

# Can Large Language Models *Unlock* Novel Scientific Research Idea?

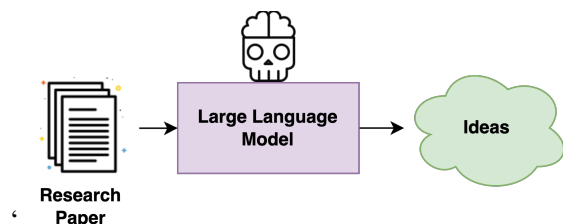


Figure 1: Large language model suggesting future research ideas after reading a research paper

## Abstract

*"An idea is nothing more nor less than a new combination of old elements"* (Young, 2019). The widespread adoption of Large Language Models and publicly available ChatGPT has marked a significant turning point in the integration of Artificial Intelligence (AI) into people's everyday lives. This study explores the capability of large language models in generating novel research ideas based on information from research papers. We conduct a thorough examination of 4 LLMs in five domains (e.g., Chemistry, Computer Economics, Medical, and Physics). We found that Claude-2 and GPT-4 generated future research ideas are more aligned with the author's perspective than GPT-3.5 and Gemini. We also found that Claude-2 generates more diverse future research ideas than GPT-4, GPT-3.5, and Gemini 1.0. We further performed a human evaluation of the novelty, relevancy, and feasibility of the generated future research ideas. This investigation offers insights into the evolving role of LLMs in idea generation, highlighting both its capability and limitations. Our work represents a first step toward evaluating and utilizing language models in generating new research ideas. We make our datasets and codes publicly available.<sup>1</sup>

"Innovation is seeing what everybody

<sup>1</sup><https://anonymous.4open.science/r/Future-Idea-Generation-DC42/README.md>

has seen and thinking what nobody has thought" —Dr. Albert Szent-Györgyi

## 1 Introduction

An *idea* can be defined as a thought or suggestion aimed at solving a problem or considering a possibility. This concept is central to fields ranging from philosophy to science and economics. According to (Plato et al., 2000), ideas are archetypal forms that represent the most accurate reality. In the context of scientific research, (Kuhn and Hawkins, 1963) in "The Structure of Scientific Revolutions" describes an idea as a realization or hypothesis that can challenge and shift paradigms within a scientific community. Therefore, an idea can be understood as a cognitive construct that arises from the human mind's ability to process information, reflect, and imagine, serving as a cornerstone for creativity, problem-solving, and innovation. Idea generation can be generally understood as a state of focused internally-directed attention involving controlled semantic retrieval (Benedek et al., 2014).

As technology improves, new capabilities emerge. Ever since the Turing Test was proposed in the 1950s, humans have explored the mastering of language intelligence by machine (Zhao et al., 2023). Technological advancements serve two key functions in innovation. Firstly, they influence the goals of generating and selecting ideas. Secondly, they impact the methodology of how ideas are generated and chosen (Kornish and Hutchison-Krupat, 2017). Large language models (LLMs) have exhibited unparalleled mastery of natural language processing (NLP). Since, these have become increasingly powerful, researchers have begun to investigate their reasoning ability in problem-solving tasks (Yao et al., 2022; Brahman et al., 2023). The concept of an idea is essentially a new combination of old elements. LLMs have access to a broad spectrum of knowledge, due to their extensive training on vast amounts of text data. However, understand-

ing how information extracted from a research paper can give rise to new ideas, which have not yet been explored. This leads us to ponder:

Can Large Language Models read a scientific paper and suggest new research ideas or directions?

Motivated by this, in this paper, we analyze the potential of LLMs in generating future research directions/ideas. As LLMs possess knowledge across various domains, we investigate five specific areas, viz. Computer Science, Physics, Chemistry, Economics, and Medicine. To address this task, we create a dataset of papers published after the year 2022 from these five domains. We annotate the papers with future research ideas. To evaluate the novelty and relevance of ideas generated by the LLMs, we propose an Idea Alignment Score (IAS-core). This score reflects how well the generated ideas align with those proposed by the authors. To study the model’s ability to generate diverse ideas, we propose an Idea Distinctness Index. We analyze and discuss the performance and limitations of four LLMs: Gemini (Anil et al., 2023), Claude-2 (Ant, 2023), GPT-3.5, and GPT-4 (OpenAI, 2023). We further conduct a human evaluation of 460 generated ideas in computer science to study the novelty, relevance, and feasibility of these research ideas. This paper demonstrates that LLMs have the potential to generate relevant, distinct, feasible, and novel ideas to some extent.

We hope that the findings of this work will unlock the potential of LLMs in idea generation and encourage further investigation in this direction. Additionally, we hope that this work will open a new topic of research focused on accelerating scientific research by automatically generating research ideas.

#### OUR CONTRIBUTIONS:

- We make the first attempt to evaluate the potential of LLMs in generating future research ideas.
- To address the task, we create a novel dataset of recent papers of five domains (Computer science, Economics, Chemistry, Physics, Medical).
- To assess the quality of generated ideas from

LLMs, we propose Idea Alignment Score and Idea Distinctness Index to evaluate the idea generation capability.

- We discuss the challenges associated with human evaluation and conduct a human evaluation on 460 generated ideas.

## 2 Related Work

Recently, LLMs have shown emergent abilities to perform tasks they were not explicitly trained for (Wei et al., 2022; Bubeck et al., 2023). This includes common sense question answering, code generation, and cross-domain problem solving, enriching their utility across unforeseen domains (Chen et al., 2021; Sarsa et al., 2022). Their capability extends to advanced scientific domains such as computer science, physics, medicine, and mathematics (Romera-Paredes et al., 2023; Huang et al., 2023). Technology Semantic Network (TechNet) was proposed to stimulate idea generation in engineering design (Sarica et al., 2021). There have been a few works in the discovery of new proteins to accelerate scientific discovery. The prior work reported in (Spangler et al., 2014) involves utilizing published studies to find new protein kinases that phosphorylate the tumor suppressor protein p53.

A hypothesis is a hunch, assumption, suspicion, assertion or an idea about a phenomenon, relationship or situation, the reality or truth of which you do not know (Kumar, 1996). There have been some works on hypothesis generation. Initial studies on automated hypothesis generation begin by constructing a corpus of distinct concepts. Subsequently, they explore the relationships between these concepts using machine learning techniques, such as analyzing the similarities among vectors representing different words (or concepts) (Tshityoyan et al., 2019), or applying link prediction methods over a graph (where concepts are nodes) (Nadkarni et al., 2021). Recently (Qi et al., 2023; Wang et al., 2023) used LLMs and extensive pre-existing knowledge of various scientific fields for hypothesis generation.

Unlike previous works, which have generally focused on the association and interpretation of two variables, our current work aims to evaluate the capabilities of LLMs in generating future research ideas.

### 3 Dataset

Our dataset creation involves three steps: (1) Dataset Collection, (2) FRI Identification and removal, and (3) FRI generation.

#### 3.1 Dataset Collection

We construct a corpus D from S2ORC collected 100 papers from the domains of Computer Science, Economics, Physics, Chemistry, Medical from (Lo et al., 2020). To ensure the quality and relevance of the data and to utilize the future research ideas mentioned in a paper, the selected papers must meet the following requirements: (1) the paper must contain the full content, and (2) the paper must include a section on future work.

#### 3.2 FRI Identification and Removal

We first identify and removed any potential research ideas mentioned in the paper. By doing this, we ensure that the large language models have no prior access to these ideas, which could otherwise affect the objectivity of the analysis.

##### 3.2.1 Annotation Guidelines

Inspired by Hao et al. (2020), we define a future research idea as a discussion that the authors believe they will conduct in the future or believe needs to be investigated in future research. We discuss more details about the annotation guidelines in Appendix A.

##### 3.2.2 Annotator Training

Given the complexity of the papers and their frequent use of technical terminology, we hired two doctoral students, each boasting over four years of experience in scientific research publishing. To facilitate their training, an expert with more than ten years of experience in scientific publishing annotated 20 random papers from each domain, adhering to our guidelines. After this initial round of annotation, we reviewed and corrected any misinterpretations with the annotators, further refining their training and enhancing the clarity of our annotation guidelines. To assess the effectiveness of the initial training, we compiled another 20 papers from each domain. From the second round onwards, the annotators demonstrated improved proficiency, accurately identifying at least 95% of the future research ideas on average.

