-Preview	0.00%	0.00%	2.94%	0.00%	3.03%	0.00%	0.49%
	InternVL3-78B	3.23%	6.90%	2.94%	0.00%	0.00%	0.00%	2.18%
Human Performance		74.19%	68.97%	55.88%	42.31%	51.52%	68.75%	60.27%

Evaluation: General Setting

Significant performance **gaps** across domains!

Agents perform much better in biochemistry and algebra compared to other fields.

Why? “Tutorial learning”



We see this as a key **opportunity** for the future development of science agents!

Table 3: Success rates on SCIENCEBOARD. We present the performance of each agent backbone across different scientific domains under various observation settings. **Proprietary Models**, **Open-Source VLMs / LLMs**, and **GUI Action Model** are distinguished by color.

Observations	Model			Success Rate (↑)				
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	Claude-3.7-Sonnet	9.67%	37.93%	2.94%	0.00%	6.06%	6.25%	10.48%
	Gemini-2.0-Flash	6.45%	3.45%	2.94%	0.00%	0.00%	6.06%	3.15%
	Qwen2.5-VL-72B	22.58%	27.59%	5.88%	0.00%	9.09%	12.50%	12.94%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	0.00%	6.25%	2.69%
	UI-TARS-1.5-7B	12.90%	13.79%	0.00%	0.00%	6.06%	0.00%	2.69%
a11ytree	GPT-4o	12.90%	20.69%	2.94%	0.00%	6.06%	0.00%	7.10%
	Claude-3.7-Sonnet	19.35%	34.48%	2.94%	3.85%	12.12%	0.00%	12.12%
	Gemini-2.0-Flash	9.68%	17.24%	0.00%	0.00%	0.00%	0.00%	4.49%
	o3-mini	16.13%	20.69%	2.94%	3.85%	15.15%	6.25%	10.84%
	Qwen2.5-VL-72B	9.68%	10.34%	2.94%	0.00%	3.03%	0.00%	4.33%
	InternVL3-78B	3.23%	3.45%	0.00%	0.00%	0.00%	0.00%	1.11%
Screenshot + a11ytree	GPT-4o	22.58%	37.93%	2.94%	7.69%	3.03%	12.50%	14.45%
	Claude-3.7-Sonnet	12.90%	41.37%	8.82%	3.85%	9.09%	18.75%	15.79%
	Gemini-2.0-Flash	16.13%	24.14%	2.94%	0.00%	18.18%	12.50%	12.32%
	Qwen2.5-VL-72B	16.13%	20.69%	2.94%	0.00%	18.18%	12.50%	11.74%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	3.03%	6.25%	3.20%
Set-of-Mark	GPT-4o	6.45%	3.45%	0.00%	0.00%	3.03%	12.50%	4.24%
	Claude-3.7-Sonnet	16.13%	31.03%	5.88%	0.00%	6.06%	12.50%	11.93%
	Gemini-2.0-Flash	3.23%	0.00%	0.00%	0.00%	3.03%	6.25%	2.09%
	Qwen2.5-VL-72B	6.45%	6.90%	2.94%	0.00%	3.03%	12.50%	6.36%
	QvQ-72B-Preview	0.00%	0.00%	2.94%	0.00%	3.03%	0.00%	0.49%
	InternVL3-78B	3.23%	6.90%	2.94%	0.00%	0.00%	0.00%	2.18%
Human Performance		74.19%	68.97%	55.88%	42.31%	51.52%	68.75%	60.27%

Evaluation: General Setting

ATP tasks remain particularly challenging.
Why?

Because agents struggle to balance normal operations, coding skills, and highly logical reasoning.

Table 3: Success rates on SCIENCEBOARD. We present the performance of each agent backbone across different scientific domains under various observation settings. Proprietary Models, Open-Source VLMs / LLMs, and GUI Action Model are distinguished by color.

