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A survey of datasets in medicine for large language models

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Abstract

With the advent of models such as ChatGPT and other models, large language models (LLMs) have demonstrated unprecedented capabilities in understanding and generating natural language, presenting novel opportunities and challenges within the medicine domain. While there have been many studies focusing on the employment of LLMs in medicine, comprehensive reviews of the datasets utilized in this field remain scarce. This survey seeks to address this gap by providing a comprehensive overview of the datasets in medicine fueling LLMs, highlighting their unique characteristics and the critical roles they play at different stages of LLMs' development: pre-training, fine-tuning, and evaluation. Ultimately, this survey aims to underline the significance of datasets in realizing the full potential of LLMs to innovate and improve healthcare outcomes.

Keywords: Large language models (LLMs), NLP, dataset in medicine, Q&A system in medicine

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if changes were made.

1. INTRODUCTION

Medicine stands as a critical field intricately connected to human well-being, where the integration of advanced technologies such as large language models (LLMs) has shown promising potential^[1]. Since the introduction of ChatGPT<su[p](#page-17-0)>[2]</sup>, numerous studies have leveraged such models for various medical applications, demonstrating their adeptn[es](#page-17-1)s at tasks ranging from biological information extraction^[3], medical advice consultation, mental health-related applications and clinical report generation. Further[m](#page-17-2)ore, LLMs have demonstrated their potential to improve patient care^[4,5]. The utilization of LLMs in medicine is often facilitated by crafting specialized prompts or instructions, [en](#page-17-3)[a](#page-17-4)bling these models to navigate the complexities of medical data effectively.

Existing LLMs can be classified into three types: encoder-only, encoder-decoder, and decoder-only LLMs. Encoder-only LLMs (e.g., BERT^[6]) are generally used for tasks that involve understanding text, such as clas-sification and sentiment analysis. [E](#page-17-5)ncoder-decoder LLMs (e.g., ChatGLM^[7]) are useful for tasks that involve both understanding and generating text, such as summarization. Decode[r-](#page-17-6)only LLMs (e.g., GPT-4^[8]) excel at generative tasks such as sentence completion and open-ended generation. LLMs in medicinea[re](#page-17-7) developed through a two-stage process: pre-training and fine-tuning. To pre-train LLMs, two common tasks are employed: language modeling and denoising autoencoding. Language modeling involves predicting the next word in a sequence, helping the model to learn language patterns and semantic relationships effectively^[9]. Denoising autoencoding, on the other hand, requires the model to recover the replaced parts of the text, [w](#page-17-8)hich aids in understanding and generating language outputs precisely^[10,11]. The pre-training phase involves training a language model on a large corpus of structured and unst[ru](#page-17-9)[ctu](#page-17-10)red text data. For LLMs in medicine, the corpus may include electronic health records (EHR)^[12], clinical notes^[13], and medical literature^[14]. Pretraining lays the foundation for the LLMs, enabling the[m to](#page-17-11) grasp the broa[d](#page-17-12) nuances of language and [ac](#page-18-0)quire generation skills^[9,15], preparing LLMs for more specialized tasks in subsequent training stages. It is important to note that [s](#page-17-8)[om](#page-18-1)e LLMs are pre-trained on general data and fine-tuned on medical data, while others are trained on medical datasets from scratch. For instance, models such as PubMedBERT^[16] are specifically pre-trained on biomedical corpora, leading to improved performance in healthcare-speci[fic](#page-18-2) tasks compared to models that are fine-tuned on medical data after general pre-training.

Having established a solid foundation through pre-training, the fine-tuning phase focuses on domain-specific adaptation. This stage involves diverse medical corpora, such as dialogue datasets, question-answer (QA) pairs, and instructional texts, ensuring the model excels in specialized tasks^[17]. Researchers have proposed some fine-tuning methods^[18-20] to develop effective medical LLMs. This p[ha](#page-18-3)se ensures that the model becomes proficient in handlin[g h](#page-18-4)[eal](#page-18-5)thcare-specific language, thereby enhancing its accuracy and efficiency. The choice of datasets for training LLMs in medicine depends on the specific task and the type of data required. Common sources of data include EHR, scientific literature, web data, and public knowledge bases. These datasets provide valuable information for training LLMs and enable them to understand and generate medical text, making them versatile tools capable of providing accurate clinical decision support $[21]$. .

Lastly, the evaluation of LLMs employs datasets as benchmarks to rigorously assess their performance, such as text classification [22], semantic understanding [23], question answering (QA) [24], and trustworthiness [25]. The application of LL[Ms](#page-18-6) in the medical field has g[ain](#page-18-7)ed significant attention inr[ec](#page-18-8)ent years. Medical LL[Ms](#page-18-9) have been evaluated and utilized in a range of medical applications, including medical queries [26], medical examin[at](#page-18-10)ions $^{[27]}$, and medical assistants $^{[28]}$. By evaluating the model performance on different datasets, researchers can ide[ntif](#page-18-11)y areas where the model [ex](#page-18-12)cels and areas where it needs improvement [29]. This feedback helps refine the model architecture and train methods. In Figure 1, we describe the constr[uc](#page-18-13)tion process of the medical Q&A LLMs.

While Q&A systems are a key application of LLMs in healthcare, they represent only a small part of their broader potential. Our survey provides an overview of datasets used for both pre-training and fine-tuning

Figure 1. Construction process of the medical Q&A LLMs.

LLMs across a variety of medical tasks. We focus on offering concise summaries and links to these datasets, which can support LLM development across diverse medical applications. Compared with the recent work by Wu *et al*., which focuses on the accessibility and characteristics of publicly available clinical text datasets, our survey encompasses a broader scope by including multimodal datasets [30]. This provides valuable resources for researchers developing LLMs for a wider range of tasks.

To further emphasize the significance of these datasets, it is crucial to recognize that comprehensive overviews of the available data are still scarce. Figure 2 illustrates a timeline of dataset development, spanning dialogue, QA, EHR, summarization, and mul[timodal d](#page-3-0)ata, which have collectively driven the advancement of medical LLMs. By offering a detailed analysis of these datasets and their applications, our survey aims to address this gap in the literature. We hope that this work not only underscores the critical role these datasets play in advancing LLM technology but also encourages further research to unlock the full potential of LLMs in medicine.