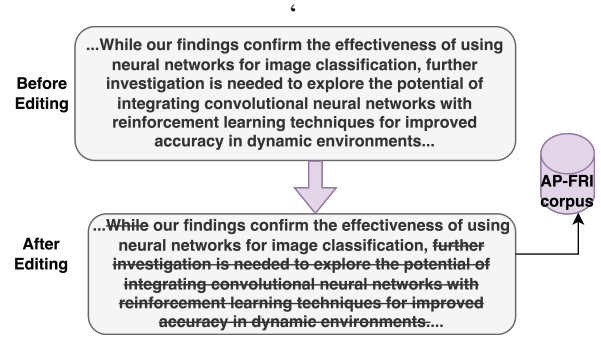


Figure 2: An example for FRI editing; Here the strike through text is removed from the paper text

##### 3.2.3 Annotation Process

We regularly monitored the annotated data, placing emphasis on identifying and rectifying inconsistencies and cases of confusion. We also implemented an iterative feedback system that continuously aimed to refine and improve the annotation process. In cases of conflict or confusion, we removed those papers as we wanted only good quality dataset. Following the annotation phase, we obtained an average inter-annotator agreement score of 0.94 using Cohen’s kappa (Cohen, 1960), indicating a substantial consensus among the annotators.

##### 3.2.4 Annotator’s Pay

We compensated each annotator according to the standard PhD salaries in India, based on the hours they worked. The appointment and salaries adhere to our university’s established practices. Payment was made per paper since the time required to read and extract future research ideas from each paper varies, depending on its complexity, technical terminology, and the annotator’s familiarity with the subject. Thus, paying based on time spent could have potentially compromised the quality of the annotations. To maintain accuracy and prevent fatigue, we imposed a daily limit of six hours for annotators.

##### 3.2.5 Future Work Removal

We observed two types of future research ideas (FRIs)<sup>2</sup> mentioned in the paper:-

- **Direct FRI:** When the sentences that mention future research idea only contains future research idea. For example *"In future*

<sup>2</sup>In this paper, we use the terms ‘ideas,’ ‘research ideas,’ ‘future research ideas,’ and ‘FRI’ interchangeably to frequently refer to future research ideas.

work, we plan to extend our approach to other code-mixed languages and evaluate its performance on more NLP tasks."

- **Mixed FRI:** We found that sometimes research papers articulate future research ideas along with other essential information of the paper in a single sentence. For example in Figure 2, this sentence not only summarizes the current research findings but also clearly outlines a direction for future work.

**AP-FRI Corpus:** We removed the sentence from the paper’s input text if it pertains to Direct FRI. However, in the case of Mixed FRI, we did not entirely remove the sentences; instead, we eliminated only parts of sentences or markers indicating future research ideas. We added the removed future ideas to a corpus, which we refer to as the AP-FRI (Author Perspective Future Research Idea Corpus). This corpus contains the future research ideas proposed by the authors of the paper. Also, before adding to the AP-FRI corpus, we merged the sentences about the same topic into a single group.

### 3.3 FRI Generation using LLM

We investigate various prompts and utilize the following prompts to generate FRIs for papers.

**System:** You are a research scientist.  
**User:** Imagine you are a research scientist. After reading the following paper, brainstorm to generate potential future research ideas:

[paper text]

Potential future research ideas from the paper in bullet points are:

Here, ‘[paper text]’ contains the full content of the paper after removal of future work sections.

### 3.4 Data Statistics

Figure 3 provides a domain-wise distribution of the average word count in academic papers, excluding discussions on future work (FWK). It can be observed that the length of papers across all fields falls within a range of 7,000 to 8,000 words. Additionally, we calculated the average word count of extracted future work within each domain, providing comparative insights into how different fields

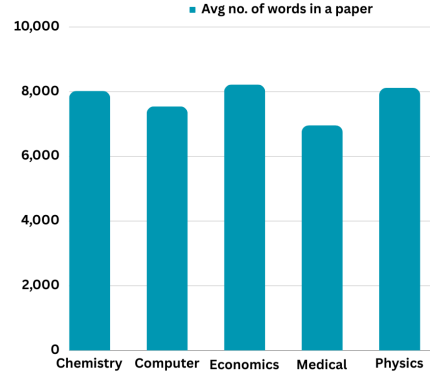


Figure 3: Domain vs Avg. number of words in a paper w/o FWK

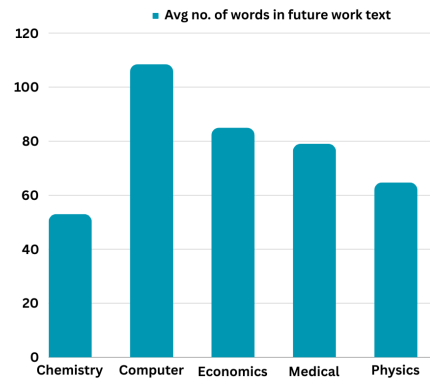


Figure 4: Domain vs Avg. number of words in FWK

prioritize discussions of future research directions. Figure 4 compares the average word count of future work text across six distinct scholarly domains. We observed that the literature in Computer Science notably prioritizes extensive discourse on future research, with an average word count significantly higher than that of other disciplines. In contrast, the literature in Chemistry demonstrates a more concise approach to discussions of future research, as evidenced by its lower average word count.

## 4 Experiments

### 4.1 Challenges

To accurately assess the novelty, relevance, and applicability of ideas generated by LLMs, evaluators must possess a high level of expertise in the specific domain and a deep understanding of the research topic to fully grasp the context. Additionally, they need knowledge of related literature to evaluate the ideas’ future potential and the broader implications of their implementation.



## 4.2 Idea Alignment Score (IAScore)

With the above challenges, the evaluation of ideas generated by LLMs is a challenging process that demands a number of high level of domain-specific experts. We, therefore, proposed an *Idea Alignment Score* (IAScore), which reflects how well the generated ideas align with those proposed by the author. The underlying idea for this score is that authors of accepted papers can be regarded as experts in their respective subjects. The reason being that they possess thorough background knowledge and have conducted deep analyses of the research topic before getting the paper accepted. Consequently, they are well-acquainted with the pertinent challenges which also may have been discussed by expert reviewers. Therefore, we propose that future ideas mentioned by the authors in the paper could be utilized as good quality of potential FRIs.

The IAScore quantifies the alignment of newly generated ideas with author’s perspectives within a specific domain, and is computed via a two-step process, detailed in Equations 1 and 2.

Initially, we compute the average alignment score  $\text{AvgScore}_j$  for each paper’s ideas. The IdeaMatcher model measures the alignment between the paper’s author *Future Research Ideas* (AP-FRI<sub>j</sub>) and its each generated idea  $I_{ij}$ . The subscript  $i$  indexes the  $i$ -th idea within the  $j$ -th paper, where  $N_j$  represents the total number of ideas proposed in that paper.

$$\text{AvgScore}_j = \frac{1}{N_j} \sum_{i=1}^{N_j} \text{IdeaMatcher}(\text{AP-FRI}_j, I_{ij}) \quad (1)$$

LLMs may generate new ideas that even the author may not have thought of. They can also generate additional future ideas, which may or may not be useful. Our goal is for this score is that the LLMs must have generated at least the author’s proposed potential future ideas. Therefore, in our formula of  $\text{AvgScore}_j$ , the sum of the alignment scores for a paper’s ideas is divided by the total number of the author’s proposed ideas,  $N_j$ , to normalize the score.

$$\text{IAScore}_{\text{domain, M}} = \frac{1}{P} \sum_{j=1}^P \text{AvgScore}_j \quad (2)$$

Subsequently, we aggregate the individual paper scores to calculate the domain-wise IAScore. This

aggregation, presented in Equation 2, averages the  $\text{AvgScore}_j$  values across all  $P$  papers within the domain. Higher the value of  $\text{IAScore}_{\text{domain}}$  signifies the more alignment of the generated ideas with author’s perspective of all papers generated by model  $M$ .

