

Advancing Molecular Graph-Text Pre-training via Fine-grained Alignment

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1

Background

2

Fine-grained Alignment

3

FineMolTex

4

Experiments

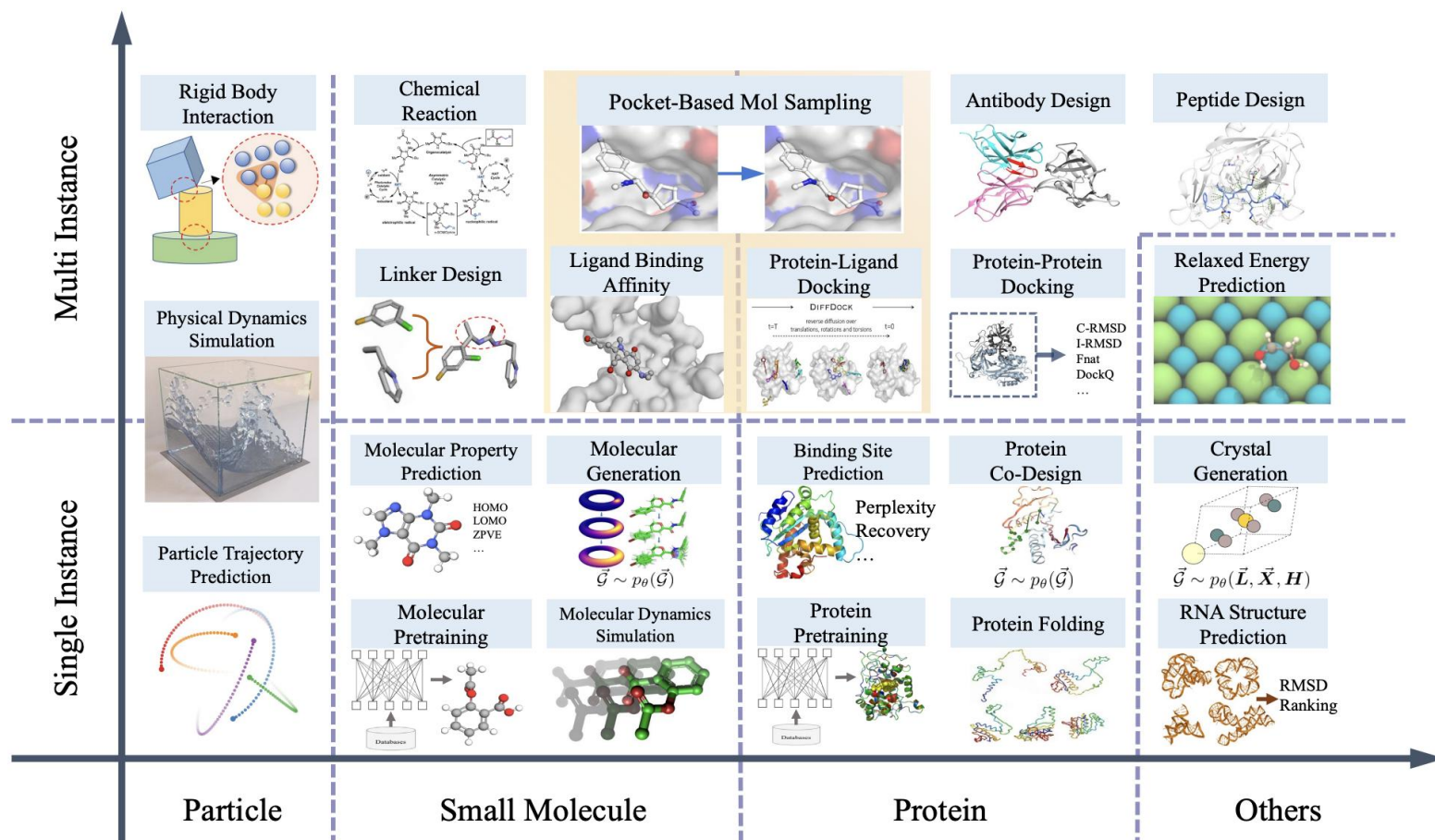
5

Conclusion

