DDxTutor: Clinical Reasoning Tutoring System with Differential Diagnosis-Based Structured Reasoning

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Abstract

Clinical diagnosis education requires students to master both systematic reasoning processes and comprehensive medical knowledge. While recent advances in Large Language Models (LLMs) have enabled various medical educational applications, these systems often provide direct answers that could reduce students' cognitive engagement and lead to fragmented learning. Motivated by these challenges, we propose DDxTutor, a framework that follows differential diagnosis principles to decompose clinical reasoning into teachable components. It consists of a structured reasoning module that analyzes clinical clues and synthesizes diagnostic conclusions, and an interactive dialogue framework that guides students through this process. To enable such tutoring, we construct DDxReasoning, a dataset of 933 clinical cases with fine-grained diagnostic steps verified by doctors. Our experiments demonstrate that fine-tuned LLMs achieve strong performance in generating structured teaching references and conducting interactive diagnostic tutoring dialogues. Human evaluation by medical educators and students validates the framework's potential and effectiveness for clinical diagnosis education. Our project is available at https://github.com/med-air/DDxTutor.

1 Introduction

Clinical diagnosis, a core task in medical practice, involves synthesizing clinical information to reach a conclusion. As this process demands both depth and precision, systematic and scientific reasoning becomes indispensable (Fauci et al., 2008). For these demands, modern medical education has placed great emphasis on fostering clinical reasoning skills in students, aiming to strengthen the rigor and scientific foundation of diagnostic thinking (Schmidt and Mamede, 2015). Recent advances

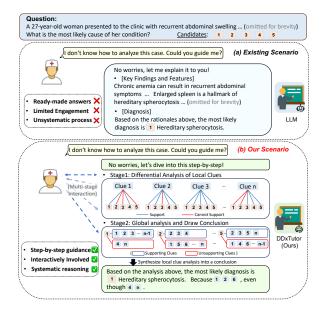


Figure 1: Comparison of clinical diagnostic training scenarios. (a) The existing scenarios usually provides direct feedback with immediate answers and explanations, where students could become passive learners with an unsystematic knowledge intaking process. (b) Our proposed framework follows the thought of Differential Diagnosis (DDx), where medical students interactively engage in two steps: (1) independent analysis of individual clinical clues, and (2) global synthesis of findings to reach a diagnostic conclusion. Our structured approach enables systematic diagnostic reasoning while maintaining active student engagement with step-by-step guidance throughout the learning process.

in Natural Language Processing (NLP), particularly Large Language Models (LLMs) like Chat-GPT (Achiam et al., 2023) and DeepSeek (Liu et al., 2024a), have enabled new possibilities for interactive clinical education. These models have been applied to various medical educational tasks, from answering medical questions (Gilson et al., 2023) to providing feedback on junior doctors' diagnostic decisions (Huang et al., 2024).

However, current medical dialogue systems and diagnostic support tools still face limitations when

Proceedings of the 63rd Annual Meeting of the Association for Computational Linguistics (Volume 1: Long Papers), pages 30934–30957 July 27 - August 1, 2025 ©2025 Association for Computational Linguistics

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applied to clinical reasoning education. Firstly, to the best of our knowledge, many existing medical LLMs are designed with a focus on providing direct diagnoses accompanied by accurate and reasonable explanations (Sviridova et al., 2024; Tian et al., 2024). While these professional explanations serve as valuable references, they often present reasoning as a complete, ready-made output rather than a step-by-step process that students can actively participate in. From an educational perspective (Sweller, 1988), this approach could reduce students' cognitive engagement, turning them into passive recipients rather than active participants in the diagnostic process. Although multi-turn dialogue systems allow students to ask follow-up questions about unclear explanations, this questiondriven approach often leads to an unsystematic learning process (Barrows, 1986). As novice learners are still developing their clinical knowledge framework, their questions tend to address immediate comprehension gaps rather than following a structured diagnostic reasoning path, potentially missing critical diagnostic elements and relationships that are essential for building comprehensive clinical thinking skills (Schmidt and Mamede, 2015).

Facing these limitations in current medical education systems, we aim to devise a new framework that leverages LLMs to promote systematic clinical reasoning education with active student engagement across multiple diagnostic stages. To structure such an educational experience, we draw inspiration from Differential Diagnosis (DDx) (Fauci et al., 2008; First, 2013), a methodical process of weighing clinical evidence to identify the most likely diagnosis among various possibilities. In the first phase, individual patient clues are analyzed sequentially, examining their meaning, indication, specificity, and any supporting or unsupporting relationships with potential diagnoses. In the second phase, each analysis is synthesized to comprehensively evaluate diagnostic possibilities, arrive at a final conclusion, and offer explanations based on core evidence.

To operationalize this framework, we curated *DDxReasoning*, a dataset containing 933 doctorverified clinical reasoning chains that capture these systematic diagnostic steps. This dataset serves dual purposes: it provides fine-tuning data to enhance LLMs' ability to generate educational reasoning chains as tutoring references. Moreover, our aims extend beyond having LLMs that merely follow these reasoning patterns. To enable interactive teaching, we further construct simulated teacher-student dialogues based on it. These dialogues demonstrate how to effectively guide students through the DDx process, enabling LLMs to conduct interactive clinical teaching while maintaining alignment with the structured reasoning objectives in *DDxReasoning*. Our main contributions are as follows:

(1) We propose *DDxTutor*, the first clinical diagnostic tutoring framework that leverages modern LLMs to implement systematic differential diagnosis teaching. By decomposing the diagnostic process into sequential clue analysis and global synthesis phases, our framework enables structured and interactive medical reasoning education.

(2) We construct *DDxReasoning*, a comprehensive dataset containing 933 expert-verified clinical cases with fine-grained diagnostic reasoning chains. This dataset provides high-quality fine-tuning data for enhancing LLMs' capability to generate structured diagnostic analyses that serve as educational references. Building upon this structured knowledge base, we further develop a dialogue generation approach that simulates teacher-student interactions aligned with the DDx process, enabling LLMs to provide guided diagnostic tutoring.

(3) Extensive experiments demonstrate the effectiveness of our approach from two complementary aspects: (a) LLMs fine-tuned on *DDxReasoning* demonstrate strong capabilities in comprehensive clinical scene analysis, providing structured analytical processes that serve as useful teaching references; (b) The derivative dialogue tutoring system effectively performs student response tracking and analysis, providing adaptive explanations that guide students through the clinical reasoning process. These results validate our framework's potential and ability to support systematic clinical reasoning education through both structured knowledge representation and interactive guidance.

2 Related Works

2.1 NLP for Medical Education

Medical education has emerged as a prominent research direction in Natural Language Processing (NLP). A notable application is the development of virtual patients powered by language models (Danforth et al., 2009; Menendez et al., 2015; Campillos-Llanos et al., 2020; Ali et al., 2021), which simulate authentic patient interactions to facilitate clinical training. These systems have demonstrated significant potential in enhancing medical education (Shi et al., 2024; Li et al., 2024b). In parallel, researchers have explored ways to improve patients' comprehension of medical instructions (Cai et al., 2023; Yao et al., 2024a). For instance, Chat-Coach (Huang et al., 2024) functions as an AI copilot, helping healthcare providers refine their patient communication skills. Beyond these specialized educational tools, general-purpose medical questionanswering systems (Yao et al., 2024b; Li et al., 2024a; Liu et al., 2024b) and applications designed for specific clinical tasks, such as medical history taking (Saley et al., 2024) and clinical note interpretation (Wang et al., 2024a), have also proven valuable as supplementary learning resources for medical students.

While these educational tools have shown promising results, they primarily focus on answering students' questions with direct explanations (as shown in Figure 1). Building upon this foundation, we propose to engage students through multiple steps of the DDx diagnostic process while tracking their fine-grained knowledge mastery.

2.2 Clinical Diagnosis Datasets

Clinical diagnosis datasets have emerged as crucial benchmarks for evaluating language models in healthcare applications. Traditional medical question-answering (QA) datasets, such as MedQA (Jin et al., 2021), MedMCQA (Pal et al., 2022), PubMedQA (Jin et al., 2019), and CMEXAM (Liu et al., 2024c), have laid the foundation for assessing medical knowledge comprehension. Recent advances have focused on developing more sophisticated diagnostic datasets (Li et al., 2024a; Wang et al.; Hou et al., 2024) that incorporate structured reasoning steps and support dynamic diagnostic processes. DDX-Plus (Fansi Tchango et al., 2022) addresses the scarcity of training data for Automatic Diagnosis (AD) and Automatic Symptom Detection (ASD) by introducing a comprehensive synthetic dataset with differential diagnoses. Dual-Inf (Zhou et al., 2024) enhances diagnostic interpretability by combining expert annotations with Large Language Models (LLMs) to create an more explainable differential diagnosis framework.

These datasets have made valuable contributions to improving LLMs' diagnostic and explanatory capabilities. Complementing these efforts, our work emphasizes the educational value in clinical process dialogues by providing more fine-grained, comprehensive annotations and explanations to impart more systematic knowledge during student guidance.

3 Problem Formulation

Our proposed DDxTutor is inspired by a key aspect of the philosophy behind differential diagnosis (DDx): to convey fine-grained, intermediate reasoning steps throughout the diagnostic process. By transparently presenting the step-by-step thought process, our framework aims to empower students to develop a comprehensive understanding of clinical reasoning and sharpen their decision-making skills. To this end, DDxTutor is built around two pivotal tasks: (1) a structured differential diagnosis reasoning component that generates detailed teaching references, and (2) an interactive teacherstudent module that provides dynamic guidance and feedback. We detail these two components below. Our framework is shown in Figure 2.

3.1 Structured DDx Reasoning

Following clinical differential diagnosis workflows (Elstein and Schwarz, 2002; Graber et al., 2005; First, 2013), we formulate a structured reasoning framework for clinical cases. Given a clinical diagnostic instance $\mathcal{I} = \{q, \mathbf{A}\},\$ where q represents the clinical presentation and $\mathbf{A} = \{a_1, a_2, ..., a_m\}$ represents candidate diagnoses, the LLM generates a structured teaching reference $\mathcal{T} = \{\mathbf{L}, \mathbf{G}\}$ through a two-stage reasoning process, where L represents the local analysis for each clinical clue and G represents the global diagnostic synthesis respectively, which are described in detail as follows. (Notations are marked at their corresponding positions in the workflow diagram of our framework (Figure 2) for better understanding and correspondence.)

