

SciClaims: An End-to-End Generative System for Biomedical Claim Analysis

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Abstract

We present SciClaims, an interactive web-based system for end-to-end scientific claim analysis in the biomedical domain. Designed for high-stakes use cases such as systematic literature reviews and patent validation, SciClaims extracts claims from text, retrieves relevant evidence from PubMed, and verifies their veracity. The system features a user-friendly interface where users can input scientific text and view extracted claims, predictions, supporting or refuting evidence, and justifications in natural language. Unlike prior approaches, SciClaims seamlessly integrates the entire scientific claim analysis process using a single large language model, without requiring additional fine-tuning. SciClaims is optimized to run efficiently on a single GPU and is publicly available for live interaction¹.

1 Introduction

Systematic Literature Review (SLR) plays a critical role in biomedical research and the pharmaceutical industry, supporting clinical decisions, regulatory submissions, and R&D pipelines. A central task in SLR is the validation of scientific claims, ensuring that assertions made in scientific texts are supported by prior peer-reviewed research. However, this task is labor-intensive, prone to errors, and increasingly difficult to scale as the number of scientific publications grows.

We present SciClaims, a fully automated system that addresses this challenge by providing an end-to-end scientific claim analysis pipeline in an interactive, user-friendly interface. The system extracts factual claims from scientific texts, retrieves evidence from a curated biomedical corpus, and verifies the validity of each claim using large language models (LLMs). To improve transparency,

SciClaims also offers natural language rationales and highlights key supporting or refuting evidence.

SciClaims is optimized for real-world deployment and operates efficiently on a single GPU, supports high-throughput processing, and handles documents of up to 10,000 characters. It is accessible through a web-based interface, allowing users to analyze both preloaded and custom input texts.

In this demonstration, we showcase SciClaims as a useful tool to enable users to validate scientific claims in real time, facilitating trustworthy knowledge discovery in high-stakes domains like biomedicine and pharmaceuticals. A how-to video² and all the code³ used in this project have been published and made publicly available.

2 Related Work

The task of analyzing scientific claims from real-world texts based on background knowledge consists of three primary components: claim extraction, evidence retrieval from a document corpus, and verifying or fact-checking the claims against the evidence (Eldifrawi et al., 2024; Vladika and Matthes, 2023). However, solving this pipeline end-to-end across all three stages remains an open challenge.

Several studies have addressed the challenge of extracting claims from scientific texts. Some frameworks based on zero-shot models (Pan et al., 2021; Wright et al., 2022) have achieved promising results, primarily focusing on generating claim datasets from raw texts to train fact-checking models in specific domains. However, these methods rely on multi-stage NLP pipelines that are prone to failure and tend to produce a large number of claims per document, making their integration into an end-to-end system difficult. In contrast, recent

¹https://labdemos.expertcustomers.ai/health_claims (user: guest / password: acldemos2025)

²www.youtube.com/watch?v=jyms_Ey0YSQ

³www.github.com/expertailab/sciclaims-backend and www.github.com/expertailab/sciclaims-frontend

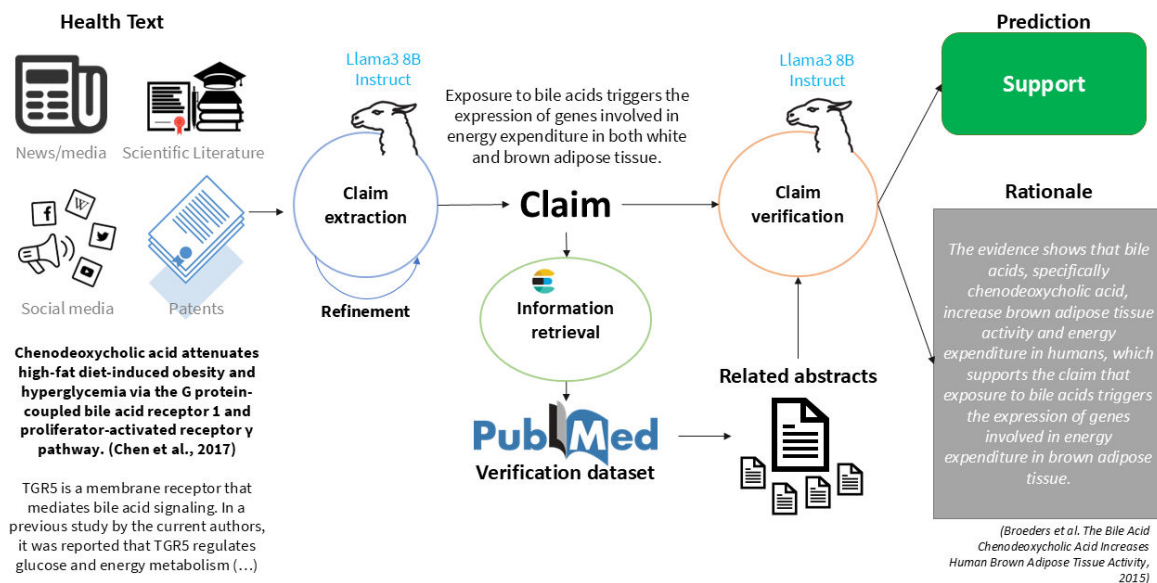


Figure 1: System Architecture.

approaches based on LLMs extract atomic factual units (Min et al., 2023; Chern et al., 2023), which serve as concise, interpretable summaries of the source texts.

For the evidence retrieval phase, dense passage retrieval methods, such as ColBERT (Khattab and Zaharia, 2020), have emerged due to their ability to retrieve highly relevant documents from large corpora with great precision. However, these methods are computationally expensive and therefore less practical for real-time, lightweight applications. Thus, simpler approaches such as BM-25 (Robertson and Zaragoza, 2009) and tools like Elasticsearch, known for their balance between retrieval quality and computational efficiency, are still preferred for deployment in production environments.

