

# DEEPMED Search: An Open-Source Agentic Platform for Medical Deep Research with Introspective Verification

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## Abstract

Navigating the deluge of heterogeneous medical data, from academic literature (PubMed) to clinical guidelines (Web) and private knowledge bases remains a critical bottleneck for evidence-based medicine. While commercial black-box tools lack transparency, standard open-source RAG implementations frequently suffer from “reasoning drift” when handling complex, long-tail queries. We present **DEEPMED SEARCH**, a fully open-source, agentic platform designed for transparent medical deep research. Built on a high-performance NEXTJS architecture, DEEPMED SEARCH features a source-adaptive router that autonomously dispatches sub-queries to PubMed, web search, or local graph-based knowledge bases based on information density. Crucially, the platform integrates an introspective verification module, powered by a causal-consistent multi-agent debate framework, to validate retrieved evidence against diagnostic logic before synthesis. To demonstrate its robustness, we showcase DEEPMED SEARCH’s ability to autonomously decompose high-difficulty rare disease queries, filter out confounding noise, and generate structured, citation-backed research reports in minutes. By open-sourcing this software, we provide the community with a robust infrastructure to democratize access to trustworthy, glass-box medical reasoning in research and prototyping settings, which is publicly available at: <https://www.deepmedsearch.cloud> and the demonstration video is available at: <https://youtu.be/4U4aok8yLpk>.

## 1 Introduction

Large Language Models (LLMs) have revolutionized clinical question answering [Singhal *et al.*, 2025; Tu *et al.*, 2023; Yang *et al.*, 2022], yet their deployment in high-stakes evidence-based medicine remains hindered by limited transparency and auditability [Ji *et al.*, 2023; Rudin, 2019]. Medical deep research requires synthesizing heterogeneous data from academic literature, constantly updating clinical

guidelines, and structured patient records. While Retrieval-Augmented Generation (RAG) is widely adopted to ground model outputs [Lewis *et al.*, 2020; Karpukhin *et al.*, 2020; Izacard and Grave, 2021], standard RAG implementations often rely on static, linear pipelines that are vulnerable to *Retrieval-Induced Reasoning Drift*, especially in complex, long-tail scenarios such as rare disease diagnosis [Mallen *et al.*, 2023]. In these contexts, standard semantic retrievers suffer from commonality bias [Kandpal *et al.*, 2023], preferentially surfacing generic evidence that supports common mimic conditions and causing the LLM to be hijacked by plausible but non-discriminative distractors. Although commercial AI research tools attempt to address this issue, their black-box nature prevents clinicians from auditing the causal logic behind retrieved evidence, which remains a non-negotiable requirement in clinical workflows.

To democratize trustworthy and auditable AI in healthcare, we introduce DEEPMED SEARCH, a fully open-source, agentic platform engineered for transparent medical deep research. Moving beyond simple RAG frameworks, DEEPMED SEARCH introduces a source-adaptive router that autonomously orchestrates sub-queries across heterogeneous endpoints, including PubMed APIs, live web search, and a local Neo4j-based knowledge graph. Furthermore, to ensure the causal alignment and faithfulness of the retrieved evidence, the platform features an introspective verification module [Kiciman *et al.*, 2024]. This engine employs contrastive hypothesis expansion to filter confounding noise and utilizes a multi-agent adversarial debate framework [Du *et al.*, 2024; Yao *et al.*, 2022] to rigorously stress-test reasoning paths before final synthesis.

In this paper, we showcase the system’s glass-box interactivity, while clinicians can observe DEEPMED SEARCH as it autonomously decomposes complex queries, routes searches, and conducts real-time multi-agent deliberations. Ultimately, the system generates structured, citation-backed research reports while maintaining complete interpretability [Xiong *et al.*, 2024].

