

Uncertainty-Aware Web-Conditioned Scientific Fact-Checking

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Abstract

Scientific fact-checking is vital for assessing claims in specialized domains such as biomedicine and materials science, yet existing systems often hallucinate or apply inconsistent reasoning, especially when verifying technical, compositional claims against an evidence snippet under source and cost/latency constraints. We present a pipeline centered on *atomic* predicate–argument decomposition and *calibrated, uncertainty-gated* corroboration: atomic facts are aligned to local snippets via embeddings, verified by a compact evidence-grounded checker, and only facts with uncertain support trigger domain-restricted web search over authoritative sources. The system supports both binary and tri-valued classification where it predicts labels from SUPPORTED, REFUTED, NEI for three-way tasks. We evaluate under two regimes, *Context-Only* (no web) and *Context+Web* (uncertainty-gated web corroboration); when retrieved evidence conflicts with the provided context, we abstain with NEI rather than overriding the context. On multiple benchmarks, our framework surpasses the strongest benchmarks. In our experiments, web corroboration was invoked for only a minority of atomic facts on average, indicating that external evidence is consulted selectively under calibrated uncertainty rather than routinely. Overall, coupling atomic granularity with calibrated, uncertainty-gated corroboration yields more interpretable and context-conditioned verification, making the approach well-suited to high-stakes, single-document settings that demand traceable rationales, predictable cost/latency, and conservative abstention under domain and evidence grounding constraints.

CCS Concepts

• **Computing methodologies** → **Search with partial observations; Information extraction; Natural language generation.**

Keywords

Natural language generation, Fact Checking, Scientific Claim Uncertainty aware verification, Evidence Grounding

1 Introduction

Scientific fact-checking is crucial in high-stakes domains such as biomedicine, public health, and materials science. However, state-of-the-art language models still struggle to verify technical statements: they often rely on surface cues, fail to correctly interpret predicate argument structure in complex sentences, and hallucinate when local evidence is sparse. As a result, their judgments can be inconsistent and weakly justified. This creates a need for methods that can verify technical claims from small, potentially noisy snippets, provide compute-efficient and explainable verdicts, and consult trusted web sources only when local evidence is insufficient.

We present a modular, context-aware, evidence-grounded pipeline. Claims are decomposed into atomic predicate argument facts, aligned to the most relevant evidence spans, and evaluated by a lightweight verifier. When uncertainty is high, a controlled corroboration step queries authoritative domains; all signals are aggregated into an interpretable verdict. This couples symbolic decomposition with probabilistic verification and targeted retrieval, balancing precision and coverage.

Empirically, the pipeline yields consistent test-time gains. On **BIONLI-300**, atomic decomposition plus web search improves factual F_1 by **+6.0** (66.7% vs. 60.7%) and Balanced Accuracy by **+7.35** points over sentence-level verification with *MiniCheck*. Uncertainty-triggered corroboration adds a further **+4.7** F_1 and raises **PubMedFact1k** Macro- F_1 by **+1.2** over the strongest benchmark, reducing ambiguous NEI↔(Supported/Refuted) cases. The same pipeline transfers to climate fact-checking, achieving strong performance on CLIMATE-FEVER under both **CONTEXT-ONLY** (text-only) and **CONTEXT+WEB** (constrained corroboration) settings with identical thresholds and prompts. Because our system is built around the same compact verifier (*MiniCheck*) and differs only in *structure* (atomic facts), *selection* (local snippets), and *confident search* (uncertainty-gated corroboration), these gains isolate the value of grounded verification rather than larger models, improving accuracy, interpretability, and cost.

We recommend our pipeline when: (i) **traceability and abstention matter** (per-atom rationales and NEI abstentions when web and local evidence disagree); (ii) **cost/latency is constrained** (a compact verifier in the inner loop and retrieval only for uncertain atoms); (iii) **source control is required** (web corroboration restricted to vetted domains such as NIH/WHO/CDC). In short, choose **ATOMIC+SEARCH** when you need *explanations and predictable cost* under domain and provenance constraints.