The release of several claim verification datasets has spurred the development of numerous models aimed at addressing this challenge. VERT5ERINI (Pradeep et al., 2021), PARAGRAPHJOINT (Li et al., 2021), and MultiVerS (Wadden et al., 2022) have achieved promising results on scientific benchmarks like SciFact (Wadden et al., 2020) and CLIMATE-FEVER (Diggelmann et al., 2021). However, these approaches still leave room for improvement and typically rely on large, labeled datasets, which limits their scalability across domains. In parallel, recent advances in LLMs have led to increased interest in verifying automatically generated content. Frameworks such as FactScore (Min et al., 2023), FacTool (Chern et al., 2023), and LLM Oasis (Scirè et al., 2024) focus on assessing the factuality of

LLM-generated text rather than existing, human-written passages. Nonetheless, their techniques, such as extracting atomic knowledge units and using zero-shot LLMs for claim extraction and verification, can be extended to real-world scientific texts, offering a promising direction for scalable, data-efficient claim analysis.

Although individual components have seen substantial progress, an integrated system capable of seamlessly connecting these steps remains an open problem. Many existing systems, such as FactDetect (Jafari and Allan, 2024), ClIVER (Liu et al., 2024), and more recently (Liang and Sonntag, 2025; Vladika et al., 2025), focus on claims that are already identified, neglecting the crucial first step of extracting relevant claims from raw, real-world texts. As a result, these systems are limited to pre-identified claims rather than addressing the full pipeline of claim extraction, retrieval, and verification.

Given the complexity of this problem, our approach aims to optimize each stage of the pipeline to ensure both efficiency and accuracy in real-world biomedical claim analysis. In this work, we tap on the strengths of modern LLMs to address previous limitations, specifically in the claim extraction and verification stages, and integrate our results as a robust online system.

3 System Description

SciClaims is an interactive, end-to-end system for scientific claim analysis, comprising three main

Select one of the sample documents below or try your own text. Then, click **ANALYZE**

Example 31 (Biomedicine-paper): Aldehyde dehydrogenase 1 (ALDH1) expression is an inde...

[source](#)

Aldehyde dehydrogenase 1 (ALDH1) expression is an independent prognostic factor in triple negative breast cancer (TNBC).

Triple negative breast cancer is a breast cancer subtype that is characterized by poor prognosis and aggressive behavior. Meanwhile, cancer stem cells are believed to contribute to tumorigenesis and metastasis. Our earlier studies in cell lines and animal models revealed that CSCs were enriched in TNBC patients. In addition, CSCs sorted from TNBC cell lines were more tumorigenic relative to non-CSCs.[3,4] In the present study, we further analyzed the relationship between CSCs and cancer development in TNBC patients using ALDH1 as a CSC marker. Since the first report by Ginestier et al[11] showing that ALDH1 expression was associated with poor clinical outcome in breast cancer, several studies have indicated that ALDH1+ breast cancer cells are related to TNBC.

891/1000%

Analyze

Triple negative breast cancer is a type of breast cancer that often has a poor prognosis and behaves aggressively.

Cancer stem cells play a key role in the development and spread of cancer.

Cancer stem cells are more common in people with triple negative breast cancer.

SUPPORT

Prediction score:

99.94%

Rationale: The evidence suggests that normal cells bearing cancer stem cell markers are associated with the triple negative receptor subtype of breast cancer, supporting the claim that cancer stem cells are more common in people with triple negative breast cancer.

<https://doi.org/10.1186/bcr3471>

Cancer stem cell markers are enriched in normal tissue adjacent to triple negative breast cancer and inversely correlated with DNA repair deficiency.

Normal cells bearing cancer stem cell markers are associated with the triple negative receptor subtype of breast cancer. This study suggests stem cell staining and gene expression signatures from normal breast tissues represent novel tissue-based risk biomarkers for triple negative breast cancer. Validation of these results in additional studies of normal tissue from cancer-free women could lay the foundation for future targeted triple negative breast cancer prevention strategies.

Figure 2: Screenshot of the SciClaims Demo.

components: claim extraction, evidence retrieval, and claim verification. It is optimized to run efficiently on a single 24GB VRAM GPU, enabling real-time performance via a web-based interface.

The system architecture (Figure 1) is built around a Llama3 8B Instruct model (Grattafiori et al., 2024) and an Elasticsearch-based retrieval engine. It processes biomedical or scientific text input and outputs a list of extracted claims, their verification status, relevant evidence, and a natural language rationale. The model is set up using vLLM (Kwon et al., 2023), which enables high-throughput processing in inference. Next, we present the main building blocks of SciClaims.

Claim extraction: This first module extracts potential claims from the source text. A claim is characterized by a specific set of properties detailed in section 5. The Claim Extraction module calls the Llama3 8B Instruct model twice: first, to generate an initial list of claims, and second, to refine and filter them, improving the quality of the resulting claims. All the prompts used in our pipeline are provided in Appendix A. Based on our evaluation, we made adjustments to the initial prompts to improve claim extraction performance. Further details on this refinement can be found in section 5.

Document retrieval: The second module retrieves potentially relevant documents from the verification dataset, using the claim as a query to the Elasticsearch index. The verification dataset contains 4.7 million abstracts from PubMed

(2000–2022) that were curated using the Semantic Scholar’s *Highly Influential Citations* metric (Valenzuela et al., 2015), ensuring that each article is backed by at least three highly influential citations. This selection criterion helps prioritize documents that have been extensively referenced in the academic community, enhancing the quality and relevance of retrieved information for verification. We chose not to filter documents based on Elasticsearch’s scoring mechanism to maximize recall. Instead, the subsequent verification module is responsible for discarding irrelevant documents.