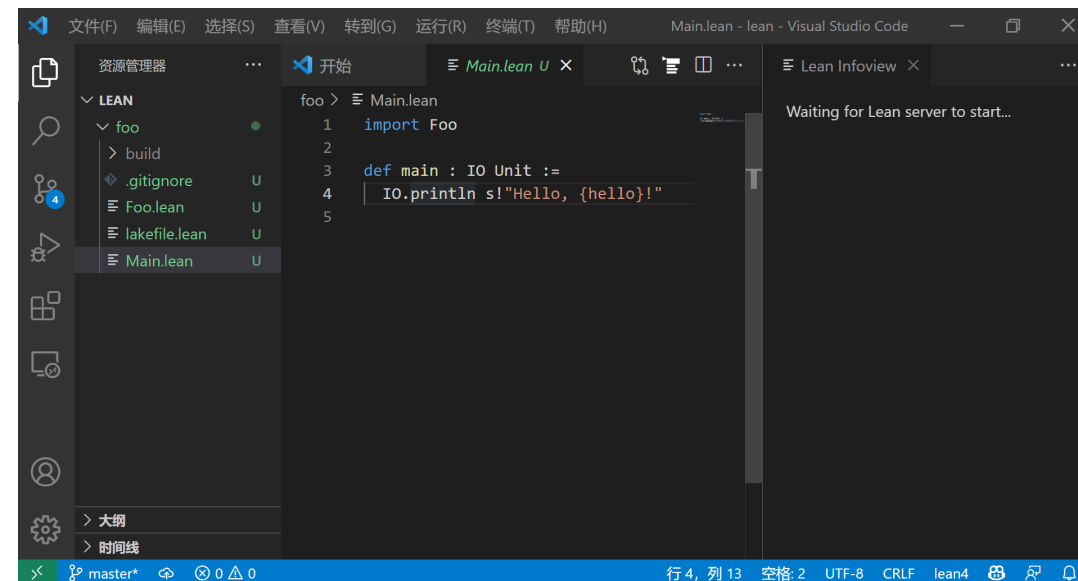
Observations	Model	Success Rate (↑)						
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	Claude-3.7-Sonnet	9.67%	37.93%	2.94%	0.00%	6.06%	6.25%	10.48%
	Gemini-2.0-Flash	6.45%	3.45%	2.94%	0.00%	0.00%	6.06%	3.15%
	Qwen2.5-VL-72B	22.58%	27.59%	5.88%	0.00%	9.09%	12.50%	12.94%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	0.00%	6.25%	2.69%
	UI-TARS-1.5-7B	12.90%	13.79%	0.00%	0.00%	6.06%	0.00%	2.69%
a11ytree	GPT-4o	12.90%	20.69%	2.94%	0.00%	6.06%	0.00%	7.10%
	Claude-3.7-Sonnet	19.35%	34.48%	2.94%	3.85%	12.12%	0.00%	12.12%
	Gemini-2.0-Flash	9.68%	17.24%	0.00%	0.00%	0.00%	0.00%	4.49%
	o3-mini	16.13%	20.69%	2.94%	3.85%	15.15%	6.25%	10.84%
	Qwen2.5-VL-72B	9.68%	10.34%	2.94%	0.00%	3.03%	0.00%	4.33%
	InternVL3-78B	3.23%	3.45%	0.00%	0.00%	0.00%	0.00%	1.11%
Screenshot + a11ytree	GPT-4o	22.58%	37.93%	2.94%	7.69%	3.03%	12.50%	14.45%
	Claude-3.7-Sonnet	12.90%	41.37%	8.82%	3.85%	9.09%	18.75%	15.79%
	Gemini-2.0-Flash	16.13%	24.14%	2.94%	0.00%	18.18%	12.50%	12.32%
	Qwen2.5-VL-72B	16.13%	20.69%	2.94%	0.00%	18.18%	12.50%	11.74%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	3.03%	6.25%	3.20%
Set-of-Mark	GPT-4o	6.45%	3.45%	0.00%	0.00%	3.03%	12.50%	4.24%
	Claude-3.7-Sonnet	16.13%	31.03%	5.88%	0.00%	6.06%	12.50%	11.93%
	Gemini-2.0-Flash	3.23%	0.00%	0.00%	0.00%	3.03%	6.25%	2.09%
	Qwen2.5-VL-72B	6.45%	6.90%	2.94%	0.00%	3.03%	12.50%	6.36%
	QvQ-72B-Preview	0.00%	0.00%	2.94%	0.00%	3.03%	0.00%	0.49%
	InternVL3-78B	3.23%	6.90%	2.94%	0.00%	0.00%	0.00%	2.18%
Human Performance		74.19%	68.97%	55.88%	42.31%	51.52%	68.75%	60.27%

Evaluation: General Setting

ATP tasks remain particularly challenging.
Why?