Figure 2. Timeline diagram of the development of medical datasets.

2. OVERVIEW OF DATASETS

2.1. Collection

Datasets for medical LLMs originate from diverse sources. Open-source platforms such as Hugging Face and GitHub provide immediate access to pre-curated datasets, while literature reviews and Google searches uncover additional resources. Together, these strategies ensure a broad and comprehensive collection covering a wide range of medical applications. A summary of these datasets is presented in Table 1.

2.2. Sources

In the realm of medical data analysis, especially for training LLMs tailored for healthcare applications, the diversity and specificity of the datasets are paramount. Unlike earlier pretrained language models (PLMs), contemporary LLMs, with their vast array of parameters, necessitate extensive training data encompassing a comprehensive spectrum of medical knowledge. To meet this requirement, a variety of specialized medical datasets have become increasingly available for research purposes. We categorize these corpora into four groups based on their sources: EHR, Scientific Literature, Web Data, and Public Knowledge Bases.

2.2.1 EHR

EHRs contain comprehensive information about patients' medical history, diagnoses, treatments, medication and allergies. They are widely used in medical research and analysis. In this category, MIMIC-III^[31], MIMIC- $\mathrm{IV}^{\left[32\right] }$ and CPRD $\left[33\right]$ are three commonly used datasets for LLM fine-tuning.

MIMIC-III^[31] is an openly available dataset featuring de-identified health data from over 40,000 patients who were admitt[ed](#page-18-14) to the intensive care units at Beth Israel Deaconess Medical Center from 2001 to 2012. This extensive dataset includes records from 58,976 hospital admissions across 38,597 patients, positioning it as a crucial resource for in-depth healthcare research. It is renowned for its substantial inclusion of 2,083,180 deidentified notes, filled with detailed patient histories and clinician observations. MIMIC-III provides a wide array of data, encompassing patient demographics, hourly vital sign measurements, lab test results, medical procedures, medication records, caregiver notes, imaging reports, and mortality data, both during hospital

Table 1. An overview of commonly used datasets in medicine for LLMs

stays and post-discharge.

Evolving from MIMIC-III, MIMIC-IV $^{[32]}$ is a relational database detailing actual patient admissions at a tertiary academic medical center in Bosto[n, M](#page-18-15)A, USA. This updated version includes modernized data spanning from 2008 to 2019, capturing a wide array of medical information such as laboratory measurements, administered medications, and recorded vital signs. Designed to support a broad spectrum of healthcare research, MIMIC-IV is instrumental for investigations in clinical decision-making, patient care optimization, and epidemiological studies, providing a rich foundation for advancing medical knowledge and improving healthcare outcomes.

In contrast to the U.S.-based datasets, CPRD^[33] contains coded and anonymized EHR data from a network of over 2000 practices in the UK. This dataset [is](#page-18-16) linked to secondary care and other health and administrative databases, providing a representative sample of the population by age, sex, and ethnicity. It includes information on demographic characteristics, diagnoses and symptoms, drug exposures, and vaccination history. Covering 60 million patients, among whom nearly 18 million are currently registered, CPRD notably provides a significant longitudinal perspective on health outcomes and trends, with 25% of these patients having been followed for at least 20 years.

2.2.2 Scientific literature

Scientific literature datasets, such as PubMed^[34], provide access to a vast collection of research papers, articles, and abstracts related to life sciences and bio[med](#page-18-17)ical topics. These datasets are valuable for training healthcare language models as they contain high-quality academic and professional text.

PubMed is a freely accessible database that supports access to several National Library of Medicine (NLM) literature resources, which contains citations and abstracts related to biomedical topics and life sciences. It provides more than 36 million citations and abstracts of biomedical literature, including content from MED-LINE, PubMed Central (PMC)^[35], and online books. These citations may include links to full-text content from other sources such as PM[C o](#page-18-18)r the publisher's website. Launched online in 1996, PubMed is maintained and upgraded by the National Center for Biotechnology Information (NCBI). The dataset contains high-quality text, making it particularly suitable for training medical LLMs.

Closely related to PubMed, PMC serves as a significant repository that archives open access full-text articles published in biomedical and life sciences journals. PMC contains over eight million full-text article records covering biomedical and life science research from the late 1700s to the present, including articles formally published in a scholarly journal, author manuscripts accepted for publication in a journal, and preprint versions of articles. It is a crucial component of the NLM collection, complementing its extensive print and licensed electronic journal holdings.

Turning to datasets tailored for systematic reviews, $\mathrm{RCT}^{\,[36]}$ comprises 4,528 reviews conducted by the Cochrane collaboration's members. These systematic reviews sou[rce](#page-18-19)d from PubMed of all trials are relevant to specific clinical questions. RCT is constructed from the abstracts of systematic reviews and the clinical trials' titles and abstracts that are summarized by these reviews. Similarly, CDSR ^[38] is a dataset of high-quality systematic reviews in various medical domains. It contains a training set of 5,1[95](#page-18-21) source-target pairs, a validation set of 500 abstract pairs, and a test set of 1,000 abstract pairs.

For summarization tasks in the biomedical field, MS^{\degree}2 (multi-document summarization of medical studies)^[37] is a multi-document summarization dataset comprising over 470 K documents and 20 K summaries in [the](#page-18-20) biomedical domain. It is constructed from papers in the Semantic Scholar literature corpus, containing a large amount of related markup.

Broadening the scope to multidisciplinary research, S2ORC^[41] is a large corpus consisting of 81.1 million English-language academic papers covering many academic [dis](#page-18-25)ciplines. It represents a significant stride in academic literature aggregation. S2ORC is meticulously constructed using data from the Semantic Scholar literature corpus, which integrates papers from a variety of sources including publishers, archives such as arXiv or PubMed and resources such as MAG.