In the first stage, the LLM performs local analysis to generate L. It begins by decomposing the clinical presentation q into a set of self-contained and relatively independent clinical clues $\mathbf{C} = \{c_1, c_2, ..., c_n\}$, where each c_i represents a complete and standalone clinical observation extracted from clinical presentation q, e.g., *Chest radiography shows signs of pulmonary hyperinflation*. For each clue c_i , the LLM conducts a detailed analysis $l_i \in \mathbf{L}$ that derives four new components: a specificity assessment of c_i (*spec*_i), an indication (*ind*_i)

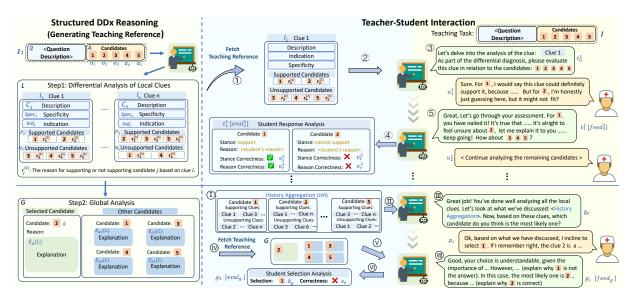


Figure 2: Our *DDxTutor* framework. Left: Teaching reference generation through structured differential diagnosis reasoning. Mid: Backend dialogue management, including teaching reference retrieval, student response analysis and record. Right: Interactive tutoring pipeline incorporating step-by-step clinical reasoning guidance and feedback.

revealing the underlying clinical conditions, and two clue-candidate relation sets \mathbf{P}_i and \mathbf{N}_i :

$$\mathbf{P}_{i} = \left\{ (a_{j}, r_{j}^{(i)}) \mid c_{i} \to a_{j}, a_{j} \in \mathbf{A} \right\}$$
$$\mathbf{N}_{i} = \left\{ (a_{j}, r_{j}^{(i)}) \mid c_{i} \not\to a_{j}, a_{j} \in \mathbf{A} \right\}$$
(1)

where $r_j^{(i)}$ represents the reasoning associated with the relationship. The symbols \rightarrow and $\not\rightarrow$ indicate whether the clue c_i supports or does not support the candidate a_j , respectively.

In the second stage, the LLM synthesizes all local analyses to generate a global synthesis G conditioning on L:

$$\mathbf{G} = \{ (\hat{a}, E_{\hat{a}}(\mathbf{L})) \} \cup \{ (a_j, E_j(\mathbf{L})) \mid a_j \in \mathbf{A} \setminus \{ \hat{a} \} \}$$

where $\hat{a} \in \mathbf{A}$ represents the selected diagnosis, and $E_j(\mathbf{L})$ denotes the explanation for candidate jbased on evidence from \mathbf{L} .

3.2 Teacher-Student Interaction

Leveraging the teaching reference \mathcal{T} generated from the *Structured DDx Reasoning* process, we design an interactive learning framework where the LLM serves as a medical educator guiding students through a systematic diagnostic reasoning process. The interaction follows a two-stage approach mirroring the structured DDx reasoning: (1) local clue analysis discussions and (2) global diagnostic synthesis discussions. (Notations are marked in Figure 2 *Teacher-Student Interaction* for better understanding and correspondence.)

In the first stage, the LLM conducts detailed analytical dialogues for each clinical clue. Given a clinical clue analysis $l_i \in \mathbf{L}$, as defined in Section 3.1, the LLM initiates a structured discussion dialogue with student D_{local}^i :

$$D_{local,0:K}^{i} = \left\langle t_{0}^{i}, (u_{1}^{i}, t_{1}^{i}), \dots, (u_{K}^{i}, t_{K}^{i}) \right\rangle$$

where t_0^i represents the initial teaching inquiry, u_k^i denotes the student's response in the k-th turn, and t_k^i represents the LLM's teaching response. Each teaching response t_k^i consists of two components: $t_k^i = (eval_k^i, feed_k^i)$, where $eval_k^i$ analyzes the student's response into a structured format: $eval_k^i = \{(a_j, stance_j, reason_j, v_s^j, v_r^j) \mid j \in$ $J_k \subseteq \{1, ..., m\}\}$. Here, J_k is the index set of candidates actually mentioned by the student in turn k. a_j denotes the *j*-th candidate diagnosis in the original option set A of the case, maintaining consistent notation with the exam choices. $stance_i$ records the student's supportive stance (whether they believe the clue supports or does not support the diagnosis) for candidate a_i , while $reason_i$ captures the student's reasoning that justifies this stance. The binary flags $v_s^j, v_r^j \in \{0, 1\}$ indicate whether the stance and reasoning, respectively, are correct with respect to the reference analysis l_i . Within the discussion of a single clue analysis l_i , we assume that students articulate their stance and reasoning for any given candidate at most once (If a candidate is revisited, only the latest response is retained.). Consequently, the variables $stance_j$, $reason_j$ and their correctness indicators v_s^j , v_r^j do not depend on the turn index k. Based on this evaluation, $feed_k^i$ delivers targeted feedback by referring to the teaching reference l_i .

The complete local analysis phase comprises dialogues for all clinical clues:

$$\mathcal{D}_{local} = \left\{ D_{local}^{i} \mid i = 1, \dots, n \right\}.$$

In the second stage, the LLM facilitates a global diagnostic synthesis discussion D_{global} that follows the structure:

$$D_{global} = \langle g_0, (p_1, g_1) \rangle$$

Here, g_0 presents a summary of the prior local clue analysis discussions (History Aggregation, HA) to provide the student with a comprehensive overview of the previous discussions. Then, p_1 stands for the student's final diagnostic decision along with their reasoning, while $g_1 = (eval_g, feed_g)$ represents the LLM's evaluation and feedback. The evaluation component $eval_g$ analyzes the student's final diagnosis: $eval_g = \{\hat{a}_g, v_d\}$, where \hat{a}_g denotes the student's selected diagnosis, and v_d is a binary indicator reflecting the correctness of their selection. Based on this evaluation, $feed_g$ delivers detailed explanatory feedback, referring to the teaching reference **G**.

4 DDxReasoning Dataset

To support the structured DDx reasoning task described above, we developed a dataset based on clinical diagnosis questions from MedQA (Jin et al., 2021). Following the formulation in Section 3.1, each case in our dataset contains a clinical scenario $\mathcal{I} = \{q, \mathbf{A}\}$ and its corresponding teaching reference $\mathcal{T} = \{\mathbf{L}, \mathbf{G}\}$. The dataset consists of 933 cases (755 for training and 178 for testing), each presenting a complex clinical scenario with multiple symptoms and signs. Following the original setting of MedQA, each question has 5 candidate diagnoses.

We develop a two-stage dataset creation pipeline that aligns with our structured reasoning process, as shown in Appendix Figure 5.

Stage 1: Local Analysis Generation and Verification

First, for each clinical case \mathcal{I} , we use the OpenAI-o1 (Jaech et al., 2024) model to the local analysis component **L** by decomposing the question description q into independent clinical clues

C. For each clue c_i , it performs the structured analysis l_i as defined in Section 3, comprising a specificity assessment $spec_i$, clinical indication ind_i , supporting relationships \mathbf{P}_i , and unsupporting relationships \mathbf{U}_i .

Three experienced doctors then independently verify these local analyses, examining: (1) Clue decomposition - ensuring each clue represents a meaningful, independent clinical observation; (2) Clinical indications - validating the correct interpretation of symptoms and signs; (3) Specificity assessments - evaluating how uniquely each clue points to specific diagnoses; (4) Supporting/unsupporting relationships - verifying the correctness of candidate categorization and the validity of medical reasoning. For each sample, doctors independently review and mark problematic aspects of the local analyses. After all three doctors complete their reviews, we merge their annotations to identify overlapping concerns and unique issues. This merged feedback serves as the basis for a focused discussion to reach consensus on necessary modifications. This process continues until the merged feedback shows no remaining issues from any doctor.

Stage 2: Global Synthesis Generation and Verification

After verification of the local analyses, we feed the refined L back to OpenAI-o1 to generate the global synthesis component G. This includes generating a final diagnostic decision, providing comprehensive reasoning for the chosen candidate, and explaining why other candidates were not selected.

The same doctor panel applies an identical review-merge-consensus process for the global analysis, focusing on (1) The logical coherence of the final decision, (2) The completeness of evidence integration, and (3) The validity of reasoning for both selected and rejected candidates. Similar to Stage 1, the annotations from all doctors are merged to identify common concerns and unique insights, followed by targeted discussions to resolve any remaining issues.

Statistics of DDxReasoning dataset, prompts used to build the data, examples, and more details are presented in Appendix A.

5 Knowledge-grounded Clinical Tutoring Dialogue Generation

Following prior work in educational systems—such as science, math, and medical tutoring (Wang et al., 2024b; Liu et al., 2024d; Chevalier et al., 2024), we propose using simulation-based approaches to construct dialogue datasets. This method enables us to operationalize the teaching framework we have designed, guiding the model to adhere to structured teaching interactions. By generating diverse, high-quality simulated dialogues, we expect this approach to better align language models with our intended educational objectives.

Our dialogue generation process consists of two stages that mirrors our framework in Section 3.2: local clue analysis dialogues and global diagnostic synthesis dialogues.

Local Analysis Dialogue Generation. For each clinical clue c_i (and its analysis l_i , the dialogue begins with the teacher presenting the clue through an initial inquiry t_0^i and requesting analysis of its relationship to the candidates. In each turn, the student randomly selects k candidates to analyze $(1 \le k \le |\mathcal{A}_t|)$, where \mathcal{A}_t represents the remaining unanalyzed candidates.

For each selected candidate, the student provides a response that exhibits one of the following patterns: (1) correct stance with aligned reasoning, (2) correct stance with misaligned reasoning, (3) incorrect stance with misaligned reasoning, (4) random guessing, or (5) complete uncertainty. The teacher then evaluates this response against the teaching reference l_i and provides targeted feedback addressing both the correctness of stance and reasoning. **Global Synthesis Dialogue Generation.**

After completing all local analysis dialogues, the simulated tutor initiates a global discussion by presenting the previously analyzed clinical clues and requesting a final diagnostic decision. The student then synthesizes the evidence from all clues to select a final diagnosis and provide comprehensive reasoning for their choice. The teacher evaluates this final decision against the teaching reference **G** and provides detailed feedback that addresses both the diagnosis selection and the quality of synthesized reasoning.

We leverage GPT-40 (Hurst et al., 2024) to generate student responses and teacher feedback through carefully crafted prompts that ensure both dialogue coherence and pedagogical effectiveness. This approach creates teacher-student interactions spanning diverse learning scenarios while remaining firmly grounded in the verified clinical knowledge from our *DDxReasoning* Dataset. The resulting dialogues comprises a Local Analysis Tutoring dialogue collection with 16,132 training and 3,760 testing turns, and a Global Synthesis dialogue collection with 1,506 training and 356 testing turns. A detailed illustration of this dialogue generation process is in Appendix B.

6 Experiments

We evaluate state-of-the-art LLMs on two core aspects of the DDxTutor framework: (1) the ability to generate comprehensive DDx reasoning chains that can serve as teaching references, and (2) the capacity to conduct tutoring dialogues that guide students through systematic diagnostic reasoning. For evaluation, we fine-tune three specialized groups of LLMs:

(1) Teaching Reference Generator (f_1) : These models are trained to generate complete teaching references $\hat{\mathcal{T}}$ given the clinical case input \mathcal{I} : $\hat{\mathcal{T}} = f_1(\mathcal{I})$.

