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Large language models for disease diagnosis: a scoping review

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Automatic disease diagnosis has become increasingly valuable in clinical practice. The advent of large language models (LLMs) has catalyzed a paradigm shift in artificial intelligence, with growing evidence supporting the efficacy of LLMs in diagnostic tasks. Despite the increasing attention in this field, a holistic view is still lacking. Many critical aspects remain unclear, such as the diseases and clinical data to which LLMs have been applied, the LLM techniques employed, and the evaluation methods used. In this article, we perform a comprehensive review of LLM-based methods for disease diagnosis. Our review examines the existing literature across various dimensions, including disease types and associated clinical specialties, clinical data, LLM techniques, and evaluation methods. Additionally, we offer recommendations for applying and evaluating LLMs for diagnostic tasks. Furthermore, we assess the limitations of current research and discuss future directions. To our knowledge, this is the first comprehensive review for LLM-based disease diagnosis.

Automatic disease diagnosis is pivotal in clinical practice, leveraging clinical data to generate potential diagnoses with minimal human input¹. It enhances diagnostic accuracy, supports clinical decision-making, and addresses healthcare disparities by providing high-quality diagnostic services². Additionally, it boosts efficiency, especially for clinicians managing aging populations with multiple comorbidities³⁻⁵. For example, DXplain⁶ analyzes patient data to generate diagnoses with justifications. Online services also promote early diagnosis and large-scale screening for diseases like mental health disorders, raising awareness and mitigating risks^{4,7-10}.

Advances in artificial intelligence (AI) have driven two waves of automated diagnostic systems¹¹⁻¹⁴. Early approaches utilized machine learning techniques like support vector machines and decision trees^{15,16}. With larger datasets and computational power, deep learning (DL) models, such as convolutional, recurrent, and generative adversarial networks, became predominant^{12,17-20}. However, these models require extensive labeled data and are task-specific, limiting their flexibility^{1,19,21}. The rise of

generative large language models (LLMs), like GPT²² and LLaMA²³, pre-trained on extensive corpora, has demonstrated significant potential in various clinical applications, such as question answering^{24,25} and information retrieval^{26,27}. These models are increasingly applied to diagnostics. For example, PathChat²⁸, a vision-language LLM fine-tuned with comprehensive instructions, set new benchmarks in pathology. Similarly, Kim et al.²⁹ reported that GPT-4 outperformed mental health professionals in diagnosing obsessive-compulsive disorder, underscoring its potential in mental health diagnostics.

Despite growing interest, several key questions remain unresolved: Which diseases and medical data have been explored for LLM-based diagnostics (Q1)? What LLM techniques are most effective for diagnostic tasks (see Box 1), and how should they be selected (Q2)? What evaluation methods best assess performance of various diagnostic tasks (Q3)? Many reviews have explored the use of LLMs in medicine³⁰⁻³⁷, but they typically provide broad overviews of diverse clinical applications rather than focusing specifically on disease diagnosis. For instance, Pressman et al.³⁸ highlighted

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introducing various clinical applications of LLMs, e.g., pre-consultation, treatment, and patient education. These reviews tend to overlook the nuanced development of LLMs for diagnostic tasks and do not analyze the distinct merits and challenges in this area, revealing a critical research gap. Some reviews^{39,40} have focused on specific specialties, such as digestive or infectious diseases, but failed to offer a comprehensive perspective that spans multiple specialties, data types, LLM techniques, and diagnostic tasks to fully address the critical questions at hand.

This review addresses the gap by offering a comprehensive examination of LLMs in disease diagnosis through in-depth analyses. First, we systematically investigated a wide range of disease types, corresponding clinical specialties, medical data, data modalities, LLM techniques, and evaluation methods utilized in existing diagnostic studies. Second, we critically evaluated the strengths and limitations of prevalent LLM techniques and evaluation strategies, providing recommendations for data preparation, technique selection, and evaluation approaches tailored to different contexts. Additionally, we identify the shortcomings of current studies and outline future challenges and directions. To the best of our knowledge, this is the first review dedicated exclusively to LLM-based disease diagnosis, presenting a holistic perspective and a blueprint for future research in this domain.

Results

Overview of the scope

This section outlines the scope of our review and key findings. Figure 1 provides an overview of disease types, clinical specialties, data types, and modalities (Q1), and introduces the applied LLM techniques (Q2) and evaluation methods (Q3), addressing the key questions. Our analysis spans 19 clinical specialties and over 15 types of clinical data in diagnostic tasks, covering modalities such as text, image, video, audio, time series, and multimodal data. We categorized existing works based on LLM techniques, which fall into four categories: prompting, retrieval-augmented generation (RAG), fine-tuning, and pre-training, with the latter three further subdivided. Table 1 summarizes the taxonomy of mainstream LLM techniques. Figure 2 illustrates the associations between clinical specialties, modalities of utilized data, and LLM techniques in the included papers. Additionally, Fig. 3 presents a meta-analysis, covering publication trends, widely-used LLMs for training and inference, and statistics on data sources, evaluation methods, data privacy, and data sizes. Collectively, these analyses comprehensively depict the development of LLM-based disease diagnosis.

Box 1 | Terms and Concepts

Disease diagnosis: receiving clinical data, such as patient symptoms, medical history, and diagnostic tests, as input and identifying which disease explains the symptoms and signs.

Diagnostic tasks: a type of tasks that generate disease diagnoses or probability estimates for specific conditions, such as differential diagnosis and conversational diagnosis.

Large language models: a type of AI models using deep neural networks to learn the relationships between words in natural language, using large datasets of text to train.

Hallucination: an AI-generated output that is plausible but factually incorrect or unrelated to the input, arising from limitations in training or reasoning.

Prompt: an input or instruction provided to an AI model to guide its response, often designed to elicit specific or task-relevant outputs.

Chain-of-thought: a technique enabling AI to generate multi-step reasoning by breaking down complex tasks into sequential steps for improved accuracy.

Self-consistency prompt: a method that samples diverse reasoning paths and selects the most consistent solution to enhance the reliability of outputs in reasoning tasks.

Study characteristics

As shown in Fig. 2, the included studies span all 19 clinical specialties, and some specialties receive particular attention, such as pulmonology and neurology. While most studies leveraged text modality, multi-modal data, such as text-image⁴¹ and text-tabular data⁴², are widely adopted for diagnostic tasks. Another observation is that various LLM techniques have been applied to diagnostic tasks, and all have been used with multi-modal data (Table 1). Additionally, we find an increasing number of LLM-based diagnostic studies all over the world, reflecting the field's growing significance (Fig. 3a). Among these studies, GPT²² and LLaMA²³ families dominate inference tasks, while LLaMA and ChatGLM⁴³ are commonly adopted for model training (Fig. 3b). Figure 3c shows that most datasets originate from North America (50.6%) and Asia (33.9%), and 50.4% of the studies used public datasets (Fig. 3e). Evaluation methods vary: 66.8% rely on automated evaluation, 28.1% on human assessment, and 5.1% on LLM-based evaluation (Fig. 3d). Figure 3f reveals that the included studies employed large datasets (e.g., 5×10^5 samples) for pre-training diagnostic models, surpassing those primarily using fine-tuning or RAG. This phenomenon aligns with another observation that over half of pre-training models used data from multiple specialties.

Prompt-based disease diagnosis

A customized prompt typically includes four components: instruction (task specification), context (scenario or domain), input data (data to process), and output indicators (desired style or role). In this review, over 60% ($N = 278$) of studies employed prompt-based techniques, categorized as hard prompts and soft prompts. Hard prompts are static, interpretable, and written in natural language. The most common methods included zero-shot ($N = 194$), Chain-of-Thought (CoT) ($N = 37$), and few-shot prompting ($N = 35$). Among them, CoT prompting excels in thoroughly digesting input clinical cues in manageable steps to make a coherent diagnosis decision. Particularly, in differential diagnosis tasks, CoT reasoning allows the LLM to sequentially analyze medical images, radiology reports, and clinical history, generating intermediate outputs that lead to a holistic decision, with an accuracy of 64%⁴⁴. Self-consistency prompting was used in a few studies ($N = 4$). For instance, a study combined self-consistency with CoT prompting to improve depression prediction by synthesizing diverse data sources through multiple reasoning paths. This hybrid approach reduced the mean absolute error by nearly 50% compared to standard CoT methods⁴⁵.