1 Background Molecular Graph-Text Pre-training

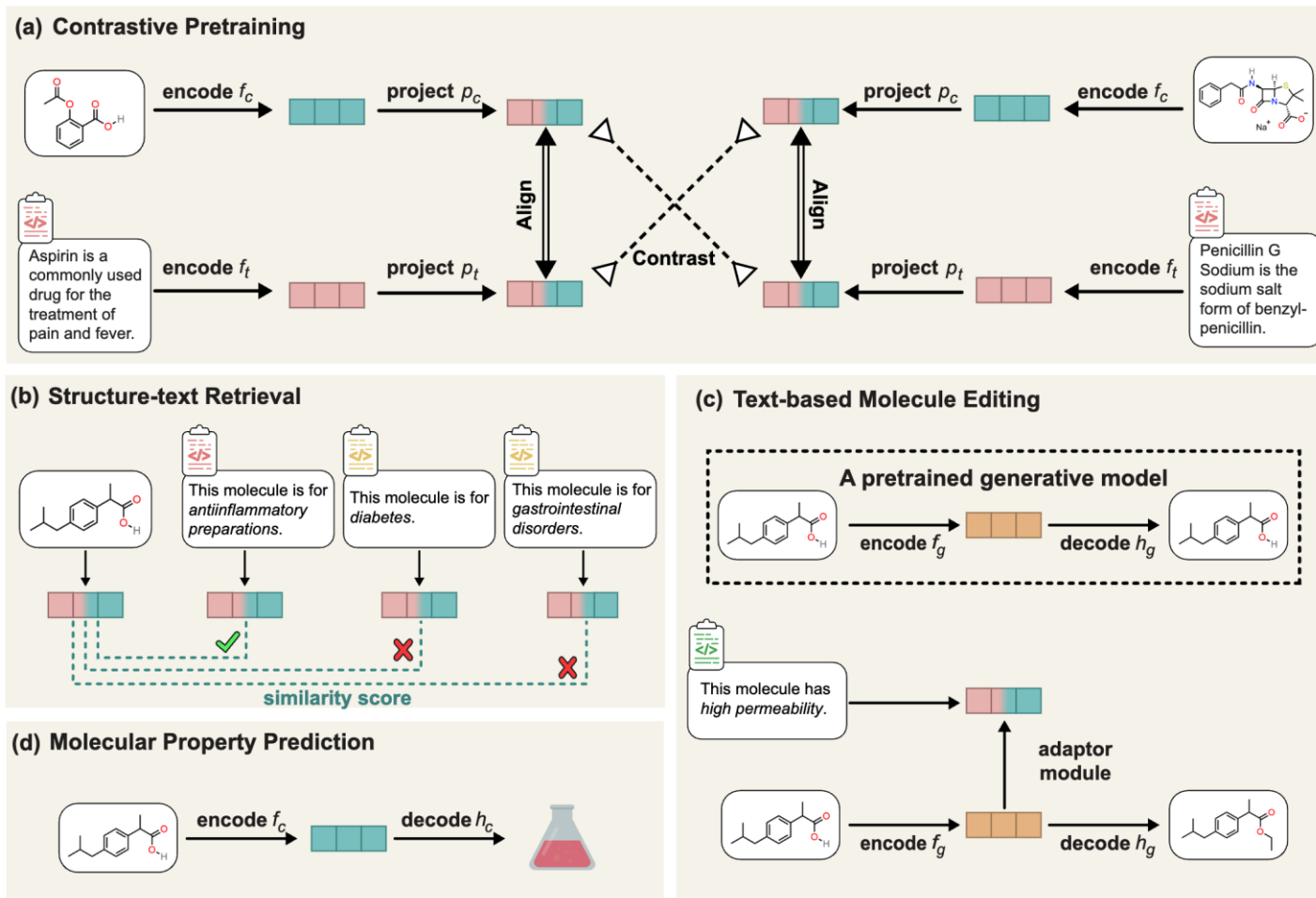


Comprehending molecular structure and related knowledge is pivotal in scientific investigations spanning diverse fields



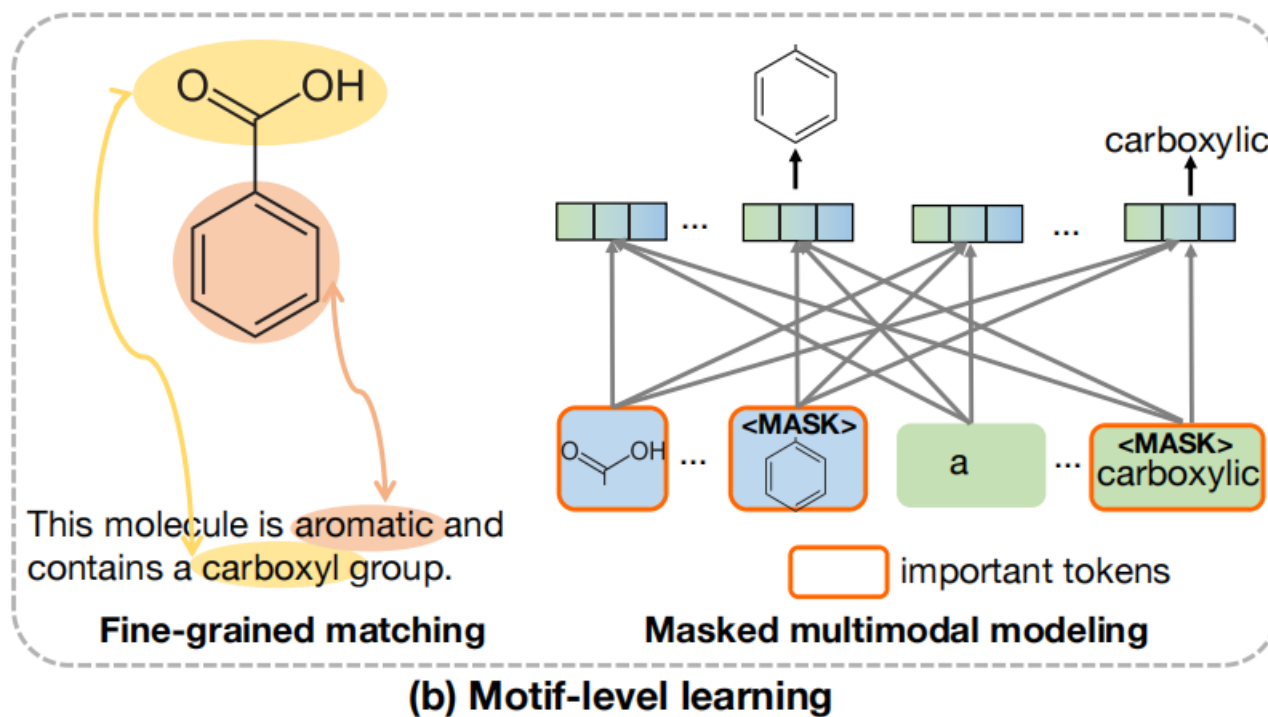
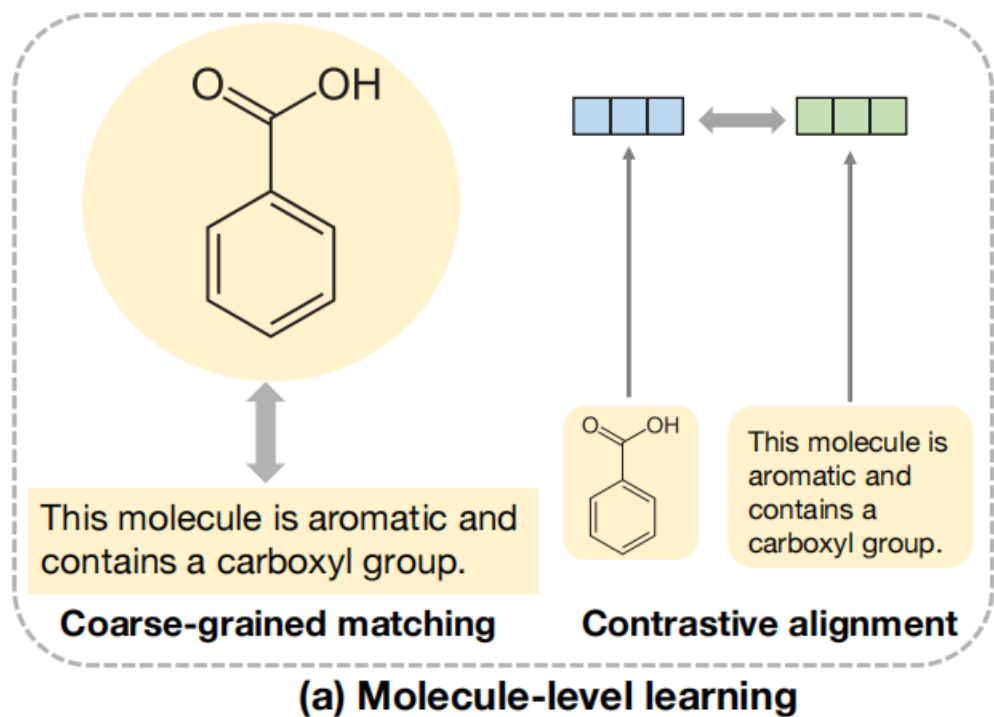