(2) Local Analysis Stage Dialogue Tutor (f_2) : These models f_2 are trained to generate teaching responses \hat{t}_k^i based on the clinical clue c_i , history dialogue $D_{local,0:(k-1)}^i$, student's last utterance u_k^i , and local teaching reference l_i : $\hat{t}_k^i = f_2(c_i, D_{local,0:(k-1)}, u_k^i, l_i)$.

(3) Global Synthesis Stage Dialogue Tutor (f_3) : These models f_3 generate final feedback g_1 based on all local analyses with aggregated information HA, student's final diagnosis p_1 , and global teaching reference G: $\hat{g}_1 = f_3(\mathbf{L}, HA, p_1, \mathbf{G})$.

We select models with parameters ranging from 3B to 32B, including 7 general open source LLMs: Qwen2.5-3B, Qwen2.5-7B, Qwen 2.5-14B, Qwen 2.5-32B (Qwen et al., 2025), LLama3.2-3B, LLama3.1-8B (Grattafiori et al., 2024), Phi4-14B (Abdin et al., 2024) and 2 Medical LLMs: Meditron-8B (Chen et al., 2023, 2024b), and Med42-8B (Christophe et al., 2024).¹

6.1 Implementation Details

All the models are finetuned with LoRA (Hu et al., 2021) using the Unsloth framework (Daniel Han and team, 2023). We set the LoRA rank r=16 and α =16, targeting key transformer modules including attention layers and feed-forward networks. The models were trained for 3 epochs using AdamW optimizer, a learning rate of 2e-4, and a batch size of 8. A linear learning rate scheduler with 20 warmup

¹For f_1 models, we further finetuned two 70B-level model: Qwen2.5-72B and Med42-70B with 4-bit Qlora (Dettmers et al., 2023).

Model	Bert_C	$Meteor_C$	Acc_{PN}	$Bert_{PN}$	$Meteor_{PN}$	$Bert_{Ind}$	Meteor _{Ind}	$Bert_{Spec}$	$Meteor_{Spec}$	Bert_E	$Meteor_E$	$\operatorname{Acc}_{\hat{a}}$
Qwen2.5-3B	0.836	0.695	0.664	0.600	0.248	0.891	0.242	0.884	0.320	0.757	0.387	0.600
Llama3.2-3B	0.884	0.760	0.681	0.620	0.284	0.896	0.290	0.888	0.357	0.803	0.434	0.706
Qwen2.5-7B	0.842	0.706	0.707	0.639	0.269	0.894	0.274	0.822	0.316	0.823	0.442	0.777
Llama3.1-8B	0.887	0.769	0.739	0.677	0.334	0.899	0.309	0.896	0.389	0.849	0.466	0.832
Meditron-8B	0.901	0.779	0.730	0.669	0.337	0.900	0.316	0.898	0.398	0.828	0.451	0.773
Med42-8B	0.895	0.790	0.729	0.668	0.331	0.901	0.315	0.899	0.414	0.849	0.476	0.836
Qwen2.5-14B	0.881	0.720	0.734	0.665	0.294	0.893	0.265	0.895	0.359	0.825	0.441	0.776
Phi4-14B	0.896	0.787	0.767	0.705	0.371	0.905	0.350	0.903	0.432	0.863	0.500	0.858
Qwen2.5-32B	0.884	0.724	0.754	0.686	0.319	0.897	0.288	0.898	0.379	0.872	0.486	0.892
Qwen2.5-72B	0.888	0.741	0.773	0.703	0.328	0.897	0.296	0.897	0.383	0.867	0.498	0.874
Med42-70B	0.912	<u>0.789</u>	0.767	0.705	0.360	<u>0.902</u>	0.339	0.904	<u>0.428</u>	0.877	0.492	0.901
Phi4-14B (2shot)	0.759	0.611	0.718	0.654	0.300	0.895	0.264	0.894	0.368	0.798	0.345	0.738
GPT-4o-mini (2shot)	0.600	0.377	0.585	0.520	0.161	0.885	0.189	0.883	0.243	0.780	0.181	0.792
GPT-4o (2shot)	0.540	0.312	0.741	0.666	0.251	0.886	0.233	0.890	0.294	0.861	0.271	0.959

Table 1: Performance of Teaching Reference Generator f_1 on the DDxReasoning Dataset.

Model	IOU _a	Acc_{stance}	$\operatorname{Bert}_{reason}$	Acc_{vs}	Acc_{vr}	$Bert_{feed}$	Meteor _{feed}	$\operatorname{Acc}_{\hat{a}g}$	Acc_{vd}	$\operatorname{Bert}_{feed_g}$	$Meteor_{feed_g}$
Qwen2.5-3B	0.994	0.987	0.949	0.985	0.867	0.895	0.415	0.978	<u>0.997</u>	0.915	0.514
Llama3.2-3B	0.996	0.992	0.950	0.991	0.875	0.900	0.454	0.983	<u>0.997</u>	0.918	0.527
Qwen2.5-7B	0.995	0.975	0.938	0.977	0.868	0.896	0.420	0.922	0.997	0.913	0.512
Llama3.1-8B	0.997	0.992	0.954	0.992	0.898	0.901	0.462	0.992	0.994	0.918	0.526
Meditron-8B	0.996	0.991	0.954	0.992	0.900	0.901	0.456	0.994	0.997	0.918	0.527
Med42-8B	0.997	0.993	0.956	0.993	0.902	0.903	0.471	0.992	0.994	0.918	0.525
Qwen2.5-14B	0.991	0.989	0.952	0.989	0.885	0.898	0.437	0.961	0.997	0.915	0.518
Phi4-14B	0.998	0.994	0.957	0.994	0.914	0.905	0.481	0.997	1.000	0.920	0.539
Qwen2.5-32B	0.997	0.993	0.953	0.993	0.903	0.900	0.451	0.994	0.997	0.916	0.523
Phi4-14B (2shot)	0.890	0.965	0.874	0.848	0.740	0.878	0.312	0.553	0.991	0.910	0.506
GPT-4o-mini (2shot)	0.886	0.893	0.880	0.878	0.704	0.881	0.360	0.919	0.997	0.901	0.476
GPT-4o (2shot)	0.981	0.955	0.881	0.919	0.789	0.880	0.335	0.800	1.000	0.896	0.469

Table 2: Performance of Local Analysis Tutor f_2 and Global Synthesis Tutor f_3 models on the simulated dialogues.

steps was employed. All the models are trained on a server with 2 A100 GPUs.

6.2 Evaluation Setting

We evaluate the f_1 models from the following aspects: (1) Local Clue Analysis Capability: the alignment of the decoupled local clues with teaching references using metrics $Bert_C$ and $Meteor_C$; the accuracy (Acc_{PN}) between predicted and ground-truth stances for each clue; the semantic and textual quality of reasoning explanations measured by $Bert_{PN}$ and $Meteor_{PN}$; the quality of clue specificity descriptions using $Bert_{Spec}$ and $Meteor_{Spec}$; the quality of indication description of the clue using $Bert_{Ind}$ and $Meteor_{Ind}$; (2) Global Synthesis and Conclusion: the final diagnosis selection accuracy $Acc_{\hat{a}}$; and the quality of final explanations using $Bert_E$ and $Meteor_E$. Here, Bert and Meteor refer to BertScore (Zhang et al., 2019) and Meteor Score (Banerjee and Lavie, 2005) respectively.

For f_2 , we evaluate their: (1) Student Response Extraction capability: measuring the intersection over union (IOU_a) between extracted student-analyzed candidates and ground truth, the accuracy of extracting students' supporting/nonsupporting stances (Acc_{stance}), and the quality of extracted students' reasons ($Bert_{reason}$); (2) Student Response Judgment capability: evaluating the accuracy of tutors' judgment on students' stances (Acc_{vs}) and the consistency between students' reasoning and teaching references (Acc_{vr}); (3) Teacher Feedback Generation capability: assessing feedback quality using $Bert_{feed}$ and $Meteor_{feed}$.

For f_3 , following a similar evaluation logic as f_2 , we assess its capability to extract students' final diagnosis selection $(Acc_{\hat{a}g})$, the accuracy of tutors' judgment (Acc_{vd}) , and the quality of teacher feedback $(Bert_{feed_q} and Meteor_{feed_q})$.

More detailed explanation of our evaluation protocol is presented in Appendix C.

6.3 **Results of Teaching Reference Generators**

The quantitative evaluation of teaching reference generators (f_1 models) reveals several findings. As shown in Table 1 and Figure 3, model performance generally improves with increasing parameter count, following established scaling laws (Kaplan et al., 2020; Zhang et al.). This trend is particularly evident in the Qwen model family, which demonstrates consistent performance gains across parameter scales.

Phi4-14B achieves optimal efficiency with the best balance between performance and computational cost among the evaluated models. Notably, some medical domain-specialized models, such as Med42-8B and Meditron-8B, fail to exhibit their expected advantages in our task, even though they are known to perform well in general medical QA tasks. This observation may be attributed to the limited representation of complex, multi-step clinical reasoning chains in their pre-training datasets, which constrains their performance to levels comparable to, or even below, that of their parent model, Llama3.1-8B.

Further analysis through correlation mapping (Figure 4) reveals that the final diagnosis accuracy $(Acc_{\hat{a}})$ is most strongly correlated with three local analysis metrics: stance accuracy (Acc_{PN}) and reasoning quality measures $(Bert_{PN})$ and $Meteor_{PN}$). This phenomenon aligns with clinical intuition that accurate stance identification and sound reasoning at each local analysis step are fundamental to reaching correct final diagnoses.

Few (two) shot results of Phi4-14B, GPT-4omini and GPT-40 demonstrate that few-shot models seem to take a different path to reach solutions, or to some extent, they are 'slacking off'. This is evidenced by their poor clue decoupling scores $(Bert_C \text{ and } Meteor_C)$ while maintaining high final selection accuracy $(Acc_{\hat{a}})$. This could indicate that few-shot models are leveraging their pretrained knowledge to make direct diagnostic leaps rather than following the systematic reasoning process we aim to teach. While this may achieve correct final diagnoses, it bypasses the step-by-step analytical process that is crucial for medical education. This observation highlights the importance of fine-tuning in ensuring models adhere to structured diagnostic reasoning patterns rather than relying on shortcuts that, while potentially effective for diagnosis, may be less valuable for teaching purposes.

6.4 Results of Dialogue Tutors

The evaluation of dialogue tutors (f_2 and f_3 models) reveals promising capabilities across multiple dimensions (Table 2). All fine-tuned models demonstrate robust performance in recognizing candidate diagnoses from student responses—a critical ability for maintaining effective dialogue flow and monitoring student progress. The models also exhibit high precision in assessing student response correctness, as evidenced by strong Acc_{vs} and Acc_{vr} scores.