### 4.2.1 IdeaMatcher

To select an effective IdeaMatcher, we create a small annotated corpus. We first manually searched for matching pairs of ideas from generated ideas and AP-FRI of the paper. After obtaining 61 matching pairs, we searched for non-matching pairs of ideas, which is straightforward as only one generated idea will match or won’t match with another one from AP-FRI while others would not match, so we picked an equal number of non-matching pairs. Then, we experimented with the idea-matching task by considering it similar to the Natural Language Inference task. In particular, we considered the generated FRIs to be hypotheses and their corresponding AP-FRIs of the paper to be premises. If the idea matches, the hypothesis should be entailed by the premise. In particular, we used a pre-trained RoBERTa MNLI model (Liu et al., 2019) for this task. We found that this technique produces many false negative cases, resulting in an accuracy of 65.5%. We also evaluated the idea-matching capability of BERTScore (Zhang\* et al., 2020), considering that both ideas will be semantically similar. We found that BERTScore performed better than the entailment technique, resulting in an accuracy of 75.4%. We also tried GPT by prompting it with various questions and found that it resulted in 91.8% accuracy when prompted with a specific question prompt below:-

**Prompt:** Your task is to examine whether a particular idea is incorporated within a set of ideas and to what degree.

**Collection of ideas:** {API-FRIs}

**Single idea:** {A generated Idea}

Is the single idea contained within the collection of ideas?

If yes, quantify its degree of presence or relevance of the single idea in the collection of ideas on a scale from 0 to 1.

We found that GPT performs better than the existing NLI (Natural Language Inference) and similarity measure such as BERTScore. Therefore, we

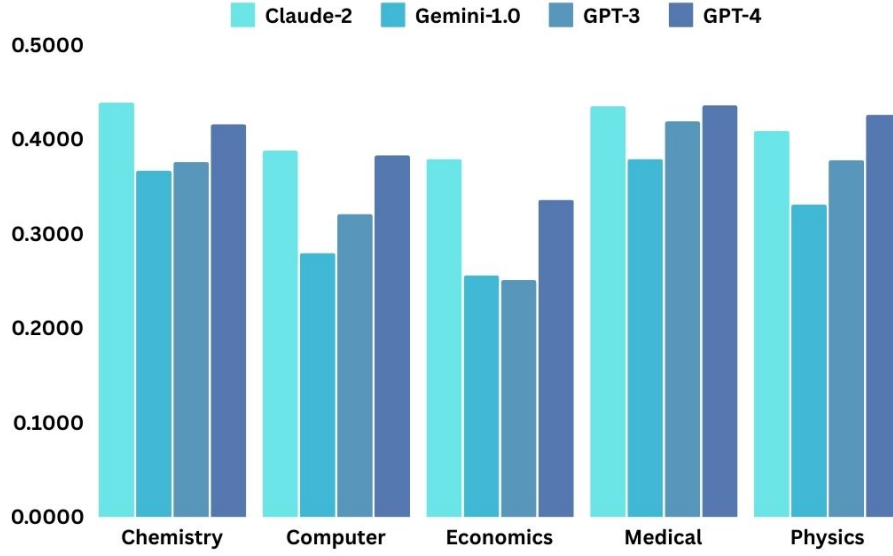


Figure 5: IAScore for each domain and model; a higher value indicates better alignment with the author.

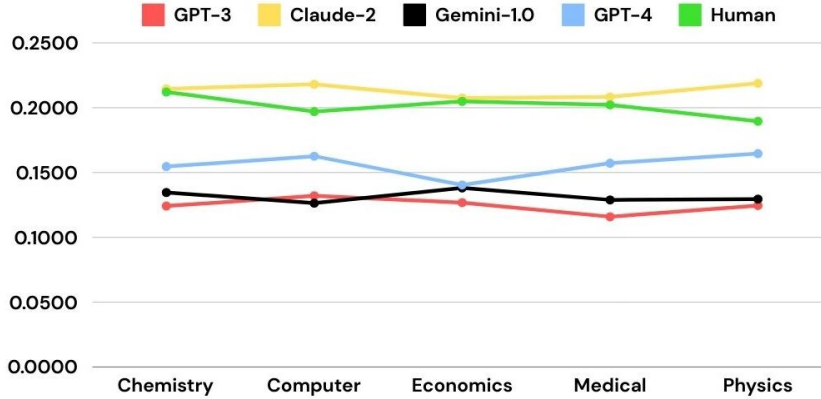


Figure 6: Idea distinctness index analysis; Here human is the authors of the paper

chose GPT for this task<sup>3</sup>.

### 4.3 Idea Distinctness Index

Distinct-N (Li et al., 2015), is a metric that measures the diversity of a sentence. It focuses on the number of distinct n-grams of a sentence and thus penalizes sentences with lots of repeated words. However comparing two ideas need semantic comparisons rather than just syntactic differences. So, we introduce a method to semantically evaluate the distinctness of the generated ideas. This method

<sup>3</sup>We used the OpenAI model GPT-3.5-turbo-0125 using OpenAI API

in particular leverages semantic embedding to capture the essence of each idea and computes their distinctness based on semantic similarity measures.

Given a set of generated ideas  $I = \{id_1, id_2, \dots, id_n\}$ , representing individual ideas, we first encode each idea into a high-dimensional vector space using a pre-trained BERT model (Devlin et al., 2019)<sup>4</sup>  $\text{BERT} : id_i \mapsto \mathbf{v}_i$ , where  $\mathbf{v}_i \in \mathbb{R}^d$  is the embedding of idea  $id_i$  and  $d$  is the dimensionality of the embedding space.

To quantify the distinctness between pairs of ideas, we compute the cosine similarity between

<sup>4</sup>bert-base-uncased

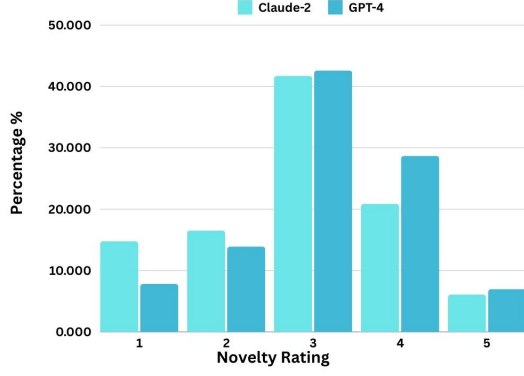


Figure 7: Novelty human evaluation for Computer Science domain

their embeddings,  $\text{sim}(\mathbf{v}_i, \mathbf{v}_j) = \frac{\mathbf{v}_i \cdot \mathbf{v}_j}{\|\mathbf{v}_i\| \|\mathbf{v}_j\|}$ , for each pair of ideas  $(id_i, id_j)$  in  $I$ . The distinctness  $D_{ij}$  between two ideas  $i$  and  $j$  is then inversely related to their similarity:  $D_{ij} = 1 - \text{sim}(\mathbf{v}_i, \mathbf{v}_j)$ .

The overall distinctness of the set  $I$  is calculated as the mean of all pairwise distinctness scores:

$$D_I = \frac{1}{n(n-1)} \sum_{i=1}^n \sum_{j=1, j \neq i}^n D_{ij} \quad (3)$$

This measure provides a single scalar value  $D_I$  that quantifies the average diverseness of ideas within a corpus of ideas, with higher values indicating a greater degree of diverseness among the ideas.

Subsequently, we aggregated the distinctness scores across all ideas in each paper to compute the mean distinctness for that paper. Let  $P = \{p_1, p_2, \dots, p_m\}$  represent the set of papers in a domain, where  $m$  is the number of papers in the domain. Finally, for a comprehensive assessment of model performance within a domain, we averaged the mean distinctness scores of all papers generated by model  $M$  as follows:

$$D_{\text{domain}, M} = \frac{1}{m} \sum_{p=1}^m D_{I_{pM}} \quad (4)$$

The resultant metric,  $D_{\text{domain}, M}$ , represents the average idea distinctness for model  $M$  in a given domain, indicating the model’s ability to generate diverse ideas.

