From Generalist to Specialist: A Survey of Large Language Models for Chemistry

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Abstract

Large Language Models (LLMs) have significantly transformed our daily life and established a new paradigm in natural language processing (NLP). However, the predominant pretraining of LLMs on extensive web-based texts remains insufficient for advanced scientific discovery, particularly in chemistry. The scarcity of specialized chemistry data, coupled with the complexity of multi-modal data such as 2D graph, 3D structure and spectrum, present distinct challenges. Although several studies have reviewed Pretrained Language Models (PLMs) in chemistry, there is a conspicuous absence of a systematic survey specifically focused on chemistry-oriented L paper, we outline methodologie rating domain-specific chemist and multi-modal information in also conceptualize chemistry Ll using chemistry tools and investigate their potential to accelerate scientific research. Additionally, we conclude the existing benchmarks to evaluate chemistry ability of LLMs. Finally, we critically examine the current challenges and identify promising directions for future research. Through this comprehensive survey, we aim to assist researchers in staying at the forefront of developments in chemistry LLMs and to inspire innovative applications in the field. $¹$ $¹$ $¹$ </sup>

1 Introduction

Recent years have witnessed remarkable advancements in daily life driven by LLMs. Competitive models like GPT-4 [\(Achiam et al.,](#page-8-0) [2023\)](#page-8-0) and Claude [\(Anthropic,](#page-8-1) [2024\)](#page-8-1) have demonstrated exceptional abilities across diverse tasks, often matching or surpassing human-level performance, marking significant progress toward Artificial General Intelligence (AGI, [Bubeck et al.](#page-9-0) [\(2023\)](#page-9-0)). In sci-

GPT40: The molecular formula for '[3-(carboxyamino)-2oxopropyl]azanium' is C4H8N2O3. x - – – – – – – –
products of the i - – – – – – – – – – –
action described in th - - - - - - - - - - - -
Question: What are the

Question: What is the molecule formula of the molecule with the the IUPAC name '[3-(carboxyamino)-2-oxopropyl]azanium'?

Question: Please synthesize aspirin.

GPT40: Synthesizing aspirin (acetylsalicylic acid) involves a common laboratory procedure using salicylic acid and acetic anhydride...

Figure 1: Three common errors in general LLMs arising from the key challenges.

showing promising results [\(Fang et al.,](#page-10-0) [2023\)](#page-10-0). Among these, chemistry LLMs, further tailored for chemical applications via additional training or advanced prompt engineering, have garnered significant attention. Before the advent of LLMs, there are lots of notable efforts towards chemistry, such as MolT5 [\(Edwards et al.,](#page-10-1) [2022\)](#page-10-1), Text2Mol [\(Edwards et al.,](#page-10-2) [2021\)](#page-10-2), MoMu [\(Su et al.,](#page-13-0) [2022\)](#page-13-0), Text+Chem T5 [\(Christofidellis et al.,](#page-9-1) [2023\)](#page-9-1). However, these models are built on PLMs like BERT [\(Devlin,](#page-9-2) [2018\)](#page-9-2) and T5 [\(Raffel et al.,](#page-12-0) [2020\)](#page-12-0), requiring fine-tuning for specific tasks and lacking emergent abilities [\(Wei et al.,](#page-14-0) [2022a\)](#page-14-0), such as Chain-of-Thought (CoT, [Wei et al.](#page-14-1) [\(2022b\)](#page-14-1)) reasoning and tool-using capabilities [\(Qin et al.,](#page-12-1) [2023\)](#page-12-1). Existing reviews [\(Xiao et al.,](#page-14-2) [2024;](#page-14-2) [Liao et al.,](#page-11-0) [2024;](#page-11-0) [Pei](#page-12-2) [et al.,](#page-12-2) [2024a\)](#page-12-2) have already discussed those PLMs in

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¹We maintain an up-to-date Github repository at: [https:](https://github.com/OpenDFM/LLM4Chemistry) [//github.com/OpenDFM/LLM4Chemistry](https://github.com/OpenDFM/LLM4Chemistry).

chemistry, such as [Liao et al.](#page-11-0) [\(2024\)](#page-11-0), which emphasize molecule encoding methods and pretraining objectives. More related works are discussed in the Appendix [A.](#page-15-0) In contrast, our survey focuses on generative models with Transformer decoder ar-

chitectures [\(Vaswani et al.,](#page-13-1) [2017\)](#page-13-1), addressing key challenges of general LLMs and reviewing existing approaches to adapt them for chemistry-specific tasks and applications.

General LLMs, such as the GPT [\(Ouyang et al.,](#page-12-3) [2022;](#page-12-3) [Achiam et al.,](#page-8-0) [2023\)](#page-8-0) and LLaMA series [\(Tou](#page-13-2)[vron et al.,](#page-13-2) [2023a](#page-13-2)[,b\)](#page-13-3), have demonstrated impressive performance. However, they tend to underperform on chemistry-related tasks as shown in Figure [1.](#page-0-1) We identify three key challenges contributing to these limitations.

Challenge 1: domain knowledge is not enough. Most LLMs are pre-trained with the objective of predicting the next token based on web data sourced from the internet [\(Ouyang et al.,](#page-12-3) [2022\)](#page-12-3), as demonstrated by open-source models like LLaMa series [\(Touvron et al.,](#page-13-2) [2023a](#page-13-2)[,b\)](#page-13-3). While some chemistry-related data exist within these datasets, the quantity is minimal, and there is a lack of data specifically tailored for chemistry. This deficiency extends to other crucial steps in the development of LLMs, such as supervised fine-tuning (SFT) and reinforcement learning from human feedback (RLHF, [Christiano et al.](#page-9-3) [\(2017\)](#page-9-3); [Stiennon et al.](#page-13-4) [\(2020\)](#page-13-4)).