Claim verification: The final module performs fact-checking by making another call to the LLM, providing the claim along with the retrieved related documents. The model assigns one of the following three labels: 1) *SUPPORT* if the claim is verified by the document, 2) *REFUTE* if the claim is refuted by the document, or 3) *NEI* (not enough information) if the document lacks sufficient evidence or is not relevant to the claim. To improve transparency and interpretability, we also request the model to provide a rationale for its decision, including identifying the most relevant sentence(s) that support its conclusion.

4 User interface

In this section, we demonstrate the functionality of our approach through a web-based claim analysis tool that allows users to easily interact with the system and analyze claims within relevant texts,

leveraging the backend architecture presented in the previous section. The demo showcases the complete workflow, from entering a scientific passage to obtaining fact-checking results with supporting evidence and explanations.

The user interface provides a drop-down menu featuring over 30 pre-selected examples from various domains, including biomedicine papers, COVID-related news, social media, and patents (see Figure 2). These examples were chosen to represent a broad spectrum of platforms and relevant disciplines, allowing users to explore diverse contexts. By selecting any example from the list, the corresponding text will be displayed in the text box below, with a hyperlink to the original source. The text box is fully editable, enabling users to modify the content and analyze their own text. While the application can process texts up to 10,000 characters, we recommend keeping the input under 2,000 characters for faster results.

As shown in Figure 2, once the user has selected or entered a text, they can click the *Analyze* button to initiate the claim analysis process. Results are presented as a list of identified claims, each of which can be expanded to reveal detailed information. For each claim, the interface shows:

Prediction label: Whether the claim is supported (green) or refuted (red). Results labeled as *Not Enough Information* by the LLM are not presented to the user.

Prediction score: A normalized probability score that represents the confidence level of the model in its prediction. This score is derived from the statistical outputs of the model, particularly the tokens representing the label string, and it is expressed as a percentage.

Evidence: The related document selected by the retrieval module, along with its DOI. The specific sentence(s) in the abstract of the paper that were most influential for the prediction according to the LLM are highlighted.

Rationale: A justification of the reasoning behind the classification, providing additional insight into the decision-making process carried out by the model.

This interactive interface provides an intuitive and user-friendly environment for testing and exploring claim analysis in various health-related texts. The analysis goes through the entire pipeline, offering a label for the claims extracted along with the retrieved evidence. When a claim receives conflicting labels from different pieces of evidence,

our system returns all relevant pairs, allowing users to assess which is more accurate. Since scientific knowledge evolves, evidence considered true at one time may later be contradicted. We recognize the value of providing all supporting and refuting evidence along with their publication dates.

5 Experimentation and Results

We evaluate our system following a three-stage approach. First, we evaluate SciClaims’ modules against some baselines using SciFact as benchmark. Then, we evaluate the performance of the system at every step of the pipeline using an LLM as a judge. Finally, we perform a human evaluation for the whole system.

5.1 Evaluation in SciFact

We use SciFact, a benchmark dataset focused on biomedical literature, to evaluate SciClaims. It includes 1,400 expert-authored scientific claims, each linked to evidence-containing abstracts and annotated with a verification label.

SciFact supports two complementary evaluations: (1) claim extraction, by comparing system-generated claims to human-written ones using shared source documents, and (2) claim verification, by assessing how accurately the system labels claim-evidence pairs.

5.1.1 Claim extraction

To evaluate the claim extraction module, we consider the expert-written claims from SciFact as the ground truth for each source paragraph.

Method	Rouge1	Rouge2	RougeL	Similarity Score
Sentence tokenizer	0.3313	0.1980	0.3030	0.7211
(Pan et al., 2021)	0.2780	0.1334	0.2555	0.6561
(Wright et al., 2022)	0.2525	0.0762	0.2220	0.6475
Noun Phrases Gen	0.2567	0.0953	0.2293	0.6693
SciClaims	0.3387	0.1896	0.3084	0.7250

Table 1: Claim quality scores in SciFact

As a first baseline, we select a sentence tokenizer, treating each sentence in the source paragraph as a claim. Next, we select two baselines (Pan et al., 2021; Wright et al., 2022) that use a pipeline with two transformer models to generate claims from the source paragraph through entity extraction. The first model generates a question-answer pair, where the answer is the entity, while the second reformulates it as a claim. We also introduce a method where we propose to build the claims around noun phrases rather than named entities.

We evaluate the quality of claim generation by matching each system-generated claim with its most similar gold claim from the same paragraph. We discard pairs with a Levenshtein similarity score below 0.3, a threshold empirically tuned to balance recall and noise. For valid matches, we compute ROUGE-1, ROUGE-2, ROUGE-L, and a semantic similarity score using a DeBERTa model fine-tuned on the STS-B dataset. As shown in Table 1, SciClaims achieves the highest scores in ROUGE-1, ROUGEL, and semantic similarity, indicating more faithful and informative claim generation.

5.1.2 Claim verification

We evaluate SciClaims on SciFact’s test set, comparing its label accuracy for claim-evidence pairs with existing approaches.

Architecture	Precision	Recall	F1-Score
Roberta-base*	0.4662	0.5963	0.5220
MultiVerS*	0.7286	0.7321	0.7303
SciClaims (LLaMA3 8B Instruct)	0.7034	0.6863	0.6788
SciClaims (Qwen2 7B Instruct)	0.6649	0.6677	0.6439
SciClaims (Phi3 Small 8K)	0.7100	0.6894	0.6525
SciClaims (OLMO2 7B Instruct)	0.6379	0.6118	0.5886

Table 2: Precision, recall and F1-Scores in Claim Verification of SciFact test set. RoBERTa-base and MultiVerS are fine-tuned models, while SciClaims is zero-shot. In parenthesis, the backbone used in each SciClaims configuration

We compare SciClaims with two strong baselines: a RoBERTa-base classifier (Liu et al., 2019) and MultiVerS (Wadden et al., 2022), a longformer-based model for claim verification. Both baselines are fine-tuned on SciFact. While MultiVerS achieves the highest F1-score, SciClaims performs competitively despite operating in a zero-shot setting. As shown in Table 2, the relatively narrow performance gap demonstrates SciClaims’ strong generalization ability without task specific training. Furthermore, as shown in Table 2, we evaluate SciClaims with different LLMs with similar sizes as backbone, being LLaMA3 8B Instruct the one which offered the best performance.