Soft prompt: a learnable embedding added to the input space of a pre-trained model to guide its behavior without modifying the model's parameters.

Retrieval-augmented generation: integrates retrieved data into LLMs, enhancing responses by leveraging external information for improved context and accuracy in content generation.

Fine-tuning: the process of adapting a pre-trained model to a specific task by training it further on a smaller, task-specific dataset.

Supervised fine-tuning: refining a pre-trained model for a task using labeled data to enhance task-specific performance.

Parameter-efficient fine-tuning: adapting pre-trained models to new tasks by updating limited parameters (e.g., adapters), reducing computational costs while preserving performance.

Reinforcement learning from human feedback: a method where models improve outputs by learning from human-provided feedback, aligning behavior with human goals through reinforcement learning.

Pre-training: the foundational training phase of a model on a large, general dataset to learn broad patterns, features, and representations, which can later be adapted to specific tasks through fine-tuning.

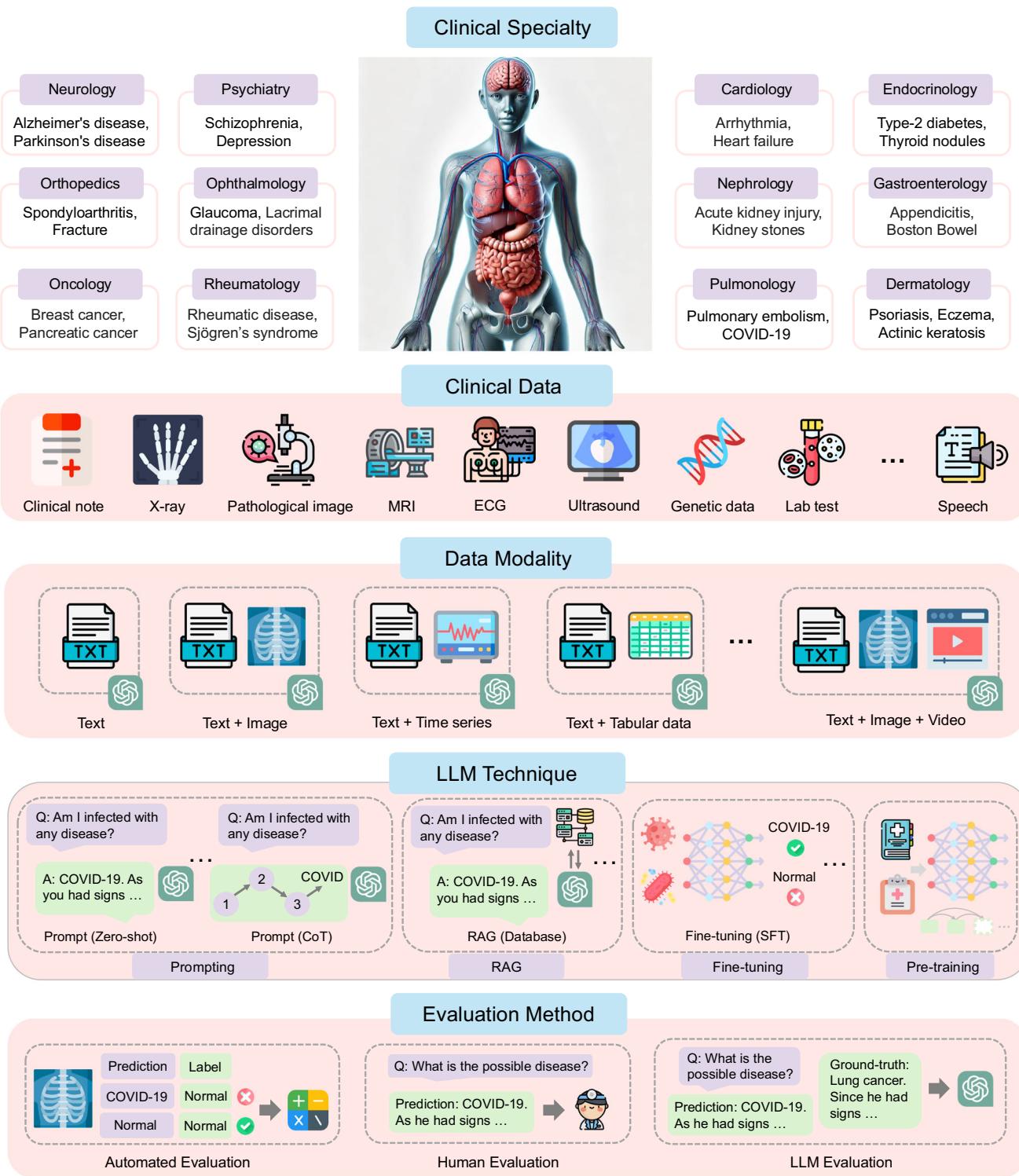


Fig. 1 | Overview of the investigated scope. It illustrated disease types and the associated clinical specialties, clinical data types, modalities of the utilized data, the applied LLM techniques, and evaluation methods. We only presented part of the clinical specialties, some representative diseases, and partial LLM techniques.

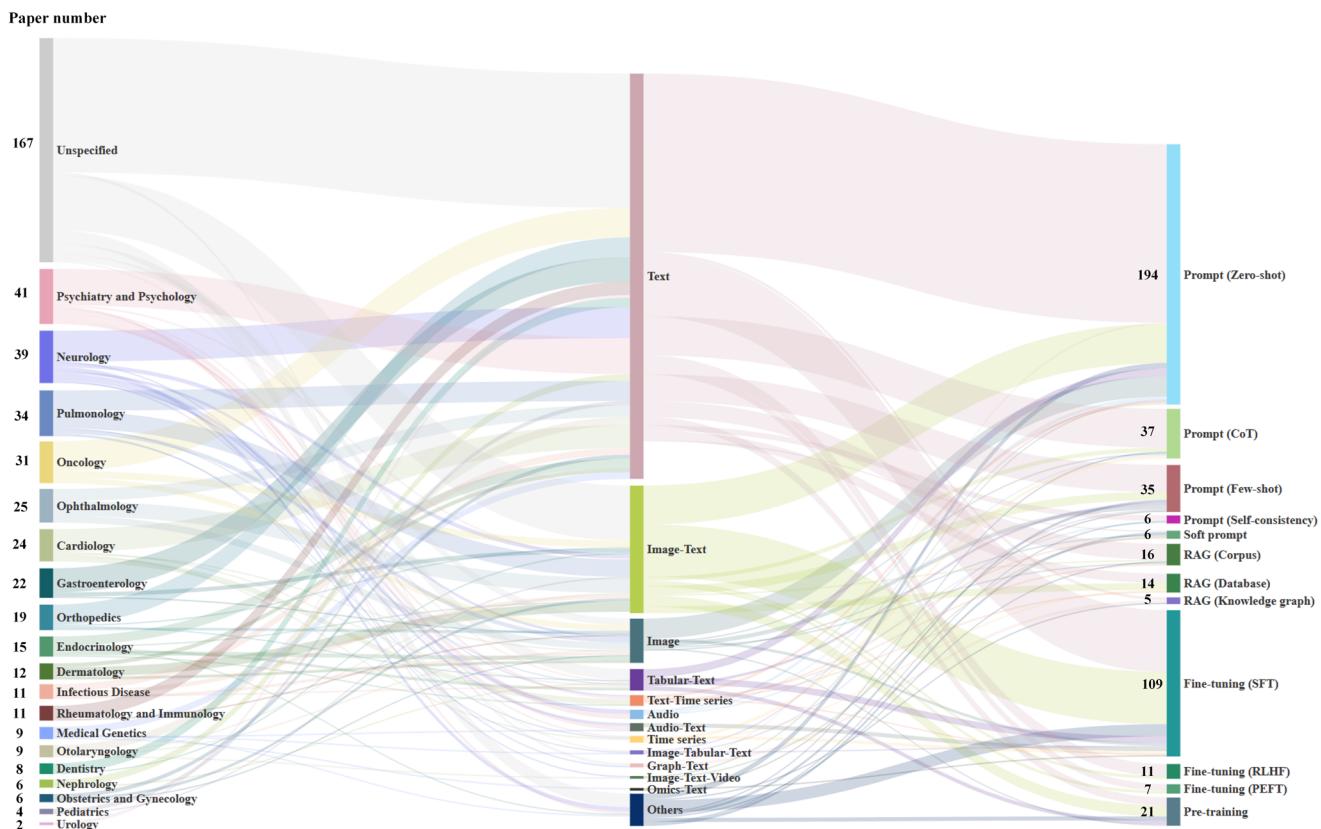
In contrast, soft prompts ($N = 6$) are continuous vector embeddings trained to adapt the behavior of LLMs for specific tasks⁴⁶. These prompts effectively integrate external knowledge, such as medical concept embeddings and clinical profiles, making them well-suited for complex diagnostic tasks requiring nuanced analysis. This knowledge-enhanced approach achieved F1 scores exceeding 0.94 for diagnosing common diseases like hypertension and coronary artery disease and demonstrated superiority in rare disease diagnosis⁴⁷.