For highly specialized research, CORD-19^[42] is a dataset containing over 1M papers on COVID-19 and related historical coronavirus research, including [ful](#page-18-24)l text content for nearly 370 K papers. This dataset is valuable for pandemic research, emphasizing the essential role of domain-specific resources in meeting urgent biomedical needs.

2.2.3 Web data

Web data includes a broad spectrum of text that can be sourced from the internet, embodying a vast array of information types and formats. Among these, social media content stands out as one of the most prevalent and rich data sources. Reddit is a popular online platform where users can submit various types of content, including links, text posts, images, and videos. These submissions can be endorsed or disapproved by others through "upvotes" or "downvotes". Content that garners a significant number of upvotes is typically regarded as valuable and can serve as a rich source for creating high-quality datasets.

WebText^[44] is a well-known corpus, which is compiled from highly upvoted links on Reddit, but it is not publicly avail[ab](#page-19-20)le. In response to the limited availability of WebText, OpenWebText $^{[45]}$, an open-source alternative, is released.

COMETA^[43] is an entity linking dataset of medical terminology. It contains 20,015 English biomedical concept mentionsf[rom](#page-19-0) Reddit expert-annotated with their corresponding SNOMED-CT links, covering a wide range of concepts such as symptoms, diseases, anatomical expressions and procedures across a range of conditions.

Colossal Clean Crawled Corpus $(C4)^{[11]}$ is a dataset containing about 750 GB clean English text scraped from the Common Crawl web dump.

2.2.4 public knowledge bases

There exist many public knowledge bases in medicine, such as $\mathrm{UMLS}^{\, [46]}$, $\mathrm{CMeKG}^{\, [47]}$ and $\mathrm{DrugBank}^{\, [48]}$.

UMLS^[46] is one of the most popular repository of biomedical vocabularies, which is developed by the US NLM. [It h](#page-19-2)as over two million names representing around 900,000 concepts sourced from over 60 families of biomedical vocabularies, as well as 12 million relations among these concepts.

CMeKG[47] is a Chinese medical knowledge graph, which is a structured description of professional medical knowled[ge.](#page-19-21) It is constructed by referencing authoritative international medical standards and a wide range of sources such as clinical guidelines, industry standards and medical textbooks. This knowledge graph lays a foundation for a medical QA system and serves as a comprehensive resource for medical information.

DrugBank^[48] is a comprehensive freely available database containing detailed drug, drug-target, drug action and drug i[nte](#page-19-3)raction information. The most recent version (5.1.12) has 16,581 drug entries including 2,769 approved small molecule drugs, 1,620 approved biologics, 135 nutraceuticals and over 6,723 experimental drugs. Additionally, 5,291 non-redundant protein sequences are linked to these drug entries. DrugBank's data is highly structured and accessible in various formats, including SMILES, SDF, MOL, PDB, InChI, and InChIKey for chemical structures, FASTA for sequence data, and XML and JSON for textual data. These standardized formats make DrugBank's data easily usable for training LLMs in tasks such as drug discovery and pharmacological research. Known for its high data quality, DrugBank curates information from peerreviewed scientific literature, patents, and reputable databases, offering extensive and comprehensive data on drug targets. Regular updates ensure the dataset remains accurate and reliable. DrugBank's structured data on drug-target interactions makes it highly suitable for LLM tasks such as QA on drug-related topics. Additionally, its detailed pharmacological data supports LLMs in generating accurate summaries of drug mechanisms, and its information on interactions and metabolic pathways can be leveraged for dialogue generation in systems assisting healthcare professionals in drug-related decision-making.

It is important to note that, while datasets such as MIMIC-III form the foundation for many LLMs, they do have certain limitations. For example, MIMIC-III comes from a single institution - Beth Israel Deaconess Medical Center in Boston, USA, which makes it less representative of healthcare practices and patient demographics in other regions or countries, limiting the generalizability of models trained on this data. The dataset spans from 2001 to 2012, and medical practices, technologies, and treatment protocols have evolved significantly since then. This makes LLMs trained on MIMIC-III potentially less effective when applied to current healthcare scenarios. Moreover, MIMIC-III primarily contains data from critical care units, focusing on patients with severe conditions. As a result, it lacks coverage of broader healthcare settings, such as outpatient care or chronic disease management, which limits the scope of models trained on this dataset. Even MIMIC-III, one of the largest clinical text datasets available, contains only 0.5 billion tokens of clinical text from a single hospital - far less than the tens of billions used in LLM training^[88]. Similarly, while CPRD provides a robust source of UK population health data, it has certain limitation[s.](#page-20-16) The dataset primarily reflects the ethnic and disease distributions specific to the UK, which may introduce regional and population biases that affect the model's generalizability. Additionally, CPRD data are primarily derived from primary care practices, making it more representative of outpatient care. Although CPRD includes linkages to hospital care and other health-related datasets, its emphasis on primary care may limit the depth of information available for studies that require detailed inpatient or intensive care data. This highlights the pressing need for more extensive and diverse clinical text datasets to advance clinical LLMs.

2.3. Datasets structure

Datasets for medical LLMs can be classified into two broad categories based on their structure: conventional text data and multimodal data.

2.3.1 Conventional text data

This category comprises datasets primarily centered on text, which are essential for training models to understand and generate human language. Within this category, QA and dialogue datasets stand out as the most widely utilized.

QA datasets

QA datasets are designed to train models capable of answering human-posed questions. These datasets typically consist of question-answer pairs, which are used to enable models to comprehend the question context and generate accurate responses based on the information available in the dataset or linked knowledge bases.

cMedQA2^[49], the extension and amendment of version 1.0, is a dataset designed specifically for Chinese community me[dic](#page-19-4)al QA. It is collected from an online Chinese medical QA forum, where users post their queries and receive answers from qualified doctors. It contains 108,000 questions and 203,569 answers, doubling the number of questions and answers compared to version v1.0, and performs some data cleaning preprocessing steps, such as eliminating greeting words and replacing English punctuation with Chinese punctuation.

webMedQA^[50] is a real-world Chinese medical QA dataset collected from professional health-related consultancy we[bsi](#page-19-5)tes. The dataset is collected through some steps including data preprocessing and removing questions with more than one best-adopted reply. It consists of 63,284 questions, covering most of the clinical departments of common diseases and health problems.