Challenge 2: multi-modal data is not perceived. Chemistry encompasses various modalities, including 1D sequences [\(Krenn et al.,](#page-11-1) [2020\)](#page-11-1), 2D molecular graphs [\(Duvenaud et al.,](#page-10-3) [2015;](#page-10-3) [Xu et al.,](#page-14-3) [2018;](#page-14-3) [Liu et al.,](#page-11-2) [2019\)](#page-11-2), and 3D structures [\(Schütt](#page-13-5) [et al.,](#page-13-5) [2018;](#page-13-5) [Satorras et al.,](#page-12-4) [2021;](#page-12-4) [Atz et al.,](#page-8-2) [2021\)](#page-8-2). Additionally, there are numerous chemical spectra, such as Nuclear Magnetic Resonance (NMR, [Simpson et al.](#page-13-6) [\(2012\)](#page-13-6)), Liquid Chromatography-Tandem Mass Spectrometry (LC-MS, [Seger](#page-13-7) [\(2012\)](#page-13-7); [Dührkop et al.](#page-10-4) [\(2015\)](#page-10-4); [Litsa et al.](#page-11-3) [\(2023\)](#page-11-3)), and Infrared Spectroscopy (IR, [Alberts et al.](#page-8-3) [\(2023\)](#page-8-3)). These spectra contain substantial information that LLMs currently fail to fully exploit.

Challenge 3: chemistry tools are not utilized. Due to the core design of LLMs, they often struggle with retaining up-to-date knowledge and performing specific chemistry operations [\(Castro Nasci](#page-9-4)[mento and Pimentel,](#page-9-4) [2023;](#page-9-4) [Schick et al.,](#page-12-5) [2024\)](#page-12-5). On the other hand, there are numerous powerful chemistry tools, such as the structure knowledge retrieval (PubChem [\(Kim et al.,](#page-11-4) [2019\)](#page-11-4), OPTIMADE [\(Evans et al.,](#page-10-5) [2024\)](#page-10-5)), and various expert-designed artificial intelligence systems tailored to address specific problems like reaction prediction [\(Pesci](#page-12-6)[ullesi et al.,](#page-12-6) [2020\)](#page-12-6), retrosynthesis planning [\(Segler](#page-13-8) [et al.,](#page-13-8) [2018\)](#page-13-8) and so on. The absence of integration with these chemistry tools significantly hinders the

performance of LLMs in the field of chemistry.

In this survey, we critically review current efforts addressing the three key challenges outlined in Figure [2.](#page-2-0) Additionally, we review the existing benchmarks used to evaluate the performance of chemistry LLMs and offer suggestions for future research directions. To the best of our knowledge, this is the first systematic survey reviewing existing approaches for transferring general LLMs to chemistry-specific LLMs in decoder architecture.

2 Domain Knowledge

Pre-training, SFT and RLHF have been the de facto way to enhance domain knowledge of LLMs. We will detail those methods in the following sections.

2.1 Pre-training

The natural of LLMs lay in language modeling, given a set of examples $(x_1, x_2, ..., x_n)$ each composed of variable length sequences of symbols (s_1, s_2, \ldots, s_m) , language model is framed as a unsupervised distribution estimation and the joint probabilities over symbols can be formulated [\(Radford](#page-12-7) [et al.,](#page-12-7) [2019\)](#page-12-7):

$$
p(x) = \prod_{i=1}^{n} p(s_n | s_1, ..., s_{n-1}),
$$
 (1)

self-attention architectures like the Transformer can be applied to compute these conditional probabilities. Training on a large-scale corpus in this manner enables LLMs to capture rich language representations, refering to pre-training.

Continue pre-training is prefered given the existence of advanced foundation models like LLaMA [\(Touvron et al.,](#page-13-2) [2023a,](#page-13-2)[b\)](#page-13-3) and Galactica [\(Taylor et al.,](#page-13-9) [2022\)](#page-13-9), which already contain some basic chemistry knowledge. In contrast, pretraining from scratch is cost-prohibitive. Chemistry knowledge is typically encoded in specific languages, such as the Simplified Molecular-Input Line-Entry System (SMILES) [\(Weininger,](#page-14-4) [1988\)](#page-14-4), which represents 3D structures as flattened sequences while preserving most structural information. Other representations include molecular formulas, SELFIES [\(Krenn et al.,](#page-11-1) [2020\)](#page-11-1), International Union of Pure and Applied Chemistry (IUPAC) names, and the Chemical Identifier (InChI) [\(Heller](#page-10-6) [et al.,](#page-10-6) [2013\)](#page-10-6). To enhance foundation models with domain-specific chemistry knowledge, it is necessary to gather pre-training corpora in these chemical languages and apply continued pre-training.

Figure 2: Taxonomy of currect approachs for transfering general LLMs to specialized chemistry LLMs.

The volume of pre-training data required for chemistry LLMs is immense, making it difficult to obtain and, in some cases, restricted by copyright. To the best of our knowledge, ChemDFM [\(Zhao](#page-15-6) [et al.,](#page-15-6) [2024b\)](#page-15-6) is the sole chemistry LLM specifically pre-trained on a chemical corpus. ChemDFM's training data comprises 34 billion tokens from 3.9 million chemical papers collected online before January 2022 and 49 million tokens from 1.4 thou-sand chemistry books sourced from LibreTexts^{[2](#page-2-2)} and Gold Books^{[3](#page-2-3)}. Through pre-training on this chemical text, ChemDFM can acquire a solid understanding of chemistry and emerge as the top open-source model [\(Feng et al.,](#page-10-12) [2024\)](#page-10-12). Another T5-based chemistry LM, Nach0 [\(Livne et al.,](#page-12-12) [2024\)](#page-12-12), collects 13 million abstracts from PubMed, 119K patent descriptions from the USPTO, and incorporates approximately 100 million documents from ZINC.