Most prompt-based studies ($N = 221$) focused on unimodal data, predominantly text ($N = 171$). Clinical text sources like clinical notes⁴⁸, imaging reports^{49–51}, and case reports^{52,53} were commonly used. These studies often prompted LLMs with clinical notes or case reports to predict potential diagnoses^{54–57}. A smaller subset ($N = 19$) applied prompt engineering to medical image data, analyzing CT scans⁵⁸, X-rays^{59,60}, MRI scans^{58,61}, and pathological images^{62,63} to detect abnormalities and provide evidence for differential diagnoses^{62,64–66}.

Table 1 | Overview of LLM techniques for diagnostic tasks

Techniques	Types	Representative studies
Prompting	Zero-shot	Text ^{196,197} , image ^{65,198} , audio ^{70,72} , text-image ⁶² , text-time series ^{73,199} , text-tabular ²⁰⁰
	Few-shot	Text ^{25,187} , image ⁵⁸ , text-image ^{41,201} , text-image-tabular ¹⁵³
	CoT	Text ^{51,202} , audio ²⁰³ , time series ¹⁵⁵ , text-image ^{44,204}
	Self-consistency	Text ⁸⁹ , audio ²⁰⁵ , text-image-tabular-time series ⁴⁵
	Soft prompt	Text ²⁰⁶ , image ²⁰⁷ , tabular-time series ^{47,208} , text-image-graph ⁵⁹
RAG	Knowledge graph	Text ⁸¹ , text-time series ⁹⁴
	Corpus	Text ^{85,87} , text-image ^{64,86} , text-time series ⁸³
	Database	Text ^{80,93} , text-image ⁹⁰
Fine-tuning	SFT	Text ^{98,209,210} , text-image ^{133,211,212} , text-video ^{102,112} , text-audio ^{111,213} , text-tabular ^{42,200}
	RLHF	Text ^{116,117,214} , text-image ¹¹⁵
	PEFT	Text ^{98,124,215} , text-image ¹⁰⁴
Pre-training	-	Text ^{124,129,131} , text-image ^{109,133,137} , text-tabular ^{135,200} , text-video ²¹³ , text-omics ¹⁰⁹

SFT supervised fine-tuning, RLHF reinforcement learning from human feedback, PEFT parameter-efficient fine-tuning.

**Fig. 2 | Summary of the association between clinical specialties (left), data modalities (middle), and LLM techniques (right) across the included studies on disease diagnosis.**

With the advancement of multimodal LLMs, 57 studies explored their application in disease diagnosis through prompt engineering. Visual-language models (VLMs) like GPT-4V, LLaVA, and Flamingo ($N = 37$) integrated medical images (e.g., radiology scans) with textual descriptions (e.g., clinical notes)^{67–69}. For example, incorporating ophthalmologist feedback and contextual details with eye movement images significantly improved GPT-4V's diagnostic accuracy for amblyopia⁶⁴.

Beyond image-text data, more advanced multimodal LLMs (e.g., GPT-4o and Gemini-1.5 Pro) have also integrated other data types to support disease diagnosis in complex clinical scenarios. Audio and video data have been used to diagnose neurological and neurodegenerative disorders, such as autism^{70,71} and dementia^{59,72}. Time-series data, such as ECG signals and

wearable sensor outputs, were used to support arrhythmia detection^{73,74}. With the integration of tabular data such as user demographics^{75,76}, and lab test results^{47,77}, the applications have been extended to depression and anxiety screening⁴⁵. Omics data has been integrated to aid in identifying rare genetic disorders⁷⁸ and diagnose Alzheimer's disease⁷⁶. Some studies further enhanced diagnostic capabilities by integrating medical concept graphs to provide a richer context for conditions such as neurological disorders⁵⁹.

Retrieval-augmented LLMs for diagnosis

To enhance the accuracy and credibility of the diagnosis, alleviate hallucination issues, and update LLMs' stored medical knowledge without needing re-training, recent studies^{79–81} have incorporated external medical