Huatuo-26M^[51], named after the ancient Chinese physician Hua Tuo, stands as the largest Chinese medical QA dataset av[ai](#page-19-6)lable today. It is collected from multiple sources through text cleaning and data deduplication methods, including an online medical consultation website, medical encyclopedias, and medical knowledge bases. It contains over 26 million QA pairs, covering various aspects such as diseases, symptoms, treatment methods, and drug information. Huatuo-26M significantly expands the scale of existing medical QA datasets and offers an unprecedented resource in the Chinese medical domain.

PubMedQA^[53] is a novel biomedical QA dataset collected from PubMed abstracts. It aims to answer research questions wi[th](#page-19-8) yes/no/maybe, using the corresponding abstracts. PubMedQA has 1K expert-annotated, 61.2 K unlabeled and 211.3 K artificially generated QA instances. It is the first QA dataset where reasoning over the contexts, especially their quantitative contents, is required to answer the questions.

HealthSearchQA^[59] is a new dataset of 3,173 commonly searched consumer medical questions. It is curated using seed medic[al](#page-19-11) conditions and their associated symptoms. The dataset diverges from other medical text QA datasets in three significant ways including question only, free text response and open domain.

Dialogue

Dialogue datasets record conversational exchanges, mirroring real human-to-human interactions. They are crucial for training models to understand conversational nuances, and provide accurate and contextually appropriate responses.

MedDialog-CN^[60] is a Chinese dataset containing conversations between doctors and patients. It contains 1.1 million dial[ogu](#page-19-12)es, covering 29 broad categories of specialties and 172 fine-grained specialties. The data is collected from an online consultation website. MedDialog-EN^[60] is an English dataset with 0.26 million dialogues, covering 51 categories of communities and 96 specialti[es](#page-19-12). The data is collected from two online platforms of healthcare services.

IMCS-21^[61] is a dialogue dataset that contains a total of 4,116 annotated samples with 164,731 utterances, coveringt[en](#page-19-22) pediatric diseases: bronchitis, fever, diarrhea, upper respiratory infection, dyspepsia, cold, cough, jaundice, constipation and bronchopneumonia.

Pubmed Causal^[63] is a dataset for causal statements in science publications, containing 2,446 annotated sentences.

Instructions

Instruction datasets comprise step-by-step directives or guidelines intended to train models to perform specific tasks or understand procedural language, which is particularly useful for instructional AI applications.

Alpaca [69] is a dataset based on the self-instruct [89] method. This dataset employs the text-davinci-003 model on the [17](#page-19-19)5 human-crafted instruction-output [pai](#page-20-17)rs from Self-Instruct to generate 52,000 new instructions along with inputs and outputs. Moreover, around 40% of the examples have an input in the final dataset.

sft-20 ${\rm k}^{[\tau_1]}$ is a dataset originating from the QiZhen medical knowledge base, which includes real medical QA data be[tw](#page-20-1)een patients and doctors, and drug text knowledge. It constructs an instructional dataset by applying specific question templates to semi-structured data. Qizhen Medical Knowledge Base collects QA data on various topics such as diseases, medications, diagnostic tests, surgeries, prognoses, and dietary information, summing up to 560,000 instructional entries.

ShenNong-TCM-Dataset $^{[72]}$ is constructed on the foundation of TCM-neo4j, an open-source medical knowledge graph. It employs an [in](#page-20-2)novative entity-centric self-instruction method, leveraging ChatGPT to generate

over 110,000 pieces of instructional data centered around TCM.

Summarization

Summarization is a concise description that captures the salient details of information. In the medical domain, summarization can be useful for helping people easily understand and address the diverse nature of questions and answers.

MeQSum^[73] is a dataset comprising 1,000 consumer health questions and their expert-crafted summaries. The questi[on](#page-20-3)s are carefully chosen from a collection provided by the U.S. NLM, ensuring a diverse and representative selection of consumer health questions. The dataset is particularly noteworthy for its method of summarization, meticulously carried out by three medical experts adhering to stringent guidelines to ensure the quality and utility of the summaries. The summarization process followed by these experts is governed by two critical principles. First, the summary must allow the retrieval of correct and complete answers to the original questions. Second, the summary cannot be shortened further without meeting the first condition.

CHQ-Summ^[74] is a CHQ summarization dataset consisting of 1,507 consumer health questions and corresponding su[mm](#page-20-4)aries. It is created from the Yahoo community QA forum that has a diverse set of users' questions. It contains additional annotations about question focus and question type of the original question, which are all annotated by domain experts.

MEDIQA-AnS^[75] is a dataset designed for question-driven, consumer-focused summarization. It contains 156 consumer h[ea](#page-20-5)lth questions, corresponding answers to these questions, and manually generated summaries of these answers.

In Table 2, we compare some medical QA datasets. A notable distinction among the datasets is their source. M[ost datas](#page-10-0)ets, such as cMedQA2, webMedQA, and Huatuo-26M, are sourced from community-driven medical consultation platforms such as Xywy Community, Baidu Doctor, and Qianwen Health. These platforms are primarily Chinese-language websites, which means the datasets predominantly represent Chinese patients and healthcare practitioners, resulting in limited ethnic and geographical diversity. Additionally, the range of diseases covered may not be fully comprehensive, and the quality and reliability of the answers can vary due to differences in expertise among the responding doctors. Despite the limitations, these datasets provide a rich repository of user-generated medical inquiries and professional responses, making them particularly useful for developing consumer-facing medical QA systems. On the other hand, PubMedQA and MeQSum leverage more formal medical literature and research databases such as PubMed and the U.S. NLM. While these datasets do include some international content, they are primarily based on U.S. sources, thus reflecting healthcare practices, patient demographics, and disease prevalence patterns that are more representative of the American population. This may limit their generalizability to other ethnic groups and geographical regions. A major limitation of datasets sourced from online communities is the variability in the accuracy and reliability of the information. These datasets may have limited coverage of diseases, and the quality of responses can vary significantly based on the expertise of the contributing doctors. Conversely, datasets such as PubMedQA offer a higher degree of reliability but may lack the diversity of inquiries that arise from everyday medical concerns.