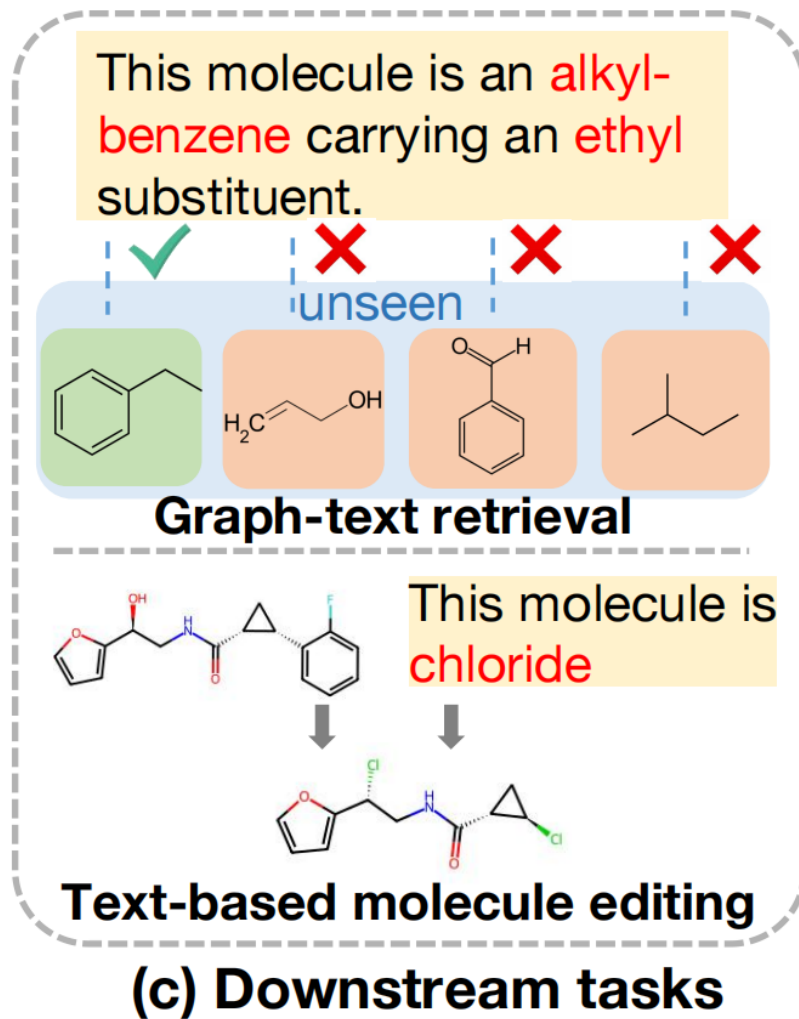
- Several studies explore molecular structures along with their corresponding descriptions.



- 1 Background
- 2 **Fine-grained Alignment**
- 3 FineMolTex
- 4 Experiments
- 5 Conclusion





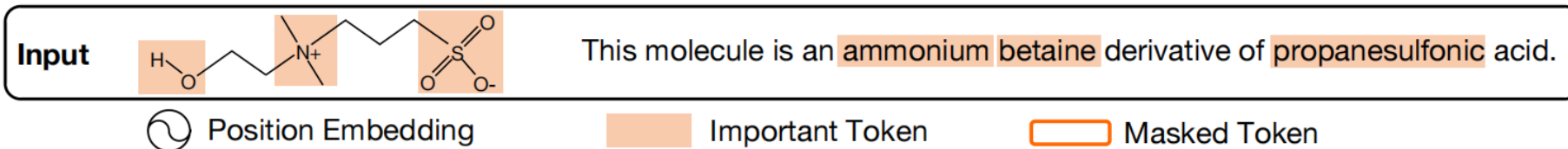
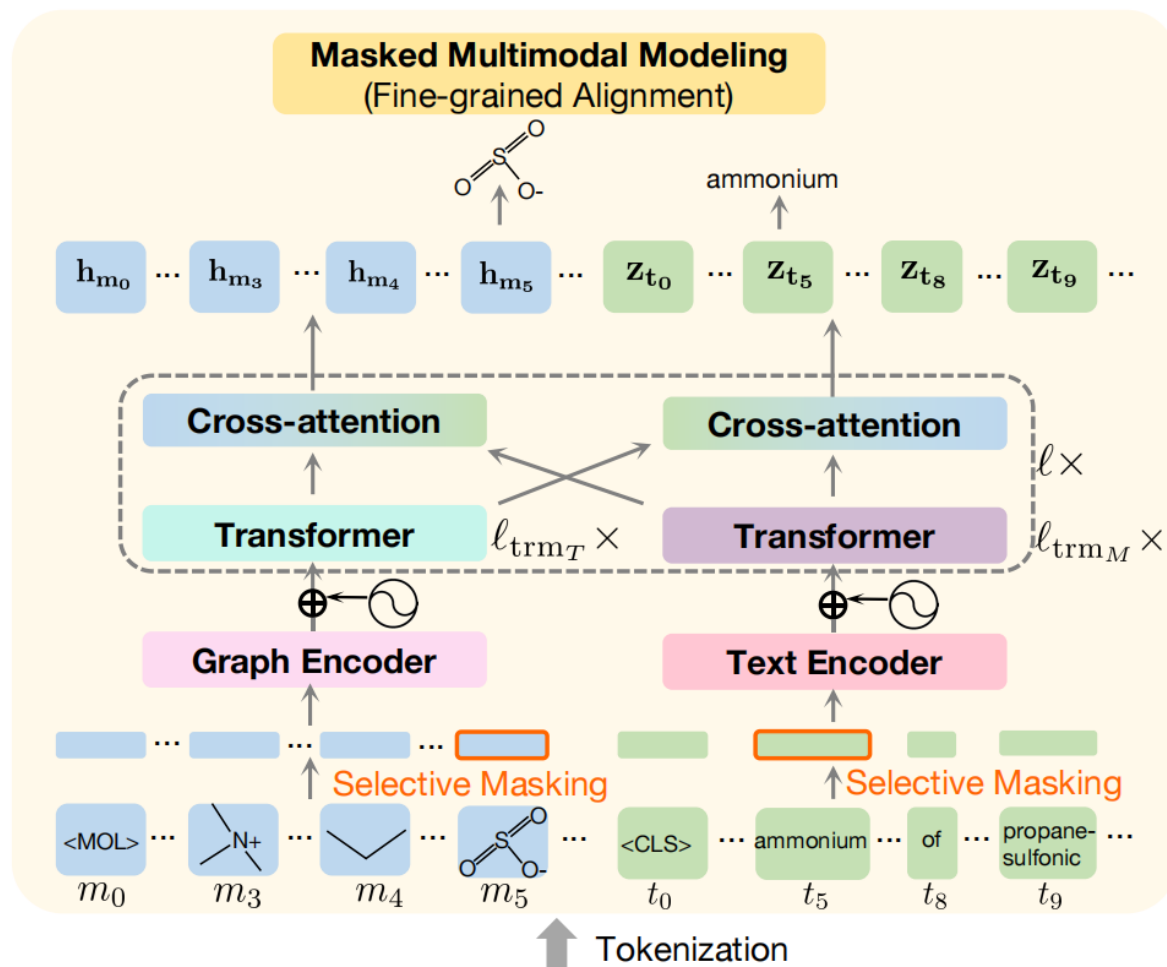
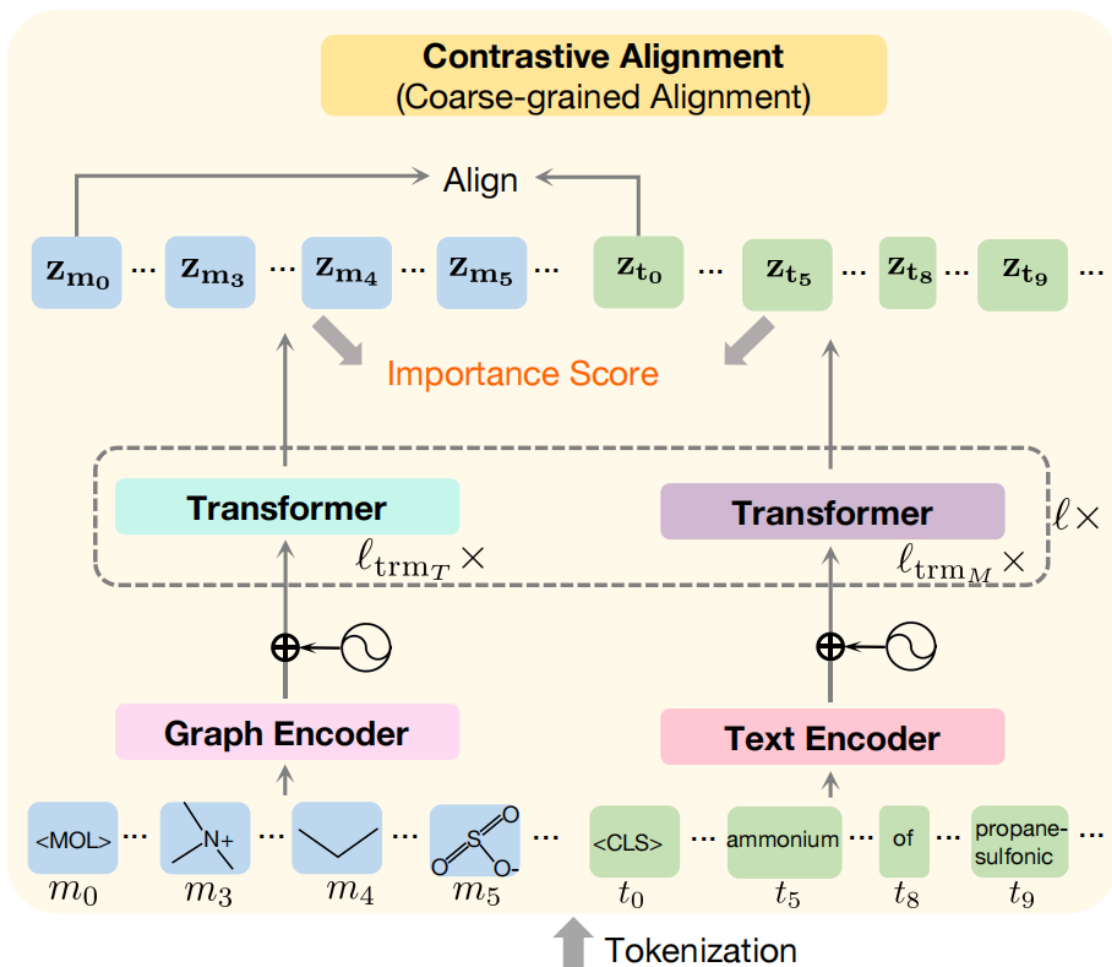