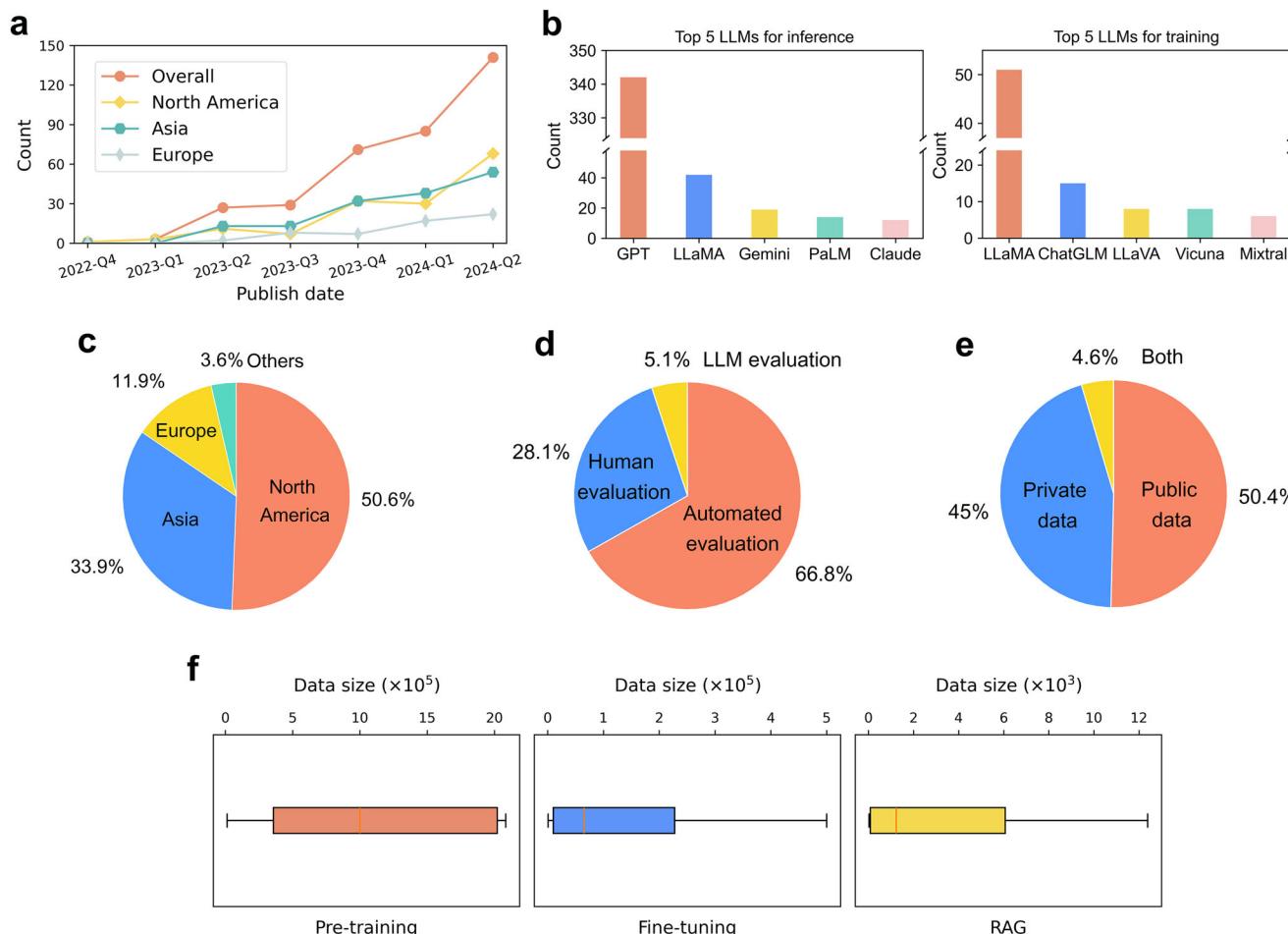


Fig. 3 | Metadata of information from LLM-based diagnostic studies in the scoping review. **a** Quarterly breakdown of LLM-based diagnostic studies. Since the information for 2024-Q3 is incomplete, our statistics only cover up to 2024-Q2. **b** The top 5 widely-used LLMs for inference and training. **c** Breakdown of the data source by regions. **d** Breakdown of evaluation methods (note that some papers utilized multiple evaluation methods). **e** Breakdown of the employed datasets by privacy status. **f** Distribution of data size used for LLM techniques. The red line

indicates the median value, while the box limits represent the interquartile range (IQR) from the first to third quartiles. Notably, pre-trained diagnostic models were often followed by other LLM techniques (e.g., fine-tuning), yet this figure only includes studies that primarily used fine-tuning or RAG. Statistics for prompting methods are not included because: (i) hard prompts generally utilize zero or very few demonstration samples, and (ii) although soft prompts require more training data, the number of relevant studies is insufficient for meaningful distribution analysis.

knowledge into diagnostic tasks. The external knowledge primarily comes from corpus^{64,79,82–88}, databases^{74,80,89–93}, and knowledge graph^{81,94}, in the included papers. Based on the data modality, these RAG-based studies can be roughly categorized into text-based, text-image-based, and time-series-based augmentations.

In text-based RAG, most studies^{80,82,84,85,91–93} utilized basic retrieval methods where external knowledge was encoded as vector representations using sentence transformers, such as OpenAI's text-embedding-ada-002. Queries were similarly encoded, and relevant knowledge was retrieved based on vector similarities. The retrieved data was then input into LLMs with specific prompts to produce diagnostic outcomes. In contrast, Li et al.⁸⁸ developed guideline-based GPT agents for retrieving and summarizing content related to diagnosing traumatic brain injury. They found that these guideline-based GPT-4 agents significantly outperformed the off-the-shelf GPT-4 in terms of accuracy, explainability, and empathy evaluation. Similarly, Thompson et al.⁷⁹ employed regular expressions to extract relevant knowledge for diagnosing pulmonary hypertension, achieving about a 20% improvement compared to structured methods. Additionally, Wen et al.⁸¹ integrated knowledge graph retrieval with LLMs to enable diagnostic inference by combining implicit and external knowledge, achieving an F1 score of 0.79.

In text-image data processing, a common approach^{87,91} involved extracting image features and text features and aligning them within a

shared semantic space. For instance, Ferber et al.⁹¹ used GPT-4V to extract crucial image data for oncology diagnostics, achieving a 94% completeness rate and an 89.2% helpfulness rate. Similarly, Ranjit et al.⁸⁷ utilized multi-modal models to compute image-text similarities for chest X-ray analysis, leading to a 5% absolute improvement in the BERTScore metric. Notably, one study fine-tuned LLMs with retrieved documents to enhance X-ray diagnostics⁸⁶, attaining an average accuracy of 0.86 across three datasets.

For time-series RAG, most studies focused on the electrocardiogram (ECG) analysis^{74,83}. For example, Yu et al.⁸³ transformed fundamental ECG conditions into enhanced text descriptions by utilizing relevant information for ECG analysis, resulting in an average AUC of 0.96 across two arrhythmia detection datasets. Additionally, Chen et al.⁹⁵ integrated retrieved disease records with ECG data to facilitate the diagnosis of hypertension and myocardial infarction.

Fine-tuning LLMs for diagnosis

Fine-tuning an LLM typically encompasses two pivotal stages: supervised fine-tuning (SFT) and reinforcement learning from human feedback (RLHF). SFT trains models on task-specific instruction-response pairs, enabling it to interpret instructions and generate outputs across diverse modalities. This phase establishes a foundational understanding, ensuring the model processes inputs effectively. RLHF further refines the model by aligning its behavior with human preferences. Using reinforcement

learning, the model is optimized to produce responses that are helpful, truthful, and aligned with societal and ethical standards⁹⁶.

In medical applications, SFT enhances in-context learning, reasoning, planning, and role-playing capabilities, improving diagnostic performance. This process integrates inputs from various data modalities into the LLM's word embedding space. For example, following the LLaVA approach⁹⁷, visual data is converted into token embeddings using an image encoder and projector, then fed into the LLM for end-to-end training. In this review, 49 studies focused on SFT using medical texts, such as clinical notes⁹⁸, medical dialogs^{99–101}, or reports^{102–104}. Additionally, 43 studies combined medical texts with images, including X-rays^{102,105–107}, MRIs^{104,107,108}, or pathology images^{109–111}. A few studies explored disease detection from medical videos^{102,112}, where video frames were sampled and converted into visual token embeddings. Generally, effective SFT requires collecting high-quality, diverse responses to task-specific instructions to ensure comprehensive training.

RLHF methods are categorized as online or offline. Online RLHF, integral to ChatGPT's success¹¹³, involves training a reward model on datasets of prompts and human preferences and using reinforcement learning algorithms like Proximal Policy Optimization (PPO)¹¹⁴ to optimize the LLM. Studies have shown its potential in improving medical LLMs for diagnostic tasks^{115–117}. For instance, Zhang et al.¹¹⁷ aligned their model with physician characteristics, achieving strong performance in disease diagnosis and etiological analysis; the diagnostic performance of their model, HuatuoGPT, surpassed GPT-3.5 in over 60% of cases of Meddialog¹¹⁸. However, online RLHF's effectiveness depends heavily on the reward model's quality, which may suffer from over-optimization¹¹⁹ and data distribution shifts¹²⁰. Additionally, reinforcement learning often faces instability and control challenges¹²¹. Offline RLHF, such as Direct Preference Optimization (DPO)¹²², frames RLHF as optimizing a classification loss, bypassing the need for a reward model. This approach is more stable and computationally efficient, proving valuable for aligning medical LLMs^{123,124}. Yang et al.¹²⁴ reported significant performance drops on pediatric benchmarks when the offline RLHF phase was omitted. A high-quality dataset of prompts and human preferences is essential for online RLHF reward model calibration¹²⁵ or the convergence of offline methods like DPO¹²⁶, whether sourced from experts¹¹³ or advanced AI models¹²⁷.