A critical factor in the utility of these datasets is the inclusion of real versus generative data. Datasets such as Huatuo-26M, Chatmed, and ShenNong incorporate both real and synthetic data. The inclusion of generative data, when carefully modeled, can significantly enhance the coverage of edge cases and underrepresented medical conditions. It enables QA systems to handle rare or hypothetical medical inquiries, which may not often arise in real-world consultations. However, generative data can also introduce noise into the training process, particularly if the synthetic data is of lower quality or inaccurately generated. This underscores the need for careful validation of generated content to prevent the dissemination of incorrect or harmful medical advice.

Each dataset is suited for specific applications in medical QA. For instance, IMCS-21 is tailored to pediatric medicine, making it highly specialized for QA systems focusing on children's health. The detailed, contextspecific nature of this dataset makes it ideal for systems that require deep, domain-specific knowledge. Similarly, CovidDialog, released in 2020 in response to the COVID-19 pandemic, offers targeted information on COVID-19 and other pneumonia-related conditions, making it an invaluable resource for QA applications focused on respiratory diseases. The release dates of these datasets are indicative of their relevance to current healthcare challenges. For example, datasets such as Huatuo-26M and Chatmed, released in 2023, reflect the latest developments in large-scale QA system training and integrate recent developments in medical knowledge, making them well-suited for modern medical applications.

In summary, each dataset within this comparative analysis offers unique strengths depending on the target application, the need for real or synthetic data, and the specific medical domain. While large datasets such as Huatuo-26M and sft-20k are indispensable for building comprehensive, large-scale QA models, smaller datasets such as PubMedQA and MeQSum remain critical for high-accuracy tasks that demand evidencebased answers. Thus, the choice of dataset should be guided by the specific needs of the QA system - whether for broad coverage, specialized knowledge, or linguistic precision.

2.3.2 Multimodal data

This category includes datasets that involve multiple modalities, such as text, images, and time series data. In the medical domain, multimodal language models offer a promising direction for further research. In Figure 3, we present a partial display of the content from four multimodal datasets.

VQA Datasets

Medical visual question answering (Med-VQA) has tremendous potential in medicine, particularly in fields such as radiology and pathology. These two fields are rich in both imaging data and textual reports, making them prime candidates for VQA applications.

VQA-RAD^[76] is a manually-crafted dataset in radiology where questions and answers are given by clinicians. It contains [3,5](#page-20-6)15 visual questions of 11 types and 315 corresponding radiological images.

 $\rm SLAKE^{\rm [77]}$ is a large bilingual dataset with comprehensive semantic labels annotated by experienced physicians and an [ext](#page-20-7)endable knowledge base for Med-VQA. It contains 642 radiology images including 12 diseases and 39 organs of the whole body, with 14,028 QA pairs and 5232 medical knowledge triplets.

PathVQA^[78] is a pathology VQA dataset containing 32,799 QA pairs of eight categories, generated from 4,998

Figure 3. Partial content display of multimodal dataset.

images. The majority of questions in PathVQA are open-ended, and the other half are "yes/no" questions.

ROCO^[80] is a multimodal image dataset, containing over 81K radiology images with several medical imaging modalit[ies](#page-20-10). It is constructed by retrieving all image-caption pairs from PMC. All images have corresponding captions, keywords extracted from the image caption, UMLS Concept Unique Identifiers and Semantic Type.

MedICaT $^{[81]}$ is a dataset that encompasses medical figures, captions, subfigures and subcaptions, and inline references [th](#page-20-11)at enable the study of these figures in context. The dataset's content is meticulously extracted from open-access articles available in PMC, ensuring figures and captions. Additionally, the corresponding reference texts are sourced from the S2ORC $^{[41]}$. It contains 217,060 figures collected from 131,410 open-access scientific papers. Moreover, the dataset inc[lud](#page-18-25)es inline references for approximately 25,000 figures from the ROCO^[80] dataset.

Among these datasets, VQA-RAD is a pioneering dataset for radiology VQA, consisting of radiological images and related medical questions. It focuses on enhancing the understanding of diagnostic images through QA, making it valuable for medical decision support systems. SLAKE extends the concept by incorporating a broader set of modalities (CT, MRI, X-rays) and both visual and textual inputs. This allows for more complex, multimodal reasoning tasks. PathVQA is specific to pathology images, such as histopathological slides, and is valuable for disease diagnosis in pathology using VQA techniques.

U-Xray, CheXpert, PadChest, and MIMIC-CXR focus on chest X-ray classification and segmentation. These datasets are widely used in developing models for automatic disease detection (e.g., pneumonia, pneumothorax) and diagnosis from X-rays. CheXpert, with over 224,000 labeled images, provides precise annotations for several lung diseases, making it a benchmark for chest disease classification. PadChest contains over 160,000 labeled images and expands its scope with Spanish-language reports, contributing to multilingual model training. ROCO and ROCOv2 are aimed at report generation from medical images. They cover a variety of medical imaging modalities, such as X-rays, CT scans, and MRIs, along with their corresponding textual descriptions.

These datasets enable models to generate human-readable medical reports, which can assist radiologists in producing structured reports efficiently.

MIMIC-CXR, with over 370,000 chest X-rays and associated clinical reports, is one of the largest and most diverse datasets available for medical imaging research. It is widely used for tasks such as disease classification, report generation, and clinical decision support. Similarly, CheXpert and PadChest offer large volumes of annotated images but focus more specifically on X-ray data, whereas MIMIC-CXR includes a broader range of accompanying clinical information, making it suitable for more complex multimodal tasks. U-Xray, though smaller in size, remains valuable for tasks such as detecting lung diseases in X-ray images. Its specialized focus on specific modalities and conditions makes it ideal for benchmarking models on chest X-ray classification tasks.