Hard to perform human-like operations

e.g., Autocomplete



Evaluation: General Setting

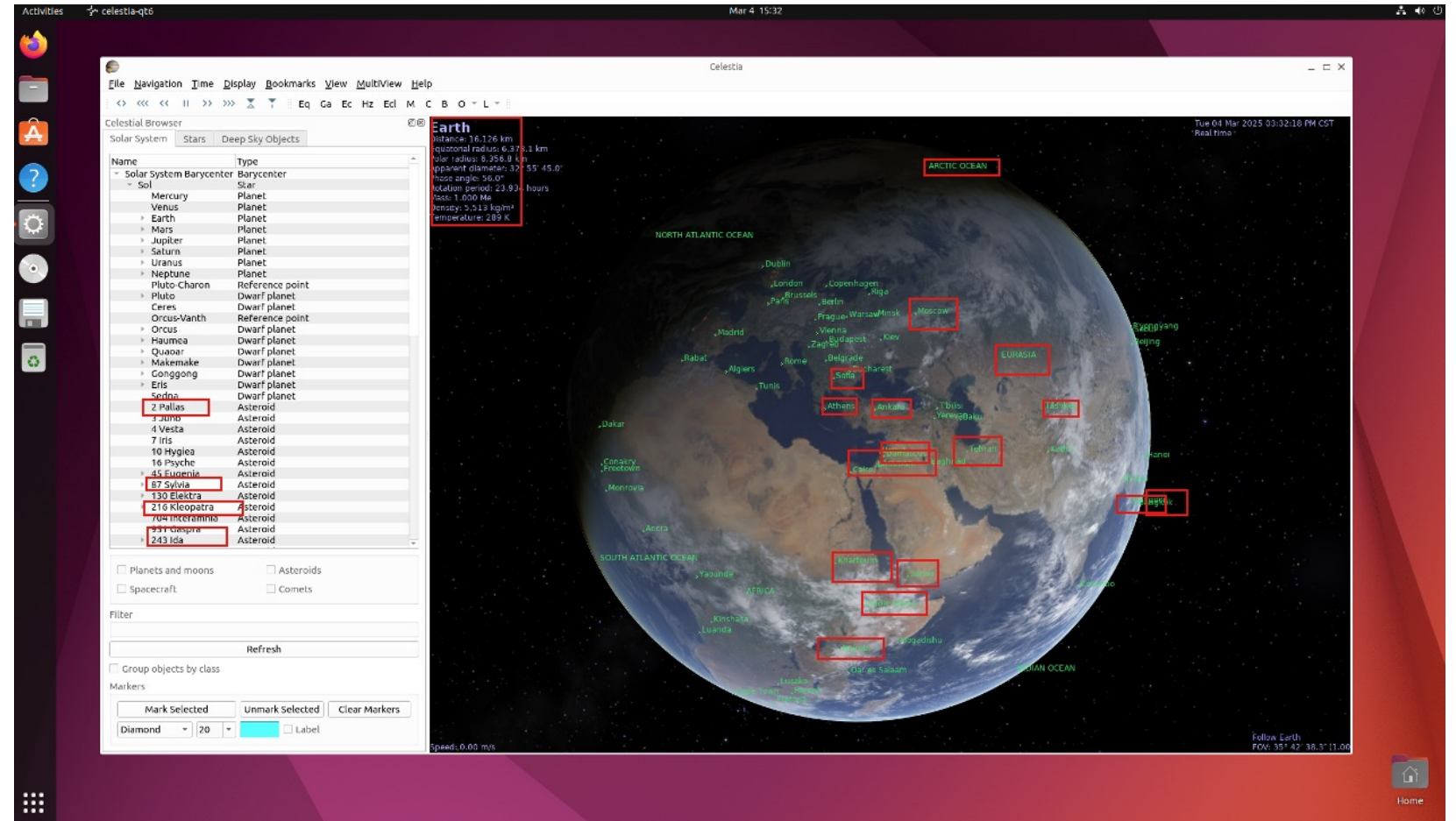
SoM? Doest not fit all scenarios!

Table 3: Success rates on SCIENCEBOARD. We present the performance of each agent backbone across different scientific domains under various observation settings. Proprietary Models , Open-Source VLMs / LLMs , and GUI Action Model are distinguished by color.

Observations	Model	Success Rate (↑)						
		Algebra	Biochem	GIS	ATP	Astron	Doc	Overall
Screenshot	GPT-4o	3.23%	0.00%	0.00%	0.00%	0.00%	6.25%	1.58%
	Claude-3.7-Sonnet	9.67%	37.93%	2.94%	0.00%	6.06%	6.25%	10.48%
	Gemini-2.0-Flash	6.45%	3.45%	2.94%	0.00%	0.00%	6.06%	3.15%
	Qwen2.5-VL-72B	22.58%	27.59%	5.88%	0.00%	9.09%	12.50%	12.94%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	0.00%	6.25%	2.69%
	UI-TARS-1.5-7B	12.90%	13.79%	0.00%	0.00%	6.06%	0.00%	2.69%
a11ytree	GPT-4o	12.90%	20.69%	2.94%	0.00%	6.06%	0.00%	7.10%
	Claude-3.7-Sonnet	19.35%	34.48%	2.94%	3.85%	12.12%	0.00%	12.12%
	Gemini-2.0-Flash	9.68%	17.24%	0.00%	0.00%	0.00%	0.00%	4.49%
	o3-mini	16.13%	20.69%	2.94%	3.85%	15.15%	6.25%	10.84%
	Qwen2.5-VL-72B	9.68%	10.34%	2.94%	0.00%	3.03%	0.00%	4.33%
	InternVL3-78B	3.23%	3.45%	0.00%	0.00%	0.00%	0.00%	1.11%
Screenshot + a11ytree	GPT-4o	22.58%	37.93%	2.94%	7.69%	3.03%	12.50%	14.45%
	Claude-3.7-Sonnet	12.90%	41.37%	8.82%	3.85%	9.09%	18.75%	15.79%
	Gemini-2.0-Flash	16.13%	24.14%	2.94%	0.00%	18.18%	12.50%	12.32%
	Qwen2.5-VL-72B	16.13%	20.69%	2.94%	0.00%	18.18%	12.50%	11.74%
	InternVL3-78B	6.45%	3.45%	0.00%	0.00%	3.03%	6.25%	3.20%
Set-of-Mark	GPT-4o	6.45%	3.45%	0.00%	0.00%	3.03%	12.50%	4.24%
	Claude-3.7-Sonnet	16.13%	31.03%	5.88%	0.00%	6.06%	12.50%	11.93%
	Gemini-2.0-Flash	3.23%	0.00%	0.00%	0.00%	3.03%	6.25%	2.09%
	Qwen2.5-VL-72B	6.45%	6.90%	2.94%	0.00%	3.03%	12.50%	6.36%
	QvQ-72B-Preview	0.00%	0.00%	2.94%	0.00%	3.03%	0.00%	0.49%
	InternVL3-78B	3.23%	6.90%	2.94%	0.00%	0.00%	0.00%	2.18%
Human Performance		74.19%	68.97%	55.88%	42.31%	51.52%	68.75%	60.27%