Since full training of LLMs is challenging due to high GPU demands, parameter-efficient fine-tuning (PEFT) reduces the number of tunable parameters. The most common PEFT method, Low-Rank Adaptation (LoRA)¹²⁸, introduces trainable rank decomposition matrices into each layer without altering the model architecture or adding inference latency. In this review, all PEFT-based studies ($N = 7$) used LoRA to reduce training costs^{98,104,124}.

Pre-training LLMs for diagnosis

Pre-training medical LLMs involves training on large-scale, unlabeled medical corpora to develop a comprehensive understanding of the structure, semantics, and context of medical language. Unlike fine-tuning, pre-training enables the acquisition of extensive medical knowledge, enhancing generalization to unseen cases and improving robustness across diverse diagnostic tasks. In this review, five studies performed text-only pretraining on the LLMs from different sources^{129–132}, such as clinical notes, medical QA texts, dialogs, and Wikipedia. Moreover, eight studies injected medical visual knowledge into multimodal LLMs via pretraining^{109,133–137}. For instance, Chen et al.¹³⁷ employed an off-the-shelf multimodal LLM to reformat image-text pairs from PubMed into VQA data points for training their diagnostic model. To improve the quality of the image encoder, pre-training tasks like reconstructing images at tile-level or slide-level¹⁰⁹, and aligning similar images or image-text pairs¹³³ are common choices.

Performance evaluation

Evaluation methods for diagnostic tasks generally fall into three categories (Table 2): automated evaluation¹³⁸, human evaluation¹³⁸, and LLM evaluation¹³⁹, each with distinct advantages and limitations (Fig. 4).

In this review, most studies ($N = 266$) relied on automated evaluation, which is efficient, scalable, and well-suited for large datasets. These metrics can be grouped into three types. (1) Classification-based metrics, such as accuracy, precision, and recall, are commonly used for disease diagnosis. For instance, Liu et al.¹³³ evaluated COVID-19 diagnostic performance using AUC, accuracy, and F1 score. (2) Differential diagnosis metrics, including top-k precision, assess ranked diagnosis lists. Tu et al.¹⁴⁰ employed top-k accuracy to evaluate the correctness of differential diagnosis predictions. (3) Regression-based metrics, such as mean squared error (MSE)¹⁴¹, quantify deviations between predicted and actual values¹⁴². Despite their efficiency, automated metrics rely on ground-truth diagnoses¹⁴³, which may be unavailable, and cannot understand contexts, such as the readability of diagnostic explanations or their clinical utility¹⁴⁴. They also struggle with complex tasks, such as evaluating the medical correctness of diagnostic reasoning¹⁴⁵.

Human evaluation ($N = 112$), conducted by medical experts^{24,138}, does not require ground-truth labels and integrates expert judgment, making it suitable for complex, nuanced assessments. However, it is costly, time-consuming, and prone to subjectivity, limiting its feasibility for large-scale evaluation. Recent studies have explored using LLM evaluation ($N = 20$), a.k.a. LLM-as-Judges¹³⁹, to replace human experts in evaluation and combine the interpretative depth of LLM judgment with the efficiency of automated evaluation. While ground-truth accessibility is not strictly necessary^{99,116}, its inclusion improves reliability¹⁴³. Popular LLMs used for this purpose include GPT-3.5, GPT-4, and LLaMA-3. However, this approach remains constrained by LLM limitations, including susceptibility to hallucinations⁹⁹ and difficulties in handling complex diagnostic reasoning¹⁴⁶. In summary, each evaluation approach has distinct advantages and limitations, with the choice dependent on the specific requirements of the task. Figure 4 guides the selection of suitable evaluation approaches for different scenarios.

Discussion

This section analyzes key findings from the included studies, discusses the suitability of mainstream LLM techniques for varying resource constraints and data preparation, and outlines challenges and future research directions.

The rapid rise of LLM-based diagnosis studies (Fig. 3a) might partially be attributed to the increased availability of public datasets¹⁴⁷ and advanced off-the-shelf LLMs⁵⁷. Besides, the top five LLMs used for training and inference differ significantly (Fig. 3b), reflecting the interplay between effectiveness and accessibility. Generally, closed-source LLMs, with their vast parameters and superior performance¹⁴³, are favored for LLM inference, while open-source LLMs are essential for developing domain-specific models due to their adaptability¹⁴⁸. These factors underscore the dual influence of effectiveness and accessibility on diagnostic applications. Additionally, the regional analysis of datasets (Fig. 3c) reveals that 84.5% of datasets originate from North America and Asia, potentially introducing racial biases in this research domain¹⁴⁹.

Most studies employed prompting for disease diagnosis (Fig. 2), leveraging its advantages, such as minimal data requirements, ease of use, and low computational demands¹⁵⁰. Meanwhile, LLMs' extensive medical knowledge allowed them to perform competitively across diverse diagnostic tasks when effectively applied^{24,143}. For example, a study fed two data samples into GPT-4 for depression detection¹⁵¹, and the performance significantly exceeded traditional DL-based models. In summary, prompting LLMs facilitates the development of effective diagnostic systems with minimal effort, contrasting with conventional DL-based approaches that require extensive supervised training on large datasets^{2,17}.

We then compare the advantages and limitations of mainstream LLM techniques to indicate their suitability for varying resource constraints, along with a discussion of data preparation. Generally, the choice of LLM technique for diagnostic systems depends on the quality and quantity of available data. Prompt engineering is particularly effective in few-data scenarios (e.g., zero or three cases with ground-truth diagnoses), requiring minimal setup^{24,152}. RAG relies on a high-quality external knowledge base,