#### 4.4 Human Evaluation

The evaluation of generated future ideas necessitates familiarity with both previous works related to

the subject and the work being evaluated. Specifically, the evaluator must be an expert in the domain and topic. Given the complexity of human evaluation, we approached authors (as the authors have the knowledge of their paper and they also have knowledge of the literature) who have published papers in reputable venues, possess over 5 years of experience in scientific publishing, and have authored more than 5 scientific papers. We collected their accepted papers and followed the dataset preparation as we discussed in Section 3 and generated FRIs. We modify the prompt slightly to specifically generate only the top five results (see Appendix B). We selected the outputs from Claude and GPT-4<sup>5</sup> models due to their better IAScore and Idea Distinction index. We adopt this approach to avoid author exhaustion and to get an accurate evaluation. We ask the following questions from each human evaluator:-

- Q1: Is the idea relevant with the research topic of the paper. (Relevant/Not relevant)
- Q2: Assess the originality/novelty of the research idea (5 scale)
- Q3: Review the research idea for factual correctness and feasibility. Is the idea impractical or too vague to be actionable? (Not Possible/Possible)

For Q2, we used Best-Worst Scaling (Louviere et al., 2015) on a 5-point scale. More details about the human evaluation are mentioned in the Appendix B.

## 5 Results and Discussion

### 5.1 Alignment Results

Figure 5 provides a comparative overview of the IAScore for four language models<sup>6</sup> Claude-2, Gemini-1.0, GPT-3, and GPT-4 across five academic domains: Chemistry, Computer Science, Economics, Medical, and Physics.

In the Chemistry and Economics domains, Claude has the highest IAScore, indicating strong alignment with the authors’ future research ideas. Claude and GPT-4 have almost similar values for the Computer, Medical, and Physics domains (with GPT-4 slightly higher). GPT-3 and Gemini have

<sup>5</sup>We used gpt-4-turbo using OpenAI API for the generation

<sup>6</sup>We set maximum token length to 512, and temperature=0 for each models

lower scores than both GPT-4 and Claude in every domain. GPT-3 has almost the same score as Gemini in the Chemistry and Economics domains. However, it scores higher than Gemini in the Computer, Medical, and Physics domains. The results underscore the advancements in language model capabilities, with each model showcasing domain-specific strengths in idea generation. This alignment of LLMs shows that large language models are able to generate relevant and novel ideas to some extent.

## 5.2 Distinctness Results

We show the comparative evaluation of idea distinctness scores in Figure 6. The line graph depicts the variation of distinctness between the generated ideas and the human-written ideas (AP-FRIs). GPT-3 shows the least distinctness among the generated ideas, except in the Computer domain, where it is slightly more distinct than Gemini. As shown in the graph, the distinctness of Gemini is also quite low; however, it is slightly better than GPT-3, except in the Computer domain.

The generated ideas of GPT-4 are more distinct than those of Gemini and GPT-3 (except for economics, whereas the distinctness of GPT-4 is the same as Gemini). However, it is lower than both Claude and Human. The Idea Distinctness Index of the generated ideas from Claude are almost the same as those of humans for Chemistry, Economics, and Medical domains. However, they are higher than even human scores in the Computer and Physics domains, which shows that it generates very distinct FRIs.

## 5.3 Human Evaluation Results

We conducted a human evaluation on 460 generated ideas for 46 papers in the computer science domain. To validate the quality of human annotation, we measure the inter-annotator agreement ratio where 20% of the generated ideas are evaluated by two different authors of the same paper. We measured Cohen’s kappa coefficient (Cohen, 1960), which was 0.83, thereby confirming the high quality of the generated research ideas.

**Novelty:** Figure 7 displays the results of the human evaluation. We observed that Claude generates 14.78% of non-novel and 16.52% generic FRIs, 41.73% moderately novel, 20.86% very novel, and 16.52% extremely novel FRIs. GPT generates 7.83% not-novel, 13.91% generic, 42.61% moderately novel, 28.70% very novel, and 6.96% ex-

tremely novel ideas. Claude generates more non-novel and generic ideas than GPT-4, while GPT-4 produces more very novel ideas and nearly the same number of excellent ideas. This demonstrates that although large language models also generate generic or already explored ideas, they are capable of producing novel ideas that have either not been explored or have been minimally explored.

**Relevance and Feasibility:** After human evaluation, we found that that 76.67% of the ideas generated by Claude and 93.34% by GPT-4 are relevant. Furthermore, 83.34% of Claude’s generated ideas and 96.64% of GPT-4’s ideas were judged to be practically feasible and factually correct. These results highlight that Claude and GPT-4 can generate relevant and feasible research ideas. However, the reason Claude generates more impractical and irrelevant research ideas may be that Claude attempts to generate more distinct research ideas than GPT-4, as we evaluated and discussed in Section 5.2.

## 6 Conclusion and Future Work

In conclusion, we present the first attempt to evaluate the potential of LLMs in generating future research ideas across five domains: Computer Science, Economics, Chemistry, Physics, and Medicine. Our results and analysis show that LLMs possess domain-specific strengths in idea generation. Furthermore, the results from the Idea Distinctness Index indicate that LLMs, such as Claude and GPT-4, generate distinct research ideas than Gemini and GPT 3.5. GPT-4 and Claude aligns better with authors written future research ideas than Gemini and GPT-4. The alignment of LLMs with the authors of generated ideas, and our human evaluations on relevance, novelty, and feasibility, reveal that although LLMs often produce non-novel and generic ideas, they have the potential to generate relevant and novel and diverse ideas to a significant extent. We hope that the findings and experiments of this work will unlock the potential of LLMs in idea generation and will open a new topic of accelerating scientific research by automatically generating ideas.

In the future, we will explore how integration of knowledge from multiple papers can help LLMs boost the novelty of the generated ideas and prevent the generation of generic and existing ideas.



## Limitations

### 6.1 Limitations of Data Collection

We extracted papers using the Semantic Scholar Academic Graph API from January 2023 to February 2024. The number of papers available is limited by the scope of our data extraction from the Semantic Scholar Academic Graph. We excluded papers that are not in English, as well as those whose abstracts could not be correctly parsed from the PDFs. Not all of these papers include sections on future work; therefore, we annotated only those that contained sections outlining future research directions. So due to such limitations, we collected 100 papers from each domain for analysis.

### 6.2 Memorization

([Carlini et al., 2022](#)) highlight that LLMs are prone to memorizing portions of their training data, a significant concern in the evaluation of contemporary LLMs. Despite this, the data used for pre-training and post-training includes "a small amount" of more recent data. Therefore, we gathered recent papers from 2023 and 2024. By focusing our evaluation on papers published in these years, the likelihood of test papers appearing in the pre-training corpora for the models is substantially reduced. In addition, we conducted a manual review of these papers to assess memorization. This involved asking various questions related to the papers, such as their titles, publishing venues, author names, etc., to see if the models could supply the missing information. Our findings showed no evidence of such memorization occurring.

## Ethics Statement

We have utilized the open source dataset for our work. Our aim for this work is to assess the potential of language models in generating ideas. Our Institutional Review Board (IRB) evaluated and approved this study. We do not encourage the use of LLMs to generate AI generated research papers (by generating new ideas) or misuse it for harmful idea generation. Notably, it is a common challenge among existing LLMs with strong creative and reasoning abilities.

## References

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Gemini Team Google Rohan Anil, Sebastian Borgeaud, Yonghui Wu, Jean-Baptiste Alayrac, Jiahui Yu, Radu

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## A Dataset Annotation Guidelines

Recognizing future research idea in a paper involves analyzing the portion of text containing directions for future research. The following steps can be followed:

**Step 1:** Begin by reading the Title and Abstract of the paper to gain an understanding of its subject matter. It is important to read these sections multiple times to grasp the paper’s main points, such as

its motivation, contributions, and other relevant aspects. If necessary, refer to the paper itself or read related material to enhance your understanding.