5.2 Evaluation with a judge model

The second stage of our evaluation provides a comprehensive assessment of the entire system using a LLM as judge. Based on the Judge Arena leaderboard⁴, we select Qwen 2.5 72B turbo as judge,

since it is the first ranked model with open weights and a higher parameter count than our system’s LLM (Llama3 8B Instruct). We chose its 4-bit quantization version (Qwen2.5 72B AWQ⁵) due to hardware limitations. For the evaluation sample, we randomly select 120 documents from the PubMed dataset presented in section 3. The evaluation is conducted in three phases: Claim quality, document retrieval, and claim verification and full-system evaluation.

5.2.1 Claim quality evaluation

In this phase, we evaluate the quality of the claims generated by SciClaims and compare it to other methods, using the same baselines mentioned in Section 5.1.

We devised a questionnaire consisting of eight yes/no questions (see Appendix B) to capture the desired properties in a correct claim and ask the judge model to answer it. The first question requires context from the source paragraph, while the remaining questions focus solely on the claim. Correct claims need to receive a *Yes* to all questions. Table 3 shows that our LLM-based system outperforms all other methods, scoring 15 points higher than the second-best, the sentence tokenizer.

To further enhance the results, we refined our claim extraction prompts with two optimizations:

Claim Definition Properties (CDP). Here we enhance the prompt by incorporating the characteristics of a claim, such as precision, conciseness, or check-worthiness. These characteristics are derived from the questionnaire used to evaluate the quality of the claims. The goal is to guide the LLM to adhere to these criteria when generating the list of claims.

Claim Refinement (CR): This upgrade involves a follow-up call to the LLM in order to refine the initial list of claims. For each candidate claim, we pair it with the source paragraph and ask the LLM to refine the claim based on the same criteria (precision, conciseness, check-worthiness, etc.). This step aims to eliminate poorly formed claims and reinforce the application of the specified criteria.

These two upgrades result in an additional 29-point increase in the percentage of correct claims generated by SciClaims while reducing the number of candidate claims generated by our approach. Notably, QA-oriented baselines, such as (Wright et al., 2022) and our noun phrase-based generation

⁴<https://huggingface.co/spaces/AtlaAI/judge-arena>

⁵huggingface.co/Qwen/Qwen2.5-72B-Instruct-AWQ

Method	Q1	Q2	Q3	Q4	Q5	Q6	Q7	Q8	Correct Claims (%)	Candidate claims
Sentence tokenizer	0.9987	0.9100	0.8436	0.8709	0.9322	0.9009	0.3703	0.9596	32.46	767
(Pan et al., 2021)	0.2449	0.1303	0.1618	0.1371	0.2112	0.1169	0.0337	0.1641	3.15	445
(Wright et al., 2022)	0.8043	0.4714	0.5040	0.3842	0.5593	0.4829	0.1641	0.6249	11.28	4175
Noun Phrases Gen	0.3001	0.1562	0.1857	0.1209	0.2388	0.1320	0.0316	0.2203	2.42	4335
SciClaims	0.9949	0.9474	0.9718	0.9089	0.9320	0.9345	0.5238	0.9756	47.37	779
SciClaims+CDP	0.9942	0.9690	0.9845	0.9380	0.9574	0.9593	0.5930	0.9903	55.43	516
SciClaims+CDP+CR	0.9923	0.9787	0.9884	0.9845	0.9806	0.9748	0.7810	0.9903	76.36	516

Table 3: Claim extraction (phase 1) evaluation results with a judge model (Qwen2.5 72B AWQ).

Claims	All Claims			Correct Claims		
	R@1	R@3	R@5	R@1	R@3	R@5
Sentence tokenizer	0.5591	0.7205	0.7795	0.6265	0.7390	0.7871
(Pan et al., 2021)	0.2970	0.4653	0.5248	0.6429	0.7143	0.7143
(Wright et al., 2022)	0.5860	0.7474	0.8038	0.7113	0.8365	0.8726
Noun Phrases Gen	0.4634	0.6456	0.7013	0.6286	0.7619	0.8095
SciClaims	0.5045	0.6645	0.7135	0.5257	0.6938	0.7344
SciClaims+CDP	0.5146	0.6940	0.7466	0.5944	0.7448	0.7867
SciClaims+CDP+CR	0.4727	0.6250	0.6777	0.5102	0.6675	0.7107

Table 4: Document retrieval (phase 2) evaluation results with a judge model (Qwen2.5 72B AWQ).

method, generate significantly larger quantities of claims compared to the other methods in Table 3.

5.2.2 Document retrieval evaluation

We assess document relevance by asking the judge model whether each retrieved paragraph aids claim verification. Recall is evaluated at $k = 1, 3, 5$ retrieved documents per claim. As shown in Table 4, claims generated by (Wright et al., 2022) retrieve more potentially relevant documents than SciClaims. However, SciClaims performs well, retrieving at least one relevant document in 75% of cases when fetching five documents per claim, a common real-world scenario. This highlights its balance between claim accuracy and document relevance. QA-oriented methods generate more compact, less specific claims than LLMs, increasing the likelihood of retrieving related documents.