SLAKE, ROCO, MedlCaT, and PMC-15M offer multimodal data, which combines medical images with textual annotations or descriptions. This allows models to tackle tasks that require understanding both modalities simultaneously, such as VQA, report generation, or image-text matching. PMC-15M, with its combination of images and full-text articles, supports complex natural language processing (NLP) tasks in medical research, while ROCO enables report generation tasks that can directly influence clinical workflow efficiency.

Multi-omics datasets

Multi-omics datasets, which integrate various types of biological data such as genomics, proteomics, and metabolomics, provide a rich and multidimensional foundation for developing LLMs.

Cancer multi-omics datasets [90] include ten datasets that contain multi-omics data from different cancer types, with each dataset correspon[din](#page-20-18)g to a specific cancer. These datasets typically include three key omics layers: gene expression, DNA methylation, and miRNA expression. The number of patients ranges from 170 for acute myeloid leukemia (AML) to 621 for breast invasive carcinoma (BIC), offering a diverse range of data for training LLMs in cancer research.

Each dataset offers unique benefits depending on the intended use case and the specific tasks in medical image processing, report generation, or VQA. CheXpert, PadChest, and MIMIC-CXR stand out in disease detection and classification from chest X-rays due to their size and rich annotations. VQA-RAD, SLAKE, and PathVQA are essential for advancing VQA tasks in radiology and pathology. Datasets such as PMC-15M and MedlCaT provide invaluable resources for multimodal tasks that combine medical images with textual data, enabling more sophisticated models for clinical decision support and medical research. What is more, the integration of multi-omics data serves as both a guide for biomedical researchers in identifying suitable deep learning-based fusion methods and an indication of promising directions for improving multi-omics data fusion techniques.

In summary, the diversity and richness of data sources lay a solid foundation for the development of medical LLMs, facilitating significant advancements in understanding medical language, processing medical data, and providing medical decision support. Each dataset's unique structure and characteristics cater to specific aspects of healthcare AI technology, ranging from basic QA systems to complex diagnostic tools and patient management systems. By leveraging these resources, LLMs demonstrate substantial potential to enhance patient care quality and accelerate medical research.

3. DATASET APPLICATION

Datasets are fundamental to the deployment of LLMs in medicine. They are mainly employed in three aspects: pre-training, fine-tuning, and evaluation. The application of datasets has been presented in Figure 4.

Figure 4. The application of datasets in developing medical Q&A LLMs.

3.1. Pre-training

In the pre-training phase, a large corpus of text data including both structured and unstructured text is used to train the LLMs. The corpus typically consists of various sources such as EHRs, clinical notes, and medical literature^[91]. Some of the most commonly used medical datasets for pre-training medical LLMs include PubMed^[34], [M](#page-20-19)IMIC-III clinical notes^[31], and PMC literature^[35]. High-quality datasets form the backbone of LLM p[re](#page-18-17)-training, with MIMIC-IIIa[nd](#page-18-14) PubMed serving as p[ivo](#page-18-18)tal resources. MIMIC-III provides valuable and extensive data for research, which has facilitated the development of several LLMs, such as ClinicalBert^[14], , GatorTron^[92], and BlueBERT^[93]. PubMed, known for its extensive biomedical literature, has been the fo[un](#page-18-0)dationfor t[rai](#page-20-20)ning models suc[h a](#page-20-21)s PubMedBERT $^{[16]}$, BioBERT $^{[94]}$ and GatorTron. These models benefit from PubMed's extensive collection of research articles [an](#page-18-2)d studies, [wh](#page-20-22)ich enables them to capture a wide range of biomedical knowledge essential for engaging with scientific texts.

A notable example is GatorTronGPT, which is trained on a massive corpus of 82 billion words of de-identified clinical text^[13] and 195 billion words of general English text from the Pile dataset^[40]. GatorTronGPT is trained fro[m sc](#page-17-12)ratch using the GPT-3^[95] architecture with five billion and 20 billion [par](#page-18-23)ameters. By leveraging such diverse datasets, models such [as](#page-20-23) GatorTronGPT gain the ability to understand both general English and domain-specific medical language, enhancing their utility in clinical applications. These datasets can be employed in combination to enrich the pre-training phase. BlueBERT combines both PubMed and MIMIC-III for pre-training; BioBERT is pre-trained on both PubMed and PMC; MEDITRON [96] is pre-trained on the GAP-REPLAY data mixture that contains papers from PubMed and PMC. Through [pre](#page-20-24)-training on these medical corpora, LLMs are equipped with rich medical knowledge and to tackle various healthcare-related tasks.

3.2. Fine-tuning

After pre-training, LLMs require further fine-tuning to enhance their abilities. This phase leverages domainspecific datasets, such as dialogue data, QA pairs, and instructional texts, enabling the models to develop a nuanced understanding of both natural language and medical terminology. It tailors the model for specific medical tasks, allowing it to interpret and generate medical texts effectively.

For LLMs in medicine, supervised fine-tuning (SFT) and instruction fine-tuning (IFT) are two commonly used methods. These methods involve training the model on specific datasets to adapt it to the desired domain or task. SFT utilizes high-quality medical corpus such as physician-patient conversations, medical QA, and knowledge graphs. IFT constructs instruction-based training datasets, typically comprising instructioninput-output triples, to enhance the ability of instruction following. An example of IFT is the training process of Med-PaLM 2^[18], where the base LLM is PaLM 2^[97]. Med-PaLM 2 is fine-tuned using the datasets including MedQA, MedM[CQ](#page-18-4)A, HealthSearchQA, LiveQA,a[nd](#page-20-25) MedicationQA. The fine-tuning followed the protocol used by Chung *et al.*, resulting in improved performance on medical Q&A benchmarks^[98]. .