**Step 2:** Identify Key Sections for Analysis Focus primarily on the Discussion and Conclusion sections of the paper, as these areas often contain explicit mentions of future research directions. Scan the Methodology section as well, as sometimes suggestions for improving future studies or addressing current study limitations are mentioned here.

**Step 3:** Distinguish Future Research Ideas from General Statements: Differentiate explicit future research suggestions from general discussion. Future research directions usually involve specific recommendations, plans, or identified gaps that require further exploration. These are often phrased using terms like "future studies should," "further research is needed," or "additional work will." Avoid confusing these with broader statements of potential relevance or applicability, which do not provide direct guidance on future work.

We offer multiple examples of papers with its future research ideas to assist and direct the annotators. We found a few text which looks like future work but is on contrary the motivation of the work. As an example, consider the following: *"The goal of this work was to direct attention to emerging and novel research involving "magnetogel nanohybrid materials" that might be relevant in future applications for the treatment of wastewater, as well as in other fields.*

The second example is: *"Our data could be useful for designing high-quality trials in the future to define the exact role of hemoabsorption in ARDS."* Here, how novel research involving magnetogel nanohybrid material will help in future application is written.

Also another example is: *"The goal of this work was to direct attention to emerging and novel research involving magnetogel nanohybrid materials that might be relevant in future applications for the treatment of wastewater, as well as in other fields."* This is the application in future, and not the future work.

**Step 4:** Separate Future Research from Limitations: Carefully examine any limitations mentioned in the paper to determine if they are explicitly linked to future research. Only consider a limitation as future work if the authors clearly indicate a direct intention to address it in subsequent studies. This helps avoid assuming that all limitations



naturally lead to future research directions.

There is also very thin line between limitation and future work, where a limitation can or cannot be a future work. There were few cases where limitations were mentioned "*One limitation of this paper is the absence of a coordinated attention structure to capture cross-channel information.*". As limitations can or cannot be a future work, we only take those limitations which is explicitly mentioned by the author to be a future work. Hence, we only considered the explicit mention of the future work by the author in their paper.

## B Human Annotation

We prepared a Google Form for each paper and provided the links to the annotators. We also specified instructions for them at the beginning of the form. We have added an example of the form for a paper in Figure 8, Figure 9, and Figure 10.

Here is the little modified form for human evaluation that generates only top 5 research ideas:-

**System:** You are a research scientist.

**User:** Imagine you are a research scientist. After reading the following paper, brainstorm to generate potential top 5 future research ideas:

[paper text]

Potential top 5 future research ideas from the paper in bullet points are:

Here, '[paper text]' contains the full content of the paper after removal of future work sections.

### B.1 Output Examples

Our LLM generated future research output can be found in Table 1, Table2, 3, Table 4, Table5.

Model	Response
Claude-2	<ol style="list-style-type: none"> <li>1. Investigate the antioxidant and oil-stabilizing effects of peptides from the soy protein hydrolysates. The paper showed the pepsin hydrolysate had better activity, likely due to its peptide composition. Specific peptides could be isolated and tested for their antioxidant effects in oil.</li> <li>2. Study the effects of the soy protein hydrolysates on the quality and shelf life of the fried food. This paper focused only on the oil stability. It would be useful to also assess how the hydrolysates affect lipid oxidation, texture, color etc. in the fried banana chips.</li> <li>3. Evaluate the antioxidant effects of the soy protein hydrolysates in other vegetable oils such as sunflower, canola, corn oil etc. during frying. Since oil composition affects stability, the effects may differ in other oils high in polyunsaturated fatty acids.</li> <li>4. Examine the impacts of frying parameters such as temperature, food type, frying time etc. on the efficacy of the soy protein hydrolysates. Varying these factors could provide insights into the optimal conditions for the hydrolysates to exert antioxidant effects.</li> <li>5. Study the sensory qualities of fried foods with the soy protein hydrolysate additions. An analysis of taste, aroma, color acceptability by consumers would provide important information for potential commercial use of the hydrolysates as natural antioxidants.</li> </ol>
Gemini-1.0	<ol style="list-style-type: none"> <li>1. Investigate the effects of different hydrolysis conditions on the antioxidant activity of soy protein hydrolysates. This could involve varying the enzyme concentration, temperature, and pH of the hydrolysis reaction.</li> <li>2. Identify the specific peptides in the pepsin hydrolysate that are responsible for its antioxidant activity. This could be done using mass spectrometry and other analytical techniques.</li> <li>3. Test the antioxidant activity of soy protein hydrolysates in other food systems. This could include using them to stabilize other types of oils or to protect food products from oxidation.</li> <li>4. Investigate the potential health benefits of soy protein hydrolysates. This could include studying their effects on oxidative stress, inflammation, and other chronic diseases.</li> <li>5. Develop new food products that incorporate soy protein hydrolysates as antioxidants. This could include developing new types of oil-based products, such as salad dressings and mayonnaise, as well as new types of food products, such as fortified cereals and snacks.</li> </ol>
GPT-4	<ol style="list-style-type: none"> <li>1. Exploration of Peptide Sequences and Amino Acid Compositions for Enhanced Oxidative Stability: Investigate the specific peptide sequences and amino acid compositions derived from different proteases besides trypsin and pepsin to understand their contribution to the oxidative stability of oils. This could involve using a variety of proteolytic enzymes to hydrolyze soy proteins and other plant-based proteins, followed by an analysis of their peptide profiles and antioxidant activities in oil and food systems.</li> <li>2. Application of Soy Protein Hydrolysates in Various Food Systems: Extend the application of soy protein hydrolysates to other food systems beyond frying oils, such as baked goods, dairy products, and meat alternatives, to assess their potential in improving oxidative stability, shelf-life, and nutritional profiles. This could involve comparative studies of hydrolysate effectiveness in different food matrices and processing conditions.</li> <li>3. Mechanistic Studies on the Antioxidant Action of Hydrolysates: Conduct in-depth mechanistic studies to understand how soy protein hydrolysates interact with lipid molecules and radical species at the molecular level. This could involve computational modeling and in vitro experiments to explore the dynamics of peptide-lipid interactions and the role of specific amino acids in scavenging free radicals and metal ions.</li> <li>4. Development of Environmentally Friendly and Health-focused Antioxidants: Given the interest in replacing synthetic antioxidants with natural alternatives, further research could focus on optimizing the production, purification, and application of soy protein hydrolysates as natural antioxidants. This could include exploring sustainable extraction methods, evaluating the safety and efficacy of hydrolysates in food systems, and assessing consumer acceptance.</li> <li>5. Comparative Analysis of Hydrolysates from Different Oilseed Proteins: Since the study focused on soy proteins, future research could explore the antioxidative potential of hydrolysates derived from other oilseed proteins, such as sunflower, canola, or hemp. This would involve a comparative analysis of their antioxidative activities, peptide compositions, and effectiveness in enhancing the oxidative stability of different types of oils and fats used in food processing and storage.</li> </ol>
GPT-3.5	<ol style="list-style-type: none"> <li>1. Investigate the impact of different proteases on the antioxidant properties of soy protein hydrolysates in stabilizing oils during frying cycles.</li> <li>2. Study the influence of peptide size and amino acid composition on the oxidative stability of oils when treated with protein hydrolysates.</li> <li>3. Explore the potential of specific amino acids, such as tryptophan, histidine, and methionine, in enhancing the antioxidant activity of hydrolyzed soy proteins in frying oils.</li> <li>4. Examine the effects of pH adjustments during enzymatic hydrolysis on the antioxidant performance of soy protein hydrolysates.</li> <li>5. Evaluate the long-term stability and effectiveness of pepsin and trypsin hydrolyzed soy protein in preventing oil oxidation in different frying conditions over extended periods.</li> </ol>

Table 1: Example of generated response for chemistry domain. Paper: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC10528889/pdf/foods-12-03498.pdf>