To support further research, we release **PUBMEDFACT1K**, a three-way (SUPPORTED/REFUTED/NEI) medical claim verification dataset derived from the **PUBMEDQA** dataset [9]. Together, these results show that combining atomic reasoning, calibrated uncertainty handling, and domain-aware corroboration enables more faithful and reproducible scientific fact-checking.

2 Related Work

Closed-book QA suggests that autoregressive transformers retain substantial factual knowledge, but whether such parametric memory alone suffices for veracity classification remains unclear. On *SciFact*, zero-shot GPT-4 yield promising results with a few in-domain exemplars [1], yet performance drops substantially on *SciFact-Open* and related COVID/fake-news settings without retrieval [6, 8, 15], motivating retrieval with grounded checking. Prior approaches such as post-hoc revision (RARR, “research→revise”) [7], broad tool-augmented checking (FACTOOL, Web/Scholar/Python) [3], and RAG

	Atomic+Search	RARR	FacTool	Provenance	Base
Explicit typed atomic facts	✓	✗	✗	✗	✗
Uncertainty-gated retrieval	✓	✗	✗	✗	✗
Domain-restricted external evidence	✓	✗	✗	n/a	n/a
Compact verifier (NLI)	✓	✗	✗	✓	✓
LLM judge aggregation	✓	✗	✓	✗	✗
Post-generation editing objective	✗	✓	✗	✗	✗
Tri-valued labels with NEI	✓	✗	✗	✗	✓

Table 1: Comparison of recent fact-checking methods. ✓ indicates the feature is present, ✗ indicates it is absent, and n/a means not applicable. “Base” represents a typical retrieve–verify pipeline.

provenance verification via cross-encoder NLI [13] do not explicitly structure verification. In contrast, our ATOMIC+SEARCH decomposes claims into ≤ 25 -word predicate–argument units inspired by OpenIE, entailment graphs, and PropBank SRL [5, 10, 11]; selects localized evidence windows; and verifies each atomic fact with a compact MiniCheck-style evidence-grounded model [14], invoking domain-restricted web corroboration only under calibrated uncertainty. This atomic, uncertainty-gated loop unifies symbolic decomposition, probabilistic verification, and controlled corroboration in a single transparent pipeline, yielding finer-grained rationales and clearer failure modes (scope, negation, numerical qualifiers) than sentence-level verifiers and post-generation editors [3, 7, 13], as summarized in Table 1.

3 Method

We propose an *Atomic+Search* pipeline for scientific fact checking that couples (i) atomic fact decomposition, (ii) semantic local-evidence selection, (iii) lightweight verification with a grounded verifier, (iv) uncertainty-triggered, domain-constrained web corroboration, and (v) a final deductive judge. Given a natural-language claim c and evidence D (e.g., a paragraph or abstract), our system outputs a binary/ Tri-Valued verdict $\{\text{TRUE}, \text{FALSE}, \text{NEI}\}$ and a short rationale.

3.1 Problem Formulation

Let c be a claim and D the associated evidence context. We decompose c into a set of atomic facts $\mathcal{F} = \{f_i\}_{i=1}^m$, where each f_i targets a single predicate–argument tuple. For each f_i , a verifier returns a support probability $p_i \in [0, 1]$ given evidence D :

$$p_i = \text{MiniCheck}(f_i | D). \quad (1)$$

Facts are labeled SUPPORTED, REFUTED and UNCERTAIN based on the grounding probabilities. Facts with $p_i \in [\alpha, \beta]$ (the *Uncertain band*) trigger targeted web corroboration before a final verdict is produced.