5.2.3 Claim verification and full-system

The verification task involves predicting the veracity of claim-evidence pairs, classifying each as *SUPPORT*, *REFUTE*, or *NOT ENOUGH INFORMATION* (*NEI*). Thus we evaluate our system’s label accuracy by asking to the judge model whether the assigned label is correct. We compare SciClaims results with MultiVerS (Wadden et al., 2022).

SciClaims demonstrates superior claim verification accuracy, outperforming the fine-tuned MultiVerS (Wadden et al., 2022) model despite operating in a zero-shot setting. As shown in Table 5, SciClaims consistently produces more accurate labels

across all claim generation methods, except (Pan et al., 2021). Notably, SciClaims returns significantly fewer *NEI* labels than MultiVerS, providing greater value to users by delivering more definitive *SUPPORT* or *REFUTE* labels. The unusually high accuracy of (Pan et al., 2021), likely stems from its excessive *NEI* labels, which simplify label selection. Furthermore, the SciClaims+CDP+CR system achieves the highest overall performance, correctly labeling 50% of generated claims, outperforming the next-best MultiVerS-based system by over five points.

Table 5 also reports the average processing time per document for each system configuration. Systems using MultiVerS as the verification module tend to be the fastest overall, with the exception of those paired with claim extraction modules like (Wright et al., 2022) and Noun Phrases Generation, which produce a high number of claims and thus increase runtime. Among all configurations, those using SciClaims+CDP+CR for claim generation—paired with either SciClaims or MultiVerS for verification—offer the best trade-off between predictive accuracy and time efficiency. Importantly, only the configuration that uses SciClaims as both the generation and verification module can run on a single 24GB GPU, making it uniquely suitable for real-world deployment.

5.3 Human evaluation

To complement our automated evaluation with LLMs, we conducted a human evaluation to assess the quality, relevance, and usability of SciClaims outputs. Five NLP-experts independently reviewed a sample of 154 PubMed documents processed by the system. The annotation tasks were divided into three phases, following the same structure as our LLM-based evaluation (see Section 5.2).

Claim quality evaluation. Annotators were shown an input paragraph and one randomly selected claim extracted by SciClaims. They answered the same eight binary questions used in the

System		All Claims		Correct Claims		System Score	Time/doc (secs.)
Extraction Module	Verification Module	Label Accuracy	Not NEI (%)	Label Accuracy	Not NEI (%)		
Sentence tokenizer	MultiVerS	0.5494	4.77	0.5127	4.82	0.1664	2.36
(Pan et al., 2021)	MultiVerS	0.7591	2.31	0.5714	11.90	0.0180	2.17
(Wright et al., 2022)	MultiVerS	0.5015	5.53	0.4268	8.35	0.0482	14.25
Noun Phrases Gen	MultiVerS	0.5965	3.08	0.4825	6.67	0.0117	15.01
SciClaims	MultiVerS	0.5944	6.02	0.5709	6.50	0.2704	2.86
SciClaims+CDP	MultiVerS	0.5867	5.71	0.5361	5.60	0.2971	1.82
SciClaims+CDP+CR	MultiVerS	0.6204	3.72	0.5922	7.44	0.4522	2.93
Sentence tokenizer	SciClaims	0.6041	61.20	0.6693	59.17	0.2173	11.72
(Pan et al., 2021)	SciClaims	0.6568	39.60	0.619	71.43	0.0195	7.61
(Wright et al., 2022)	SciClaims	0.6397	69.44	0.7098	74.66	0.0801	65.28
Noun Phrases Gen	SciClaims	0.6608	53.02	0.6762	70.48	0.0164	68.01
SciClaims	SciClaims	0.6361	58.49	0.6567	59.71	0.3111	12.38
SciClaims+CDP	SciClaims	0.6296	58.87	0.6364	62.47	0.3527	8.13
SciClaims+CDP+CR	SciClaims	0.6589	53.06	0.6574	54.23	0.5020	9.24

Table 5: Verification evaluation (phase 3) results with a judge model (Qwen2.5 72B AWQ). The table shows the label accuracy of each combination of extraction and verification module. The Correct Claims column counts only those considered correct in the Phase 1 questionnaire. Not NEI represents the proportion of claim–evidence pairs labeled as SUPPORT or REFUTE, rather than NEI (Not Enough Information). The last two columns summarize system-level performance: Processing Time per Document is the average time (in seconds) each system takes to process a single document, and System Score reflects the ratio of correctly labeled and well-formed claims across the system.

LLM-based evaluation (see Appendix B), assessing conciseness, precision, and other key dimensions. Table 6 shows that human judgments closely align with the judge model, with 70.5% of the claims meeting all eight criteria.

Claim Quality	Score	Rationale Quality	Score
Q1 (grounding)	0.9048		
Q2 (grammar)	0.9810		
Q3 (completeness)	0.9810	RQ1 (justification)	0.8571
Q4 (precision)	0.8667	RQ2 (relevance)	0.9206
Q5 (relevance)	0.8952	RQ3 (completeness)	0.7937
Q6 (conciseness)	0.8571		
Q7 (self-contained)	0.9238		
Q8 (contribution)	0.8857		
Correct Claims (%)	70.47	Correct Rationales (%)	74.61

Table 6: Claim and rationale evaluation of SciClaims by human annotators. Score indicates the overall ratio of ‘yes’ responses to the question along the annotators.

Document retrieval evaluation. Annotators were presented with a claim and its corresponding retrieved paragraph from SciClaims. They were asked whether the retrieved evidence provided sufficient information to verify the claim. In 60% of cases, annotators judged the paragraph as informative enough for claim verification, indicating moderate effectiveness of the retrieval module in real-world scenarios.