Table 2 | Overview of evaluation metrics for diagnostic tasks

Type	Evaluation metric	Purpose	Scenario	Representative task
Automated evaluation	Accuracy ²¹⁶	The ratio of all correct predictions to the total predictions	G	DD ¹⁵⁴ , DDX ²¹⁷ , CD ²¹⁸ , RP ²¹⁹ , DRG ¹⁰⁵ , MHD ²²⁰
Precision ⁵⁵		The ratio of true positives to the total number of positive predictions	G	DD ⁵⁵ , CD ²²¹ , MIC ⁴⁴ , RP ¹⁹ , DRG ¹⁰⁵
Recall ⁵⁵		The ratio of true positives to the total number of actual positive cases	G	DD ⁴⁵ , CD ²²¹ , RP ¹⁹ , DRG ¹⁰⁵
F1 ¹³³		Calculated as the harmonic mean of precision and recall	G	DD ⁵⁵ , DDX ²²² , CD ²²¹ , MIC ²²³ , RP ²¹⁹ , DRG ¹⁰⁵
AUC ²²⁴		The area under the Receiver Operating Characteristic curve	G	DD ⁴⁹ , CD ²²⁵ , MIC ²²⁶ , RP ²¹⁹ , DRG ¹⁰⁵ , MHD ²²⁷
AUPR ²²⁸		The area under the precision-recall curve	G	DD ²²⁹ , MIC ²²⁸ , RP ²³⁰ , DRG ²²⁹
Top-k accuracy ¹⁴⁰		The ratio of instances with the true label in the top k predictions to total instances	G	DD ¹⁴⁰ , DDX ¹⁶⁸
Top-k precision ⁶⁰		The ratio of true positives to total positive predictions within the top k predictions	G	DD ¹⁴⁰ , DDX ²²²
Top-k recall ²³¹		The ratio of true positives within the top k predictions to actual positive cases	G	DD ¹⁴⁰ , DDX ¹⁶⁸
Mean square error ¹⁴²		The average of the squared differences between predicted and actual values	G	DD ¹⁴² , RP ¹⁴¹
Mean absolute error ¹⁴¹		The average of the absolute differences between predicted and actual values	G	DD ¹⁴² , RP ¹⁴¹
Cohen's κ ²³²		Measure the agreement between predicted score and actual score	G	DD ³³²
BLUE ¹¹⁵		Calculate precision by matching n-grams between reference and generated text	T	DD ²³³ , CD ²³⁴ , MIC ²³⁵ , DRG ¹¹⁵
ROUGE ¹⁸⁷		Calculate F1-score by matching n-grams between reference and generated text	T	DD ²³³ , CD ¹⁸⁷ , MIC ²³⁵ , DRG ¹¹⁵
CIDEr ¹⁰²		Evaluate n-gram similarity, emphasizing alignment across multiple reference texts	T	CD ¹⁰² , MIC ²³⁶ , DRG ²³⁷
BERTScore ⁸¹		Measure similarity by comparing embeddings of reference and generated text	T	DD ²³⁸ , DDX ¹⁴³ , CD ¹⁸⁷ , DRG ⁸⁷
METEOR ²³⁴		Evaluate text similarity by considering precision, recall, word order, and synonym matches	T	DDX ¹⁴³ , CD ²³⁴ , MIC ²³⁶ , DRG ¹¹⁵
Human evaluation				
Necessity ¹⁸⁷		Whether the response or prediction assists in advancing the diagnosis	T	CD ⁸⁷
Acceptance ²³⁹		The degree of acceptance of the response without any revision	T	DD ¹⁴⁴ , CD ²⁴⁰
Reliability ¹⁷⁶		The trustworthiness of the evidence in the response or prediction	T	DD ¹⁴⁴ , CD ¹⁷⁶
Explainability ⁹⁸		Whether the response or prediction is explainable	T	DD ²⁴¹ , CD ²¹⁸
Human or LM evaluation				
Correctness ²⁴²		Whether the response or prediction is medically correct	T	DD ¹³⁴ , DDX ²¹⁷ , CD ¹⁸⁷ , DRG ²⁴³ , MHD ¹⁷⁶
Consistency ⁹⁹		Whether the response or prediction is consistent with the ground-truth or input	T	DD ¹⁰⁸ , DDX ²⁴¹ , CD ⁹⁹ , MHD ¹⁷⁶
Clarity ⁸⁰		Whether the response or prediction is clearly clarified	T	DD ¹⁴⁹ , CD ²⁴⁴
Professionality ¹⁷⁶		The rationality of the evidence based on domain knowledge	T	CD ⁴⁹ , MHD ¹⁷⁶
Completeness ¹⁸⁷		Whether the response or prediction is sufficient and comprehensive	T	DDX ¹⁴³ , CD ²¹⁸ , DRG ²⁴³
Satisfaction ²⁴⁵		Whether the response or prediction is satisfying	T	CD ¹⁴⁰ , DRG ²³⁷
Hallucination ⁹⁹		Response contains inconsistent or unmentioned information with previous context	T	DDX ²²² , CD ²¹⁸ , DRG ²⁴⁶
Relevance ⁸⁰		Whether the response or prediction is relevant to the context	T	CD ⁹⁰ , DRG ²⁴⁶
Coherence ²⁴⁷		Assess logical consistency with the dialog history	T	CD ¹⁰⁰ , DRG ¹⁹⁰

Since diagnostic tasks might include explanations alongside the predicted diagnosis, existing studies also evaluated these explanatory descriptions. We categorized the metrics based on their application scenarios: **G** denotes that the metric requires ground-truth diagnosis for evaluation, while **T** indicates those applicable to textual descriptions (e.g., generated explanations). Notably, we only present a selection of representative diagnostic tasks from the included papers: disease diagnosis (DDx), differential diagnosis (DD), and diagnostic report generation (DRG), conversational diagnosis (CD), medical image classification (MIC), risk prediction (RP), mental health disorder detection (MHD), and diagnostic report generation (DRG).

Evaluation Approach	Accuracy & Stability	Cost-Effectiveness	Comprehension & Handling Complex Tasks	Representative Papers
Automated Evaluation	Accurate. Strong reproducibility. Quantifiable metrics.	Minimal evaluation cost. The one-time cost of ground-truth preparation. Scalable.	Inability to understand context. Inadaptable to complex tasks and scenarios.	55, 133, 140, 143, 224, 228
Human Evaluation	Accurate. Subject to individual bias.	Costly. Labor-intensive.	Remarkable context understanding. Adaptable to complex tasks and scenarios.	54, 144, 145, 176, 187, 239
LLM Evaluation	LLM-dependent accuracy. Risk hallucination. LLM-dependent stability.	Moderate cost. Scalable. Require expert design.	Moderate context understanding. Limited adaptability for complex tasks and scenarios.	80, 99, 116, 143, 176, 217

Fig. 4 | Summary of the evaluation approaches for diagnostic tasks.

such as databases⁸⁰ or corpora⁸², to retrieve accurate information during inference. Fine-tuning requires well-annotated datasets with sufficient labeled diagnostic cases¹³³. Pre-training, by contrast, utilizes diverse corpora, including unstructured text (e.g., clinical notes, literature) and structured data (e.g., lab results), to establish a robust knowledge foundation via unsupervised language modeling^{42,153}. Although fine-tuning and pre-training facilitate high performance and reliability¹³³, they demand significant resources, including advanced hardware and extensive biomedical data (see Fig. 3f), which are costly and often hard to obtain²⁴. In practice, not all diagnostic scenarios require expert-level accuracy. Applications such as large-scale screenings¹⁵⁴, mobile health risk alerts¹⁵⁵, or public health education³⁰ prioritize cost-effectiveness and scalability. Overall, balancing accuracy with resource constraints depends on the specific use case.

Despite advances in LLM-based methods for disease diagnosis, this scoping review highlighted several barriers to their clinical utility (Fig. 5). One limitation lies in information gathering. Most studies implicitly assume that the available patient information is sufficient for diagnosis, which often fails¹⁵⁶, especially in initial consultations or with complex diseases, increasing the risk of misdiagnosis¹⁵⁷. In practice, clinical information gathering is iterative, starting with initial data (e.g., subjective symptoms), refining diagnoses, and conducting further tests or screenings¹⁵⁸. This process relies heavily on experienced clinicians¹⁴⁰. To reduce this dependence, recent studies have explored multi-round diagnostic dialogs to collect relevant information^{159,160}. For example, AIME¹⁴⁰ uses LLMs for clinical history-taking and diagnostic dialog, while Sun et al.¹⁶⁰ utilized reinforcement learning to formulate disease screening questions. Future efforts could further embed awareness of information incompleteness into models or develop techniques for automatic diagnostic queries¹⁶¹. Another limitation arises from the reliance on single data modalities, whereas clinicians typically synthesize information from multiple modalities for accurate diagnosis⁴⁴. Additionally, real-world health systems often operate in isolated data silos, with patient information distributed across institutions²⁶. Addressing these issues will require efforts to collect and integrate multi-modal data and establish unified health systems that facilitate seamless data sharing across institutions¹⁶².

Barriers also exist in the information integration process. Some studies utilized clinical vignettes for diagnostic tasks without fulfilling the SOAP standard¹⁶³. While adhering to clinical guidelines is crucial¹⁴², limited studies have incorporated this factor into diagnostic systems¹⁶⁴. For example, Kresevic et al.⁸² sought to enhance clinical decision support systems by accurately explaining guidelines for chronic Hepatitis C management. Besides, the integration and interpretation of lab test results pose significant value in healthcare¹⁶⁵. For example, Bhasuran et al.¹⁶⁶ reported that incorporating lab data enhanced the diagnostic accuracy of GPT-4 by up to 30%. A future direction is the effective integration of lab test results into LLM-based diagnostic systems.

Exploring clinician-patient-diagnostic system interactions offers a promising research direction¹⁶⁷. Diagnostic systems are desired to assist clinicians by providing Supplementary information to improve accuracy and efficiency^{58,168}, incorporating expert feedback for continuous refinement. A user-friendly interface is essential for effective human-machine interaction, enabling clinicians to input data and engage in discussions with the system. Human language interaction further enhances usability by allowing natural conversation with LLM-based diagnostic tools¹⁶⁸, reducing cognitive load. Additionally, LLM-aided explanations improve transparency by providing rationales for suggested diagnoses¹⁴⁵, fostering trust, and facilitating informed decision-making among clinicians and patients.