CONTENTS

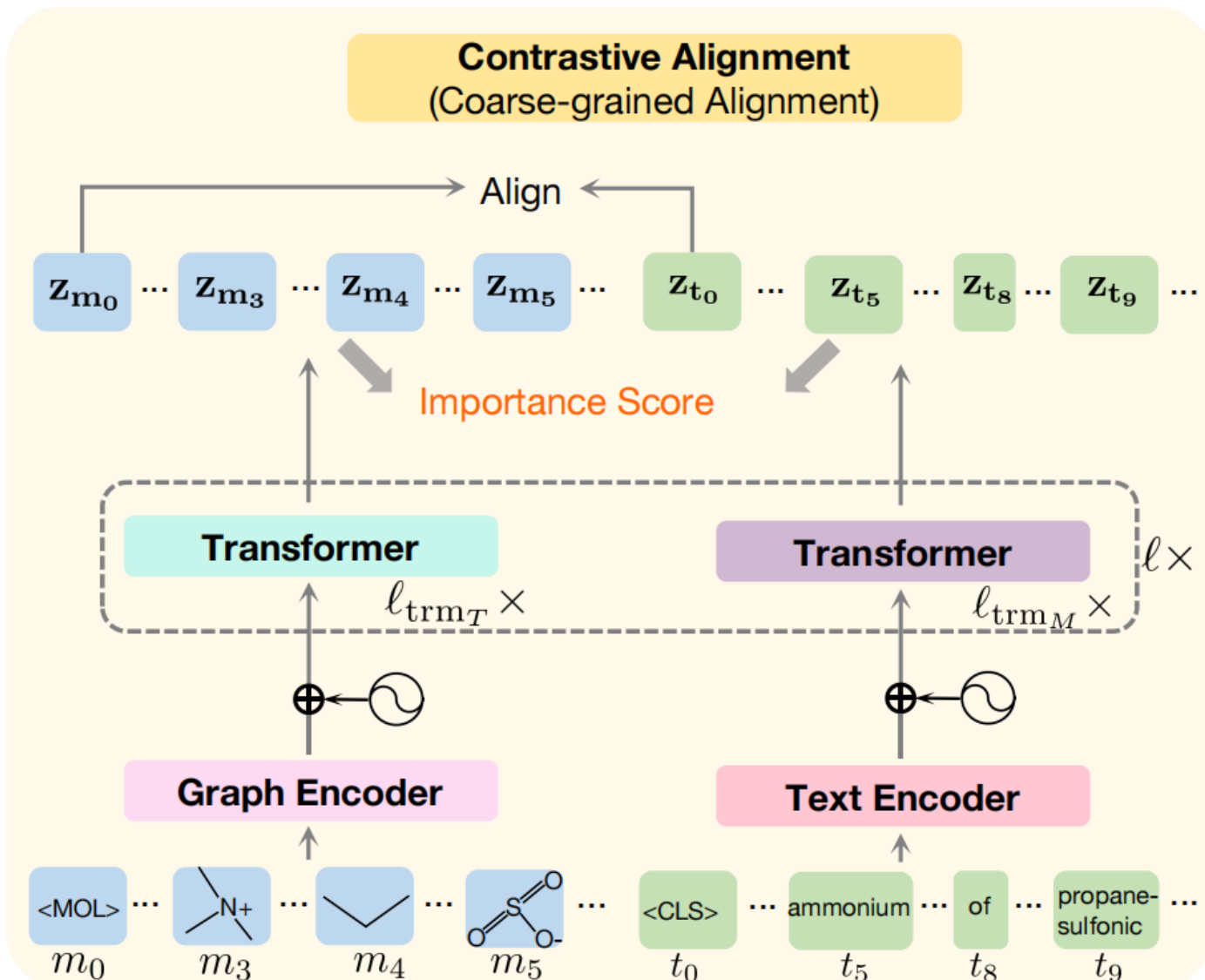
- 1 Background
- 2 Fine-grained Alignment
- 3 **FineMolTex**
- 4 Experiments
- 5 Conclusion