Evaluation: General Setting

“Cognitive burden”



Evaluation: Modular Setting

Planner + Action Model



Step 1: To create a new folder in Markor, I need to first open the Markor app.

```
action: {"action_type": "open_app", "app_name": "Markor"}
```

Step 2: To create a new folder, I need to click on the "Create a new file or folder" button, which is indicated by the plus icon.

```
action: {"action_type": "click", "x": 964.5, "y": 2074.5}
```

Step 3: I need to change the folder name to folder_20241224. The current text field for the folder name is visible and editable.

```
action: {"action_type": "type", "text": "folder_20241224", "x": 373.5, "y": 552.0}
```

Evaluation: Modular Setting



GPT-4o as the planner + GUI model

Clear performance improvement (up to ~20% SR)

Separating planning and action offers a promising direction!

Table 4: Success rates of different VLM agent combinations under the planner + grounding model setting on SCIENCEBOARD. The observation setting used in this experiment is screenshot. Colors denote Proprietary Models , Open-Source VLMs and GUI Action Models.

Planner	Grounding Model	Success Rate (↑)				
		Algebra	Biochem	GIS	Astron	Overall
GPT-4o	OS-Atlas-Pro-7B	6.25%	10.34%	0.00%	3.03%	4.92%
	UGround-V1-7B	0.00%	3.45%	0.00%	3.03%	1.62%
	Qwen2.5-VL-72B	12.50%	34.48%	11.76%	9.09%	16.96%
	UI-TARS-72B	3.23%	10.34%	5.88%	6.06%	6.38%
	GUI-Actor-7B	21.88%	44.83%	2.94%	12.12%	20.44%
GPT-4o		3.23%	0.00%	0.00%	0.00%	0.81%

Analysis

CLI is very helpful effective.

Finding: (V)LMs tend to prefer completing tasks **via CLI** when possible.

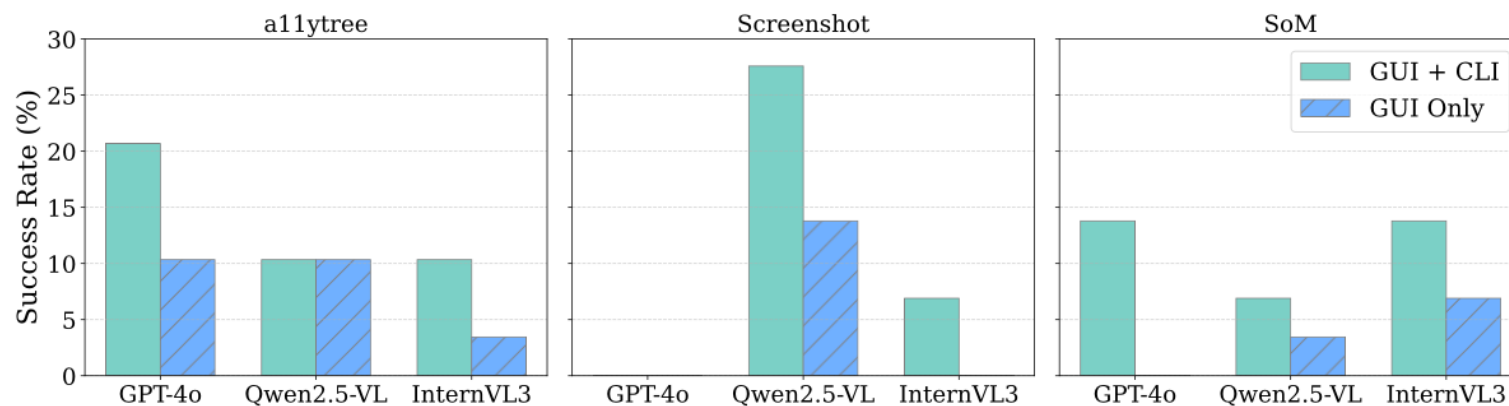



















Figure 8: Extended analysis of Vision-Only vs. Hybrid Interface.

More analysis available in the paper!