## 2 System Architecture

DEEPMED SEARCH is engineered as a modular, agentic workflow designed to transform high-level clinical inquiries into structured research insights. As illustrated in Figure 1, the platform transitions from autonomous data orchestration

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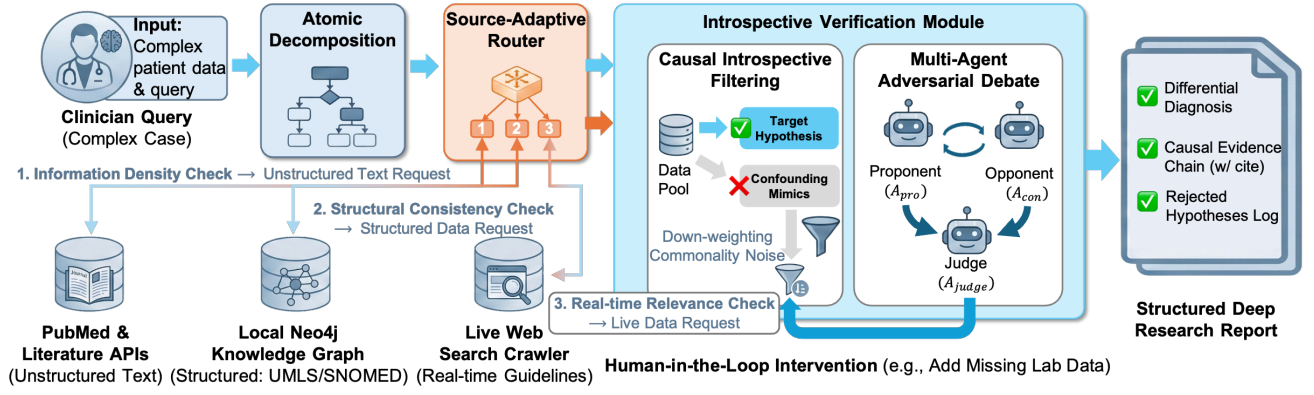


Figure 1: System architecture of DEEPMED SEARCH. The platform integrates a source-adaptive router for multi-endpoint evidence retrieval, causal introspective filtering to mitigate commonality bias, and a multi-agent debate engine to verify reasoning and synthesize reports.

to rigorous causal verification across three primary stages: Source-Adaptive Routing, Causal Introspective Filtering, and Multi-Agent Synthesis.

### 2.1 Source-Adaptive Routing

A core pillar of DEEPMED SEARCH is its ability to navigate heterogeneous data landscapes. Unlike monolithic RAG systems, our platform incorporates a **Source-Adaptive Router** that treats the information retrieval process as a sequential decision-making task.

Upon receiving a clinical query, the system first performs *Atomic Decomposition*, breaking the input into distinct clinical constraints (e.g., symptomatic triad, temporal progression, and treatment resistance). The Router then dynamically evaluates the information density of each sub-query to dispatch them to optimal endpoints:

- **Scientific Evidence:** Direct API integration with **PubMed** for fetching peer-reviewed literature and clinical trial abstracts [Qiu *et al.*, 2025].
- **Pathophysiological Logic:** Querying a local NEO4J knowledge graph (integrating UMLS and SNOMED-CT) to identify multi-hop causal links between symptoms [Pearl, 2009].
- **Real-time Guidelines:** Utilizing a **Web Search** crawler to retrieve the latest consensus statements and medical news.

### 2.2 Causal Introspective Filtering

To mitigate the *Commonality Bias* identified in clinical LLMs [Kandpal *et al.*, 2023], the platform utilizes a *Causal Introspective Filtering* mechanism. This stage aligns the retrieved evidence with diagnostic logic via two steps: **1) Contrastive Hypothesis Expansion:** The system proactively formulates “confounding hypotheses”, prevalent diseases that mimic the target long-tail condition (e.g., generating “Essential Hypertension” when analyzing potential “Pheochromocytoma”) [Park and Lee, 2024]. **2) Evidence Down-weighting:** Chunks are filtered based on a discriminative score  $S_i = \frac{\text{sim}(c_i, H_{\text{target}})}{\text{sim}(c_i, H_{\text{mimic}}) + \epsilon}$ . Here,  $\text{sim}(\cdot, \cdot)$  denotes the cosine similarity between BGE-M3 sentence embeddings, and  $\epsilon$

is set to  $10^{-6}$  to avoid division by zero. A candidate chunk is regarded as discriminative when  $S_i$  is larger than the threshold  $\tau_s$ ; chunks failing to meet a dynamic threshold are tagged as “non-discriminative noise” and visually dimmed in the interface, ensuring the model’s attention remains on the long-tail signals.