3 FineMolTex Framework

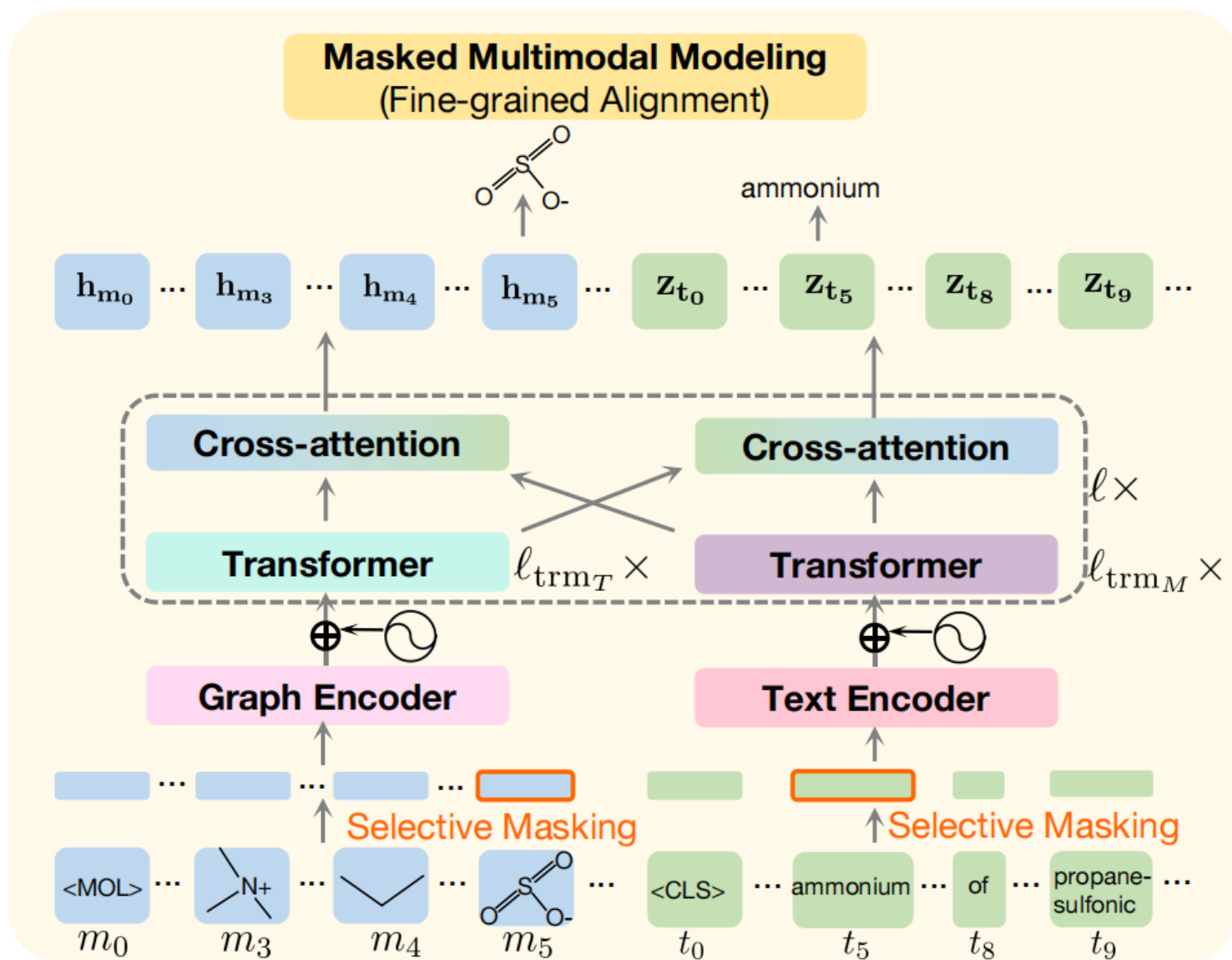


3 FineMolTex Framework



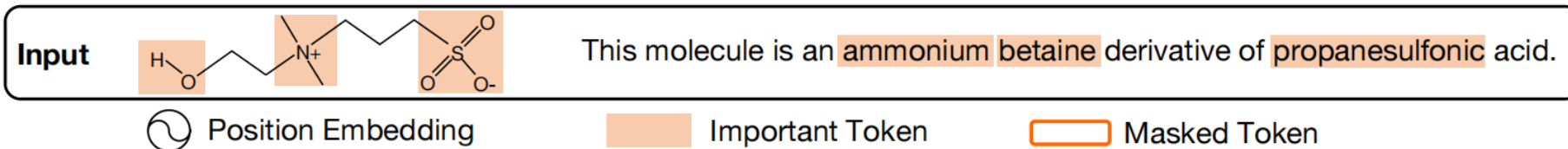
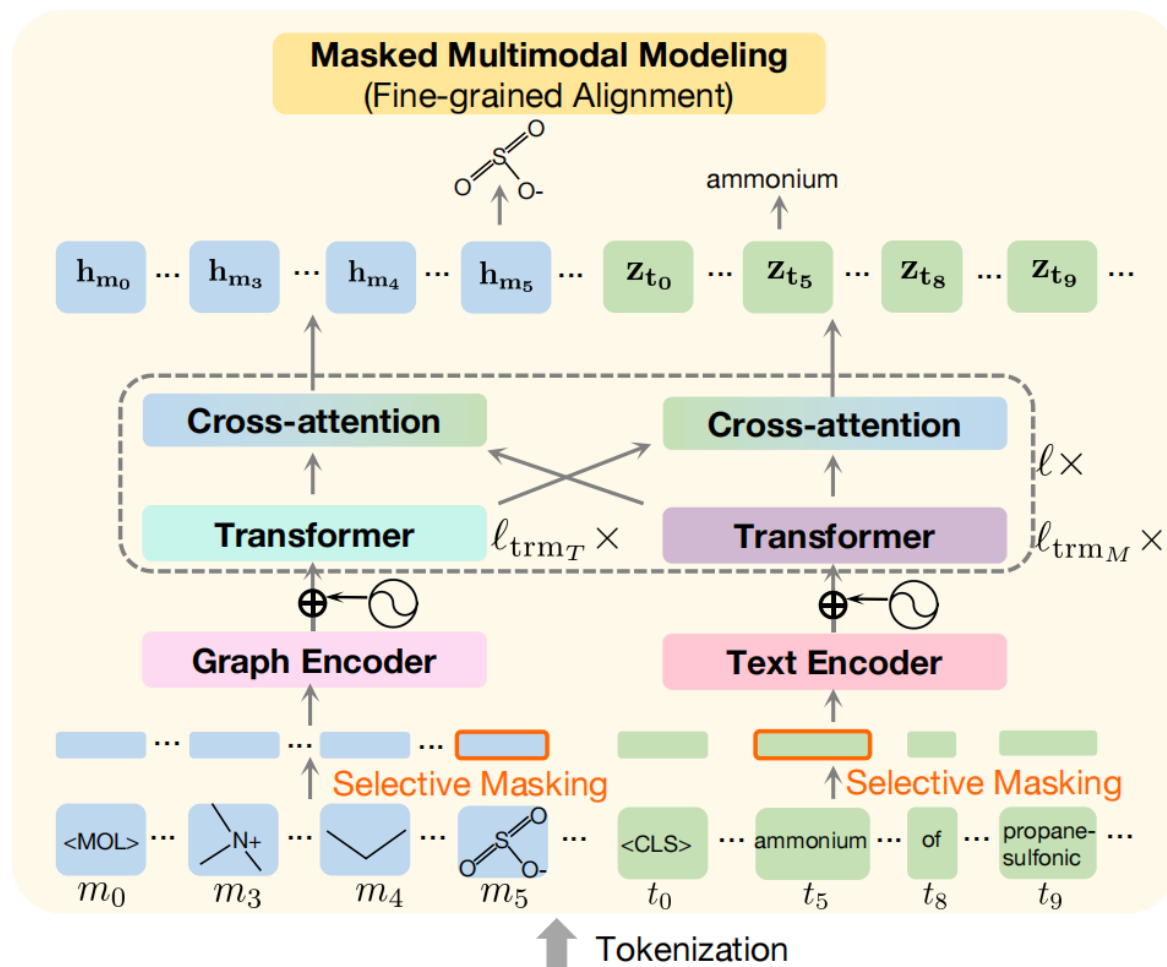
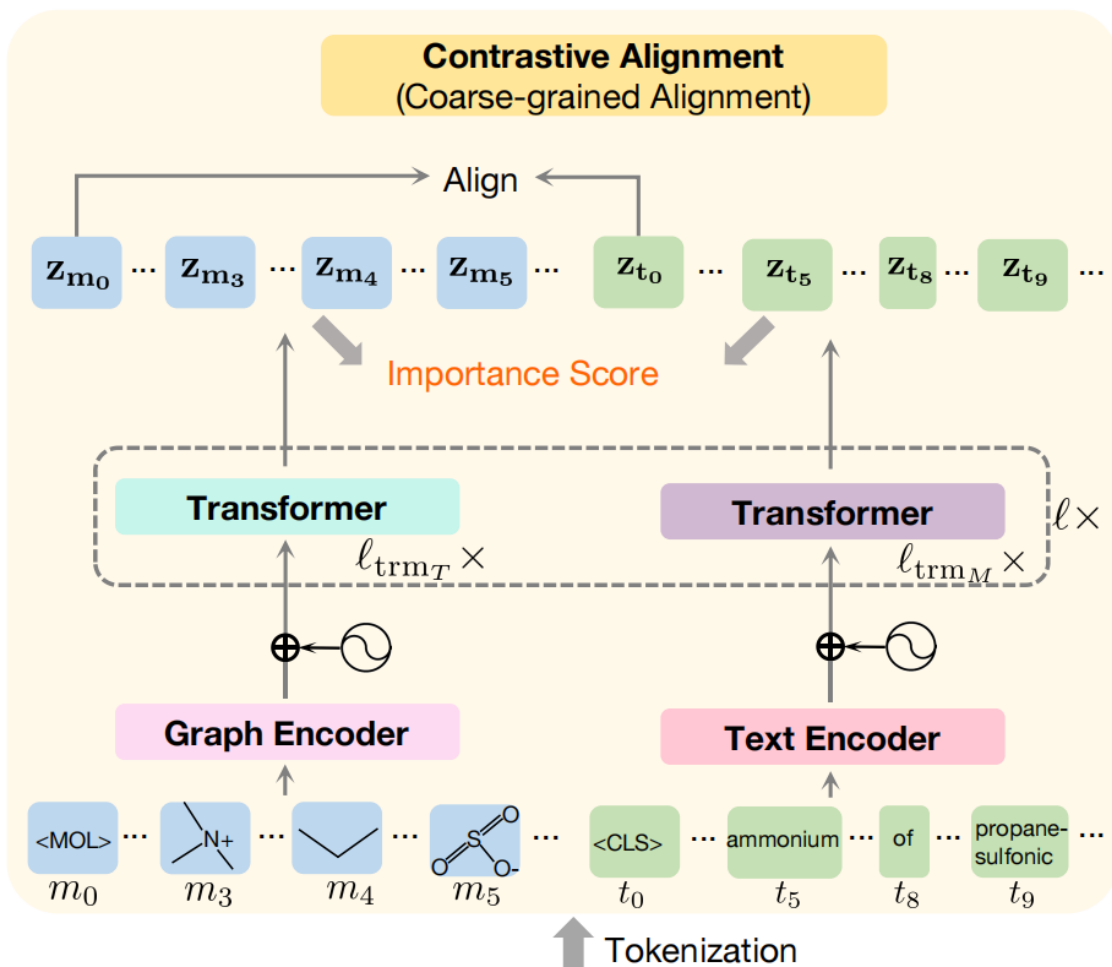
$$L_{\text{con}} = -\frac{1}{2} \mathbb{E}_{m_0, t_0} \left[\log \frac{\exp(\cos(\mathbf{z}_{m_0}, \mathbf{z}_{t_0})/\tau)}{\exp(\cos(\mathbf{z}_{m_0}, \mathbf{z}_{t_0})/\tau) + \sum_{t'_0} \exp(\cos(\mathbf{z}_{m_0}, \mathbf{z}_{t'_0})/\tau)} + \log \frac{\exp(\cos(\mathbf{z}_{t_0}, \mathbf{z}_{m_0})/\tau)}{\exp(\cos(\mathbf{z}_{t_0}, \mathbf{z}_{m_0})/\tau) + \sum_{m'_0} \exp(\cos(\mathbf{z}_{t_0}, \mathbf{z}_{m'_0})/\tau)} \right],$$