Throughout, we adopt a *single-document* verification regime: for each claim we assume exactly one evidence document D is available. In our intended use cases, D is either (i) the artifact supplied by the task, or (ii) the user’s *top retrieved evidence* the highest-ranked page/abstract they have already opened after a standard search. This isolates the contribution of atomic decomposition and calibrated verification from open-domain retrieval, mirrors how people assess claims while reading a single page, and avoids mixing potentially conflicting sources. When D is insufficient, our uncertainty-gated

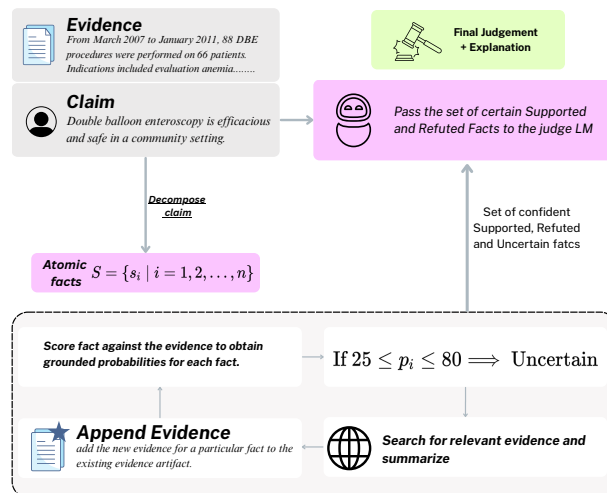


Figure 1: Atomic+Search pipeline

corroboration step may consult additional sources; if those conflict with D , we abstain with UNCERTAIN rather than overwrite the provided context.

3.2 Datasets

We evaluate our framework on three fact-checking datasets where each claim is paired with scientific evidence and a gold label, emphasizing grounded justification and traceable provenance. Unlike standard fact-checking benchmarks, scientific claim verification often requires decomposing claims and corroborating sub-claims across multiple sources to establish true evidential support. **PubMedFact1k**¹ is a 1,000 example biomedical claim verification set derived from the human-annotated PubMedQA PQA-Labeled split [9], where each question is rewritten as a declarative claim paired with its abstract and *yes/no/maybe* are mapped to SUPPORTED/REFUTED/NEL. **BIONLI-300** is a 300-example subset of BIONLI that treats each hypothesis as the claim and its source abstract as context, mapping *entailment* to SUPPORTED and *contradiction* to REFUTED [2]. **CLIMATE-FEVER** provides 1,535 climate-related claims, each linked to five Wikipedia evidence sentences; we merge the five sentences into a single document and evaluate only on the SUPPORTED and REFUTED subsets [4].

3.3 Uncertainty-Aware Web Corroboration

Our pipeline (Fig. 1) takes a claim c and document D and produces a structured, claim-level verdict through uncertainty-aware stages. First, a reasoning-capable LLM performs **atomic fact decomposition**, converting (c, D) into short atomic facts $\{f_i\}$ (each ≤ 25 words, expressing a single predicate–argument tuple) and returning strict JSON of the form `facts[$\{id, text, targets\}$]`. This yields consistent, fine-grained units for downstream verification.

Next, **semantic snippet selection** identifies local evidence for each fact. The document is chunked into overlapping ≈ 420 -character windows $\{x_j\}$. For each fact f_i , we form a query from its text and targets, compute embeddings for all queries and chunks

¹<https://huggingface.co/datasets/umbc-scify/PubmedFact1k>

using text-embedding-3-large, and select the window with highest cosine similarity; if embeddings are unavailable, we fall back to a deterministic token-overlap heuristic. Given fact-snippet pairs, a lightweight verifier, *MiniCheck-7B*, produces a calibrated support probability for each fact. Facts with probability ≥ 0.80 are labeled SUPPORTED, those ≤ 0.25 as REFUTED, using binary calibration instead of multi-class NLI to simplify aggregation and reduce label drift across biomedical domains.

Uncertainty-triggered web corroboration is applied only to facts with intermediate scores between 0.25 and 0.80. For these, we issue concise web queries that restate the fact and include its local snippet as context, restricting search to authoritative biomedical sources (e.g., PubMed, WHO, CDC, FDA, NIH, ClinicalTrials.gov, Wikipedia). Retrieved snippets are summarized with inline citations into an auxiliary text, which is concatenated with the snippet to form augmented evidence; *MiniCheck-7B* is re-run on this augmented evidence, and we record both the original and updated probabilities and whether rescoring occurred. The 0.25–0.80 band is chosen from a held-out calibration study of *MiniCheck* [12], where most ambiguous cases concentrate.