2.2 SFT

Pre-training on large corpus with next token prediction does not align well with users' objective,

as users expect models to "follow their instructions helpfully and safely" [\(Zhang et al.,](#page-14-8) [2023b\)](#page-14-8). SFT effectively aligns LLMs with user expectations by training them on datasets consisting of (INSTRUC-TION, OUTPUT) pairs, where INSTRUCTION refers to specific chemistry tasks and OUTPUT represents the desired responses. Given the variety of chemistry tasks in the SFT dataset, it can be further categorized as follows:

1. *Multi-task SFT*: We categorize commonly used chemistry tasks into four types: SMILES understanding, reaction understanding, notation alignment and chemistry-related QA, as detailed in Appendix [B.](#page-15-7) The most significant distinction among different SFT models [\(Yu et al.,](#page-14-6) [2024;](#page-14-6) [Fang et al.,](#page-10-0) [2023;](#page-10-0) [Zhao](#page-15-6) [et al.,](#page-15-6) [2024b;](#page-15-6) [Zhang et al.,](#page-14-7) [2024a\)](#page-14-7) lie in their data sources and the volume of data used, and the detailed data distribution is shown in Appendix [B.](#page-15-7) The total dataset volume ranges from 1.5M to 3M, although [Zhang et al.](#page-14-7) [\(2024a\)](#page-14-7) does not provide exact figures, it is likely of a similar magnitude. The distribution of tasks within the SFT dataset determines the model's chemistry capabilities, as identified

²https://libretexts.org/

³ https://goldbook.iupac.org/

by [\(Feng et al.,](#page-10-12) [2024\)](#page-10-12). [Zhao et al.](#page-15-6) [\(2024b\)](#page-15-6); [Zhang et al.](#page-14-7) [\(2024a\)](#page-14-7) focus more on chemistryrelated QA, gathering major data from sources such as chemistry exams and existing datasets, which enhances the model's ability to answer user questions more naturally.

2. *Task-specific SFT*: Task-specific finetuning of LLMs has demonstrated effective prediction performances, often surpassing traditional machine learning models, particularly in low-data scenarios[\(Jablonka et al.,](#page-10-11) [2024\)](#page-10-11). [Jablonka et al.](#page-10-11) [\(2024\)](#page-10-11) finetune GPT-3 for classification, regression, and inverse design tasks, achieving competitive results in three case studies (polymers, metal-organic frameworks, and photoswitches). More recently, [Liu et al.](#page-12-11) [\(2024d\)](#page-12-11) propose hybrid instruction tuning on more than 1000 property tasks with LLaMA2- 7b-chat [\(Touvron et al.,](#page-13-3) [2023b\)](#page-13-3), reporting up to a 16.6% average improvement over leading LLM baselines across all classification tasks. Additionally, [Chen et al.](#page-9-9) [\(2023\)](#page-9-9) also fine-tune LLaMA2-7B-chat with 13,878 pieces of structured material knowledge data to predict inorganic material synthesis pathways.

In addition to these chemistry tasks, chemical text mining is also a crucial foundation in chemical research, as much scientific knowledge is dispersed across the text, tables, and figures in millions of academic papers [\(Dagdelen et al.,](#page-9-10) [2024\)](#page-9-10). [Dagdelen](#page-9-10) [et al.](#page-9-10) [\(2024\)](#page-9-10) focus on joint named entity recognition and relation extraction, enabling the generation of simple English sentences or more structured formats, such as JSON object, from individual sentences or entire paragraphs. [Zhang et al.](#page-14-9) [\(2024c\)](#page-14-9) extend these efforts to more chemical text mining tasks, achieving the best performance across all tasks, with exact accuracy ranging from 69% to 95% using minimal annotated data.

2.3 RLHF

While pre-training and SFT provide chemistry LLMs with domain-specific knowledge and enable them to perform specific tasks, these models are still prone to hallucination. RLHF is the most effective method to alleviate hallucinations and build a truthful, helpful and harmless LLM [\(Ouyang et al.,](#page-12-3) [2022\)](#page-12-3). There are many detail algorithms to utilize human feedback, such as PPO [\(Schulman et al.,](#page-13-11) [2017\)](#page-13-11), DPO [\(Rafailov et al.,](#page-12-13) [2024\)](#page-12-13). Beyond human feedback, other methods for collecting preference

feedback include AI feedback [\(Lee et al.,](#page-11-11) [2023;](#page-11-11) [Bai](#page-8-5) [et al.,](#page-8-5) [2022\)](#page-8-5) and environment feedback [\(Cao et al.,](#page-9-11) [2024;](#page-9-11) [Dong et al.,](#page-9-12) [2024\)](#page-9-12).

Existing research on human alignment for chemistry LLMs primarily focuses on molecular generation tasks. [Fang et al.](#page-10-9) [\(2024b\)](#page-10-9) first pre-trains LLM on SELFIES [\(Krenn et al.,](#page-11-1) [2020\)](#page-11-1), enabling the generation of syntactically correct molecules; however, the model also produces undesirable molecules, referred as molecular hallucinations. To mitigate these hallucinations and better align with actual chemical contexts, they apply a rank loss [\(Liu et al.,](#page-11-12) [2022\)](#page-11-12) by assigning higher probabilities to molecule candidates with desired properties. [Zholus et al.](#page-15-5) [\(2024\)](#page-15-5) finetunes a GPT-based model for 3D molecular design, and utlizes external feedback from docking software using REINFORCE algorithm [\(Williams,](#page-14-10) [1992\)](#page-14-10). [Hu et al.](#page-10-10) [\(2024\)](#page-10-10) further investigates multiple GPT agents to generate desirable molecules in diverse directions, with the reward function estimated by docking software. The objective is to maximize the average reward while simultaneously improving molecular diversity.

