

Automated Generative Experimental Design via Relational Learning-Guided Large Language Models

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Introduction

Large Language Models (LLMs) have revolutionized AI, achieving impressive results on tasks traditionally requiring human intelligence. At the forefront of applying Al in science is the automation of research, offering faster discoveries and new insights. By using traditional models and logic to generate hypotheses, LLMs can then be given a logical scaffold to reason around, reducing hallucinations and enhancing the reliability of its outputs. Here we demonstrate a way to leverage the use of LLMs to automate experimental design, given relational learning derived hypotheses and physical laboratory constraints. We integrate this methodology with an automated laboratory cell, demonstrating a way of flexibly doing automated scientific discovery in a predefined hypothesis space.





Automated analysis

7

Experimental outcome is analyzed³ and compared to expected outcomes from (4) using automated hypothesis testing. Results are added to knowledge-base and a report is generated.

> Automated experiment execution through standardized lab protocols

> > **Running the experiment**



6

5

Experiment Formalization Experimental design is **automatically** formalized in adherence to the interface of the automated laboratory, producing a standardized, machine readable,

experimental protocol in JSON

Hypothesis Filtering & Weighting

Learn an association between logic programs and a biological observable through regularized linear models, creating a descriptive hypothesis concerning a measurable biological observable (e.g. metabolite)².

Hardware availability constrains design-space

Generative Experimental Design



3

The hypothesis is passed to an **LLM** along with hardware and feasibility constraints. From that, automatically generate an experimental design to prove or disprove generated hypothesis

4

Control software determines formalization

Experiment is automatically performed in an automated laboratory cell along with integrated phenotyping, sampling and metabolomics readouts based on ion mobility mass spectrometry³



Conclusion

Generative experimental design is a promising avenue for enabling the complete automation of science. By integrating traditional modeling and relational learning alongside Large Language Models, we can guide this process to not only generate valid scientific hypotheses and creative experimental designs without excessive hallucinations, but also automatically incorporate insights from complex interconnected datasets into the scientific process.

Image generated using GPT40 with the prompt: Isometric view of a stylized automated laboratory cell performing biology experiments 1. Ross D. King, Ashwin Srinivisan, Luc Dehaspe. WARMR: A Data Mining Tool for Chemical Data (2001). Journal of Computer-Aided Molecular Design. 2. Daniel Brunnsåker, Filip Kronström, levgeniia A. Tiukova, Ross D. King. Interpreting protein abundance in Saccharomyces cerevisiae through relational learning (2024). Bioinformatics. 3. Gabriel K. Reder, Erik Y. Bjurström, Daniel Brunnsåker et al.. AutonoMS: Automated Ion Mobility Metabolomic Fingerprinting (2024). Journal of the American Society for Mass Spectrometry.

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