Leaderboard

ScreenshotA1ly TreeScreenshot + A1ly TreeSet of Marks

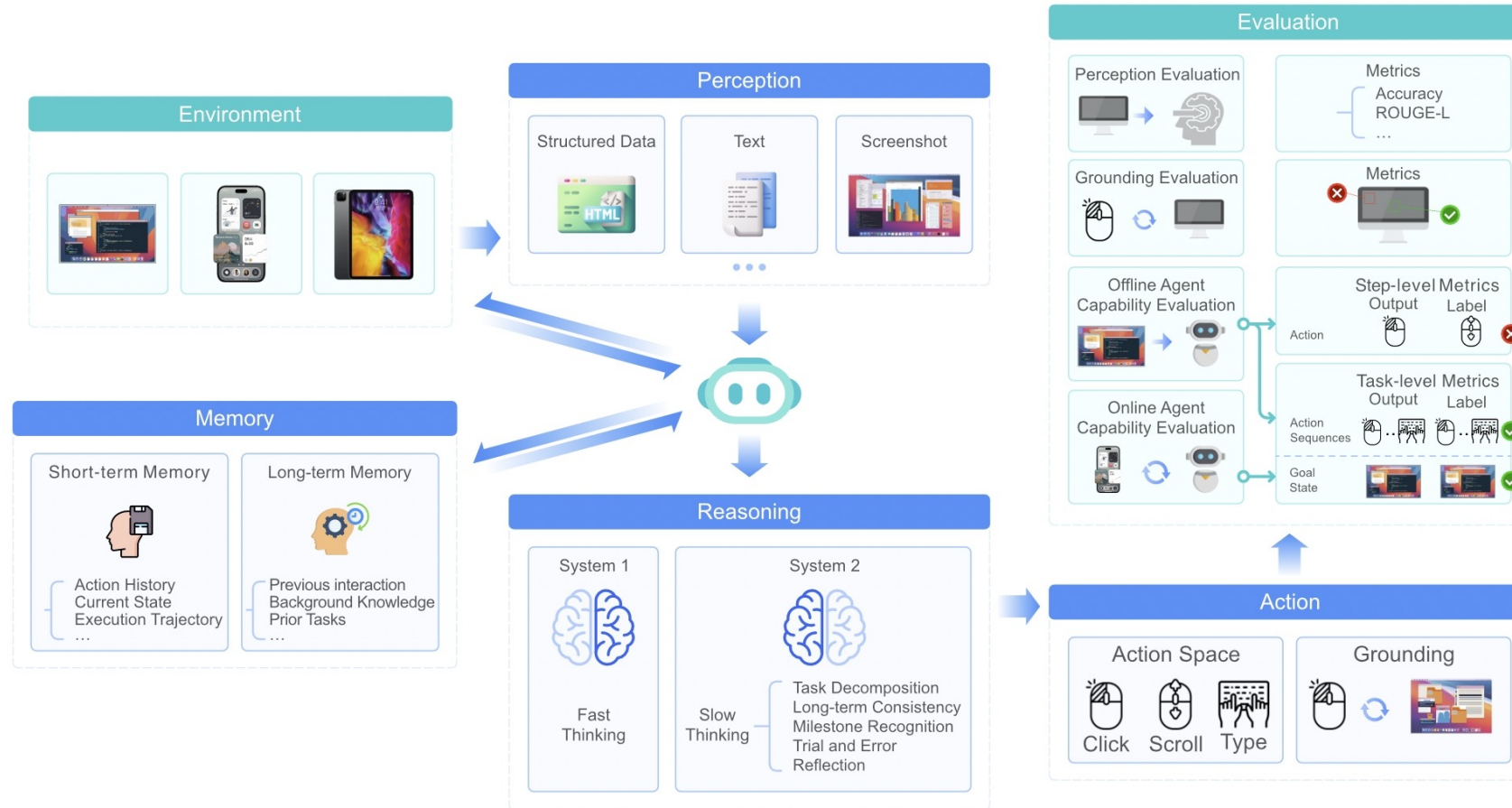
Search by keywords

O..	Settings	% Acc ↓	% Alg	% Biochem	% GIS	% ATP	% Astron	% Doc
	Calude-3.7-Sonnet w/ screenshot...	15.79	12.90	41.37	8.82	3.85	9.09	18.75
	GPT-4o (2024-08-06) w/ screensh...	14.45	22.58	37.93	2.94	7.69	3.03	12.50
	GPT-4o (2024-08-06) w/ set_of_m...	14.45	6.45	3.45	0.00	0.00	3.03	12.50
	Qwen2.5-VL-72B w/ screenshot	12.94	22.58	27.59	5.88	0.00	9.09	12.50
	Gemini-2.0-Flash w/ screenshot+a...	12.32	16.13	24.14	2.94	0.00	18.18	12.50
	Calude-3.7-Sonnet w/ a1ly_tree	12.12	19.35	34.48	2.94	3.85	12.12	0.00
	Calude-3.7-Sonnet w/ set_of_marks	11.93	16.13	31.03	5.88	0.00	6.06	12.50
	Qwen2.5-VL-72B w/ screenshot+a...	11.74	16.13	20.69	2.94	0.00	18.18	12.50
	o3-mini (2025-01-31) w/ a1ly_tree	10.84	16.13	20.69	2.94	3.85	15.15	6.25
	Calude-3.7-Sonnet w/ screenshot	10.48	9.67	37.93	2.94	0.00	6.06	6.25
	GPT-4o (2024-08-06) w/ a1ly_tree	7.10	12.90	20.69	2.94	0.00	0.00	6.06
	Qwen2.5-VL-72B w/ set_of_marks	6.36	6.45	6.90	2.94	0.00	3.03	12.50
	UI-TARS-1.5 w/ screenshot	5.92	12.90	13.79	0.00	0.00	6.06	0.00
	Gemini-2.0-Flash w/ a1ly_tree	4.49	9.68	17.24	0.00	0.00	0.00	0.00
	Qwen2.5-VL-72B w/ a1ly_tree	4.33	9.68	10.34	2.94	0.00	3.03	0.00
	InternVL3-78B w/ screenshot+a11...	3.20	6.45	3.45	0.00	0.00	3.03	6.25
	Gemini-2.0-Flash w/ screenshot	3.15	6.45	3.45	2.94	0.00	0.00	6.06