Model	Response
Claude-2	<ol style="list-style-type: none"> <li>1. Investigate other machine learning models like recurrent neural networks for multi-round transfer learning in lung cancer detection. The paper focused on convolutional neural networks, but RNNs may capture temporal patterns in medical imaging data.</li> <li>2. Explore different loss functions and regularization techniques to further minimize negative transfer. The paper proposed custom loss functions, but there is room to experiment with other formulations.</li> <li>3. Evaluate the multi-round transfer learning approach on diverse medical imaging datasets beyond lung cancer, such as brain MRI scans. This could demonstrate the generalizability of the method.</li> <li>4. Develop more advanced generative models like CycleGAN to synthesize intermediate domains between source and target. This could create higher quality bridging distributions.</li> <li>5. Perform multi-round transfer learning in a low-data regime by leveraging data augmentation and semi-supervised learning techniques. This could show the approach is viable when target data is extremely scarce.</li> </ol>
Gemini-1.0	<ol style="list-style-type: none"> <li>1. Investigate the impact of different prioritization algorithms on the performance of MTL-MGAN. The paper uses a hybrid approach that merges modified 2D dynamic warping (M2DW) and Silhouette coefficient to prioritize source domains. Future research could explore other prioritization algorithms, such as mutual information, transfer entropy, or graph-based methods, to determine if they can further improve the performance of MTL-MGAN.</li> <li>2. Develop new loss functions to minimize negative transfer.** The paper proposes loss functions that are designed to minimize negative transfer in the aspects of domains, instances, and features. Future research could investigate other loss functions that can further reduce negative transfer, such as adversarial loss, maximum mean discrepancy, or knowledge distillation.</li> <li>3. Explore different architectures for the MGAN.** The paper uses a modified GAN (MGAN) to create two intermediate domains as bridges between the source domain and the target domain. Future research could explore other GAN architectures, such as deep convolutional GAN (DCGAN), Wasserstein GAN (WGAN), or StyleGAN, to determine if they can further improve the performance of MTL-MGAN.</li> <li>4. Apply MTL-MGAN to other medical diagnosis tasks.** The paper evaluates the performance of MTL-MGAN on 10 benchmark datasets related to lung cancer detection. Future research could apply MTL-MGAN to other medical diagnosis tasks, such as breast cancer detection, skin cancer detection, or diabetic retinopathy detection, to determine its generalizability.</li> <li>5. Investigate the use of MTL-MGAN for other types of data.** The paper applies MTL-MGAN to medical images. Future research could investigate the use of MTL-MGAN for other types of data, such as text, audio, or time series data, to determine its applicability to a wider range of problems.</li> </ol>
GPT-4	<ol style="list-style-type: none"> <li>1. Extension to Other Forms of Cancer and Diseases: Investigating the applicability and effectiveness of the MTL-MGAN approach for the detection and diagnosis of other types of cancer, such as breast, prostate, or skin cancer, and potentially other diseases that have distinct imaging characteristics. This would involve adapting the transfer learning and MGAN components to handle different imaging modalities and disease signatures.</li> <li>2. Real-Time Diagnostic Systems Development: Developing real-time, AI-driven diagnostic systems using the MTL-MGAN framework that can be integrated into clinical workflows. This could significantly reduce diagnosis times and improve patient outcomes by providing immediate, accurate diagnostic information during patient examinations or surgeries.</li> <li>3. Integration with Electronic Nose Technology for Early Detection: Combining the MTL-MGAN approach with emerging technologies like electronic nose devices, which can detect volatile organic compounds associated with lung cancer from breath samples. Research could focus on creating a hybrid detection system that leverages imaging data and breath analysis for earlier detection of lung cancer.</li> <li>4. Exploration of Unsupervised and Semi-supervised Learning Techniques: Given the challenge of obtaining large, annotated medical imaging datasets, exploring unsupervised and semi-supervised learning methods within the MTL-MGAN framework could be valuable. This might involve developing algorithms that can learn from unlabelled data or very small labelled datasets, potentially increasing the model's applicability in low-resource settings.</li> <li>5. Improvement of Transfer Learning Techniques for Heterogeneous Data Sources: Investigating advanced transfer learning techniques that can effectively handle more significant variations between source and target domains, particularly when dealing with heterogeneous data sources. This research could include the development of novel algorithms for better domain adaptation and intermediate domain creation to facilitate the transfer of knowledge across diverse medical datasets and imaging modalities.</li> </ol>
GPT-3.5	<ol style="list-style-type: none"> <li>1. Exploring Multiround Transfer Learning in Other Healthcare Applications: Investigate the applicability of multiround transfer learning in various healthcare domains beyond lung cancer detection, such as cardiovascular disease diagnosis, diabetic retinopathy detection, or infectious disease prediction.</li> <li>2. Advanced Loss Function Design for Negative Transfer Avoidance: Develop novel loss functions to further minimize the impact of negative transfer in transfer learning models by incorporating more sophisticated domain, instance, and feature-based penalties.</li> <li>3. Enhanced Generative Adversarial Networks for Bridging Domain Gaps: Investigate the development of advanced generative adversarial networks that can create more effective intermediate domains to bridge the gap between diverse source and target domains in transfer learning scenarios.</li> <li>4. Automated Prioritization Algorithm for Source Domains: Develop an automated algorithm that can dynamically prioritize source domains based on their similarities to the target domain, thereby enhancing the efficiency and accuracy of the transfer learning process.</li> <li>5. Evaluation of MTL-MGAN in Real-World Clinical Settings: Conduct a comprehensive evaluation of the proposed multiround transfer learning and modified generative adversarial network algorithm in real-world clinical settings to assess its performance, scalability, and generalizability in diverse healthcare environments.</li> </ol>

Table 2: Example of generated response for computer domain. Paper: <https://www.hindawi.com/journals/ijis/2023/6376275/>