### 2.3 Multi-Agent Synthesis

The final reasoning phase is handled by an introspective verification module, instantiated as a multi-agent adversarial debate framework [Chan *et al.*, 2023]. The system deploys three distinct agents: a **Proponent** ( $A_{pro}$ ) that constructs reasoning chains from evidence to diagnosis, an **Opponent** ( $A_{con}$ ) that identifies missing causal links or inconsistencies, and a **Judge** ( $A_{judge}$ ) that delivers the final verdict once reasoning converges (Information Gain  $< \tau$ ).

Crucially, rather than producing a single-line answer, the synthesis engine generates a structured research report. These reports are citation-backed and organized into sections: Differential Diagnosis, Supporting Evidence (from both Graph and Text), and Rejected Hypotheses. This “glass-box” output allows clinicians to audit the platform’s logic in real-time.

## 3 Implementation Details

The DEEPMED SEARCH backend is engineered for high-throughput concurrent processing to support real-time clinical deep research workflows. **Frontend & Orchestration:** Built on NEXT.JS, the frontend integrates a *WebSocket*-driven UI for streaming real-time agent deliberations. A dedicated orchestration layer handles the concurrent dispatch of sub-queries via the Source-Adaptive Router. **Knowledge Engines:** Structured multi-hop reasoning is powered by a NEO4J graph database integrating UMLS and SNOMED-CT. For unstructured evidence, we use MILVUS to index the MIRAGE corpus (Textbooks and PubMed) [Xiong *et al.*, 2024], utilizing BGE-M3 for dense vector similarity coupled with a lightweight cross-encoder for precise re-ranking. **LLM Serving:** To meet the computational demands of deploying three concurrent agents (Proponent, Opponent, Judge), we serve models like QWEN2.5-14B and DEEPSEEK-V3 via VLLM,

**Query:** “Patient presents with paroxysmal hypertension, headache, palpitations, and sweating. Standard anti-hypertensive drugs ineffective.”

× **Vanilla RAG Response:**

Diagnosis: **Essential Hypertension**. Treatment involves lifestyle changes...

[Error: Fixates on common disease; ignores paroxysmal & sweating cues]

✓ **DEEPMED SEARCH (Ours):**

Diagnosis: **Pheochromocytoma**.

**Reasoning:** 1. *Contrastive Filter:* ‘Essential Hypertension’ down-weighted (cannot explain sweating). 2. *Debate Verdict:* Symptom triad strongly indicates catecholamine-secreting tumor.

Table 1: Case study: DEEPMED SEARCH actively filters common disease noise (Hypertension) to correctly identify the rare condition.

leveraging *PagedAttention* to optimize memory footprint and maximize throughput. **Report Generation Engine:** A custom synthesis module compiles the verified evidence into a structured markdown/PDF report, automatically mapping reasoning steps to specific reference nodes (PubMed IDs or Graph edges) to ensure full traceability.

## 4 Demonstration Scenarios

We present using real-world rare disease cases from the MedR-Bench dataset, as exemplified in Table 1. The web interface allows clinicians to input complex case reports and interactively audit the deep research process.

### 4.1 Scenario 1: Autonomous Routing & Causal Filtering

Consider a patient presenting with episodic hypertension, headaches, and palpitations. A standard RAG system typically retrieves generic guidelines for “Primary Hypertension”, leading to reasoning drift. In DEEPMED SEARCH: **1) Routing in Action:** The Router identifies the temporal cue (“episodic”) and dispatches parallel queries to Neo4j and PubMed. **2) Noise Filtering:** The “Evidence Panel” marks generic hypertension evidence with a *Low Relevance* tag (visualized as dimmed text), explicitly stating it was down-weighted due to overlap with common confounders. Conversely, evidence pointing to *Pheochromocytoma* is dynamically highlighted.