<https://qiushisun.github.io/ScienceBoard-Home/>

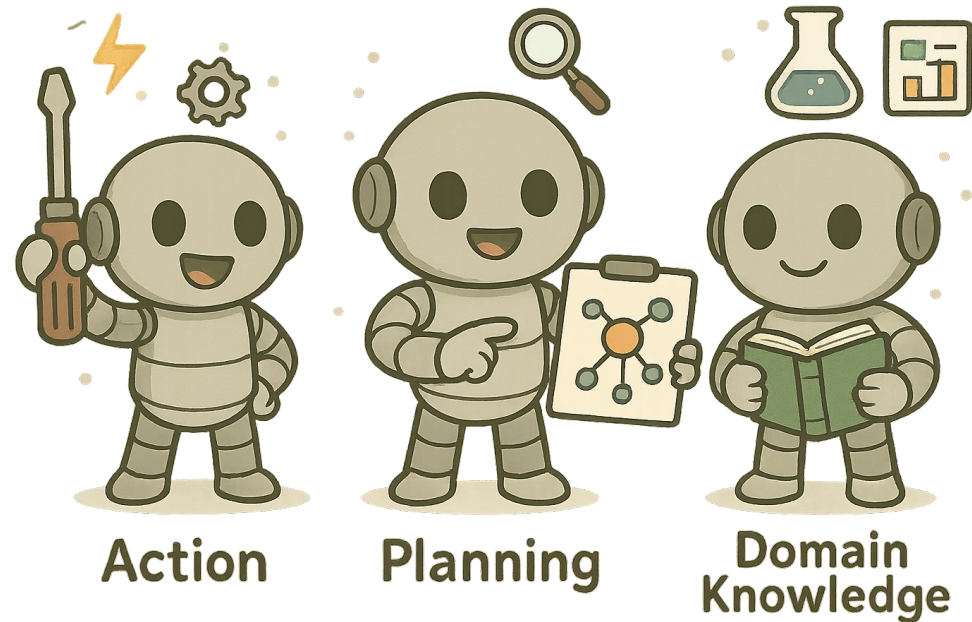
Towards High Value Area

Recap: Core capabilities and evaluation for GUI agents



Towards High Value Area

For science agents, we need to strike a balance!



Some Limitations

1. The current evaluation uses binary (0/1) scoring; allowing **partial credit** for intermediate steps may better reflect real-world scenarios.

Challenge: exploration space!

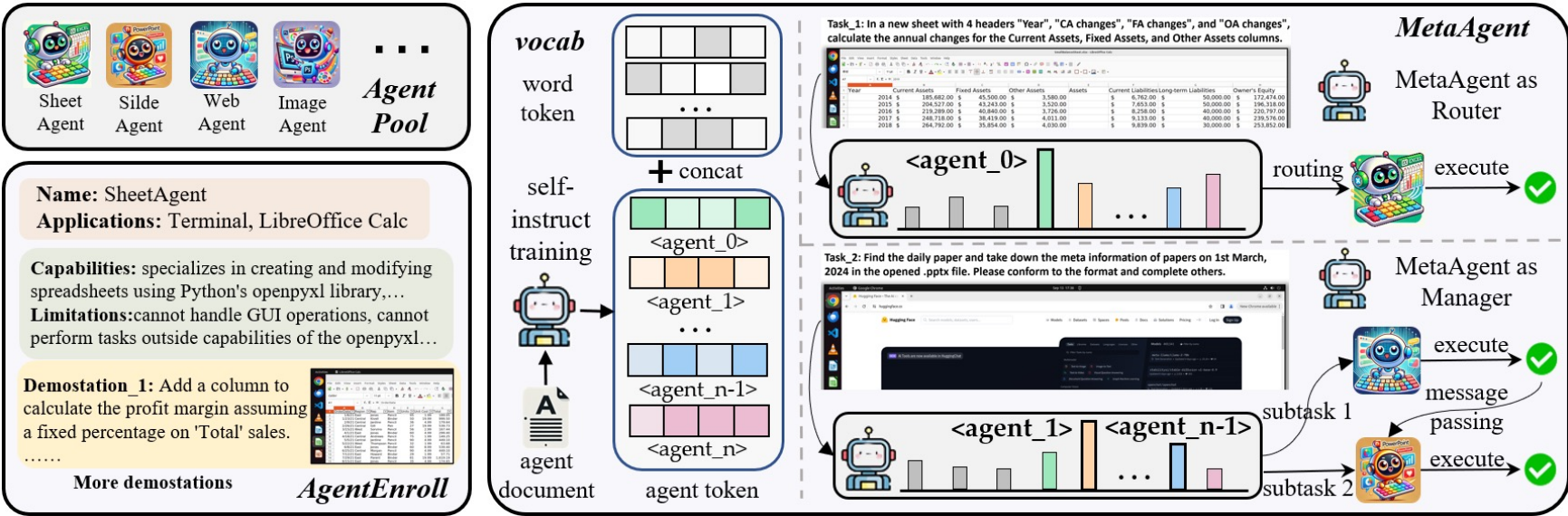
2. All evaluated software is open-source distribute; incorporating commercial software remains challenging.