Most of the studies focused on diagnostic accuracy, but overlooked ethical considerations, like explainability, trustworthiness, privacy protection, and fairness¹⁶⁹. Providing diagnostic predictions alone is insufficient in clinical scenarios, as the black-box nature of LLMs often undermines trust⁹⁹. Designing diagnostic models with explainability is desired¹⁴⁵. For example, Dual-Inf is a prompt-based framework that offers potential diagnoses while explaining its reasoning¹⁴³. Besides, since LLMs suffer from hallucinations, how to enhance users' trustworthiness toward LLM-based diagnostic models is worth exploring¹⁷⁰. Potential solutions include using fact-checking tools to verify the output's factuality¹⁷¹. Regarding privacy, adherence to regulations like HIPAA and GDPR, including de-identifying sensitive data, is essential^{26,172}. For example, SkinGPT-4, a dermatology diagnostic system, was designed for local deployment to ensure privacy protection¹⁷³. Fairness is another concern, as patients should not face discrimination based on gender, age, or race¹⁶⁹, but research on fairness in LLM-based diagnostics remains scarce¹⁷⁴.

In the context of modeling, building superior models for accurate and reliable diagnosis remains an exploration. While pre-training on extensive medical datasets benefits diagnostic reasoning¹⁷⁵, many medical LLMs generally lag behind general-domain counterparts in parameter scale^{148,176}, underscoring the potential of developing large-scale generalist models for disease diagnosis. Besides, LLMs are prone to catastrophic forgetting¹⁷⁷, where previously acquired knowledge or skills are lost when learning new information. Addressing this issue facilitates the development of generalist diagnostic models but requires incorporating robust continuous learning capabilities¹⁷⁸. One alternative approach for accurate diagnosis involves coordinating multiple specialized models, simulating interdisciplinary clinical discussions to tackle complex cases¹⁷⁹. For example, Med-MoE¹⁸⁰ is a mixture-of-experts framework leveraging medical texts and images and achieved an accuracy of 91.4% in medical image classification. Additionally, hallucinations in LLMs undermine diagnostic reliability¹⁷⁰, necessitating solutions such as knowledge editing¹⁸¹, external knowledge retrieval⁸², and novel model architectures or pre-training strategies¹⁷⁵. Another promising avenue is longitudinal data modeling, as clinicians routinely analyze EHRs

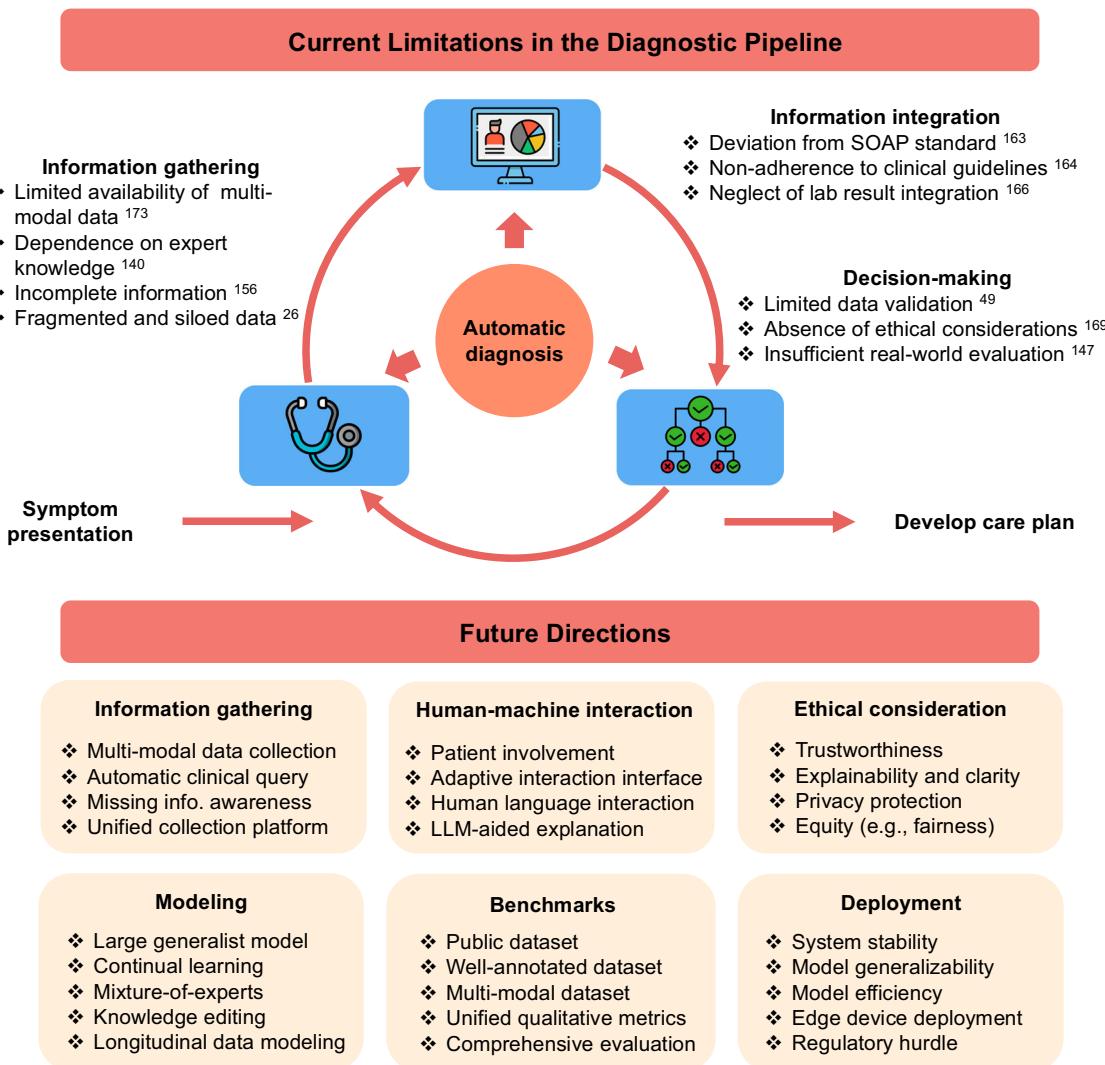


Fig. 5 | Summary of the limitations and future directions for LLM-based disease diagnosis.

spanning multiple years to inform decision-making^{182,183}. Besides, modeling temporal data helps with early diagnosis^{56,184} to improve patient outcomes. For example, early detection of lung adenocarcinoma might increase the 5-year survival rate to 52%¹⁸⁵. However, challenges like irregular sampling intervals and missing data persist¹⁸⁶, necessitating advanced methodologies to effectively capture temporal dependencies²⁵.