Claim verification evaluation. In this phase, annotators were shown a claim, its evidence, the system’s predicted label, and the corresponding rationale. They rated the label’s accuracy on a five-point scale, from 1 (completely inaccurate) to 5 (highly accurate). SciClaims achieved a strong

average score of 4.40, reflecting high alignment between system predictions and human judgment. Additionally, annotators evaluate the quality of the generated rationale using three yes/no questions: (RQ1) Is the label justified by the rationale? (RQ2) Does the rationale focus on relevant information? (RQ3) Does the rationale provide enough context to understand the label?. As shown in Table 6, the responses from annotators indicate high satisfaction with the relevance and justification of rationales, though they also noted that completeness could be improved.

6 Conclusion

We presented SciClaims, a practical, end-to-end system for scientific claim analysis in the biomedical domain. Through an interactive web-based interface, users can extract, verify, and explore scientific claims with evidence-backed rationales, all powered by LLMs and a curated biomedical corpus. SciClaims is designed for usability and speed, and it is already being explored in real-world settings such as pharmaceutical patent analysis and systematic literature reviews. By combining explainable outputs with efficient infrastructure, it provides a robust tool for researchers, clinicians, and analysts who need to validate scientific information quickly and reliably. The system is openly accessible and ready for demonstration, offering an engaging experience to showcase the capabilities of modern LLM-based scientific reasoning.

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- ichtenhofer, Cynthia Gao, Damon Civin, Dana Beaty, Daniel Kreymer, Daniel Li, David Adkins, David Xu, Davide Testuggine, Delia David, Devi Parikh, Diana Liskovich, Didem Foss, Dingkan Wang, Duc Le, Dustin Holland, Edward Dowling, Eissa Jamil, Elaine Montgomery, Eleonora Presani, Emily Hahn, Emily Wood, Eric-Tuan Le, Erik Brinkman, Esteban Arcaute, Evan Dunbar, Evan Smothers, Fei Sun, Felix Kreuk, Feng Tian, Filippos Kokkinos, Firat Ozgenel, Francesco Caggioni, Frank Kanayet, Frank Seide, Gabriela Medina Florez, Gabriella Schwarz, Gada Badeer, Georgia Swee, Gil Halpern, Grant Herman, Grigory Sizov, Guangyi, Zhang, Guna Lakshminarayanan, Hakan Inan, Hamid Shojanazeri, Han Zou, Hannah Wang, Hanwen Zha, Haroun Habeeb, Harrison Rudolph, Helen Suk, Henry Aspegren, Hunter Goldman, Hongyuan Zhan, Ibrahim Damla, Igor Molybog, Igor Tufanov, Ilias Leontiadis, Irina-Elena Veliche, Itai Gat, Jake Weissman, James Geboski, James Kohli, Janice Lam, Japhet Asher, Jean-Baptiste Gaya, Jeff Marcus, Jeff Tang, Jennifer Chan, Jenny Zhen, Jeremy Reizenstein, Jeremy Teboul, Jessica Zhong, Jian Jin, Jingyi Yang, Joe Cummings, Jon Carvill, Jon Shepard, Jonathan McPhie, Jonathan Torres, Josh Ginsburg, Junjie Wang, Kai Wu, Kam Hou U, Karan Saxena, Kartikay Khandelwal, Katayoun Zand, Kathy Matosich, Kaushik Veeraraghavan, Kelly Michelena, Keqian Li, Kiran Jagadeesh, Kun Huang, Kunal Chawla, Kyle Huang, Lailin Chen, Lakshya Garg, Lavender A, Leandro Silva, Lee Bell, Lei Zhang, Liangpeng Guo, Licheng Yu, Liron Moshkovich, Luca Wehrstedt, Madian Khabza, Manav Avalani, Manish Bhatt, Martynas Mankus, Matan Hasson, Matthew Lennie, Matthias Reso, Maxim Groshev, Maxim Naumov, Maya Lathi, Meghan Keneally, Miao Liu, Michael L. Seltzer, Michal Valko, Michelle Restrepo, Mihir Patel, Mik Vyatskov, Mikayel Samvelyan, Mike Clark, Mike Macey, Mike Wang, Miquel Jubert Hermoso, Mo Metanat, Mohammad Rastegari, Munish Bansal, Nandhini Santhanam, Natascha Parks, Natasha White, Navyata Bawa, Nayan Singhal, Nick Egebo, Nicolas Usunier, Nikhil Mehta, Nikolay Pavlovich Laptev, Ning Dong, Norman Cheng, Oleg Chernoguz, Olivia Hart, Omkar Salpekar, Ozlem Kalinli, Parkin Kent, Parth Parekh, Paul Saab, Pavan Balaji, Pedro Rittner, Philip Bontrager, Pierre Roux, Piotr Dollar, Polina Zvyagina, Prashant Ratanchandani, Pritish Yuvraj, Qian Liang, Rachad Alao, Rachel Rodriguez, Rafi Ayub, Raghotham Murthy, Raghu Nayani, Rahul Mitra, Rangaprabhu Parthasarathy, Raymond Li, Rebekkah Hogan, Robin Battey, Rocky Wang, Russ Howes, Ruty Rinott, Sachin Mehta, Sachin Siby, Sai Jayesh Bondu, Samyak Datta, Sara Chugh, Sara Hunt, Sargun Dhillon, Sasha Sidorov, Satadru Pan, Saurabh Mahajan, Saurabh Verma, Seiji Yamamoto, Sharadh Ramaswamy, Shaun Lindsay, Shaun Lindsay, Sheng Feng, Shenghao Lin, Shengxin Cindy Zha, Shishir Patil, Shiva Shankar, Shuqiang Zhang, Shuqiang Zhang, Sinong Wang, Sneha Agarwal, Soji Sajuyigbe, Soumith Chintala, Stephanie Max, Stephen Chen, Steve Kehoe, Steve Satterfield, Sudarshan Govindaprasad, Sumit Gupta, Summer Deng, Sungmin Cho, Sunny Virk, Suraj Subramanian, Sy Choudhury, Sydney Goldman, Tal Remez, Tamar Glaser, Tamara Best, Thilo Koehler, Thomas Robinson, Tianhe Li, Tianjun Zhang, Tim Matthews, Timothy Chou, Tzook Shaked, Varun Vontimitta, Victoria Ajayi, Victoria Montanez, Vijai Mohan, Vinay Satish Kumar, Vishal Mangla, Vlad Ionescu, Vlad Poenaru, Vlad Tiberiu Mihailescu, Vladimir Ivanov, Wei Li, Wenchen Wang, Wenwen Jiang, Wes Bouaziz, Will Constable, Xiao Cheng Tang, Xiaojian Wu, Xiaolan Wang, Xilun Wu, Xinbo Gao, Yaniv Kleinman, Yanjun Chen, Ye Hu, Ye Jia, Ye Qi, Yenda Li, Yilin Zhang, Ying Zhang, Yossi Adi, Youngjin Nam, Yu, Wang, Yu Zhao, Yuchen Hao, Yundi Qian, Yunlu Li, Yuzi He, Zach Rait, Zachary DeVito, Zef Rosnbrick, Zhaoduo Wen, Zhenyu Yang, Zhiwei Zhao, and Zhiyu Ma. 2024. [The llama 3 herd of models](#). *Preprint*, arXiv:2407.21783.
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A Prompts

