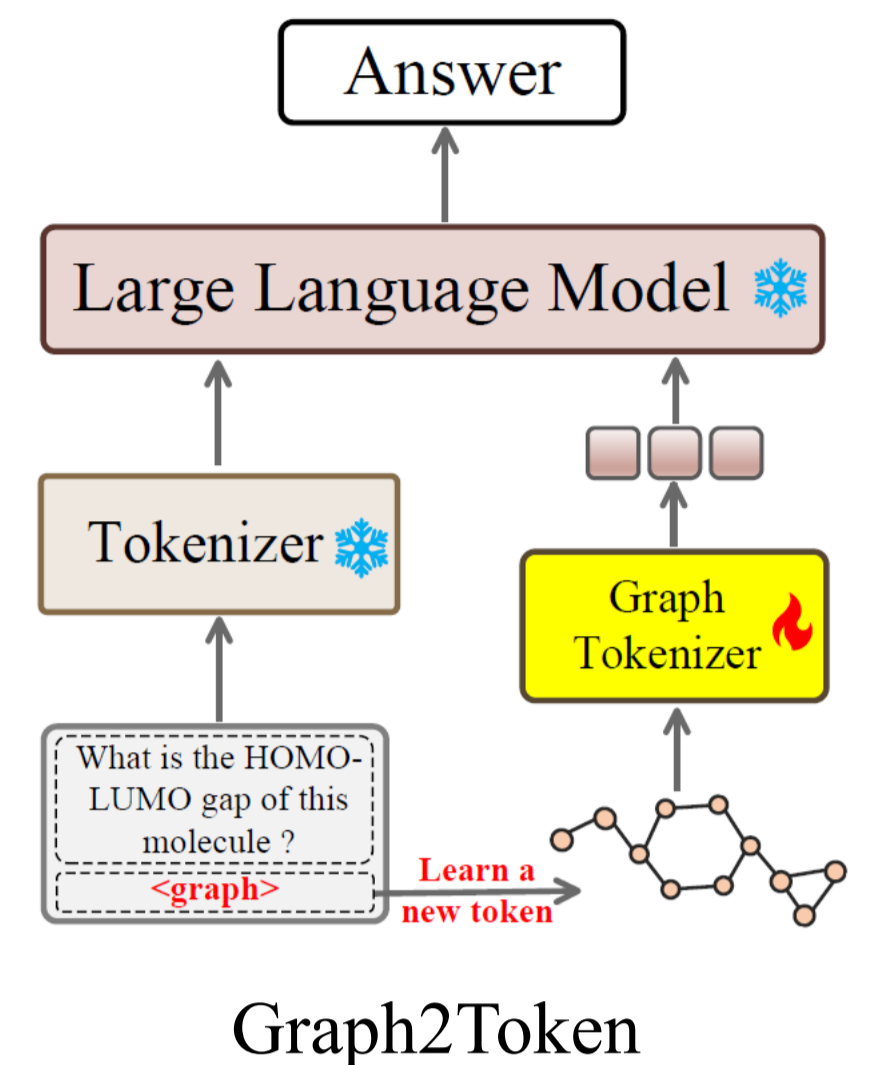
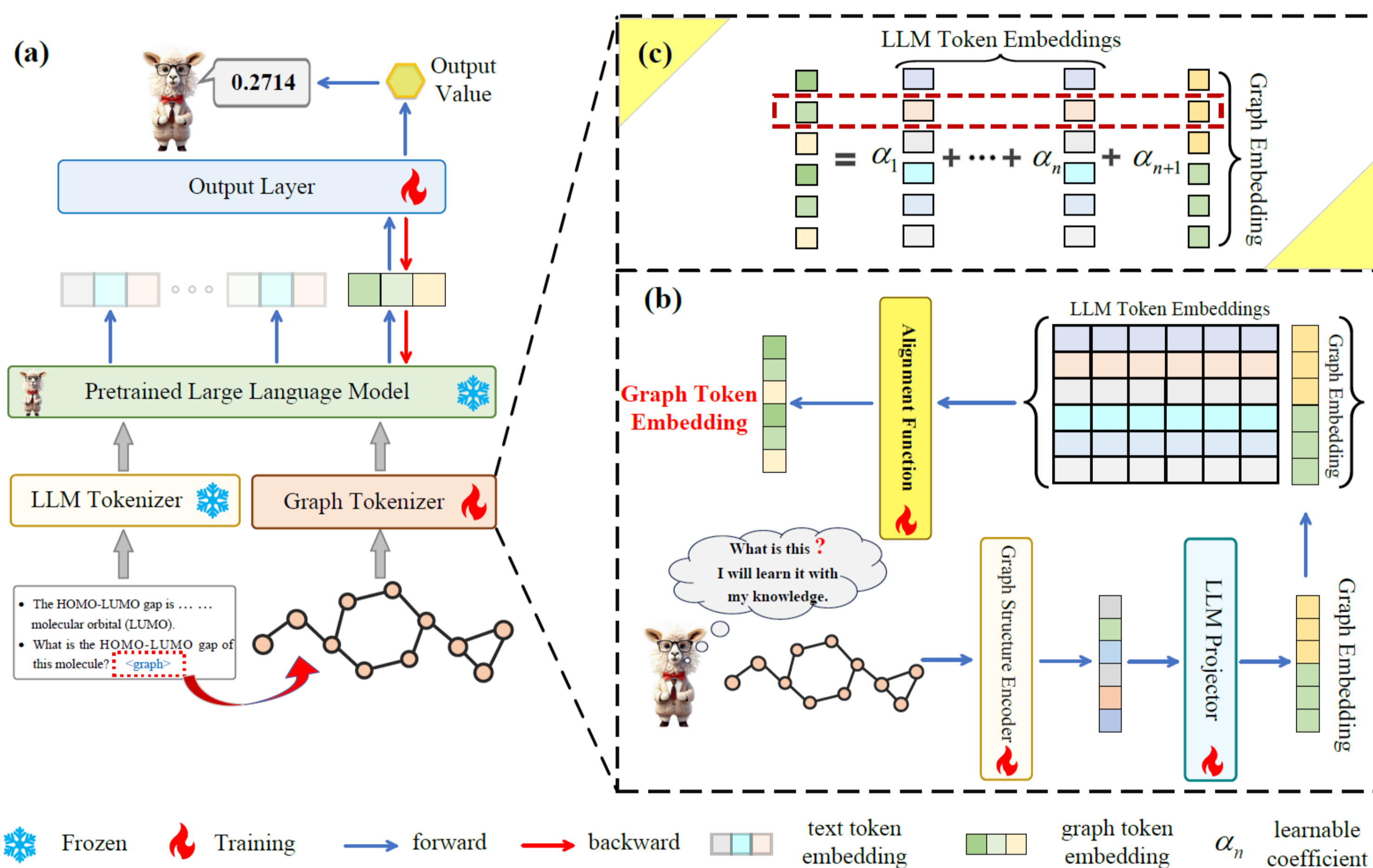