Finally, a **Judge LLM** aggregates fact-level outputs into a claim-level decision. It receives the original claim and document along with two fact sets after corroboration: \mathcal{S} , facts labeled SUPPORTED (probability ≥ 0.80), and \mathcal{R} , facts labeled REFUTED (probability ≤ 0.25). Facts with intermediate probabilities are omitted for two-way decisions but reported as UNCERTAIN in a three-way setting. Conditioned on these inputs, the Judge outputs strict JSON with fields `final_verdict` (SUPPORTED, REFUTED, or NEI), `explanation` (a short rationale referencing fact IDs), and `used_facts`.

3.4 Implementation Details

We split D into sentence groups with a maximum length of approximately 420 characters per chunk, trading off locality against coverage. For both decomposition and judging we use a chat LLM in JSON mode, while a high-capacity text-embedding model is employed for embeddings and *MiniCheck-7B* is used for verification. We fix the high-confidence threshold at $\tau_{hi}=0.80$, define an uncertainty band of $[0.25, 0.80]$, and cap the maximum fact length at 25 words. Web-based corroboration is restricted to reputable scientific domains, including PubMed, WHO, CDC, FDA, NIH and ClinicalTrials.gov, with the domain list configurable to match the target area (e.g., materials science). For each example we record the baseline verdict on c , per-fact probabilities before and after corroboration, the synthesized web evidence and citations, and the final verdict together with an explanation.

4 Experimental Setup

We evaluate ATOMIC+SEARCH in two claim verification regimes: a binary setting (SUPPORTED/REFUTED) and a three-way setting that additionally includes NEI. We compare against four families of approaches: (i) **compact verifiers and sentence-level baselines**: *MiniCheck* (sentence-level), a calibrated NLI-style checker; (ii) **closed-book and tool-augmented LLMs**: *GPT-5 Mini*, *Qwen-32B (Instruct/MAD)*, and *GPT-5 Mini + Search* (generic web-augmented prompting); (iii) **recent retrieval- and verification-centric systems**, *RARR* (retrieve-revise attribution editing), and

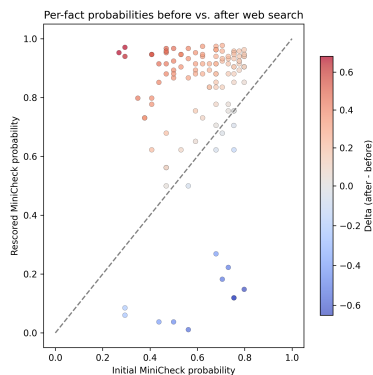


Figure 2: Distribution of per-fact support p_i before and after web corroboration on BIONLI-300.

(iv) **reasoning-focused LMs: o1** (reasoning model with long-context verification). Our variants include the full **Atomic+Search** pipeline and ablations (Table 3).

4.1 Main Results

Table 2 reports results on **BIONLI-300** (2-way), **PubMedFact1k** (3-way), and **CLIMATE-FEVER** (2-way). On **BIONLI-300**, Our framework performs best overall, markedly improving Balanced Accuracy and F_1 over sentence-level *MiniCheck* and tool-augmented LLM baselines; since *MiniCheck* is our underlying verifier, these gains isolate the impact of atomic decomposition and *confident search*. On **PubMedFact1k**, it achieves the top Macro- F_1 , outperforming *GPT-5 Mini + Search* and *GPT-o1*, and reducing borderline NEI vs. SUPPORTED/REFUTED errors. On **CLIMATE-FEVER**, it again leads in Balanced Accuracy and F_1 , with consistent gains over both compact verifiers and larger LLMs, indicating strong out-of-domain generalization.