3 FineMolTex Framework



$$L_{\text{pre}} = \beta \sum_i \text{CE}(\hat{y}_{m_i}, y_{m_i}) + \alpha \sum_j \text{CE}(\hat{y}_{t_j}, y_{t_j}).$$

3 FineMolTex Framework



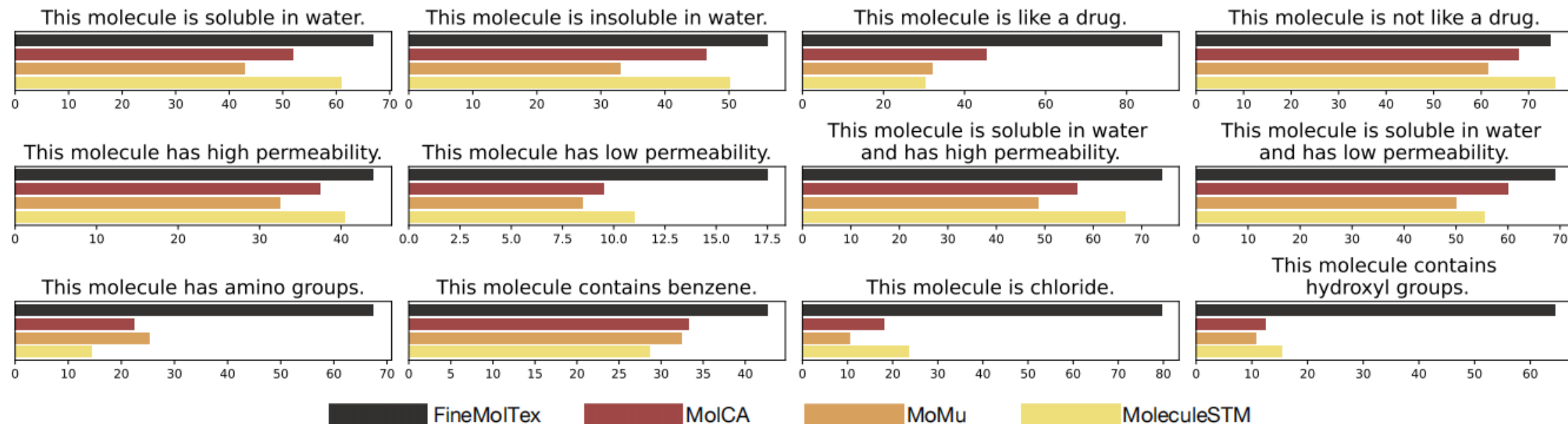
CONTENTS

- 1 Background
- 2 Fine-grained Alignment
- 3 FineMolTex
- 4 **Experiments**
- 5 Conclusion





<i>T</i>	Given Molecular Graph			Given Text		
	4	10	20	4	10	20
KV-PLM	68.38±0.03	47.59±0.03	36.54±0.03	67.68±0.03	48.00±0.02	34.66±0.02
MolCA	83.75±0.54	74.25±0.26	66.14±0.21	81.27±0.33	69.46±0.17	62.13±0.16
MoMu-S	70.51±0.04	55.20±0.15	43.78±0.10	70.71±0.22	54.70±0.31	44.25±0.43
MoMu-K	69.40±0.11	53.14±0.26	42.32±0.28	68.71±0.03	53.29±0.05	43.83±0.12
3D-MoLM	81.35±0.14	73.65±0.13	64.79±0.15	79.78±0.22	62.38±0.16	53.43±0.11
MV-Mol	92.24±0.26	85.38±0.19	79.41±0.43	91.28±0.13	85.32±0.15	80.37±0.22
MoleculeSTM	92.14±0.02	86.27±0.02	81.08±0.05	91.44±0.02	86.76±0.03	81.68±0.03
FineMolTex	96.78±0.05	92.48±0.02	87.94±0.14	96.29±0.12	91.65±0.15	85.07±0.11





Model	BBBP	Tox21	ToxCast	Sider	ClinTox	MUV	HIV	Bace	Avg
AttrMask	67.8±2.6	75.0±0.2	63.6±0.8	58.1±1.2	75.4±8.8	73.8±1.2	75.4±0.5	80.3±0.0	71.2
ContextPred	63.1±3.5	74.3±0.2	61.6±0.5	60.3±0.8	80.3±3.8	71.4±1.4	70.7±3.6	78.8±0.4	70.1
InfoGraph	64.8±0.6	76.2±0.4	62.7±0.7	59.1±0.6	76.5±7.8	73.0±3.6	70.2±2.4	77.6±2.0	70.0
MolCLR	67.8±0.5	67.8±0.5	64.6±0.1	58.7±0.1	84.2±1.5	72.8±0.7	75.9±0.2	71.1±1.2	71.3
GraphMVP	68.1±1.4	77.1±0.4	65.1±0.3	60.6±0.1	84.7±3.1	74.4±2.0	77.7±2.5	80.5±2.7	73.5
GraphCL	69.7±0.7	73.9±0.7	62.4±0.6	60.5±0.9	76.0±2.7	69.8±2.7	78.5±1.2	75.4±1.4	70.8
KV-PLM	70.5±0.5	72.1±1.0	55.0±1.7	59.8±0.6	89.2±2.7	54.6±4.8	65.4±1.7	78.5±2.7	68.2
MoMu-S	70.5±2.0	75.6±0.3	63.4±0.5	60.5±0.9	79.9±4.1	70.5±1.4	75.9±0.8	76.7±2.1	71.6
MoMu-K	70.1±1.4	75.6±0.5	63.0±0.4	60.4±0.8	77.4±4.1	71.1±2.7	76.2±0.9	77.1±1.4	71.4
MolCA	70.0±0.5	77.2±0.5	64.5±0.8	63.0±1.7	89.5±0.7	72.1±1.3	77.2±0.6	79.8±0.5	74.2
MoleculeSTM	70.0±0.5	76.9±0.5	65.1±0.4	61.0±1.1	92.5±1.1	73.4±2.9	77.0±1.8	80.8±1.3	74.6
FineMolTex	73.5±1.6	77.1±1.2	68.6±0.9	64.8±1.4	92.5±0.8	76.3±1.2	79.0±1.4	84.0±1.5	76.9