Motivation

- Large language models (LLMs) excel at various text-related tasks.
- It is still challenging for LLMs to process molecular graph data.
- Textualized graph as input \longrightarrow **insufficient** graph reasoning
- Instruction fine-tuning \longrightarrow **change the semantics** of LLM backbone
- **Graph2Token**, an efficient solution that aligns a graph token to LLM tokens.



Aligning a graph token with the LLM token vocabulary



- a) The architecture of Graph2Token with **frozen LLM tokenizer** and **trainable graph tokenizer**.
- b) The trainable graph tokenizer learns the unknown graph token representation using **LLM token vocabulary**.
- c) Alignment function utilizes a **learnable combination of tokens** pre-trained by LLM to represent the graph tokens.

Finetune and few-shot learning performance on molecular datasets

Method Type	Method	BBBP \uparrow	BACE \uparrow	HIV \uparrow	TOX21 \uparrow	Avg \uparrow
<i>Supervised Learning</i>	GIN	67.8	76.8	76.5	73.9	73.8
	GT	68.7	77.2	74.2	75.5	73.9
<i>Graph Pretrain Finetuning</i>	GraphMVP-C	72.4	81.2	77.0	74.4	76.3
	Mole-BERT	70.8	79.3	76.0	75.9	75.5
	MolFM	<u>72.9</u>	83.9	<u>78.8</u>	<u>77.2</u>	<u>78.2</u>
	SimSGT	72.3	83.6	77.7	75.7	77.3
<i>LLM-Based Tuning</i>	Llama-2-7B-chat	65.6	74.8	62.3	-	67.6
	Vicuna-v1.3-7B	60.1	68.3	58.1	-	62.6
	MolCA-S	70.8	79.3	-	76.0	75.4
	MolCA-GS	70.0	79.8	-	<u>77.2</u>	75.7
	InstructMol-G	64.0	85.9	74.0	-	74.6
	InstructMol-GS	70.0	82.3	68.9	-	73.7
	Graph2Token	73.5	<u>85.0</u>	79.4	79.2	79.3

Ratio	Dataset	Graph2Token	GIN	GCN
5%	BBBP	64.7	61.8	64.4
	BACE	73.2	64.4	65.1
	HIV	68.5	66.2	62.7
	TOX21	70.6	62.6	58.4
10%	BBBP	69.5	66.9	67.0
	BACE	74.6	68.1	64.6
	HIV	69.7	66.9	60.0
	TOX21	71.2	66.7	68.4

- Few-shot learning using 5% and 10% training data on different molecular datasets.

- Results (ROC-AUC) of finetune learning on molecular classification tasks on different datasets compared with LLM-based methods and graph learning methods.