AI and environment feedback are the most commonly used rewards for chemistry LLMs, as the more valuable human feedback is often unavailable due to the need for strong domain knowledge and the lack of effective tools to collect chemistryspecific feedback. [Hu et al.](#page-10-10) [\(2024\)](#page-10-10) design a Pythonbased open-source graphical user interface (GUI) to explore and evaluate molecules, and capture chemist's implicit knowledge and preferences more efficiently. This tool provides a promising approach for collecting chemistry-specific feedback to better align chemistry LLMs with human expertise.

3 Multi-Modal Data

Domain knowledge training is a standard approach for developing domain-specific LLMs, as demonstrated in fields like geoscience [\(Deng et al.,](#page-9-13) [2024\)](#page-9-13), law [\(Zhou et al.,](#page-15-8) [2024\)](#page-15-8), and medicine [\(Zhang et al.,](#page-14-11) [2023a\)](#page-14-11). However, chemical data is highly fragmented across multiple modalities [\(Mirza et al.,](#page-12-14) [2024\)](#page-12-14), such as 2D graphs, 3D structures, and spectra, as shown in Figure [3,](#page-5-3) which cannot be directly processed by vanilla LLMs. Inspired by recent advances in multi-modal and vision LLMs [\(Liu et al.,](#page-11-13) [2024a;](#page-11-13) [Li et al.,](#page-11-14) [2024a;](#page-11-14) [Huang et al.,](#page-10-13) [2024a\)](#page-10-13), numerous studies have focused on integrating chemical modalities with vanilla LLMs through the design of alignment components. We provide a compre-

hensive review of these works based on the modalities they support: *1D Sequences*, *2D Graphs*, *3D Structures*, and *Other Modalities*.

3.1 1D Sequences

SMILES [\(Weininger,](#page-14-4) [1988\)](#page-14-4) is a widely used molecular representation, but it is generally processed as text using a byte-pair encoding tokenizer [\(Sen](#page-13-12)[nrich,](#page-13-12) [2015\)](#page-13-12), which fails to capture its inherent chemical information. To address this limitation, MolX [\(Le et al.,](#page-11-10) [2024\)](#page-11-10) treats SMILES as a distinct modality and proposes a pre-trained BERTlike [\(Devlin,](#page-9-2) [2018\)](#page-9-2) SMILES encoder to extract features, which are then aligned with other modalities through projection. MoleculeGPT [\(Zhang et al.,](#page-15-3) [2023c\)](#page-15-3) also adapt ChemBerta [\(Ahmad et al.,](#page-8-6) [2022\)](#page-8-6) for SMILES encoding. However, SMILES lacks robustness and does not fully capture spatial information, leading to the development of other 1D sequence representations, such as SELFIES [\(Krenn](#page-11-1) [et al.,](#page-11-1) [2020\)](#page-11-1), IUPAC names, molecular fingerprints [\(Morgan,](#page-12-15) [1965\)](#page-12-15), and InChI [\(Heller et al.,](#page-10-6) [2013\)](#page-10-6). These 1D sequences are generally processed similarly to text but can be further refined using specialized encoders, such as SELFormer [\(Yüksel et al.,](#page-14-12) [2023\)](#page-14-12) for SELFIES and variational autoencoders (VAE, [Kingma](#page-11-15) [\(2013\)](#page-11-15)) for molecular fingerprints.

3.2 2D Graphs

Compared to 1D sequences, 2D graphs offer a more intuitive representation of molecular structures and chemical bonds. To process 2D graphs, an encoder is required to convert them into vector representations, followed by a projector to align these vectors with LLMs. Graph neural networks (GNNs, [Hu et al.](#page-10-14) [\(2019\)](#page-10-14); [Xiao et al.](#page-14-13) [\(2022\)](#page-14-13)) are widely used as 2D graph encoders and have been adopted by most multimodal chemistry LLMs [\(Liu](#page-12-10) [et al.,](#page-12-10) [2024e;](#page-12-10) [Li et al.,](#page-11-9) [2024b;](#page-11-9) [Zhang et al.,](#page-15-3) [2023c;](#page-15-3) [Le et al.,](#page-11-10) [2024;](#page-11-10) [Zhang et al.,](#page-15-4) [2024e\)](#page-15-4). For instance, MolTC [\(Fang et al.,](#page-10-8) [2024a\)](#page-10-8) train two GNNbased encoders and representation projectors by freezing the LLM and backpropagating the generation loss. InstructMol[\(Cao et al.,](#page-9-7) [2023\)](#page-9-7) employs MoleculeSTM's graph encoder [\(Liu et al.,](#page-11-16) [2023a\)](#page-11-16), which is trained through molecular-textual contrastive learning. MolCA [\(Liu et al.,](#page-12-9) [2023b\)](#page-12-9) utilze a more expressive GNN model - Graph isomorphism network (GIN, [Hu et al.](#page-10-14) [\(2019\)](#page-10-14)), which pre-trained on 2 million molecules from the ZINC15 [\(Ster](#page-13-13)[ling and Irwin,](#page-13-13) [2015\)](#page-13-13). HIGHT[\(Chen et al.,](#page-9-8) [2024b\)](#page-9-8) further introduce a hierarchical graph tokenizer

which em Vector Quantized-Variational AutoEncoder (VQVAE, [\(Zang et al.,](#page-14-14) [2023\)](#page-14-14)) to extract highorder structural information and then feed them into LLMs.