A.1 SciClaims claim extraction first step

```
<| begin_of_text |><| start_header_id |>system<|
end_header_id|>
```

Your task is to generate a list with the main factual claims stated in a text. A factual claim makes an assertion about something regarding the subject matter that can be proved or contradicted with factual evidence. Factual claims must be expressed as meaningful, self-contained sentences. Do not include narrative context and disregard absolutely ALL self-referential parts.

Arrange your output using the format:

```
-- claim
-- claim
-- claim.
```

```
<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
```

TEXT: {text}

A.2 SciClaims+CDP

```
<| begin_of_text |><| start_header_id |>system<|
end_header_id|>
```

Identify and list the main scientific claims stated in a passage. Each claim must satisfy the following criteria :

- ****Convey an insight, interpretation, or conclusion drawn from the passage that is testable and generalizable****: The claim should assert an outcome, capability, or effect rather than merely describing a method, aim, or process.
 - Example (Good): "Neural networks outperform decision trees in image classification tasks." (Testable outcome)
 - Example (Bad): "The proposed method aims to use neural networks for better image classification." (Descriptive, not assertive)
- ****Be expressed as a meaningful, self-contained statement****: Each claim should be fully understandable on its own, without needing context from the passage or other claims. It must convey a complete, independent idea. If referencing a study, survey, result, or process, phrase it as a general, verifiable claim.
 - Example (Good): "The Amazon rainforest is home to over 10 million species."
 - Example (Bad): "As mentioned earlier, the Amazon is one of the most biodiverse places in the world." (Requires prior context and doesn't stand alone.)
- ****Emphasize generalization and scientific assertion****: Avoid descriptive or narrative conclusions.
 - Example (Good): "Exposure to blue light before sleep can reduce melatonin production." (Generalized, testable assertion)
 - Example (Bad): "The study investigates how exposure to blue light before sleep affects melatonin production." (Descriptive of method, not a general claim)
- ****Be clear and concise****: Use straightforward language without unnecessary words.
 - Example (Good): "The Eiffel Tower is in Paris."
 - Example (Bad): "The Eiffel Tower, which is one of the most iconic landmarks in Europe and

attracts millions of tourists every year, is located in Paris, France."

- ****Exclude narrative context****: Focus on the factual assertion itself, not the surrounding story or background information.
 - Example (Good): "Water boils at 100C under normal atmospheric pressure."
 - Example (Bad): "In many cultures, people have believed for centuries that water boils at 100C, as scientists confirmed in the 18th century." (Includes unnecessary background information.)
- ****Disregard all self-referential content****: Ignore any statements referring to the passage itself or the author intentions.
 - Example (Good): "The Earth orbits the Sun."
 - Example (Bad): "The study explains how the Earth orbits the Sun." (This is self-referential and refers to the passage itself.)
- ****Be precise and objective****: Avoid ambiguity, subjective interpretation, or vague statements. Present claims as clear, verifiable facts.
 - Example (Good): "The Great Wall of China stretches approximately 13,000 miles."
 - Example (Bad): "The Great Wall of China is pretty long." (Vague and subjective.)
- ****Be relevant to the broader debate or public discourse****: Focus on verification –worthy claims that introduce new information rather than merely restating common knowledge.
 - Example (Good): "The global temperature has increased by about 1C since the late 19th century."
 - Example (Bad): "The Earth is a planet." (Common knowledge and not contributing new, verifiable information.)

Present the output using the following format:

```
-- claim
-- claim
-- claim
```

```
<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
```

PASSAGE: {text}

A.3 SciClaims claim extraction second step (refinement)

Now, using the information given by the passage, reformulate each individual claim to be fully understandable by itself, even without having the context from the passage or the rest of the claims from the list. Change the terminology or add context information to each claim if necessary.

A.4 SciClaims+CDP+CR (refinement)

```
<| begin_of_text |><| start_header_id |>system<|
end_header_id|>
```

Given a claim and the passage it was extracted from, reformulate the claim to fully adhere to ****ALL**** the following criteria.

- ****Convey an insight, interpretation, or conclusion drawn from the passage that is testable and generalizable****: The claim should assert an outcome, capability, or effect rather than merely describing a method, aim, or process.