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In addition, building upon the CMeKG $^{[47]}$, BenTsao $^{[99]}$ utilizes diverse instructional data for its instruction tuning process. MedAlpaca^[19], built u[po](#page-19-21)n the LLa[MA](#page-20-26)^[100], is fine-tuned using over 160,000 medical QA pairs sourced from [M](#page-18-26)edical Meadow^[19]. Similarly, Cha[tDo](#page-21-0)ctor^[64] is obtained by fine-tuning the LLaMA model^[100] on HealthCareMagic-100k^{[\[64](#page-18-26)]} following the Stanford [Alp](#page-19-15)aca^[69] training method. It is first finetuned [with](#page-21-0) Alpaca's data and further r[efin](#page-19-15)ed on HealthCareMagic-100k t[o im](#page-19-19)prove its medical knowledge accuracy. Other models such as Qilin-Med^[101] and Zhongjing^[68] are obtained by incorporating the knowledge graph from ChiMed^[101] and CMtMedQ[A](#page-21-1)^[68] to perform fi[ne-](#page-19-18)tuning on the Baichuan^[102] and LLaMA^[100] respectively to enhan[ce t](#page-21-1)heir medical reaso[nin](#page-19-18)g capabilities.

3.3. Evaluation

Evaluation is critical to the success of LLMs, which helps us better understand their strengths and weaknesses. Evaluating LLMs for medical applications typically involves using Q&A benchmarks where the models answer questions from a dataset, and their responses are scored based on predefined metrics, such as accuracy, precision, and recall, which are highly relevant to clinical applications. For text generation tasks, BLEU and ROUGE metrics are commonly applied to assess how closely the generated output matches the ground truth. For example, MultiMedQA $^{[59]}$ is designed to evaluate the capabilities of LLMs in answering medical questions across various formats, incl[ud](#page-19-11)ing multiple-choice and long-form answers. This benchmark compiles datasets from diverse sources, such as professional medical exams, medical research, and consumer health inquiries, providing a more thorough assessment of LLM performance beyond traditional multiple-choice accuracy or standard natural language generation metrics such as BLEU. The evaluation process tests LLMs not only on their factual accuracy but also on their medical reasoning capabilities and ability to handle both open-domain and closed-domain questions.

In addition, metrics such as Recall@K for retrieval tasks and AUC for classification tasks are frequently employed. For instance, PMC-CLIP^[82], pretrained on the PMC-OA dataset, is evaluated using Recall@K for image-text retrieval on ROCO^[80] [an](#page-20-12)d AUC and accuracy metrics for image classification, where it showed strong capabilities in these tas[ks.](#page-20-10) Clinical prediction with LLMs (CPLLM)^[103] is evaluated on four prediction tasks: patient hospital readmission prediction, along with three specific [diag](#page-21-2)nosis predictions for Chronic Kidney Disease, Acute and Unspecified Renal Failure, and Adult Respiratory Failure. The first two diagnoses are derived from the MIMIC-IV dataset, while the last diagnosis is derived from the eICU-CRD dataset $[104]$. . MIMIC-IV provides the start time for admission and discharge times, and eICU-CRD associates each dia[gno](#page-21-3)sis with a timestamp, making them similarly applicable for patient readmission prediction tasks.

Models such as Med-PaLM $^{[59]}$ and Med-PaLM 2 $^{[18]}$ are tested on MultiMedQA. USMLE $^{[105]}$, PubMedQA $^{[53]}$, and MedMCQ[A](#page-19-8) $^{[106]}$ are thr[ee](#page-19-11) popular datasets to [ev](#page-18-4)aluate their effectiveness. Codex-Med $^{[107]}$ $^{[107]}$ $^{[107]}$, PMC-LLaMA $^{[108]}$ Galactica^[109], G[ator](#page-21-5)TronGPT^[12] and Med-PaLM 2^[18] are evaluated on these three data[sets](#page-21-6). USMLE is als[o](#page-21-7) used to e[valu](#page-21-8)ate the performa[nce](#page-17-11) of MedAlpaca in a [ze](#page-18-4)ro-shot setting. Additionally, iCliniq $^{[64]}$ is used to test ChatDoctor's performance for a quantit[ati](#page-19-15)ve evaluation. HuatuoGPT^[56] undergoes evaluation using three Chinese QA datasets: cMedQA2^[49], webMedQA^[50], and Huatuo26M^{[\[51](#page-19-9)]} with GPT-4 and doctors comparing the responses from HuatuoG[PT](#page-19-4) and making e[va](#page-19-5)luations. The cMe[dQ](#page-19-6)A2 dataset is also used to evaluate ClinicalGPT^[110], which is conducted using automated evaluation metrics, with GPT-4 serving as the refer-ence model. [Oth](#page-21-9)er models such as LLaVA-Med $^{[111]}$ and Med-Flamingo $^{[112]}$ are evaluated on VQA datasets. VQA-RAD^[76], SLAKE^[77] and Path-VQA^[78] ar[e us](#page-21-10)ed to evaluate LLaV[A-M](#page-21-11)ed. VQA-RAD, Path-VQA and Visual USM[LE](#page-20-6) are used [to](#page-20-7) evaluate Med-Fl[am](#page-20-8)ingo to measure their performance in medical-related tasks. Recently, MMedBench, covering 21 medical fields, has been designed to assess the accuracy of multiple-choice QA tasks and the ability to generate rationales across multiple languages^[113]. The evaluation of eleven LLMs shows that MMed-Llama 3, built on the foundation of LLaMA 3, demons[trat](#page-21-12)es strong performance compared with models such as LLaMA, ChatDoctor, and MedAlpaca. The linguistic diversity of datasets used in evaluations provides deeper insights into the models' capabilities, not only across various medical domains but also in their adaptability to multilingual healthcare contexts.

4. CHALLENGES

Several important factors such as data availability, data curation, quality, and deserve careful consideration.