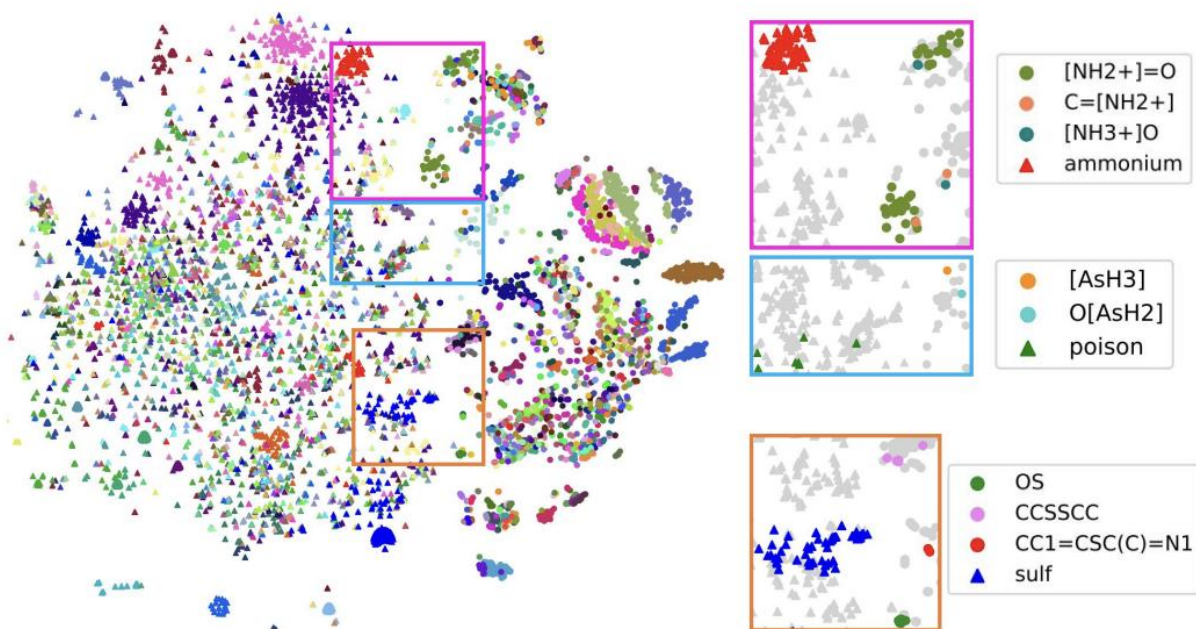


Figure 1: Visualization of motif tokens and word tokens using *t*-SNE. Triangles denote word tokens; circles denote motif tokens.

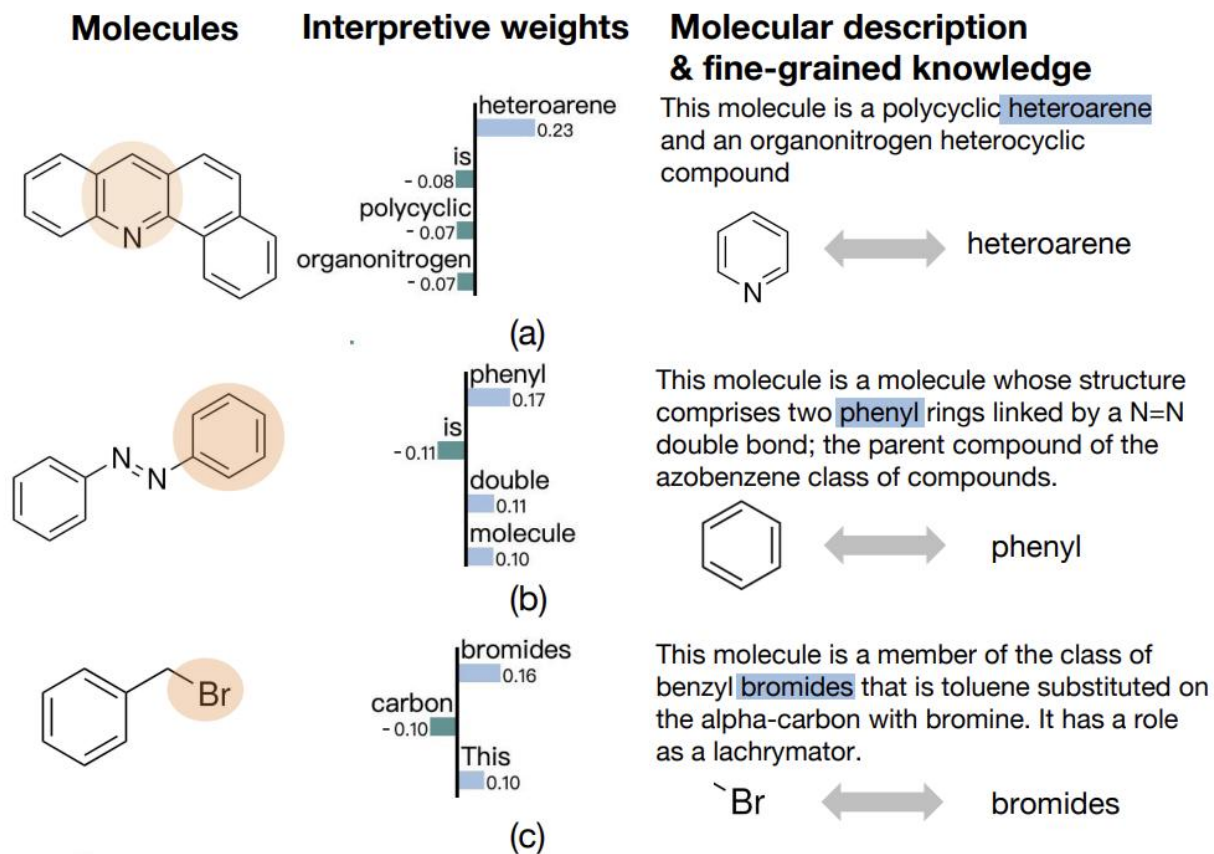


Figure 2: Explanation of the prediction of certain masked motifs based on text tokens utilizing LIME.

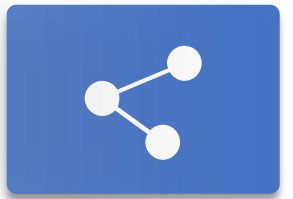
CONTENTS

- 1 Background
- 2 Fine-grained Alignment
- 3 FineMolTex
- 4 Experiments
- 5 **Conclusion**





- We reveal that fine-grained motif-level knowledge is crucial for molecular representation learning.
- We propose FineMolTex to jointly learn both coarse- and fine-grained knowledge through a contrastive alignment task and a masked multimodal learning task, respectively.
- By selectively masking the important motif/word tokens and predicting their labels using tokens from the other modality, we can effectively learn fine-grained alignment between motifs and words.
- Experimental results on three downstream tasks and two case studies demonstrate the effectiveness of FineMolTex.



Thank you !

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