4.2 Per component Importance

On BIONLI-300 (Table 3), removing web corroboration (NO-SEARCH) reduces F_1 from 66.7% to 62.0% (−4.7); removing atomic decomposition (NO-ATOMIC) further drops F_1 to 60.3% (−6.4), indicating decomposition is the larger contributor. Replacing the Judge with majority vote (MAJORITYVOTE) yields 52.1% F_1 (−14.6), showing that calibrated fact-level aggregation is crucial. Overall, decomposition, corroboration, and principled aggregation are complementary.

4.3 Retrieval and Corroboration Analysis

Figure 2 plots per-fact support probabilities before and after corroboration on BIONLI-300. The distribution shifts rightward for many facts, reflecting successful evidence augmentation when local context is insufficient. Most of the helpful external evidence comes from authoritative medical sources (e.g., PubMed/NIH/NCBI), with Wikipedia contributing recall for broad biomedical background.

System	BIONLI-300			PubMedFact1k (3-way)	CLIMATE-FEVER (2-way)		
	Bal. Acc. \uparrow	Rec. \uparrow	F1 \uparrow	Macro F1 \uparrow	Bal. Acc. \uparrow	Rec. (S) \uparrow	F1 (S) \uparrow
GPT-o1	65.8%	60.2%	64.9%	71.2%	69.70%	56.10%	65.05%
MiniCheck	61.35%	59.8%	60.7%	—	69.10%	55.00%	64.00%
GPT-5 Mini	62.9%	58.7%	61.8%	68.5%	67.90%	54.20%	63.10%
Qwen 32B MAD	62.5%	59.1%	61.3%	61.8%	67.30%	53.60%	62.60%
RARR	66.4%	62.3%	65.3%	72.3%	70.40%	57.80%	66.30%
GPT-5 Mini + Search	66.9%	62.7%	65.8%	72.5%	71.20%	58.50%	67.00%
Atomic+Search (ours)	68.7%	65.4%	66.7%	73.7%	73.83%	62.14%	70.04%

Table 2: Main results. Balanced Accuracy, Recall, and F1 on BIONLI-300; Macro-F₁ on PUBMEDFACT1K; and Balanced Accuracy, Recall (Supports), and F1 (Supports) on CLIMATE-FEVER, evaluated only on the SUPPORTED/REFUTED subsets.

BIONLI-300	F ₁ \uparrow	Δ vs. full
Atomic+Search (full)	66.7%	—
NO-SEARCH	62.0%	− 4.7
NO-ATOMIC	60.3%	− 6.4
MAJORITYVOTE (NO-JUDGE)	52.1%	− 14.6

Table 3: Ablations on BIONLI-300. Each line removes one component from the full pipeline.

5 Conclusion

We introduced an uncertainty-aware, web-conditioned pipeline for scientific fact-checking at predicate–argument (“atomic”) granularity: claims are decomposed into minimal facts, each aligned to a local snippet, verified with a compact evidence-grounded checker, and, only under calibrated uncertainty, supplemented by domain-restricted corroboration before a final judge combines high-confidence atoms into an interpretable claim-level verdict. Across biomedical and out-of-domain evaluations, ATOMIC+SEARCH consistently outperforms sentence-level verifiers, tool-augmented LLMs, and recent retrieval+verification benchmarks, while ablations show that atomic decomposition, uncertainty-gated corroboration, and the judge are complementary, corroboration is needed for only a minority of atoms, and the system abstains with NEI when retrieved evidence conflicts with the provided context (Tables 2–3). Our approach still relies on fixed calibration thresholds, curated domain allow-lists, and heuristic snippet selection. Future work includes adaptive calibration; tighter integration of decomposition, selection, and verification; richer temporal and quantitative reasoning; improved source-quality modeling; and human-in-the-loop workflows that surface atomic rationales. We hope that combining atomic structure with calibrated corroboration, together with our released components and PUBMEDFACT1K, will spur more robust, context-aware scientific fact-checking.

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