- Example (Good): "Neural networks outperform decision trees in image classification tasks ." (Testable outcome)
- Example (Bad): "The proposed method aims to use neural networks for better image classification ." (Descriptive , not assertive)
- ****Be expressed as a meaningful, self-contained statement****: Each claim should be fully understandable on its own, without needing context from the passage or other claims. It must convey a complete, independent idea. If referencing a study, survey, result, or process, phrase it as a general, verifiable claim.
 - Example (Good): "The Amazon rainforest is home to over 10 million species."
 - Example (Bad): "As mentioned earlier, the Amazon is one of the most biodiverse places in the world." (Requires prior context and doesn't stand alone.)
- ****Emphasize generalization and scientific assertion****: Avoid descriptive or narrative conclusions.
 - Example (Good): "Exposure to blue light before sleep can reduce melatonin production." (Generalized, testable assertion)
 - Example (Bad): "The study investigates how exposure to blue light before sleep affects melatonin production." (Descriptive of method, not a general claim)
- ****Be clear and concise****: Use straightforward language without unnecessary words.
 - Example (Good): "The Eiffel Tower is in Paris."
 - Example (Bad): "The Eiffel Tower, which is one of the most iconic landmarks in Europe and attracts millions of tourists every year, is located in Paris, France."
- ****Exclude narrative context****: Focus on the factual assertion itself, not the surrounding story or background information.
 - Example (Good): "Water boils at 100C under normal atmospheric pressure."
 - Example (Bad): "In many cultures, people have believed for centuries that water boils at 100C, as scientists confirmed in the 18th century." (Includes unnecessary background information.)
- ****Disregard all self-referential content****: Ignore any statements referring to the passage itself or the author intentions.
 - Example (Good): "The Earth orbits the Sun."
 - Example (Bad): "The study explains how the Earth orbits the Sun." (This is self-referential and refers to the passage itself.)
- ****Be precise and objective****: Avoid ambiguity, subjective interpretation, or vague statements. Present claims as clear, verifiable facts.
 - Example (Good): "The Great Wall of China stretches approximately 13,000 miles."
 - Example (Bad): "The Great Wall of China is pretty long." (Vague and subjective.)
- ****Be relevant to the broader debate or public discourse****: Focus on verification-worthy claims that introduce new information rather than merely restating common knowledge.
 - Example (Good): "The global temperature has increased by about 1C since the late 19th century."
 - Example (Bad): "The Earth is a planet." (Common knowledge and not contributing new, verifiable information.)

Present the output using the following format:
{"original_claim ": <str>, "refined_claim ": <str>, "rationale ": <str>}

```
<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
CLAIM: {claim}
PASSAGE: {text}
```

A.5 SciClaims claim verification

```
<| begin_of_text |><| start_header_id |>system<|
end_header_id|>
```

You are a claim analyst. Upon receiving a claim and an evidence, your task is to figure out if the claim is either supported, contradicted or unrelated based exclusively on the evidence.

- If you are confident that the claim is supported by the evidence your answer will be "SUPPORT".
- If you are certain that the evidence directly contradicts the claim, your answer will be "CONTRADICT". Please note that if the claim is just not mentioned in the evidence, or if it is unrelated to the evidence, it does not mean it is contradicted. For that cases, the answer will be "NEI".
- If the evidence does not contain enough information or if it is not related to the claim, your answer will be "NEI", which stands for Not Enough Information.

Arrange the output as a JSON dictionary with the keys "response" and "evidence" and ensure your output is JSON-valid.

- The "response" values can only be "SUPPORT", "CONTRADICT" or "NEI".
- The "evidence" value must be a list of sentences from the evidence which are more related to your decision. If the decision is "NEI", this field will be empty.

```
<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
CLAIM: {claim}
EVIDENCE: {evidence}
```

A.6 Phase 1 evaluation with judge model (Q1)

```
<| begin_of_text |><| start_header_id |>system<|
end_header_id|>
```

Given a sentence and a paragraph, answer the following question. Use exclusively the content of the paragraph to answer the question.

Is the sentence supported by the paragraph?

Return your response as a json dictionary, following this structure : {"answer":<Yes/No>, "rationale ": <str>}

```
<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
SENTENCE: {claim}
PARAGRAPH: {text}
```

A.7 Phase 1 evaluation with judge model (Q2-8)

Given a claim, answer the following question .

{QUESTION}

Return your response as a json dictionary , following
this structure : {"answer":<Yes/No>, "rationale ":
<str>}

<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
CLAIM: {claim}

A.8 Phase 2 evaluation with judge model

<| begin_of_text |><| start_header_id |>system<|
end_header_id|>

Given a claim and a paragraph, answer the following
question .

Is the information contained in the paragraph useful
to verify the claim?

Return your response as a json dictionary , following
this structure : {"answer":<Yes/No>, "rationale ":
<str>}

<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
CLAIM: {claim}
PARAGRAPH: {text}

A.9 Phase 3 evaluation with judge model

<| begin_of_text |><| start_header_id |>system<|
end_header_id|>

Given a claim and a paragraph, answer the following
question .

Is the claim {SUPPORTED/REFUTED} by the paragraph
?

Return your response as a json dictionary , following
this structure : {"answer":<Yes/No>, "rationale ":
<str>}

<| begin_of_text |><| start_header_id |>user<|
end_header_id|>
CLAIM: {claim}
PARAGRAPH: {text}

B Claim Quality Questionnaire

Id	Question
Q1	Is the claim grounded by the original text?
Q2	Is the claim grammatically correct?
Q3	Does the claim have all the necessary components (subject, predicate, and relevant qualifiers) to form a complete thought?
Q4	Is the claim precise and specific rather than vague?
Q5	Does the claim introduce new information rather than just restating common knowledge?
Q6	Is the claim concise without losing essential information?
Q7	Does the claim provide enough information to be understood independently?
Q8	Would verifying the claim add value to public knowledge?

Table 7: Questions asked to the LLM judge and human annotators to evaluate the quality of the generated claims.