There are various projectors to map graph features into the LLM embedding space, such as crossattention [\(Alayrac et al.,](#page-8-7) [2022\)](#page-8-7), Q-Former [\(Li et al.,](#page-11-17) [2023\)](#page-11-17), position-aware vision language adapters [\(Bai et al.,](#page-8-8) [2023\)](#page-8-8), and light-weight Multi-layer Perceptron (MLP). Q-Former is the most widely adopted projector [\(Liu et al.,](#page-12-9) [2023b;](#page-12-9) [Fang et al.,](#page-10-8) [2024a;](#page-10-8) [Zhang et al.,](#page-15-3) [2023c\)](#page-15-3), maintaining a set of learnable query tokens to interact with the graph encoder and extract features. However, Instruct-Mol [\(Cao et al.,](#page-9-7) [2023\)](#page-9-7) argues that Q-Former requires a large number of paired data for pretraining, making alignment inefficient, and instead employs a lightweight MLP for alignment. DeCo [\(Yao et al.,](#page-14-15) [2024\)](#page-14-15) also find that Q-Former tends to lose finegrained visual attributes and spatial locality in visual LLMs.

3.3 3D Structures

The 3D structures of molecules is crucial because it contains spatial information essential for understanding molecular dynamics, protein-ligand interactions, enzymatic functions, and other biomolecular phenomena [\(Li et al.,](#page-11-8) [2024d\)](#page-11-8). Unlike 1D sequences or 2D graphs, 3D structures provide a complete geometric representation of the molecule, allowing models to take into account the threedimensional arrangement of atoms and the distances between them. MolLM [\(Tang et al.,](#page-13-14) [2024\)](#page-13-14) and Uni-Mol [\(Zhou et al.,](#page-15-9) [2023\)](#page-15-9) demotarte performance enhancement in downstream tasks when incorporating 3D information. 3D-MoLM [\(Li et al.,](#page-11-8) [2024d\)](#page-11-8) utilizes Uni-Mol [\(Zhou et al.,](#page-15-9) [2023\)](#page-15-9) to encode 3D conformations generated from SMILES and employs Q-Former [\(Li et al.,](#page-11-17) [2023\)](#page-11-17) for crossmodal alignment. This approach outperforms baseline models that rely on 1D or 2D molecular perceptions in tasks such as molecule-text retrieval, molecule captioning, and open-text question answering, particularly when addressing 3Ddependent properties. In contrast, 3D-MolT5 [\(Pei](#page-12-16) [et al.,](#page-12-16) [2024b\)](#page-12-16) contends that the modality alignment approach employed by 3D-MoLM [\(Li et al.,](#page-11-8) [2024d\)](#page-11-8) is inefficient and introduces a specialized 3D vocabulary to train 1D, 3D, and text modalities within a unified architecture, demonstrating significant improvements over 3D-MoLM [\(Li et al.,](#page-11-8) [2024d\)](#page-11-8) in various downstream tasks.

Figure 3: For example, the compound $C_8H_{11}NO$ can be represented across various modalities. 1D sequeues include SMILES, IUPAC name and so on. Molecular structure consist of 2D graphs and 3D structures, 2D graphs encompass three matrices: atomic features, atom connection, chemical bonds features, 3D strutures compromise the coordinate of every atom. Other modalities consist of mass spectra, images, and so on.

3.4 Other Modalities

2D graphs or 3D structures generated by RDKit are often represented as matrices, which are not humanreadable. In contrast, chemical images are more intuitive and frequently used to represent chemical structures in a human-friendly format. At the same time, numerous efficient image algorithms, such as the Vision Transformer (ViT) [\(Dosovitskiy,](#page-9-14) [2020\)](#page-9-14) and Swin Transformer [\(Liu et al.,](#page-12-17) [2021\)](#page-12-17), can be directly employed as modality encoders. GIT-Mol [\(Liu et al.,](#page-11-18) [2024c\)](#page-11-18) utilizes Swin Transformer [\(Liu et al.,](#page-12-17) [2021\)](#page-12-17) from SwinOCSR for image ecoding, and adopt cross-attention for modal alignment. ChemVLM [\(Li et al.,](#page-11-7) [2024c\)](#page-11-7) adopts InternViT-6B [\(Chen et al.,](#page-9-15) [2024d\)](#page-9-15) as the vision encoder, following the LLaVA [\(Liu et al.,](#page-11-13) [2024a\)](#page-11-13) architecture in the "ViT-MLP-LLM" style. Additionally, ChemVLM introduces three new chemical image datasets — ChemOCR, MMCR-Bench, and MMChemBench, However, these datasets are not open-source at this time. To facilitate future research on chemical images, we provide a summary of existing chemical image datasets in Appendix [C.](#page-15-10)

Another important chemistry-specific modality is spectral , which can be obtained through simulations (CFMID 4.0, [Wang et al.](#page-13-15) [\(2021\)](#page-13-15)) and experiments. This data is rich in structural information and plays a vital role in determining molecular structures. For example, MSNovelist [\(Stravs et al.,](#page-13-16) [2022\)](#page-13-16) utilizes an encoder-decoder neural network to generate molecular structures de novo from tandem mass spectrometry, but its accuracy is less than 50%. Comprehensive exploration of the diverse information embedded in these spectral modalities is crucial for advancing research in this domain.