Significantly, even models with relatively modest parameters (3B) achieve satisfactory performance in workflow control, suggesting the framework's viability across various model scales. The minimal performance differential between models of different sizes indicates that dialogue tutoring, when supported by teaching references, may not demand the same model capacity as reference generation. To validate this hypothesis, we conducted an ablation study without teaching references (detailed in Appendix Table 8). The results align with our expectations, showing a predictable performance decline when Teaching material input is removed.

7 Human Evaluation

To validate the effectiveness of our framework, we adopt the human evaluation methodology outlined in NoteChat (Wang et al., 2024a), which centers on simulating patient-doctor dialogues. Our evaluation incorporates both expert assessments and user experience analyses from two distinct perspectives.

For evaluating the pedagogical value of Teaching References, we enlisted three experienced educators, each with over a decade of medical teaching experience. These experts evaluated 60 Teaching References, comprising 20 samples each from Phi4-14b, GPT-4o (2-shot), and the ground truth from *DDxReasoning* (GT). Their assessment focused on the materials' potential as both immediate teaching resources and structured draft materials.

From the learning perspective, we engaged five medical students to assess the quality of local and global dialogues generated by our f_2 and f_3 models respectively. The evaluation covered 20 cases per dialogue type, comparing outputs from LLaMA-3.2-3B, Phi4-14B, and GPT-40 (2-shot).

Participants in both evaluations ranked outputs on a 1-to-3 scale, with 1 being most preferred. We quantified preferences using Mean Reciprocal Rank (MRR), as detailed in Table 3. The results demonstrate that our fine-tuned Phi4-14B model achieves comparable or superior performance to GPT-40 (2-shot) mode within our tutoring framework, highlighting its potential for practical applications.

8 Conclusion

This paper presents DDxTutor, a novel framework that leverages LLMs to provide systematic clinical diagnostic education through structured rea-

Teaching Reference f_1											
GT Phi4-14B GPT4o (2sh											
0.850	0.600	0.383									
Local Analysis Dialogue f_2											
llama3.2-3B	Phi4-14B	GPT40 (2shot)									
0.425	0.758	0.650									
Global	Synthesis Di	alogue f3									
llama3.2-3B	Phi4-14B	GPT40 (2shot)									
0.483	0.708	0.642									

Table 3: MRR scores for human evaluation.

soning and interactive tutoring. By decomposing the diagnostic process into sequential clue analysis and global synthesis phases, our approach enables active student engagement while maintaining scientific rigor in clinical reasoning education. The framework is supported by DDxReasoning, a comprehensive dataset of expert-verified clinical reasoning chains that serves both as fine-tuning data and evaluation benchmark.

Our experimental results demonstrate that finetuned LLMs can effectively generate structured teaching references and conduct pedagogically sound tutoring dialogues. The strong performance of mid-sized models like Phi4-14B suggests that effective clinical tutoring systems are achievable with reasonable computational resources. Human evaluation from both medical educators and students validates the educational value of our framework, with experts highlighting the quality of generated teaching materials and students confirming the effectiveness of the interactive tutoring approach.

Limitations

While DDxTutor demonstrates promising potential in clinical education, several important limitations need to be acknowledged:

First, DDxTutor's educational scope is relatively focused. Although differential diagnosis (First, 2013) represents a classical approach to clinical reasoning, it is not the sole objective of medical education. Traditional medical teaching encompasses various pedagogical formats, including didactic lectures (Alaagib et al., 2019), Problem-Based Learning (PBL) (Trullàs et al., 2022), case-based discussions (Weidenbusch et al., 2019), and bedside teaching (Peters and Ten Cate, 2014). Each format serves distinct educational goals and develops different aspects of clinical competency. In this context, DDxTutor primarily addresses structured diagnostic reasoning training and may need adaptation to support broader educational objectives and teaching modalities.

Second, our framework makes several assumptions about student-teacher interactions, particularly in dialogue construction. For instance, during local clue analysis, we assume students will provide structured responses containing both their analysis and supporting rationale. While this assumption aligns with conventional medical discussions and proved effective during human evaluation, it may not fully capture the diversity of student responses in real educational settings. As a pioneering work in this domain, DDxTutor currently has limited capability to handle off-pattern or unexpected student responses. Future iterations could potentially integrate agent workflow frameworks to enhance robustness in real-world applications, allowing more flexible and adaptive responses to diverse student interaction patterns.

Acknowledgement

This research work was supported in part by the Research Grants Council of the Hong Kong Special Administrative Region, China under Project No. T45-401/22-N, in part by the National Natural Science Foundation of China under Project No. 62322318, and in part by the Hong Kong Innovation and Technology Fund under Project No. GHP/167/22SZ.

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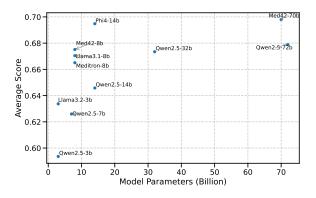


Figure 3: Performance vs Parameters of Teaching Reference Generator f_1 models.

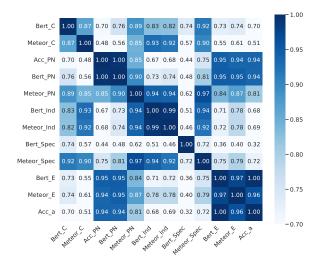


Figure 4: The correlation matrix of the performance metrics of Teaching Reference Generators, as presented in Table 1.

A More Details on DDxReasoning Dataset

Our DDxReasoning dataset was developed through a systematic two-stage annotation process combining LLM generation with expert medical verification. Initially, we selected 1,000 clinical cases from the MedQA dataset (Jin et al., 2021), with an intended split of 800 training and 200 testing cases. However, through our rigorous verification process, some cases were eliminated due to quality concerns, resulting in a final dataset of 933 cases (755 training and 178 testing cases).

The first stage focuses on generating and validating detailed local analyses for individual clinical clues. We utilize OpenAI o1 with Local Analysis Prompt (PromptL, presented in Table 4) to generate initial structured analyses for each clinical presentation, breaking down complex cases into discrete, analyzable components. Each generated

Prompt L

Question: {question}

Please extract the clues from the question for differential analysis. For each clue, analyze it systematically using the following structure:

 Clue Name: Clearly label the clue using concise and descriptive language.
 Description: Provide a detailed description of the clue, retaining as much

Description: Provide a detailed description of the clue, retaining as much of the original wording as possible.

3. Specificity Description: Assess whether the clue is diagnostically specific and explain its relevance to particular conditions (e.g., "The presence of xxx is highly specific for the xxx disease"). 4. Indication: Analyze what the clue suggests based on medical knowledge and its implications (e.g., "This finding could suggest xxx because of xxx").

5. Supported Candidates: Identify which candidate diagnoses the clue could support and explain why (e.g., "This clue supports xxx because xxx"). If there are no supported diagnoses, leave this empty.

6. Unsupported Candidates: Identify which candidate diagnoses the clue could not support and explain why (e.g., "This clue refutes xxx because xxx" or "This clue is unrelated to xxx because xxx"). If all the candidates are supported, leave this empty.

Note: Please always ensure that the sum of supported and unsupported diagnoses is equal to the number of all candidates.

Here is a JSON format of the expected output: {Local Clue Analysis Json}.

1.100.001.1.0

{question}

Below is a clue-by-clue analysis for each candidate in the options: Clue_Differential_Analysis

Please generate a final decision with a detailed differential diagnosis explanation based on the given clues. Your task is to determine the most likely diagnosis and explain why each candidate is selected or rejected.

Requirements:

Citation Format: Cite specific clues using the format [Clue X] (e.g., [Clue 1], [Clue 3]). Avoid combining multiple clue citations (e.g., [Clue 1,2] or [Clue 3,5,10]). When discussing multiple related clues, clearly state each finding with its corresponding clue index.

Example citation: "Although scleroderma is more common in middleaged women [Clue 1] and presents with fatigue [Clue 2] and difficulty swallowing [Clue 4], it typically does not present with upper esophageal webs [Clue 9], koilonychia [Clue 7], or iron deficiency anemia [Clue 10]. Additionally, the presence of neck pain [Clue 3] and leukocytosis [Clue 11] is inconsistent with typical scleroderma features, making it a less likely diagnosis."

Evidence Evaluation: Prioritize strong, decisive clues over weaker or peripheral evidence. Focus on clues that are sufficient to support diagnostic decisions. Avoid overemphasis of non-specific or irrelevant findings.

Rejected Candidates: Provide clear, evidence-based explanations for why each rejected candidate is less likely. Base rejections on specific clue citations. Explain any missing key features expected for these diagnoses.

Balanced Analysis: Use concessive reasoning (although-style) when analyzing candidates with mixed evidence. Explicitly weigh supporting versus conflicting evidence. Clearly justify final decisions when evidence is mixed.

Example reasoning: "Although Candidate A shows feature X [Clue 3], which supports the diagnosis, the absence of feature Y [Clue 5], a hallmark finding, makes it less likely."

Important Note: Your analysis will serve as a teaching resource for identifying strong and sufficient evidence in diagnostic reasoning. Therefore: Ensure all cited clues are accurate and directly relevant. Select only strong, definitive clues to support or reject diagnoses. Exclude weak or ambiguous evidence from core supporting arguments.

Here is a JSON format of the expected output: {Global Analysis Json}

Table 4: The PromptL and PromptG we used to utilized OPENAI o1 to extract a initial local clues analysis and global analysis when we curated the DDxReasoning dataset.

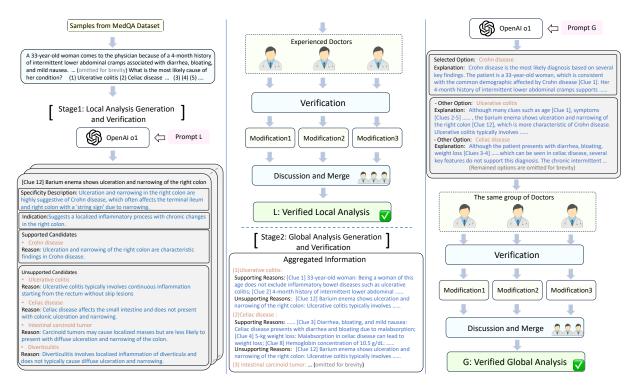


Figure 5: The two-stage generation and verification pipeline of the proposed DDxReasoning dataset

analysis comprises a specificity description, clinical indication, and detailed relationships with candidate diagnoses, classifying them into supported and unsupported categories. During this stage, approximately 49 cases were removed due to unclear or ambiguous clinical presentations that could not be effectively decomposed into distinct clues.