### 4.2 Scenario 2: Glass-Box Debate & Report

For complex diagnoses, the clinician can activate the introspective debate mode. **1) Real-Time Deliberation:** As illustrated in Figure 2, a streaming panel opens where  $A_{pro}$  argues for Pheochromocytoma citing paroxysmal symptoms, while  $A_{con}$  challenges the evidence density. **2) Human-in-the-Loop:** Crucially, the clinician can inject missing parameters (e.g., “Add lab result: elevated metanephrines”) directly into the debate stream. **Structured Output:** The  $A_{judge}$  finalizes the verdict, and the system instantly compiles a deep research report, linking the diagnosis to verified graph nodes.

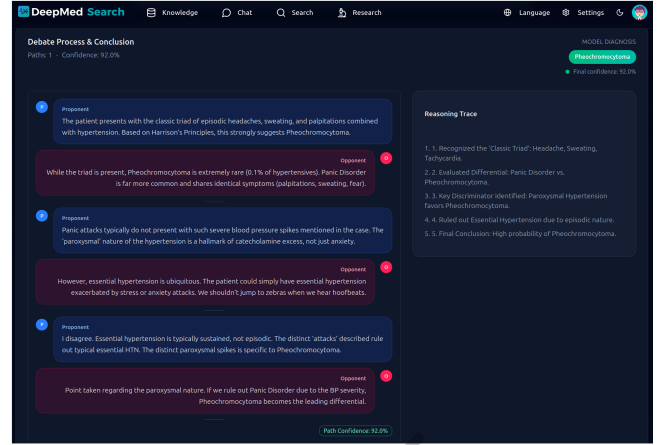


Figure 2: The Multi-Agent Debate Panel. Users can monitor agents as they argue over the diagnosis, providing transparent reasoning logic.

## 5 Evaluation

As summarized in Table 2, DEEPMED uniquely integrates source-adaptive routing and multi-agent verification, bridging the gap between vanilla RAG and commercial black-box tools.

**Performance Gains:** We evaluated on the MIRAGE benchmark [Xiong *et al.*, 2024]. DeepMed Search consistently outperforms standard RAG baselines. On the challenging PubMedQA subset, our method improves accuracy by approximately 9.0% over vanilla RAG (using Qwen3-8B). For rare diseases, experiments on the MedR-Bench dataset show that our approach achieves an accuracy of 76.43% (using DeepSeek-v3.2), bridging the gap between automated systems and expert-level reasoning.

| Capabilities                 | Vanilla RAG | Commercial | Ours |
|------------------------------|-------------|------------|------|
| Open-Source Transparency     | ✓           | ×          | ✓    |
| Adaptive Routing (Web/Graph) | ×           | ✓          | ✓    |
| Causal-Consistent Filtering  | ×           | ×          | ✓    |
| Multi-Agent Verification     | ×           | ×          | ✓    |
| Structured Deep Reports      | ×           | ✓          | ✓    |

Table 2: Feature comparison. DEEPMED SEARCH bridges glass-box transparency with practical research-oriented system capabilities.

## 6 Conclusion

DEEPMED SEARCH provides a transparent, agentic platform for medical deep research. It combines source-adaptive routing, causal-consistent filtering, and multi-agent verification to mitigate retrieval-induced reasoning drift in medical LLMs [Park and Lee, 2024] and generate structured, citation-backed reports. Current limitations include additional computational and latency overhead and the lack of a formal clinician user study. Future work will improve efficiency, involve clinicians in evaluation, and extend the framework to multimodal data such as medical imaging [Wang *et al.*, 2025].

## Acknowledgments

This work was supported by the National Natural Science Foundation of China under Grant No. 62306173.

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