Another challenge in developing diagnostic models is benchmark availability¹⁴⁷. In this review, 49.6% of the included studies relied on private datasets, which were often inaccessible due to privacy concerns⁸². Additionally, the scarcity of annotated data limits progress, as well-annotated datasets with ground-truth diagnosis enable automated evaluation, reducing reliance on human assessment¹⁴³. Hence, constructing and releasing annotated benchmark datasets would greatly support the research community¹⁴⁷. Regarding performance evaluation, some studies either used small-scale data⁵⁷ or unrealistic data, such as snippets from college books¹⁴⁵ and LLM-generated clinical notes¹⁴⁷, for disease diagnosis, while large-scale real-world data can truly validate diagnostic capabilities¹⁸². Besides, the lack of unified qualitative metrics is another issue. For example, the evaluation of diagnostic explanation varies in different studies^{143,187}, including necessity¹⁸⁷, consistency¹⁰⁸, and completeness¹⁴³. Unifying qualitative metrics foster a fair comparison. Additionally, many included studies failed to compare with conventional diagnostic models, while recent studies reported that traditional models, e.g., Transformer¹⁸⁸, might beat LLM-based counterparts in clinical

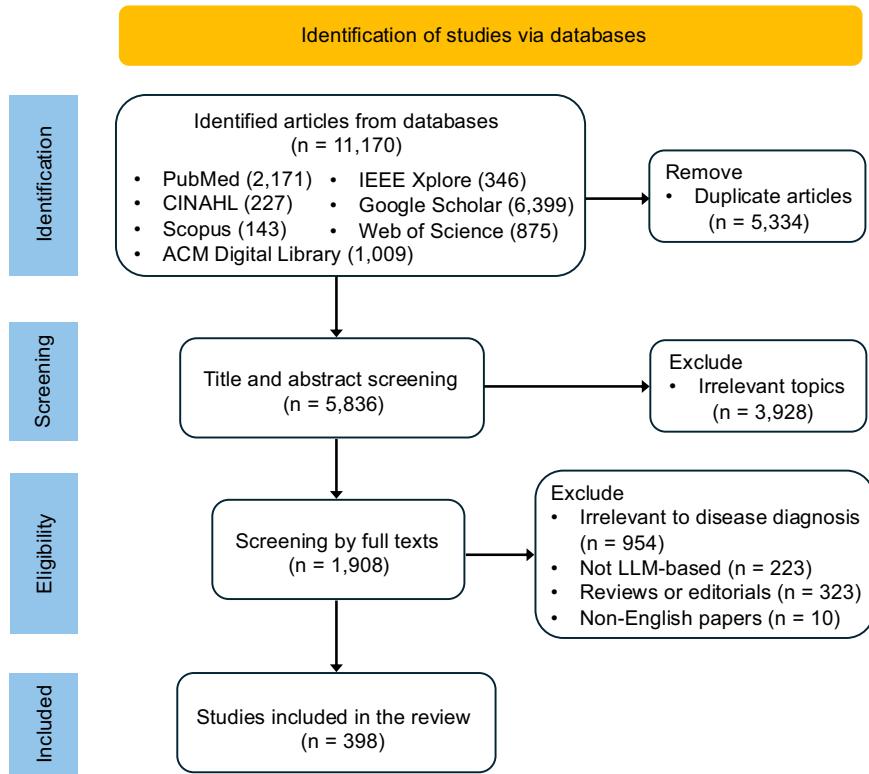
prediction¹⁸⁹. Therefore, future studies should compare with traditional baselines for comprehensive evaluation.

Regarding the deployment of diagnostic systems, several challenges warrant further investigation, including model stability, generalizability, and efficiency. Current studies have highlighted that LLMs often struggle with diagnosis stability¹⁸², fail to generalize well across data from different institutions¹⁹⁰, and encounter efficiency limitations¹⁹¹. For instance, even minor variations in instructions, such as from asking “final diagnosis” to “primary diagnosis”, can drop the accuracy 10.6% on cholecystitis diagnosis¹⁸². Addressing these limitations will advance the reliability and applicability of diagnostic models. Another promising avenue is deploying diagnostic algorithms on edge devices¹⁹². Such systems could enable the real-time collection of health data, such as ECG rhythms¹⁹, to support continuous health monitoring⁹⁵. However, regulatory barriers, including the stringent approval standards imposed by agencies such as the U.S. Food and Drug Administration (FDA) and the European Union’s Medical Device Regulation (MDR)¹⁹³, remain a significant obstacle to clinical adoption. Overcoming these challenges will be vital to ensure the safe and effective integration of LLM-based diagnostics into clinical practice.

In conclusion, our study provided a comprehensive review of LLM-based methods for disease diagnosis. Our contributions were multifaceted. First, we summarized the disease types, the associated clinical specialties, clinical data, the employed LLM techniques, and evaluation methods within this research domain. Second, we compared the advantages and limitations

Fig. 6 | PRISMA flowchart of study records.

PRISMA flowchart showing the study selection process.



of mainstream LLM techniques and evaluation methods, offering recommendations for developing diagnostic systems based on varying user demands. Third, we identified intriguing phenomena from the current studies and provided insights into their underlying causes. Lastly, we analyzed the current challenges and outlined the future directions of this research field. In summary, our review presented an in-depth analysis of LLM-based disease diagnosis, outlined its blueprint, inspired future research, and helped streamline efforts in developing diagnostic systems.

Methods

Search strategy and selection criteria

This scoping review followed the PRISMA guidelines, as shown in Fig. 6. We conducted a literature search for relevant articles published between January 1, 2019, and July 18, 2024, across seven electronic databases: PubMed, CINAHL, Scopus, Web of Science, Google Scholar, ACM Digital Library, and IEEE Xplore. Search terms were selected based on expert consensus (see Supplementary Data 1).

A two-stage screening process focused on LLMs for human disease diagnosis. The first stage involved title and abstract screening by two independent reviewers, excluding papers based on the following criteria: (a) articles unrelated to LLMs or foundation models, and (b) articles irrelevant to the health domain. The second stage was full-text screening, emphasizing language models for diagnosis-related tasks (Supplementary Data 2), excluding non-English articles, review papers, editorials, and studies not explicitly focused on disease diagnosis. The scope included studies that predicted probability values of diseases (e.g., the probability of depression) and the studies in which the foundation models involved text modalities (e.g., vision-language models) and utilized non-text data (e.g., medical images) as input. Our review excluded the foundation models without text modality, such as vision foundation models, because the scope highlighted “language” models. Following related works¹⁹⁴, we further excluded studies purely built on non-generative language models, like BERT¹⁸⁸ and RoBERTa¹⁹⁵, since the generative capability is a critical characteristic of LLMs to facilitate the development of the diagnostic system in the era

of generative AI^{30,31}. Final eligibility was determined by at least two independent reviewers, with disagreements resolved by consensus or a third reviewer.

Data extraction

Information from the articles was categorized into four groups: (1) Basic information: title, publication venue, publication date (year and month), and region of correspondence. (2) Data-related information: data sources (continents), dataset type, modality (e.g., text, image, video, text-image), clinical specialty, disease name, data availability (private or public), and data size. (3) Model-related information: base LLM type, parameter size, and technique type. (4) Evaluation: evaluation scheme (e.g., automated or human) and evaluation metrics (e.g., accuracy, precision). See Supplementary Table 1 for the data extraction form.

Data synthesis

We synthesized insights from the data extraction to highlight key themes in LLM-based disease diagnosis. First, we presented the review scope, covering disease-associated clinical specialties, clinical data, data modalities, and LLM techniques. We also analyzed meta-information, including development trends, the most widely used LLMs, and data source distribution. Next, we summarized various LLM-based techniques and evaluation strategies, discussing their strengths and weaknesses and offering targeted recommendations. We categorized modeling approaches into four areas (prompt-based methods, RAG, fine-tuning, and pre-training), with detailed subtypes. Additionally, we examined challenges in current research and outlined potential future directions. In summary, our synthesis covered data, LLM techniques, performance evaluation, and application scenarios, in line with established reporting standards.

Data availability

The analyzed data are included in this article. Aggregate data analyzed in this study is available at <https://github.com/betterzhou/Awesome-LLM-Disease-Diagnosis>

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Author contributions

S.Z. conceptualized the study and led the work. Z.Z., S.Z., J.Y., and M.Z. searched papers. S.Z., Z.X., M.Z., C.X., Y.G., Z.Z., S.D., J.W., K.X., Y.F., L.X., and J.Y. conducted paper screening and data extraction. S.Z., Z.X., M.Z.,

and C.X. performed data synthesis and contributed to the writing. S.Z., Z.X., M.Z., C.X., D.Z., G.M., and R.Z. revised the manuscript. R.Z. supervised the study. All authors read and approved the final version.

Competing interests

The authors declare no competing interests.

Additional information

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