4 Chemistry Tools

Although domian knowledge training and multimodal enhancement can encode a certain amount of domain-specific knowledge into LLMs, it is constrained by scalability and intrinsic memory capacity [\(Chiang et al.,](#page-9-6) [2024\)](#page-9-6). In this section, We emphasize improving the capability of LLMs to tackle complex chemistry and embodied problems through the use of chemistry tools, such as operating experimental equipment for scientific research. We categorize these chemistry tools into three types: structured knowledge retrieval, machine learning (ML) models, and embodied robots.

4.1 Structured Knowledge Retrieval

Structured knowledge retrieval, or retrievalaugmented generation (RAG, [\(Lewis et al.,](#page-11-19) [2020\)](#page-11-19)), has been proposed to alleviate hallucinations in both chemistry-specific and general LLMs [\(Xu](#page-14-16) [et al.,](#page-14-16) [2024\)](#page-14-16). The key component of knowledge retrieval is the knowledge source, and the retrieval method is typically determined by the source. We categorize common knowledge sources as follows:

1. *Database*: There are many famous chemistry database, such as, Materials Project (MP, [Jain et al.](#page-10-15) [\(2013\)](#page-10-15)), OPTIMADE [\(Evans et al.,](#page-10-5)

[2024\)](#page-10-5). These databases cannot be accessed through direct web searches; instead, data retrieval requires following specific API documentation. LLaMP [\(Chiang et al.,](#page-9-6) [2024\)](#page-9-6) design hierarchical ReAct [\(Yao et al.,](#page-14-17) [2022\)](#page-14-17) agents that can dynamically and recursively interact with MP to ground LLMs on highfidelity materials informatics.

- 2. *Scientific Literature*: Peer-reviewed research articles are the most accurate and authoritative data source, and there are many Scholarly engines can help us find the related papers. [Zheng et al.](#page-15-1) [\(2023\)](#page-15-1) propose to use ChatGPT for text mining the synthesis conditions of metal-organic frameworks (MOFs) and develop a ChatGPT Chemistry Assistant (CCA) chatbot base on the systhesis dataset and bibliographic context (such as authors and DOI), to alleviate hallucinatory errors.
- 3. *Knowledge Graph*: A knowledge graph is a structured representation that allows for complex queries and provides insights that traditional databases cannot easily offer [\(Ye et al.,](#page-14-18) [2024\)](#page-14-18). [Liu et al.](#page-11-6) [\(2024b\)](#page-11-6) propose KG-driven Knowledge Injection (DRAK-K) by retrieving the top-k most relevant pieces of knowledge and transforming the related knowledge into structured background context for LLMs.

4.2 ML Models

LLMs are prone to worse than existing ML baselines [\(Guo et al.,](#page-10-16) [2023\)](#page-10-16) in reaction-related tasks, and this tasks are difficult to be solved by knowledge retriveal. On the other hand, LLMs can interact with various tools (APIs) to accomplish complex tasks [\(Qin et al.,](#page-12-1) [2023\)](#page-12-1) in ReAct [\(Yao et al.,](#page-14-17) [2022\)](#page-14-17) style , and we can boost chemistry LLMs performance with SOTA ML models. ChemCrow [\(M. Bran et al.,](#page-12-8) [2024\)](#page-12-8) design reacttion tool set consist of NameRXN, ReactionPredict and ReactionPlanner provied by RXN4Chemistry API from IBM Research, and plan the syntheses of an insect repellent and three organocatalysts. ChatChemTS [\(Ishida et al.,](#page-10-7) [2024\)](#page-10-7) develop a user frendly chatbot named ChatChemTS which utilize AI-based molecule generators such as ChemTSv2 [\(Ishida](#page-10-17) [et al.,](#page-10-17) [2023\)](#page-10-17) for molecular design. ChatMOF [\(Kang](#page-11-5) [and Kim,](#page-11-5) [2024\)](#page-11-5) foucs on generating new metal organic frameworks (MOFs, [Kitagawa et al.](#page-11-20) [\(2014\)](#page-11-20)) which are useful in many chemical applications

due to large porosity, high surface area,and exceptional tunability [\(Deng et al.,](#page-9-16) [2012\)](#page-9-16), and they also predict the properties of generated MOFs. They adopt MOFTransformer [\(Kang et al.,](#page-11-21) [2023\)](#page-11-21) for the universal prediction of MOF properties and genetic algorithm [\(Park et al.,](#page-12-18) [2022\)](#page-12-18) to generate new MOFs, and achieve high accuracy of 95.7% for predicting, and 87.5% for generating tasks with GPT-4.

ML models can also help discover new catalyst by just giving feedback, ChemReasoner [\(Sprueill](#page-13-10) [et al.,](#page-13-10) [2024\)](#page-13-10) use atomistic graph neural networks (GNNs) trained from quantum chemistry simulations for structure-based scoring, the GNNs are used to yeild reward and drive LLM towards catalysts with specific properties. This novel idea suggest that ML models not only can be used as tools aid in specific task, but also can be used as feeback to guide and stimulate the LLMs to fulfill the tasks by themselfs.

4.3 Embodied Robots

Chemistry experiments are often resoure- and laborintensive, and automated experiments canattain higher throughput and precision [\(Tom et al.,](#page-13-17) [2024\)](#page-13-17). However, the discovery of new material requires not only automation but autonomy—the ability of an experimental agent to interpret data and make decisions based on it [\(Szymanski et al.,](#page-13-18) [2023\)](#page-13-18), where LLMs are excellent at planing and reasoning, showing promise of sought-after system that autonomously designs and executes scientific experiments [\(Boiko et al.,](#page-8-4) [2023\)](#page-8-4).