Model	Response
Claude-2	<ol style="list-style-type: none"> <li>1. Conduct qualitative research through interviews or focus groups to understand the specific barriers and facilitators to COVID-19 vaccination uptake in different socioeconomic groups within a country. This could provide insights to inform targeted interventions.</li> <li>2. Examine the impact of COVID-19 vaccination policies and public health messaging on vaccination uptake across socioeconomic groups. For example, study how vaccination requirements or incentives affect different populations.</li> <li>3. Investigate strategies to improve global collaboration and coordination in vaccine distribution and access. Research equitable pricing models, technology transfers, and policies to balance local needs with global supply.</li> <li>4. Develop mathematical models to forecast COVID-19 vaccination coverage over time under different policy scenarios. Simulate the impact on infection rates, healthcare burden, and emergence of new variants.</li> <li>5. Conduct intervention studies to test the effectiveness of community-based participatory approaches in increasing COVID-19 vaccine acceptance among disadvantaged populations. Evaluate culturally-tailored messaging and outreach.</li> </ol>
Gemini-1.0	<ol style="list-style-type: none"> <li>1. Investigate the impact of socioeconomic factors on vaccine hesitancy and refusal. This could be done through qualitative studies to understand the underlying reasons for vaccine hesitancy and refusal, and quantitative studies to examine the association between socioeconomic factors and vaccine uptake.</li> <li>2. Examine the role of social networks and community engagement in promoting vaccination. This could involve studying the influence of social networks on vaccination decisions, and exploring the effectiveness of community-based interventions in increasing vaccination rates.</li> <li>3. Assess the effectiveness of different strategies for addressing vaccine inequality. This could include evaluating the impact of financial incentives, targeted outreach programs, and mobile vaccination clinics on vaccination rates among underserved populations.</li> <li>4. Develop and evaluate interventions to improve the equitable distribution of vaccines. ** This could involve working with governments and international organizations to develop policies and programs that ensure that vaccines are distributed fairly and equitably to all populations.</li> <li>5. Conduct longitudinal studies to track the long-term impact of socioeconomic factors on vaccination rates. This could help to identify trends and patterns in vaccination uptake over time, and inform the development of policies and interventions to address persistent disparities.</li> </ol>
GPT-4	<ol style="list-style-type: none"> <li>1. Impact of Vaccine Acceptance and Hesitancy on Socioeconomic Disparities: Investigate how vaccine acceptance and hesitancy, influenced by cultural, historical, and information dissemination factors, contribute to the observed socioeconomic disparities in vaccination rates. This research could further dissect how misinformation and trust in healthcare systems intersect with socioeconomic status to affect vaccine uptake.</li> <li>2. Longitudinal Studies on the Evolution of Vaccine Inequality: Conduct longitudinal studies to track changes in vaccine inequality over time, particularly as new variants emerge and booster vaccinations become more common. This research would provide insights into how socioeconomic disparities in vaccination rates evolve with the pandemic's progression and the introduction of new vaccine technologies or strategies.</li> <li>3. Role of International and Public-Private Partnerships in Addressing Vaccine Inequity: Explore the effectiveness of international collaborations and public-private partnerships in improving vaccine distribution equity. This study could examine case studies of successful collaborations and identify best practices for leveraging these partnerships to reduce disparities in vaccine access.</li> <li>4. Comparative Analysis of Vaccination Strategies and Socioeconomic Impact: Compare the vaccination strategies of different countries or regions with similar socioeconomic profiles to identify what strategies are most effective in mitigating inequalities in vaccination rates. This research could include the role of mobile vaccination units, community engagement initiatives, and targeted communication strategies in improving vaccine uptake among disadvantaged populations.</li> <li>5. Impact of Socioeconomic Factors on Post-Vaccination Outcomes: Investigate how socioeconomic factors influence outcomes after vaccination, such as breakthrough infection rates, severity of breakthrough cases, and long-term immunity. This research could help identify whether socioeconomic disparities persist even after vaccination and how these disparities can be addressed through booster doses, tailored public health strategies, and improved healthcare access.</li> </ol>
GPT-3.5	<ol style="list-style-type: none"> <li>1. Investigating the Impact of Socioeconomic Factors on Access and Uptake of COVID-19 Booster Vaccines: A study could focus on analyzing how socioeconomic indicators, such as GDP, human development index, and corruption perception index, influence access to and acceptance of booster doses of COVID-19 vaccines. This research could help identify disparities in vaccine distribution and uptake among different population groups.</li> <li>2. Comparative Analysis of Inequality in COVID-19 Vaccination Across Continents: Future research could compare and contrast the levels of inequality in COVID-19 vaccination across different continents, especially focusing on the factors contributing to variations in vaccination rates and disparities in vaccine distribution. This comparative analysis could provide valuable insights for designing targeted interventions to address global vaccine inequality.</li> <li>3. Longitudinal Study on the Impact of Socioeconomic Factors on COVID-19 Vaccination: A longitudinal study could be conducted to track changes in vaccination rates over time and assess how socioeconomic factors impact the progress of COVID-19 vaccination campaigns. By analyzing trends and patterns, researchers could gain a deeper understanding of the evolving dynamics of vaccine distribution and uptake.</li> <li>4. Exploring Public Perception and Vaccine Hesitancy Among Different Socioeconomic Groups: Future research could delve into public perceptions and attitudes towards COVID-19 vaccination among diverse socioeconomic groups, investigating factors such as education level, income, and access to healthcare. This study could help identify barriers to vaccination and inform targeted communication strategies to address vaccine hesitancy.</li> <li>5. Evaluation of Intervention Strategies to Reduce Inequality in COVID-19 Vaccination: Researchers could conduct a study to evaluate the effectiveness of various intervention strategies aimed at reducing inequality in COVID-19 vaccination. By implementing targeted initiatives and measuring their impact on vaccine distribution and uptake, this research could provide evidence-based recommendations for policy interventions to promote equitable access to vaccines.</li> </ol>

Table 3: Example of generated response for economics domain. Paper: <https://www.mdpi.com/2076-393X/11/7/1147>

Model	Response
Claude-2	<ol style="list-style-type: none"> <li>1. Further optimize the minimal active H2 relaxin analog B9-31 through additional amino acid substitutions and modifications to improve binding affinity and activity.</li> <li>2. Explore alternative methods to induce helicity and stabilize the minimal H2 relaxin analogs, such as cyclization or incorporation of other helix-inducing amino acids.</li> <li>3. Develop longer-acting analogs by modifying B9-31 with polyethylene glycol, albumin binding moieties, or other half-life extending strategies.</li> <li>4. Test the in vivo efficacy of optimized minimal analogs in animal models of fibrosis, heart failure, and other conditions to assess therapeutic potential.</li> <li>5. Design biased analogs to selectively activate certain signaling pathways downstream of RXFP1 to optimize beneficial effects and reduce side effects.</li> </ol>
Gemini-1.0	<ol style="list-style-type: none"> <li>1. Investigate the potential benefits of using ECMO in combination with other therapies for severe respiratory failure. For example, ECMO could be used in combination with lung transplantation, mechanical ventilation, or extracorporeal membrane oxygenation (ECMO). This could help to improve patient outcomes by providing more comprehensive support for the lungs and other organs.</li> <li>2. Develop new strategies for managing complications associated with ECMO. Common complications of ECMO include bleeding, infection, and thrombosis. New strategies for managing these complications could help to improve patient safety and outcomes.</li> <li>3. Study the long-term effects of ECMO on patients. There is limited data on the long-term effects of ECMO on patients. Studies are needed to investigate the potential risks and benefits of ECMO in the long term.</li> <li>4. Develop new technologies to improve the performance of ECMO. New technologies could help to make ECMO more effective, safer, and easier to use. For example, new oxygenators could be developed that are more efficient and less likely to cause complications.</li> <li>5. Conduct research to better understand the mechanisms of respiratory failure. This could help to identify new targets for therapy and improve the overall management of respiratory failure.</li> </ol>
GPT-4	<ol style="list-style-type: none"> <li>1. Comparative Effectiveness of ECMO Configurations: Conduct randomized controlled trials comparing different ECMO cannulation strategies (e.g., dual-lumen cannula via the right internal jugular vein vs. traditional multi-cannula approaches) to assess their impact on patient outcomes, including survival rates, complication rates, and the ability to mobilize and rehabilitate during ECMO support. This research could also explore the use of image guidance for cannula placement and its impact on patient outcomes.</li> <li>2. ECMO in the Management of COVID-19-Induced ARDS: Given the limited randomized controlled trial data on the efficacy of ECMO for COVID-19-associated respiratory failure, further research is needed to define the role of ECMO in this context. Future studies could focus on identifying which patients benefit most from ECMO, the timing of initiation, and how ECMO is best integrated with other supportive therapies (e.g., antivirals, anti-inflammatory agents).</li> <li>3. Long-term Outcomes and Quality of Life Post-ECMO: Investigate the long-term outcomes of patients who have undergone ECMO support for severe respiratory failure, including physical, cognitive, and psychological recovery. This research could also assess the quality-adjusted life years (QALYs) gained through ECMO support and identify factors that predict better long-term outcomes.</li> <li>4. Optimization of Anticoagulation Strategies: Develop and test new anticoagulation protocols or agents that minimize the risks of bleeding and thrombotic complications in ECMO patients. This could include comparative studies of direct thrombin inhibitors versus unfractionated heparin, as well as the exploration of novel anticoagulants and biocompatible circuit coatings that may reduce the need for systemic anticoagulation.</li> <li>5. Integration of ECMO with Rehabilitation: Explore the feasibility, safety, and benefits of early rehabilitation protocols in ECMO-supported patients, including those on "awake ECMO." This research could assess the impact of rehabilitation on outcomes such as muscle strength, functional status at discharge, duration of hospital stay, and overall mortality. Studies could also investigate the best practices for physical therapy, nutritional support, and psychological support during ECMO.</li> </ol>
GPT-3.5	<ol style="list-style-type: none"> <li>1. Investigating the long-term outcomes and quality of life of patients who have undergone VV-ECMO support for respiratory failure, including factors influencing long-term recovery and rehabilitation post-ECMO.</li> <li>2. Exploring new technologies and techniques to improve the efficiency and safety of VV-ECMO, such as developing novel anticoagulation strategies or improving cannulation methods to reduce complications and enhance patient outcomes.</li> <li>3. Conducting comparative studies to evaluate the effectiveness of different ventilation strategies in conjunction with VV-ECMO support, aiming to optimize patient management and potentially improve survival rates in severe respiratory failure cases.</li> <li>4. Examining the impact of ECMO utilization during disease outbreaks, such as COVID-19, on healthcare resource allocation, patient outcomes, and healthcare system preparedness, to inform future pandemic response strategies.</li> <li>5. Investigating the role of personalized medicine in VV-ECMO patient selection and management, including the potential use of biomarkers, genetic profiling, or other advanced techniques to tailor ECMO therapy to individual patient characteristics and optimize outcomes.</li> </ol>