These local analyses then undergo thorough verification by a panel of three experienced doctors. Each doctor independently proposes modifications (Modification 1, Modification 2, Modification 3), followed by a collaborative discussion and merge phase. This verification process continues iteratively until all three doctors reach consensus, resulting in the Verified Local Analysis L.

In the second stage, we aggregate the verified local analyses and use OpenAI o1 again with Global Analysis Prompt (Prompt G, presented in Table 4) to generate comprehensive diagnostic reasoning, integrating all verified clues to form final diagnostic conclusions with detailed supporting and unsupporting reasons for each candidate. This global analysis undergoes the same rigorous verification process by the same group of doctors, leading to the final Verified Global Analysis G. Through this twostage process, an additional 18 cases were eliminated due to inconsistencies in clinical reasoning or lack of consensus among experts.

The statisics of the DDxReasoning Dataset is

presented in Table 5. An sample of the dataset is

Split	Num	Num of Clues	Max Tokens	Mean Tokens
Train		7047	12325	5223
Test	178	1643	9353	5043

Table 5: The statistics of the DDxReasoning dataset. Num of Clues refers to the total number of clues analyzed during the dataset's creation. Max Tokens and Mean Tokens indicate the maximum and average token counts within the dataset, respectively.

presented in Figure 6.

All our annotators are experienced doctors with PhD degrees, and they are compensated at rates satisfying local market guidelines.

B Knowledge-grounded Clinical Tutoring Dialogue Generation

B.1 Dialogue Generation of Local Analysis

To create a knowledge-grounded dialogue dataset that enables LLMs to effectively perform clinical tutoring, we propose a structured dialogue generation framework (Figure 7). The framework aims to develop three key capabilities in the fine-tuned LLM: (1) student response awareness - understanding what clinical elements the student analyzed in their last utterance, (2) stance recognition - identifying whether the student believes a discussed clue supports specific candidates, and (3) judgment

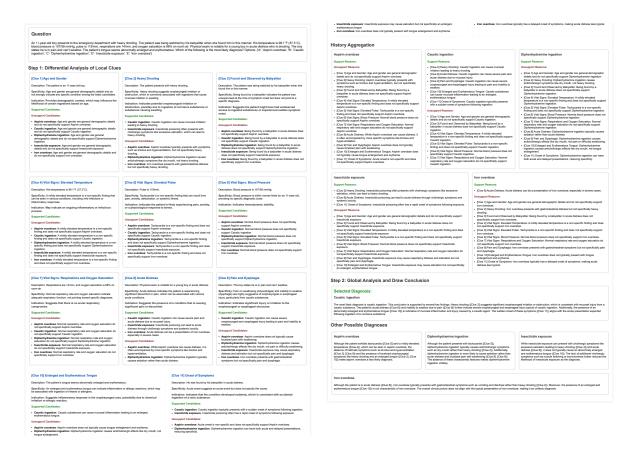


Figure 6: A sample of our DDxReasoning Dataset (Best viewed when zoomed in).

capability - assessing the correctness of student stances and reasoning.

Response State Design

For each clinical clue $c_i \in \mathbf{C}$, the student response should analyze its relationship with candidate diagnoses. Each analysis contains two key components:

- A stance *stance_j* indicating whether the clue supports or does not support the candidate *a_j*
- A reasoning text *reason_j* explaining the stance

The teacher evaluates these responses against the teaching reference $l_i \in \mathbf{L}$, assessing both stance correctness (v_s^j) and reasoning quality (v_r^j) .

Generation Process

For a given clinical case with question q, candidates set **A**, and a specific clue c_i , our dialogue generation follows a structured process that simulates a teacher-student dialogue analyzing how the clue supports or cannot support each candidate.

First, the teacher simulator combines q, c_i , and **A** to generate the initial teaching inquiry t_0^i .

Then we enter the student-teacher dialogue loop. At the beginning of each loop k, the student obtains unanswered candidates from the system state:

$$A_{remaining} = \mathbf{A} \setminus \bigcup_{k=1}^{t} J_k \tag{2}$$

where J_k represents the set of candidates discussed in turn k.

The student's response state is simulated by randomly choosing from five possible scenarios for each candidate:

- 1. Correct stance with correct reasoning
- 2. Correct stance with flawed reasoning
- 3. Incorrect stance with incorrect reasoning
- 4. Random guess (stance correct/incorrect, with planned reason as None)
- 5. Complete uncertainty

For each candidate to be addressed, we randomly select one response state to form the student's response plan. We then use GPT40 to generate

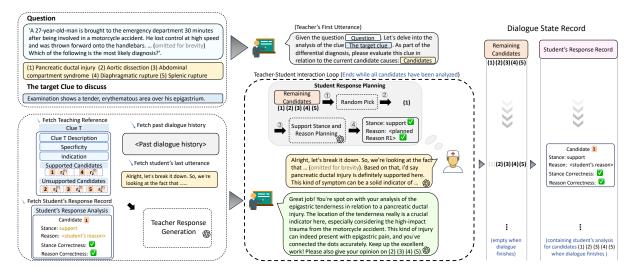


Figure 7: The Generation pipeline of Local Analysis Dialogues

simulated reasoning based on the response state, grounded in the teaching reference l_i .

For example, consider a scenario where the student's response plan includes: - For candidate a_1 : correct stance and correct reasoning (scenario 1) -For candidate a_2 : correct stance but flawed reasoning (scenario 2).

In this case, GPT40 will: (1) Generate $reason_1$ that closely aligns with the correct reasoning from l_i . (2) Generate $reason_2$ that intentionally deviates from the correct reasoning in l_i .

This process yields a structured response evaluation:

$$eval_k^i = \{(a_j, stance_j, reason_j, v_s^j, v_r^j) \mid j \in J_k\}$$
(3)

We then prompt GPT40 to generate a complete student response u_k^i following this structured plan. The generated response is expected to closely align with the evaluation structure, allowing us to use the components as ground truth labels for the dialogue output.

Teacher Response Framework

The teacher's turn t_k^i consists of two parts: the *evaluation* $(eval_k^i)$ and the *feedback* $(feed_k^i)$. The feedback policy F compares the student's answer with the reference label l_i and assigns one of three tags:

$$F(a_j) = \begin{cases} \text{Affirm} & \text{if } v_s^j \wedge v_r^j \\ \text{Refine} & \text{if } v_s^j \wedge \neg v_r^j \\ \text{Redirect} & \text{if } \neg v_s^j \end{cases}$$
(4)

• Affirm : both stance and reasoning are correct. The teacher briefly confirms the answer and may add concise supportive evidence.

- **Refine**: stance is correct but reasoning is incomplete or flawed. The teacher first acknowledges the correct stance, then amends or extends the reasoning.
- **Redirect**: the stance itself is wrong (reasoning quality is therefore moot). The teacher states the stance is incorrect, supplies the correct stance, and provides a clear, referencegrounded explanation.

In deployment, the LLM is expected to produce $eval_k^i$ and $feed_k^i$ end-to-end, $feed_k^i$ is conditioned on the internally generated $eval_k^i$ so that feedback is adaptive.

The local-analysis dialogue D_{local}^{i} proceeds until every candidate has been covered. Each turn contains:

- 1. Teacher: request analysis of the remaining candidates.
- 2. Student: supply stances and reasoning for one or more candidates.
- 3. Teacher: generate evaluation and tagconditioned feedback via the policy above.
- 4. System: update J_k with the newly discussed candidates.

This structured loop guarantees systematic yet natural coverage of clinical reasoning. The resulting dataset captures both correct and incorrect reasoning patterns, enabling the LLM to learn robust evaluation and feedback behaviours.

B.2 Global Synthesis Dialogue Generation

After completing all local clue analyses $\{D_{local}^i\}_{i=1}^n$, we generate the global diagnostic synthesis dialogue D_{global} . For each case, we generate two variants of D_{global} : one with correct diagnosis selection ($v_d = 1$) and another with incorrect selection ($v_d = 0$). This approach enables the model to learn appropriate feedback strategies for both successful and unsuccessful diagnostic reasoning.

The dialogue generation process consists of three main components:

Teacher's First Utterance: The LLM generates the initial prompt by combining:

- A summary of all analyzed clinical clues C and their analysis L.
- A request for final diagnostic decision.

Student Response Generation: For each dialogue variant, we simulate the student's response p_1 through:

1. Diagnosis selection:

$$\hat{a}_g = \begin{cases} \hat{a} & \text{if } v_d ==1\\ a_j \in \mathbf{A} \setminus \{\hat{a}\} & \text{if } v_d ==0 \end{cases}$$
(5)

where \hat{a} is the correct diagnosis from **G**.

- 2. Reasoning generation: GPT40 generates student-like reasoning that:
 - References relevant analyses from L
 - Employs natural language patterns
 - Demonstrates diagnostic conviction while maintaining educational tone

Teacher's Response Generation: The teacher's response $g_1 = (eval_g, feed_g)$ is generated conditionally:

For correct diagnosis ($v_d = 1$):

- Confirmation of correct selection
- · Validation of key reasoning points
- Supplementary supporting evidence from G
- Integration of clinical elements from L

For incorrect diagnosis ($v_d = 0$):

- · Analysis of reasoning flaws
- Presentation of correct diagnosis \hat{a}

- Comprehensive justification based on G
- References to relevant evidence from L

This structured approach generates diverse training examples for both successful and unsuccessful diagnostic scenarios, enabling the LLM to learn appropriate response strategies while maintaining educational effectiveness.

B.3 The Statistics of the Simulated Dialogues

Following the dataset design of Task-Oriented Dialogue systems (Valizadeh and Parde, 2022; Wei et al., 2018), we convert our generated dialogues into Supervised Finetuning datasets (SFT). The Statistics of the dataset is presented in Table 6.

Split	Num	Mean Total Tokens	Mean Gen Tokens								
	Local Clue Analysis Dialogue										
Train	16132	2064	418								
Test	3760	2507	641								
	Global Synthesis Dialogue										
Train	1506	6764	428								
Test	356	6574	434								

Table 6: The statistics of the simulated tutoring dialogue datasets, Local Clue Analysis Dialogue and Global Synthesis Dialogue. Here the Mean Gen Tokens means the average number of tokens in generated responses, while Mean Total Tokens represents the average length of the entire dialogue including both input and output.

C More details of the Evaluation Protocol

This section provides detailed explanations of our evaluation metrics for assessing the three types of fine-tuned models: Teaching Reference Generator (f_1) , Local Analysis Dialogue Tutor (f_2) , and Global Synthesis Dialogue Tutor (f_3) .

C.1 Content Alignment Strategy

When evaluating structured outputs from LLMs against ground truth references, a key challenge is that the order of analyses (e.g., candidate diagnoses, clinical clues) may differ between the prediction and reference, while the content remains semantically equivalent. To address this, we employ the Hungarian algorithm for optimal content matching before computing evaluation metrics.