Coscientist [\(Boiko et al.,](#page-8-4) [2023\)](#page-8-4) is a GPT-4 driven AI system which can autonomously designs, plans and performs complex experiments, it demonstrate the versatility and performance across six tasks. CLAIRify [\(Yoshikawa et al.,](#page-14-5) [2023\)](#page-14-5) also leverage robots and LLM to automate chemistry experiments, and they pay more attention to how to generate syntactically valid programs in a data-scarce domain-specific language that incorporates environmental constraints. ORGANA [\(Darvish et al.,](#page-9-5) [2024\)](#page-9-5) further extend CLAIRify with visual perception of the environment and support complex experiments between multiple robots.

5 Benchmarks

Benchmarks are essential for evaluating the performance of chemistry LLMs on chemistry-related tasks and can be broadly categorized into two categories: science benchmarks and moleculespecific benchmarks. Chemistry is a subset of science, and existing science benchmarks evaluate LLMs' ability to solve scientific problems, including those related to chemistry. Existing science benchmarks, such as SciQ [\(Welbl et al.,](#page-14-19) [2017\)](#page-14-19), Sci-Code [\(Tian et al.,](#page-13-19) [2024\)](#page-13-19), ScienceQA [\(Lu et al.,](#page-12-19) [2022\)](#page-12-19), AGIEval [\(Zhong et al.,](#page-15-11) [2023\)](#page-15-11), SciEval [\(Sun](#page-13-20) [et al.,](#page-13-20) [2024\)](#page-13-20), SciBench [\(Wang et al.,](#page-13-21) [2023\)](#page-13-21), and VisScience[\(Jiang et al.,](#page-11-22) [2024\)](#page-11-22), typically cover a wide range of scientific disciplines, including biology, earth science, physics, chemistry, and even social science. Although these science benchmarks include chemistry-related tasks, they are not specifically designed for chemistry and fail to address many chemistry-specific problems.

In contrast, molecule-specific benchmarks are designed to assess knowledge in molecule-related sciences (e.g., chemistry, materials science, biochemistry). ChemLLMBench [\(Guo et al.,](#page-10-16) [2023\)](#page-10-16) first adapts traditional chemistry tasks to a language model setting, evaluating the performance of contemporary LLMs in zero-shot and few-shot prompts. SciKnowEval [\(Feng et al.,](#page-10-12) [2024\)](#page-10-12) expands the chemistry domain to molecules by introducing a large dataset of 50,000 problems that assess various LLM abilities, including knowledge coverage, reflection and reasoning, and application. MassSpecGym [\(Bushuiev et al.,](#page-9-17) [2024\)](#page-9-17) focuses on characterization techniques, such as Tandem Mass Spectrometry (MS/MS), and evaluates the ability of LLMs to elucidate molecular structures from MS/MS data. Notably, there are several other important chemistry benchmarks, including ScholarChemQA [\(Chen et al.,](#page-9-18) [2024a\)](#page-9-18), SCIASSESS [\(Cai](#page-9-19) [et al.,](#page-9-19) [2024\)](#page-9-19), SciKnowEval [\(Feng et al.,](#page-10-12) [2024\)](#page-10-12), ChemEVal [\(Huang et al.,](#page-10-18) [2024b\)](#page-10-18), [Alberts et al.](#page-8-9) [\(2024\)](#page-8-9), and MolPuzzles [\(Guo et al.,](#page-10-19) [2024\)](#page-10-19). Due to page limitations, we provide a brief overview of these benchmarks in Table [3.](#page-17-0)

6 Future Directions

Although current approaches have made steady progress towards chemistry LLMs, there remains significant room for improvement. Future research directions can be categorized into three main aspects: data, model, and application.

6.1 Data

Data Diversity Training data is the foundation of LLMs. However, most existing datasets are built from pre-existing sources, such as MoleculeNet [\(Wu et al.,](#page-14-20) [2018\)](#page-14-20), and cover a limited range of chemistry tasks. Future work should aim to create more diverse and comprehensive datasets to enhance the training of chemistry LLMs and broaden their capabilities.

CoT Reasoning Chain-of-Thought (CoT, [Wei](#page-14-1) [et al.](#page-14-1) [\(2022b\)](#page-14-1)) reasoning is one of the most notable emergent abilities of LLMs, involving the generation of a sequence of intermediate steps leading to the final answer. However, existing chemistry LLMs often lack this critical reasoning capability due to simple training instruction pairs. Developing training data with explicit reasoning paths to effectively elicit the CoT ability in chemistry LLMs is a crucial direction for future research.

Chemical Modality As described in Section [3.4,](#page-5-2) many chemistry-specific spectra are not yet fully exploited in in chemistry LLMs. However, these spectra contain rich structural information that can be valuable for various chemical tasks. For example, tandem mass spectrometry (MS/MS) can provide detailed insights into the molecular structure, allowing for the identification and characterization of compounds and elucidation of reaction mechanisms.

6.2 Model

Multi-Modal Alignment Most works towards multi-modal chemistry LLMs always invole a single pair of modalities, limiting their representations ability. Align multiple N (\geq 3) modalities is a promising direction as different modalites are complementary and can provide more comprehensive understanding of chemistry molecules.

RLXF RLHF is a crucial step in training powerful LLMs. Although obtaining human feedback is challenging, especially in chemistry where data annotation requires specialized domain knowledg, we can leverage advanced LLMs as assistants to guide this process. Additionally, we can also utilize results from professional chemistry software as a form of reward to align chemistry LLMs.

6.3 Application

Research Assistants Chemistry LLMs have the potential to serve as powerful research assistants, aiding chemists by automating routine tasks such as literature review, data analysis, and hypothesis generation. For future development, these models can be designed to understand complex scientific queries, provide insights from vast amounts of chemical literature, suggest experimental protocols, and even propose novel research directions.