Table 4: Example of generated response for medical domain. Paper: <https://link.springer.com/article/10.1007/s41030-023-00214-2>



Model	Response
Claude-2	<ol style="list-style-type: none"> <li>1. Investigate the antioxidant and oil-stabilizing effects of specific peptide sequences identified in the pepsin and trypsin hydrolysates. The paper showed the pepsin hydrolysate had higher activity but did not identify the key active peptides. Bioinformatics could be used to predict promising peptides.</li> <li>2. Test the soy protein hydrolysates in stabilizing other vegetable oils like canola, sunflower, etc. during deep frying. The effects may differ for oils with varying fatty acid compositions.</li> <li>3. Evaluate the sensory qualities of foods fried in palm olein stabilized with the soy protein hydrolysates compared to unstabilized oil. This could include assessing odor, taste, and textural changes.</li> <li>4. Study the effects of the hydrolysates on the nutritional quality of the palm olein during repeated deep frying cycles. Parameters could include vitamin E content, formation of trans fats, and retention of essential fatty acids.</li> <li>5. Investigate combining the soy protein hydrolysates with other antioxidants like rosemary extract or green tea to evaluate synergistic effects on stabilizing frying oils. Mixtures may provide greater protection than individual additives.</li> </ol>
Gemini-1.0	<ol style="list-style-type: none"> <li>1. Investigate the effect of different hydrolysis conditions on the antioxidant activity of soy protein hydrolysates. This could involve varying the enzyme concentration, temperature, pH, and hydrolysis time to determine the optimal conditions for producing hydrolysates with the highest antioxidant activity.</li> <li>2. Evaluate the antioxidant activity of soy protein hydrolysates in different food systems. This could involve adding hydrolysates to edible oils, fried foods, or other food products and measuring the oxidative stability of the products over time. This would help to determine the potential applications of soy protein hydrolysates as natural antioxidants in food products.</li> <li>3. Study the mechanism by which soy protein hydrolysates exert their antioxidant activity.** This could involve using in vitro and in vivo assays to investigate the ability of hydrolysates to scavenge free radicals, chelate metal ions, or inhibit lipid peroxidation. This information would help to elucidate the molecular mechanisms underlying the antioxidant activity of soy protein hydrolysates.</li> <li>4. Explore the potential health benefits of soy protein hydrolysates. This could involve conducting animal or human studies to investigate the effects of hydrolysates on oxidative stress, inflammation, and other health markers. This would help to determine the potential role of soy protein hydrolysates in promoting human health.</li> <li>5. Develop novel food products containing soy protein hydrolysates. This could involve incorporating hydrolysates into functional foods, such as beverages, snacks, or supplements, to improve their nutritional value and antioxidant content. This would help to increase the consumption of soy protein hydrolysates and promote their potential health benefits.</li> </ol>
GPT-4	<ol style="list-style-type: none"> <li>1. DComparative Analysis of Different Plant-Based Proteins on Oil Stability: Extending the research to other plant-based proteins, such as pea, hemp, and rice proteins, to compare their effectiveness in improving oil oxidative stability during frying cycles. This could provide a broader understanding of the potential of plant-based proteins in enhancing the nutritional and safety profiles of fried foods.</li> <li>2. Mechanism of Action at Molecular Level: Delving deeper into the molecular mechanisms behind the antioxidant activities of different hydrolysates. This could involve investigating the interaction between specific peptide sequences and oxidative compounds within the oil, using computational modeling and in vitro experiments to elucidate the pathways for their antioxidant actions.</li> <li>3. Impact of Hydrolysate Concentration and Frying Duration: Exploring the effect of varying concentrations of the most effective hydrolysates (e.g., from pepsin digestion) over extended frying cycles beyond 12, to determine the optimal concentration for maximum oxidative stability without affecting the sensory qualities of the oil and the fried product.</li> <li>4. Application in Industrial Frying Processes: Conducting pilot-scale studies in industrial settings to assess the feasibility and economic viability of incorporating hydrolysates into commercial frying oils. This research could include the development of guidelines for the use of hydrolysates in different types of frying operations and their impact on the quality and shelf-life of fried foods.</li> <li>5. Development of Natural Antioxidant Blends: Formulating blends of hydrolysates with other natural antioxidants (e.g., extracts from spices, herbs, or other food by-products rich in phenolic compounds) to synergistically enhance the oxidative stability of frying oils. This research could also investigate the sensory impact of these blends on fried products and consumer acceptance.</li> </ol>
GPT-3.5	<ol style="list-style-type: none"> <li>1. Investigating the impact of different proteases on the generation of antioxidant peptides from soy protein hydrolysates and their effectiveness in stabilizing oils during frying cycles.</li> <li>2. Exploring the structural features of peptides derived from soy protein hydrolysates to determine the key amino acids responsible for their antioxidant properties.</li> <li>3. Comparing the oxidative stability of different types of oils (e.g., sesame oil, canola oil, corn oil) when stabilized with soy protein hydrolysates under varied frying conditions.</li> <li>4. Examining the influence of hydrolyzed protein residues on the formation of secondary oxidation products in oils during frying cycles and their impact on food quality and safety.</li> <li>5. Utilizing bioinformatics tools to predict and select specific peptide sequences from soy protein hydrolysates that exhibit the highest antioxidant capacity and stability-enhancing properties in fried oils..</li> </ol>

Table 5: Example of generated response for physics domain. Paper: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC10528889/pdf/foods-12-03498.pdf>

Please read the research idea carefully and select the appropriate response.

We explain each questions below:-

*Q1: Is the idea relevant with the research topic of the paper. (Relevant/Not relevant)*

- **Not Relevant:** If the idea seems unrelated to the research topic of the paper.
- **Relevant:** If the idea aligns with the research topic of the paper.

*Q2: Assess the originality/novelty of the research idea. (Not Novel/Novel)*

- **Not Novel:** If you believe the idea is generic or already exists.
- **Slightly novel:** The idea is somewhat generic
- **Moderately novel:** This is a promising research direction but has already been explored by a significant number of researchers.
- **Very novel:** The idea represents a good research direction and has been pursued by only a few researchers.
- **Extremely novel:** The idea is highly innovative and, to my knowledge, has not been explored by anyone.

*Q3: Review the research idea for factual correctness and feasibility. Is the idea impractical or too vague to be actionable? (Not Possible/Possible)*

- **Not Possible:** If you feel that the idea is impractical/doesn't make any sense/ not realistic
- **Possible:** If you believe the idea is practical and realistic.

Figure 8: Instructions on Human Feedback form

**Future Work 4:** Add lexical constraints during decoding to prevent minimal word changes. This could enforce more semantic changes for style transfer.

**Future Work 5:** Evaluate the model on other styles like formality, gender, political slant etc. This could test the generalizability of the approach to other text style transfer tasks.

	Not novel	Slightly novel: The idea is somewhat generic	Moderately novel: This is a promising research direction but has already been explored by a significant number of researchers.	Very novel: The idea represents a good research direction and has been pursued by only a few researchers.	Extremely novel: The idea is highly innovative and, to my knowledge, has not been explored by anyone.
Future Work 1	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Future Work 2	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Future Work 3	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Future Work 4	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
Future Work 5	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>

Figure 9: Human Feedback form for novelty

	Relevant	Possible
Future Work 1	<input type="checkbox"/>	<input type="checkbox"/>
Future Work 2	<input type="checkbox"/>	<input type="checkbox"/>
Future Work 3	<input type="checkbox"/>	<input type="checkbox"/>
Future Work 4	<input type="checkbox"/>	<input type="checkbox"/>
Future Work 5	<input type="checkbox"/>	<input type="checkbox"/>

Figure 10: Human Feedback form for relevance, feasibility