Specifically, for any two sets of content that need alignment (e.g., predicted vs. reference candidates), we:

1. Construct a cost matrix M where $M_{ij} = 1 - sim(p_i, r_j)$

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- 2. $sim(p_i, r_j)$ computes the textual similarity ratio between prediction p_i and reference r_j
- 3. Apply the Hungarian algorithm to find the optimal matching that minimizes total matching cost

This matching process ensures accurate evaluation by properly aligning corresponding contents regardless of their order in the structured output. The matched pairs are then used for computing various evaluation metrics detailed below.

C.2 Teaching Reference Generator (*f*₁) Evaluation

The evaluation of f_1 models focuses on two main aspects:

Local Clue Analysis Capability:

• Clue Decomposition Quality:

- $Bert_C(\hat{c}, c)$: BertScore measuring semantic similarity between predicted clues \hat{c} and ground-truth clues c.
- $Meteor_C(\hat{c}, c)$: Meteor score assessing textual alignment between predicted and ground-truth clues.

• Clue-Candidate Relationship Analysis:

- *Acc_{PN}*: Accuracy of predicted support/non-support stances

$$Acc_{PN} = \frac{|\{(c_i, a_j)|\hat{s}_{ij} = s_{ij}\}|}{|\{(c_i, a_j)\}|} \quad (6)$$

where \hat{s}_{ij} and s_{ij} are predicted and ground-truth stances.

- $Bert_{PN}(\hat{r}, r)$ and $Meteor_{PN}(\hat{r}, r)$: Quality metrics comparing predicted reasoning explanations \hat{r} with ground-truth reasoning r in \mathbf{P}_i and \mathbf{N}_i .
- Clue Property Analysis:
 - $Bert_{Spec}(s\hat{pec}, spec)$ and $Meteor_{Spec}(s\hat{pec}, spec)$: Quality metrics comparing predicted specificity assessment $s\hat{pec}_i$ with ground truth $spec_i$.
 - $Bert_{Ind}(\hat{ind}, ind)$ and $Meteor_{Ind}(\hat{ind}, ind)$: Quality metrics comparing predicted indication description \hat{ind}_i with ground truth ind_i .

Global Synthesis Capability:

• Diagnostic Accuracy:

$$Acc_{\hat{a}} = \frac{|\{\mathcal{I}|\hat{a} = a^*\}|}{|\{\mathcal{I}\}|}$$
(7)

where a^* is the ground-truth diagnosis.

• Explanation Quality:

- $Bert_E(\hat{E}, E)$ and $Meteor_E(\hat{E}, E)$: Quality metrics comparing predicted diagnosis explanations $\hat{E}_j(\mathbf{L})$ with ground truth $E_j(\mathbf{L})$

C.3 Local Analysis Dialogue Tutor (*f*₂) Evaluation

We evaluate f_2 models on three aspects: Student Response Extraction:

• Candidate Coverage:

$$IOU_a = \frac{|\hat{J}_k \cap J_k|}{|\hat{J}_k \cup J_k|} \tag{8}$$

where \hat{J}_k and J_k are predicted and groundtruth discussed candidate sets.

Stance Extraction:

$$Acc_{stance} = \frac{|\{j \in J_k | stance_j = stance_j\}|}{|J_k|}$$
(9)

• Reasoning Extraction: $Bert_{reason}(\hat{r}, r)$ measures quality between predicted reasoning \hat{r} and ground-truth reasoning r.

Response Judgment:

• Stance Judgment:

$$Acc_{vs} = \frac{|\{j \in J_k | \hat{v_s^j} = v_s^j\}|}{|J_k|} \quad (10)$$

where $\hat{v_s^j}$ and v_s^j are predicted and groundtruth judgments on student's stance

• Reasoning Judgment:

$$Acc_{vr} = \frac{|\{j \in J_k | v_r^j = v_r^j\}|}{|J_k|}$$
(11)

where v_r^j and v_r^j are predicted and groundtruth judgments on student's reasoning

Feedback Generation: $Bert_{feed}(\hat{f}, f)$ and $Meteor_{feed}(\hat{f}, f)$ measuring quality between predicted feedback \hat{feed}_k^i and ground-truth feedback $feed_k^i$

C.4 Global Synthesis Dialogue Tutor (f₃) Evaluation

The evaluation of f_3 models follows similar principles:

Diagnosis Extraction:

$$Acc_{\hat{a}g} = \frac{|\{\mathcal{I}|\hat{a}_g = a_g\}|}{|\{\mathcal{I}\}|} \tag{12}$$

where \hat{a}_g and a_g are predicted and ground-truth student's diagnosis selections **Diagnosis Judgment:**

$$Acc_{vd} = \frac{|\{\mathcal{I}|\hat{v_d} = v_d\}|}{|\{\mathcal{I}\}|} \tag{13}$$

where $\hat{v_d}$ and v_d are predicted and ground-truth judgments on student's final diagnosis

Global Feedback Quality: $Bert_{feed_g}(f_g, f_g)$ and $Meteor_{feed_g}(\hat{f}_g, f_g)$ measuring quality between predicted global feedback \hat{feed}_g and groundtruth feedback $feed_g$

Unless otherwise noted, we compute BertScore with the pretrained RoBERTa-large model (Liu, 2019). For the typically longer feedback produced by the f_2 and f_3 tutors, we instead use Longformer (Beltagy et al., 2020), whose extended context window better accommodates lengthy texts. All metrics are calculated by comparing model outputs with the corresponding expert-annotated ground-truth references.

D More Experimental Studies

D.1 Finetuning models to perform single clue analysis.

Model	Acc _{PN}	$Bert_{PN}$	$Meteor_{PN}$	$\operatorname{Bert}_{\operatorname{Spec}}$	$Meteor_{Spec}$	$Bert_{Ind}$	Meteor _{Ind}
Qwen2.5-3b	0.700	0.636	0.279	0.899	0.353	0.896	0.281
Llama3.2-3b	0.722	0.661	0.329	0.906	0.406	0.900	0.314
Qwen2.5-7b	0.747	0.68	0.311	0.903	0.376	0.897	0.296
Llama3.1-8b	0.754	0.692	0.351	0.908	0.426	0.902	0.328
Med42-8b	0.759	0.698	0.363	0.909	0.428	0.903	0.33
Qwen2.5-14b	0.753	0.684	0.313	0.905	0.388	0.899	0.305
Phi4-14b	0.785	0.724	0.391	0.916	0.467	0.907	0.362
Qwen2.5-32b	0.768	0.702	0.341	0.909	0.410	0.902	0.321

Table 7: Experiental results of finetuned models to perform single clue analysis.

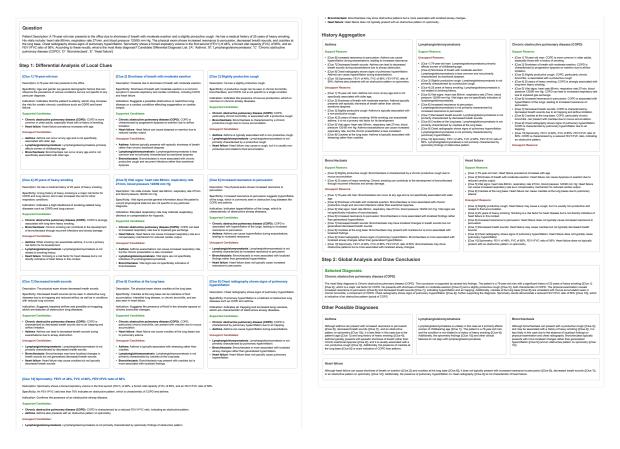
In this section, we investigate models' capability to perform focused analysis on individual clinical clues. To this end, we fine-tuned a specialized model f_4 that takes a single clue description c_i as input and produces its comprehensive analysis \hat{l}_i , i.e., $\hat{l}_i = f_4(c_i)$. Such single-clue analysis is intended for smaller-unit teaching scenarios—for example, when students encounter an unfamiliar symptom

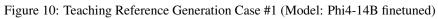


Figure 8: Teaching Reference Generation Case #1 (Model: GPT-40 2-shot In-context Learning)

Question Patient Description: A one-day-old mails is evaluated in the hospital nume	ry for bilous vomiting. The patient ha	s uninated multiple times since birth b	ut has not yet passed meconium. He was born at 38 weeks gestation to a
36 year-old gravida 3 via vaginal delivery. The programcy was complicate quadrupte screen it 15 wests gestation that demonstrated denormated monast of labor demonstrated an annotatic fuel index (VHT) of 20 cm. The prespirations are 30 km/s. On physical exam, the patient is in moderate data presentation? Candidate Othermital Diagnose Lat: (X: "Duodenal atreal	nd by gestational diabetes. The patter aternal serum alpha-feloprotein (AFP atient's two older siblings are both the tress. His abdomen is firm and disten a', '0': 'Intestinal mainstation', 'C': 'Mee	It in other received roctine prenatal (), increased beta-NCG, decreased un adhy. The patient's temperature is 92 ded with hypoactive bowel sounds. W contum lieus', 10°: Necrolizing enterod	are and declined first timester screening. The subsequently underwards conjugated esticity and increased thiblins. A time test ultrascound prior to 17° (26C°C), blood pressure is 58/37 mmHg, pulse is 16Grein, and blord of the following is the most likely esticology of this fetuar's estim, T:: "Plotic stances")
Step 1: Differential Analysis of Local Clues			
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History Aggregation			
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Step 2: Global Analysis and Draw Conclusion			
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Other Possible Diagnoses			
Interstinal matrotation $Whe billow control (Line 1) can suggest matrotation, the task of other supporting findings, such as an association with theory 21 (Clue 3) or polyhydramics (Clue 9), makes this diagnosis less likely. $	Meconium ileus Meconium ileus could explain the symp failure to pass meconium (JClue 2), but syndrome (JClue 2) and polyhydramic	tems of bilous vomiling (Clue 1) and the absence of a link to Down is (Clue R) makes it less likely.	Necrotizing enterocolitis NGC is tess likely because it typically presents later, often is preterm infants, and is not associated with the pretented findings of tissamy 21 ((Clue 20) or polyhydramics ((Clue E)).
Pytoric stemosis Pytoric stemosis is unlikely because it causes non-billous vomiting later in infancy, in	ot immediately after birth.		