Challenge: evaluation!

Part4 | Future Direction



Future Directions



Example: Heterogeneous Agents As Specialized Generalist Computer Assistant

AGENTSTORE: SCALABLE INTEGRATION OF HETEROGENEOUS AGENTS AS SPECIALIZED GENERALIST COMPUTER ASSISTANT

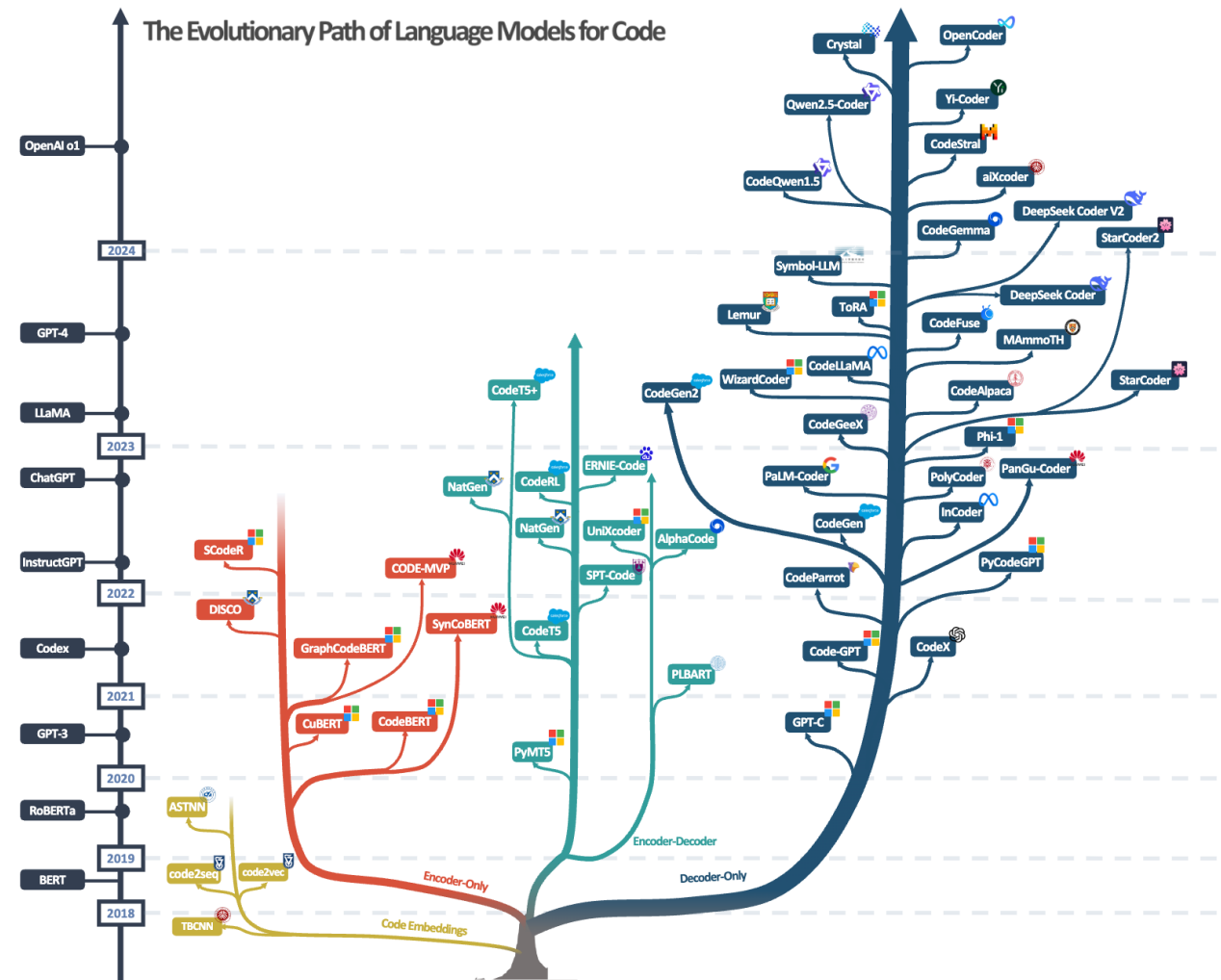
Chengyou Jia^{1,2*}, Minnan Luo¹, Zhuohang Dang¹, Qiushi Sun^{2,3}, Fangzhi Xu^{1,2}, Junlin Hu², Tianbao Xie³, Zhiyong Wu²
¹Xi'an Jiaotong University, ²Shanghai AI Lab, ³The University of Hong Kong
 cp3jia@stu.xjtu.edu.cn, wuzhiyong@pjlab.org.cn

“通专融合的Agent系统”

Future Directions

Integration with CodeLLMs?