Automated Experimentation Automated experimentation is another promising direction for advancing chemistry LLMs. Integrating these models with automated laboratory systems can enable them to not only predict molecular properties or suggest potential chemical reactions but also design, execute, and analyze experiments in real-time. Future research should explore how chemistry LLMs can be trained and aligned to interact with automated experimental setups, ensuring reliability, safety, and compliance with scientific standards.

7 Conclusion

In this survey, we systematically investigate the current approaches to adapting general LLMs for chemistry LLMs. We highlight key challenges, including domain knowledge, multi-modal data, and the integration of chemistry-specific tools, and review existing efforts to address these challenges. While significant progress has been made, achieving chemical general intelligence remains a distant goal, and we identify promising future directions. We hope this survey will inspire further innovative research in the field.

Limitations

In this paper, a comprehensive review of existing methods for constructing chemistry-focused LLMs is presented, with an emphasis on three key aspects for enhancing general LLMs: domain-specific knowledge, multi-modal data, and chemistry tools. This survey aims to provide researchers with a concise understanding of chemistry LLMs and suggest potential directions for future research. However, certain limitations may be present.

References. Due to page limitations and the rapid development of the field, we may not include all relevant references and detailed technical information. However, we strive to keep our work up-to-date on our GitHub repository.

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A Related Work

The intersection of LLMs and chemistry is an urgent and rapidly growing field. Numerous works and reviews have addressed this topic, which can be broadly categorized into:

A.1 General Science

Several surveys focus on general science, including chemistry. [Zhang et al.](#page-15-12) [\(2024d\)](#page-15-12) explore LLM applications across mathematics, physics, biology, medicine, geography, geology, environmental science, and chemistry. However, the broad scope limits the depth of discussion on chemistry-specific LLMs. [Zhang et al.](#page-14-21) [\(2024b\)](#page-14-21) focus more on the chemical domain but still include biological LLMs and BERT-style models, without discussing the emergent applications of chemistry-specific agents.

A.2 Chemistry-Specific Surveys

Chemistry's significance has drawn considerable attention, leading to various efforts summarizing current trends. [Xia et al.](#page-14-22) [\(2022\)](#page-14-22) review Chemical Pre-trained Models (CPMs) based on GNNs or Transformers but focus little on LLMs. [Janakara](#page-10-20)[jan et al.](#page-10-20) [\(2024\)](#page-10-20) emphasize the role of language models in molecular discovery but offer limited insights on training chemistry-specific LLMs. [Liao](#page-11-0) [et al.](#page-11-0) [\(2024\)](#page-11-0) concentrate on molecule encoding and pretraining objectives, while [Pei et al.](#page-12-2) [\(2024a\)](#page-12-2) discuss progress from a multi-modal perspective, neglecting LLMs' tool-using potential. [Ramos et al.](#page-12-20) [\(2024\)](#page-12-20) review chemistry LLM agent applications in literature analysis, experiment planning, and hypothesis generation, but overlook multi-modal capabilities. Notably, these surveys categorize BERTstyle LMs as LLMs, despite their need for taskspecific fine-tuning and lack of emergent abilities.

B SFT Tasks Description

The most frequently used chemistry tasks for SFT and their description are shown in Table [1.](#page-16-0) In accordance with the task division presented in Table [1,](#page-16-0) we illustrate in Figure [4](#page-16-1) the data distribution of the commonly used SFT dataset.

C Molecule Image Dataset

We describe the existing molecule image dataset in Table [2.](#page-17-1)

D Benchmarks

We briefly introduce the existing benchmarks in Table [3,](#page-17-0) covering aspects such as subject, task type, dynamics, source, and modality.

| Type | Chemistry Tasks | Description |
|--------------------------------|-------------------------------|--|
| SMILES Understanding | Molecule description | Given a molecule SMILES, generating text descrip- tion illuminating the structure, properties , biological activity, and applications. |
| | Text-based molecule design | Inverse task of molecule description, given a text description, generating the molecule SMILES. |
| | Molecular property prediction | Molecular property prediction focus on drawn from Mquantum mechanics properties of molecules drawn from MoleculeNet. |
| Reaction Understanding | Reagent prediction | Reagent prediction generate suitable catalysts, sol- vents, or ancillary substances required for a specific chemical reaction. |
| | Forward reaction prediction | Forward reaction prediction generate probable prod- $uct(s)$ of a chemical reaction. |
| | Retrosynthesis | Inverse task of forward reaction prediction, generate the synthesis routes and precursor molecules given target molecule. |
| Notation Alignment | SMILES and IUPAC names | Given SMILES, generate IUPAC name, and reverse transformation. |
| | SMILES and Formulas | Given SMILES, generate formulas, and reverse trans- formation. |
| Chemistry-Related QA | OA | Chemical QA extracted from existing dataset or exam. |

Table 1: The most frequently used chemistry tasks for SFT.

Figure 4: The compositional structure of representative SFT dataset. The definition of tasks above the the horizontal lines is shown in Table [1,](#page-16-0) the source and size of the different tasks are indicated below the horizontal lines, and percentages on the pie charts are present to show the difference of different dataset.

Table 2: Overview of molecular image datasets, categorized into synthetic and realistic groups with details on their scale and descriptions. Synthetic datasets are primarily RDKit-generated or derived from large collections, while realistic datasets include handwritten and reaction images. Some datasets are closed-source or only provide evaluation data.

Table 3: A brief introduction to the existing benchmarks. "MCQ" refers to Multi-Choice Questions, while "DA" denotes Direct-Answer tasks. "Samples" refers to the number of test set examples. The "Spectra" modality is distinctive, as spectra can be represented either as images or text.