Figure 9: Teaching Reference Generation Case #2 (Model: GPT-40 2-shot In-context Learning)





			Support Reasons:	Support Reasons:		Support Reasons:	
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(Clus 7) Two older siblings are both healthy Description: The patient's two older ablings are both healthy. Specificity: Harding healthy ablings does not provide specific historation specified the current exactle cutofics.	[Clue 8] Vital signs: temperature 97.8'7, blood pressure 8313 mmHg, putale 166/sinn, respirations 38/min Descriptor: The Jackets Inspective 80 57 F 06 C/2, blood pressere 8 3037 mmHg, patale 1650min, and patalente see 38/min Paralleline - Teaming and American Inspections are 38/min	(Clue 5) Physical exam: moderate distress, firm and distinted abdoms with hypocachive bowel sounds Dearlyton: Chysical exam, he asked in in modera distance the abdoms is firm and distinct abdoms he bowel ands. Secrifich: After and distance abdoms he hypocachis barel	Although trianical methodox can present of this black working [Cas 10] and dobrind deterations with project in best associal [Cas 12] is less Baily in this case because the detercor of mecorian passage in or specifically associated with trianistation [Cas 2]. Although the second second second second second second second second Down protome, which is more clearly lowed is duplered attem alternative methods. The 8].	[Clue 2] and can present with billou commonly associated with cystic fit The abnormal maternal serum maternal	us does not braically present with the	Although nucertaining entertacelistic car present with Nations vortiking () [] and abstrations damasing and hypotechis beauti instandi ()Car (4), less likely in this case because the patient is a term rescale born way applied delawing ()	
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Figure 11: Teaching Reference Generation Case #2 (Model: Phi4-14B finetuned)

during a case discussion, instructors can query f_4 to generate a concise, clue-specific mini-lecture, thus preserving the natural flow of the session while immediately filling knowledge gaps.

As shown in Table 7, models demonstrate strong performance in analyzing individual clinical clues, such as stance identification (Acc_{PN}) and specificity description ($Bert_{Spec}$). The Phi4-14B model attains the best overall results, with notably high scores in stance accuracy (0.724) and clinical-indication analysis ($Bert_{Ind} = 0.907$). Interestingly, although larger models still outperform smaller ones, the performance gap narrows for single-clue analysis compared with the comprehensive teaching-reference task discussed earlier. This suggests that accurate reasoning over an isolated manifestation appears less sensitive to model scale than integrating multiple pieces of evidence for a final diagnosis.

This observation complements our previous findings for the teaching-reference generator f_1 : while multi-clue reasoning benefits markedly from additional parameters, single-clue analysis remains relatively stable across model sizes. Such robustness underscores the practicality of deploying f_4 in educational settings, as even moderately sized models can deliver dependable focused analyses for on-the-spot teaching.

D.2 Dialogue Tutoring without Teaching Reference

We also study the effect of removing Teaching Reference for f_2 and f_3 models, the result are presented in Table 8.

This study reveals nuanced impacts of removing teaching references across different components of the DDxTutor framework. For local analysis dialogue tutors (f_2), the absence of teaching references leads to a notable decline in student judgment capabilities, as evidenced by decreased Acc_{vs} and Acc_{vr} scores. This degradation demonstrates the crucial role of teaching references in supporting effective dialogue-based instruction at the individual clue analysis stage.

When removing the global teaching reference **G** from the global dialogue tutor (f_3) , the impact on its judgment capability remains relatively minimal. We hypothesize that this robustness stems from the presence of complete Local Clue Analysis **L** in f_3 's context, which continues to serve as a structured foundation for information integration. This suggests that f_3 can inherently derive

accurate diagnostic reasoning by leveraging precise local analyses, even without explicit global teaching guidance.

This observation underscores an insight about our framework: the accuracy of initial local clue analysis significantly influences the quality of subsequent global synthesis. The relative stability of f_3 's performance, contingent on accurate L, validates our framework's emphasis on building strong foundations through precise local analysis before proceeding to global integration.

D.3 Qualitative results

We present some qualitative results for Teaching Reference Generators f_1 , Local Analysis Dialogue Tutor f_2 , and Global Synthesis Dialogue Tutor f_3 .

f1 models: Two Teaching Reference result from GPT-40 (2 shot) results are presented in Figure 8 and Figure 9. At the same time, the comparison results on the same cases, which are generated by finetuned Phi4-14B model, are presented in Figure 10 and 11.

 f_2 and f_3 models: A local clue analysis dialogue case and a global synthesis dialogue case, with results from Llama3.2-3B (finetuned), Phi4-14B (finetuned) and GPT-4o (2 shot In-Context Learning), are presented in Table 10 and 11.

D.4 Validity of Automatic Evaluation Metrics

Although BERTSCORE (Zhang et al., 2019) and METEOR (Banerjee and Lavie, 2005) are widely used in general NLP and medical-NLP researches (Huang et al., 2024; Yao et al., 2024a; Chen et al., 2024a). In this section, we further study how these semantic similarity scores could align with human judgment.

Experimental setup. We randomly sampled 100 predictions produced by our models. Two doctors independently rated each prediction on a five-point Likert scale (1 = very poor, 5 = excellent), judging how well the generated texts matched the ground-truth answers from a medical perspective. We then computed Kendall's τ between the doctors' scores and four common text-generation metrics—BLEU (Papineni et al., 2002), ROUGE-L (Lin, 2004), METEOR, and BERTSCORE.

Results and discussion. Table 9 shows that ME-TEOR and BERTSCORE exhibit stronger correlation with expert judgment ($\tau \approx 0.4-0.5$) than metrics such as BLEU and ROUGE-L. The magnitude of these correlation coefficients is comparable

Model	IOU _a	Acc_{stance}	$\operatorname{Bert}_{reason}$	Acc_{vs}	Acc_{vr}	$Bert_{feed}$	Meteor _{feed}	$\operatorname{Acc}_{\hat{a}g}$	Acc_{vd}	$Bert_{feed_g}$	Meteor _{feed_g}
Qwen2.5-3b	0.995	0.989	0.947	0.828	0.804	0.891	0.397	0.994	0.963	0.907	0.482
Llama3.2-3b	0.996	0.990	0.951	0.842	0.819	0.896	0.430	0.989	0.949	0.908	0.467
Qwen2.5-7b	0.994	0.990	0.947	0.844	0.812	0.892	0.408	0.889	0.969	0.906	0.477
Llama3.1-8b	0.997	0.992	0.953	0.863	0.835	0.898	0.446	0.997	0.975	0.909	0.473
Meditron-8b	0.996	0.991	0.953	0.853	0.830	0.898	0.442	0.997	0.972	0.909	0.475
Med42-8b	0.997	0.993	<u>0.955</u>	0.868	0.838	0.899	<u>0.456</u>	0.989	0.966	0.909	0.475
Qwen2.5-14b	0.982	0.991	0.948	0.866	0.836	0.894	0.421	0.966	0.980	0.909	0.490
Phi4-14b	0.997	0.993	0.955	0.888	0.857	0.901	0.468	0.997	0.989	0.912	0.484
Qwen2.5-32b	0.996	0.993	0.950	<u>0.878</u>	<u>0.846</u>	0.897	0.437	0.997	<u>0.980</u>	<u>0.911</u>	0.498

Table 8: Performance of Local Dialogue Mode f_2 and Global Dialogue Model f_3 after removing Teaching Reference L and G on the simulated dialogue datasets.

$Human \setminus Metric$	BLEU	ROUGE-L	METEOR	BERTScore	D1/D2
D1	0.39***	0.45***	0.49***	0.50***	0.57***
D2	0.26**	0.37***	0.39***	0.41***	0.57***

Table 9: Kendall τ correlations between automatic metrics and human ratings. ** p < 0.01, *** p < 0.001.

to what (Zhang et al., 2019) reported for machinetranslation benchmarks, supporting the appropriateness of using the two semantic metrics in our specialized scenario.

As METEOR and BERTSCORE align more closely and consistently with doctors' assessments, we retain them as the primary automatic semantic metrics for our experiments.

E Implementation Details and Risk Considerations

All results are obtained from a single training run per model, a choice dictated by computational limits but sufficient to demonstrate the method's promise; future work should repeat the experiments with multiple seeds to establish variance. We relied on commercial LLMs, such as GPT4o and Claude, only for polishing language use, with all scientific design and analysis performed by the authors. Although the reported metrics are promising, generated teaching references and dialogues can still contain errors. DDxTutor should therefore be used under the supervision of qualified educators, who can correct inaccuracies and emphasize that the system is a supplementary aid, not a replacement for independent hypothesis-driven medical thinking.

TEACHER

Given the question: 'A 44-year-old man presents to the family medicine clinic for some small bumps on his left thigh. The lesions are non-pruritic and have been present for the last 3 weeks. He reports feeling fatigued and malaise for the past few months. The patient has no known medical problems and takes no medications. He smokes one pack of cigarettes per day and uses intravenous street drugs. His heart rate is 82/min, the respiratory rate is 14/min, the temperature is 36.7°C (98.1°F), and the blood pressure is 126/80 mm Hg. Auscultation of the heart is without murmurs. Lungs are clear to auscultation bilaterally. Three 2-3 mm, dome-shaped, hardened papules are noted on the left thigh. Central umbilication can be observed in each papule. There is a non-tender cervical lymphadenopathy present. Which of the following is the most likely diagnosis?'.Let's delve into the analysis of the clue: 'Three 2-3 mm, dome-shaped, hardened papules are noted on the left thigh. Central umbilication can be observed in each papule.' As part of the differential diagnosis, please evaluate this clue in relation to the current candidate causes: 'A: Kaposi sarcoma, B: Bed bug bite, C: Molluscum contagiosum, D: Atopic dermatitis, E: Acute urticaria', determining whether it supports or cannot support each of them, and provide a clear explanation of your reasoning."

STUDENT

Hmm, okay, so starting with Kaposi sarcoma... honestly, I'm not sure. The description of the bumps and the other clues don't immediately scream Kaposi sarcoma to me, but I kind of feel like I don't know enough about how it might show up to totally rule it out either. So, yeah, I'd have to say 'I don't know" for now bed bug bites... again, I can't say I'm confident here either. The timing and the appearance of the bumps don't immediately make me think of bed bugs, but at the same time, I feel like I don't know their exact presentation well enough to make a clear call. So I'd stick with "I don't know" on this one too.'

LOCAL DIALOGUE TUTOR 1 (LLAMA3.2 3B FINETUNED)

[Student's Response Extraction and Analysis] ["candidate_name": "(A) Kaposi sarcoma", "student's support stance": "N.A.", "student's reasoning": "I don't know.", "student's support stance correctness": false, "student's reasoning correctness": false, "candidate_name": "(B) Bed bug bite", "student's support stance": "N.A.", "student's reasoning: "I don't know.", "student's support stance correctness": false, "student's reasoning correctness": fals

[Teacher's Response] Sure, let's go through your responses together and see how they match up with the ground truth. Here's my feedback:

Candidate: Kaposi sarcoma - **Your stance:** You said you didn't know whether the clue supports or opposes Kaposi sarcoma. - **Ground truth:** The clue does not support Kaposi sarcoma. - **Feedback:** That's totally okay—sometimes it's better to admit when you're not sure than to guess incorrectly. That said, here's the deal: the clue doesn't support Kaposi sarcoma because the specific lesion characteristics described (dome-shaped, hardened papules with central umbilication) aren't consistent with how Kaposi sarcoma presents. Kaposi sarcoma typically shows up as violaceous, irregular plaques or nodules rather than the dome-shaped, umbilicated bumps we're seeing here. So, even though you didn't take a stance, you're right to feel uncertain because this clue doesn't fit Kaposi sarcoma.