4.1. Availability and open access

Accessibility of datasets is a key factor in determining their usability for external researchers. They can generally be categorized into three groups based on their accessibility $^{[114,115]}$. Open access datasets are easily available to external researchers, often requiring only simple registrat[ion](#page-21-13) [or](#page-21-14) an email request. These datasets provide valuable resources without the need for complex approvals. In contrast, regulated access datasets require formal agreements, such as institutional approvals, ethical clearance, or payments, due to the sensitive nature of the data. While these safeguards ensure compliance with regulations, they also create barriers that can slow down access. Lastly, inaccessible datasets are publicly listed as available but are often difficult to obtain due to issues such as non-responsiveness or outdated access links. Many medical datasets fall into the regulated access category, requiring formal approvals to protect sensitive information, while some may be inaccessible despite being listed as available^[30]. These barriers highlight the challenges researchers face when trying to access medical data for LLM de[ve](#page-18-27)lopment.

4.2. Data curation and quality

Data curation tasks, including discovering, extracting, transforming, cleaning, and integrating data, remain critical yet resource-intensive efforts for organizations^[116]. Data scientists often spend over 80% of their time on these tasks^[117], as generic tools are rarely sufficien[t fo](#page-21-15)r the diverse and domain-specific requirements encountered in p[rac](#page-21-16)tice. The quality of available datasets also varies significantly, as many datasets are poorly curated, with incomplete or unlabeled data, making it difficult to train effective models. The process of human labeling is resource-intensive and requires domain expertise, while unsupervised learning methods face challenges due to the need for high accuracy $^{\rm [118]}$. Improved data curation practices, such as better structuring and labeling, are essential to enhance the qu[ality](#page-21-17) and usability of medical datasets for LLM training.

4.3. Data scarcity and fragmentation

Data scarcity and fragmentation remain significant obstacles in medical research, as medical data is frequently siloed across different institutions and stored in various formats. Multimodal biomedical data fusion has become essential in modern healthcare research, integrating diverse data sources such as medical images, biomarkers, and physiological signals to provide a more comprehensive understanding of biological systems $[119]$. . This approach enhances decision-making in key areas such as disease diagnosis, treatment planning, and [pa](#page-21-18)tient monitoring by leveraging the strengths of each data modality.

4.4. Ethical considerations

Ethical concerns about using LLMs in the medical domain are significant, particularly around patient privacy, safety, and sensitive data use. A key issue is the collection and potential exposure of protected health information input into LLM application programming interfaces (APIs), which could be accessed by unauthorized parties. The lack of transparency from companies on how they store and use this data raises ethical questions about submitting sensitive information. Thus, strict controls for de-identification and informed consent must be implemented when handling protected health information in LLM APIs. Another major issue is the leakage of personally identifiable information (PII)^[120,121]. LLMs trained on large datasets may inadvertently expose sensitive PII, such as email addresses or othe[r c](#page-21-19)[onfi](#page-21-20)dential details, through vulnerabilities such as prompt injection attacks. So, it is essential to implement rigorous safeguards and data protection measures when deploying LLMs in clinical settings.

Moreover, data that disproportionately represents certain populations may result in biased models that perform poorly for underrepresented groups, exacerbating health disparities. Researchers must consider these ethical factors and, where possible, implement bias mitigation strategies and uphold the highest standards of data privacy and protection. There are also concerns about bias in medical datasets, which can result in LLMs that benefit certain populations while marginalizing others. In particular, the datasets used to train these models typically come from well-funded institutions in high-income, English-speaking countries. This leads to a significant under-representation of perspectives from other regions of the world, causing LLMs to adopt views that are biased toward the healthcare processes of high-income countries^[122]. As a result, additional training could be integrated into the model development process to ensure that LL[Ms s](#page-21-21)erve diverse patient populations equitably, and global datasets should be incorporated to reduce geographical and socioeconomic biases.

4.5. Limitations and future directions

Current medical datasets are relatively smaller than those used for general LLMs, covering a limited portion of the medical knowledge domain^[59]. LLMs trained on these datasets may perform well on benchmarks, but they often struggle with real-world [ta](#page-19-11)sks such as differential diagnosis and personalized treatment planning $[18]$. . While generating high-quality synthetic datasets for training could help broaden the model's knowledge, it ri[sks](#page-18-4) causing LLMs to experience forgetting^[123]. Further research is necessary to validate the effectiveness of synthetic data for medical LLMs and to de[velo](#page-21-22)p techniques that mitigate such risks.

For evaluation, existing medical Q&A benchmarks often rely on metrics such as classification accuracy or natural language generation scores (e.g., $\text{BLEU}^{\{124\}}$), which may not cover the full breadth of clinical scenarios and decision-making processes that occur in re[al-w](#page-21-23)orld medical practice. Multiple-choice tasks, often featured in these benchmarks, are much easier than real-world medical decisions that require synthesizing patient information and formulating individualized treatment plans, as they are grounded by experts. Although the MultiMedQA benchmark addresses some of these gaps by offering a diverse set of questions from medical exams, research, and consumer health queries, it is not exhaustive enough. It currently lacks coverage across all medical and scientific domains and is limited to English-language datasets, which restricts its applicability in global healthcare settings. To effectively evaluate LLMs, it is crucial to expand these datasets to include multilingual evaluations and more comprehensive clinical tasks, such as open-ended assessments that mirror actual clinical workflows. This expansion will enable models to be tested on their ability to reason through medical complexities, and provide accurate responses that are essential in real-world clinical environments.

5. CONCLUSIONS

This survey presents a comprehensive overview of the datasets in medicine and their pivotal role in developing LLMs. Datasets serve not only as a foundation for training LLMs but also as benchmarks for evaluating their performance. Each stage of datasets' application is critical for ensuring that the models are practically effective in real-world medical settings. Looking forward, the continued expansion and refinement of these datasets will be essential. Future research should focus on enhancing dataset transparency and quality, addressing privacy concerns, and integrating multimodal data to enrich model training. The development of medical datasets is a dynamic and evolving field that holds the key to unlocking the full potential of LLMs in medicine.

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Made substantial contributions to the research, reviewed and summarized the literature, wrote and edited the original draft: Zhang D, Xue X, Hu M, Ying X Investigation, data analysis and interpretation: Zhang D, Xue X, Gao P Supervision: Hu M, Jin Z, Wu Y, Ying X

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