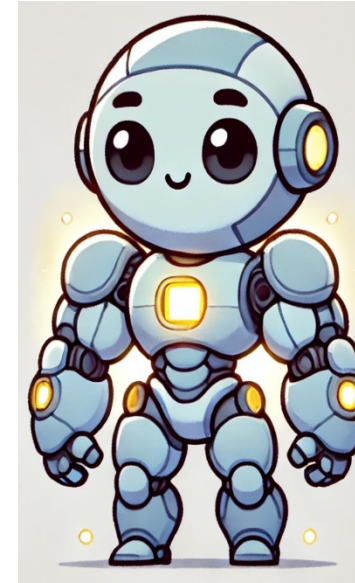
This enables the adoption of more **data-driven** methods, like ScienceAgentBench.



Future Directions



Mid-Training



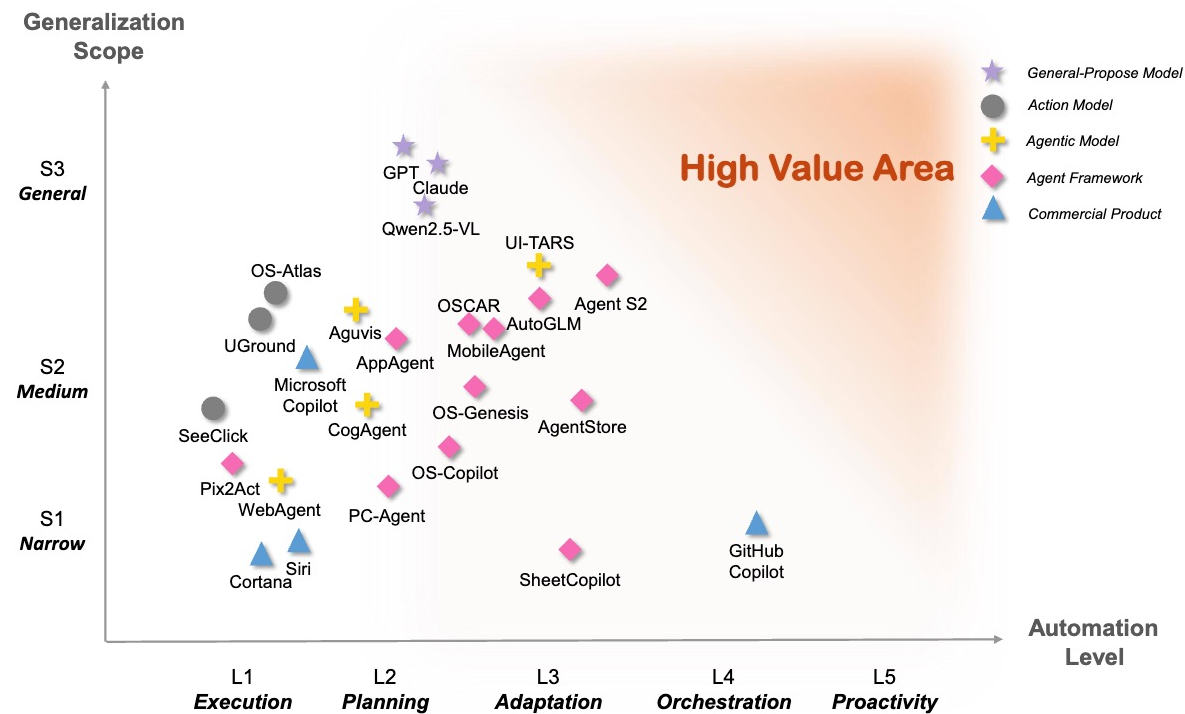
Potential solution: Mid-training?

Breaking the Data Barrier – Building GUI Agents Through Task Generalization

Junlei Zhang^{*◇☆} Zichen Ding^{*♣} Chang Ma[♣] Zijie Chen^{◇☆} Qiushi Sun[♣]
Zhenzhong Lan[☆] Junxian He[★]
[◇]Zhejiang University [☆]Westlake University [♣]Shanghai AI Laboratory
[♣]The University of Hong Kong [★]HKUST

Future Directions

Towards High Value Area



Stay tuned!

OS-MAP: How Far Can Computer Use Agents Go in Breadth and Depth?

Towards High Value Area

1. Operating robotic arms?
2. Controlling exoskeletons?
3. Utilizing highly specialized scientific software?
4. ...

We are just standing at the dawn of a long journey!

Our Project

ScienceBoard

Evaluating Multimodal Autonomous Agents in Realistic Scientific Workflows

Introducing ScienceBoard, a first-of-its-kind evaluation platform for multimodal agents in *scientific workflows*. ScienceBoard is characterized by the following core features:



Pioneering Application: ScienceBoard is the first to bring computer-using agents into the domain of scientific discovery, enabling autonomous research assistants across disciplines.



Realistic Environment: We provide a dynamic, visually grounded virtual environment integrated with professional scientific software, supporting both GUI and CLI interaction in real-time workflows.



Challenging Benchmark: A new benchmark of 169 rigorously validated tasks across 6 core domains is introduced, capturing real-world challenges.



Comprehensive Evaluations: We presents systematic evaluations across a wide range of agents powered by LLMs, VLMs, and GUI action models.

arXiv

Code

Data

VM Snapshot





上海人工智能实验室
Shanghai Artificial Intelligence Laboratory



SCHOOL OF
COMPUTING &
DATA SCIENCE
The University of Hong Kong

We are just standing at the dawn of a long journey!

Thanks for listening!

Contact: qiushisun@connect.hku.hk