Candidate: Bed bug bite* - **Your stance: You also said you didn't know whether the clue supports or opposes bed bug bites. - **Ground truth:** The clue does not support bed bug bites. - **Feedback:** Again, it's totally fine to feel unsure here. But just to clarify, the specific lesion characteristics described (dome-shaped, hardened papules with central umbilication) aren't consistent with bed bug bites. Bed bug bites typically appear as erythematous, itchy papules or bumps, often in a linear or clustered pattern, rather than the dome-shaped, umbilicated bumps we're seeing here. So, this clue doesn't fit bed bug bites.

Final Thoughts: You're doing great—recognizing when you're unsure is a key part of clinical reasoning, and it's better to admit uncertainty than to guess incorrectly. Keep practicing, and these patterns will become more intuitive over time! [/Teacher's Response]

LOCAL DIALOGUE TUTOR 2 (PHI4-14B FINETUNED)

[Student's Response Extraction and Analysis] ["candidate_name": "(A) Kaposi sarcoma", "student's support stance": "N.A.", "student's reasoning": "I don't know.", "student's support stance correctness": false, "student's reasoning correctness": false, "candidate_name": "(B) Bed bug bite", "student's support stance": "N.A.", "student's reasoning": "I don't know.", "student's support stance correctness": false, "student's reasoning correctness": fal

[Teacher's Response] Alright, let's go through your responses one by one and see how they stack up. I'll give you feedback based on what you said and the ground truth reasoning.

1. Kaposi sarcoma You said: *''I'm not sure. The description of the bumps and the other clues don't immediately scream Kaposi sarcoma to me, but I kind of feel like I don't know enough about how it might show up to totally rule it out either. So, yeah, I'd have to say 'I don't know' for now."*

Feedback: No worries about not being sure here—this is a tricky one! Let me clarify: the clue about the "3 2-3 mm, dome-shaped, hardened papules with central umbilication" does **not** support Kaposi sarcoma. Here's why: Kaposi sarcoma lesions are typically **flat or slightly elevated**, **purplish or violaceous**, and **not umbilicated**. So, this clue doesn't fit with Kaposi sarcoma at all.

You're absolutely right to feel unsure, and it's okay to admit when you don't know something. The important thing is to keep learning, and now you know how to rule this one out based on the lesion description!

2. Bed bug bite You said: *"Again, I can't say I'm confident here either. The timing and the appearance of the bumps don't immediately make me think of bed bugs, but at the same time, I feel like I don't know their exact presentation well enough to make a clear call. So I'd stick with 'I don't know' on this one too."*
Feedback: You're on the right track here! The clue about the "3 2-3 mm, dome-shaped, hardened papules with central umbilication" does **not** support bed bug bites. Bed bug bites typically present as **red, itchy welts** or **swollen bumps**, and they don't have the **dome shape or central umbilication** described in this case.

So, while you didn't take a firm stance, your instinct to question this as a possibility is spot on. And now you know that bed bug bites don't match the lesion description here. Great job!

Final Thoughts: You're doing a great job working through these! It's okay to feel unsure sometimes—clinical reasoning takes practice, and you're building the skills to analyze clues more confidently. Keep it up! [/Teacher's Response]

LOCAL DIALOGUE TUTOR 3 (GPT40 (2-SHOT))

[Student's Response Extraction and Analysis] "'json ["candidate_name": "Kaposi sarcoma", "student's support stance": "N.A.", "student's reasoning": "The description of the bumps and the other clues don't immediately scream Kaposi sarcoma to me, but I kind of feel like I don't know enough about how it might show up to totally rule it out either.", "student's support stance correctness": false, "student's reasoning correctness": false, "candidate_name": "Bed bug bite", "student's support stance ": "N.A.", "student's reasoning": "The timing and the appearance of the bumps don't immediately make me think of bed bugs, but at the same time, I feel like I don't know their exact presentation well enough to make a clear call.", "student's support stance correctness": false, "student's reasoning": "The timing and the appearance of the bumps don't immediately make me think of bed bugs, but at the same time, I feel like I don't know their exact presentation well enough to make a clear call.", "student's support stance correctness": false, "student's reasoning correctness": false, "student's Response Extraction and Analysis]

[Teacher's Response] Let's review these candidates you've mentioned.

Kaposi Sarcoma: You expressed uncertainty about this condition and acknowledged a lack of knowledge to either support or rule it out. The truth is, the appearance of the lesions you've described—dome-shaped papules with central umbilication—are not characteristic of Kaposi sarcoma. Kaposi sarcoma typically presents as purple, red, or brown lesions, often in plaques or patches, not dome-shaped papules with central umbilication. So, it's understandable to be unsure, but knowing the hallmark differences can guide you more accurately.

Bed Bug Bite: Similar to Kaposi Sarcoma, you felt unsure about bed bug bites due to a lack of detailed knowledge on presentation. Bed bug bites often appear as small, itchy wheals and usually don't have a central umbilication, which is a characteristic clue for Molluscum contagiosum, not bed bug bites. This understanding could help you differentiate in practice, even when feeling uncertain.

Considering you've now evaluated Kaposi Sarcoma and Bed Bug Bite, let's move forward. Could you analyze the same clue in relation to the remaining candidates: "Molluscum contagiosum," "Atopic dermatitis," and "Acute urticaria"?

Table 10: A Local Analysis Tutoring Dialogue Case.

QUESTION

An 8-year-old girl presents to her pediatrician with intensely pruritic lesions over her buttocks, as shown. These lesions occur intermittently but have worsened over time. Her medical history is remarkable for iron deficiency, for which she is on ferrous sulfate. The patient has also experienced intermittent episodes of mild diarrhea, previously diagnosed as lactose intolerance. Her height is at the 30th percentile and weight is at the 25th percentile for age and sex. An immunoglobulin A (IgA) tissue transglutaminase titer is 5 times the upper limit of normal. Which of the following is the most likely cause of this patient's condition? 'A': 'Crohn's disease', 'B': 'Giardiasis', 'C': 'Celiac disease', 'D': 'Common variable immune deficiency', 'E': 'Henoch-Schonlein purpura' TEACHER

Great job! You've done well analyzing all the local clues. Let's look at what we've discussed: <An Aggregation of Local Clue Analysis>

Now, based on these clues, which candidate do you think is the most likely one?

STUDENT

Based on what we discussed, the clues seem to really point towards it. First, there are the intensely pruritic lesions over her buttocks that could be dermatitis herpetiformis, which is like, strongly linked to celiac disease. Then, there's the iron deficiency; since celiac disease leads to malabsorption, it makes sense it would cause iron deficiency anemia. Also, the patient has intermittent mild diarrhea and growth at lower percentiles, which can also be explained by the malabsorption happening in celiac disease. But I think what really seals it for me is the elevated IgA tissue transglutaminase titer, which is pretty specific for celiac disease. From what I understand, that clue is pretty significant. So yeah, I think Celiac disease fits best here!

GLOBAL DIALOGUE TUTOR 1 (LLAMA3.2 3B FINETUNED)

[Students' Response Analysis]'s Selection: (C) Celiac disease; Selection Correctness: True[/Students' Response Analysis]

[Teacher's Response]Great job! You've correctly identified celiac disease as the most likely cause of this patient's condition. Let's go over why this diagnosis fits so well with the clinical clues we have.Firstly, the intensely pruritic lesions over the buttocks are a significant clue. These are characteristic of dermatitis herpetiformis, which is closely linked to celiac disease. This is a key point that helps us narrow down the diagnosis.Additionally, the elevated IgA tissue transglutaminase titer is a very strong indicator of celiac disease. This test is highly specific for the condition, and its elevation is a crucial finding in this case. The history of iron deficiency amenia is another important clue. Celiac disease leads to malabsorption, which can cause iron deficiency. This aligns perfectly with the patient's medical history.Moreover, the intermittent mild diarrhea and the growth percentiles at the 30th percentile for height and 25th percentile for weight further support the diagnosis of celiac disease. These symptoms suggest malabsorption, which can affect nutrient absorption, leading to these clinical manifestations.You've done an excellent job in piecing together these clues to arrive at the correct diagnosis. Keep up the great work in synthesizing clinical information![/Teacher's Response]

GLOBAL DIALOGUE TUTOR 2 (PHI4-14B FINETUNED)

[Students' Response Analysis]S tudent's Selection: (C) Celiac disease; Selection Correctness: True[/Students' Response Analysis]

[Teacher's Response]Great job! You've correctly identified celiac disease as the most likely diagnosis for this patient. Let's reinforce why this is the correct choice by reviewing the key clinical features that support it.Firstly, the intensely pruritic lesions over the buttocks are highly suggestive of dermatitis herpetiformis, which is a skin manifestation strongly associated with celiac disease. This is a significant clue that points us in the direction of celiac disease. Secondly, the elevated IgA tissue transglutaminase titer is a crucial finding. This test is highly specific for celiac disease, making it a strong indicator that the patient has this condition.Additionally, the history of iron deficiency anemia and intermittent mild diarrhea suggests malabsorption, which is a hallmark of celiac disease. This malabsorption can also explain the patient's growth percentiles being at the lower end, as celiac disease can impair nutrient absorption and affect growth.You've done an excellent job piccing together these clues to arrive at the correct diagnosis. Keep up the great work in analyzing clinical scenarios![/Teacher's Response]"

GLOBAL DIALOGUE TUTOR 3 (GPT40 (2-Shot))

[Students' Response Analysis]Student's Selection: (C) Celiac disease; Selection Correctness: True[/Students' Response Analysis][Teacher's Response]Excellent analysis! You've accurately identified "Celiac disease" as the most likely diagnosis, and your reasoning is well-founded.Let's reinforce the key diagnostic elements that led you to this conclusion. The intensely pruritic lesions over the buttocks you mentioned [Clue 2] are indeed characteristic of dermatitis herpetiformis, which is closely associated with celiac disease. Furthermore, the elevated IgA tissue transglutaminase titer [Clue 6] is highly specific for celiac disease, making this clue particularly significant. Your observation regarding iron deficiency [Clue 3] is critical, as malabsorption due to celiac disease can lead to iron deficiency anemia. The patient's intermittent mild diarrhea [Clue 4] and suboptimal growth percentiles [Clue 5] further support malabsorption, a hallmark of celiac disease. This diagnostic conclusion is a great synthesis of understanding the implications of IgA tissue transglutaminase test results in conjunction with recognizing the dermatological and systemic manifestations of celiac disease in children. Excellent work in putting together these clinical pieces to arrive at the correct diagnosis. Keep honing your diagnostic reasoning skills!/Teacher's Response]

Table 11: A Global Synthesis Tutoring Dialogue Case.