



# GraphGPT: Generative Pre-trained Graph Eulerian Transformer Alibaba Group

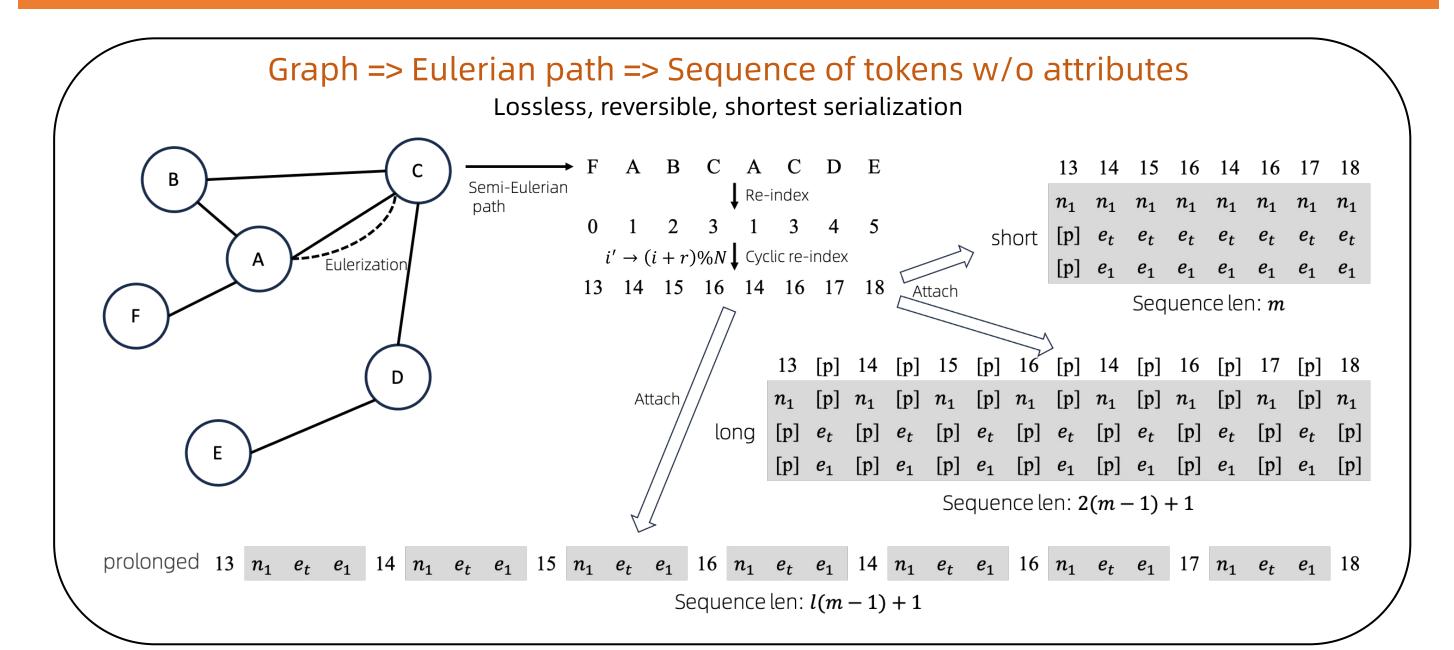


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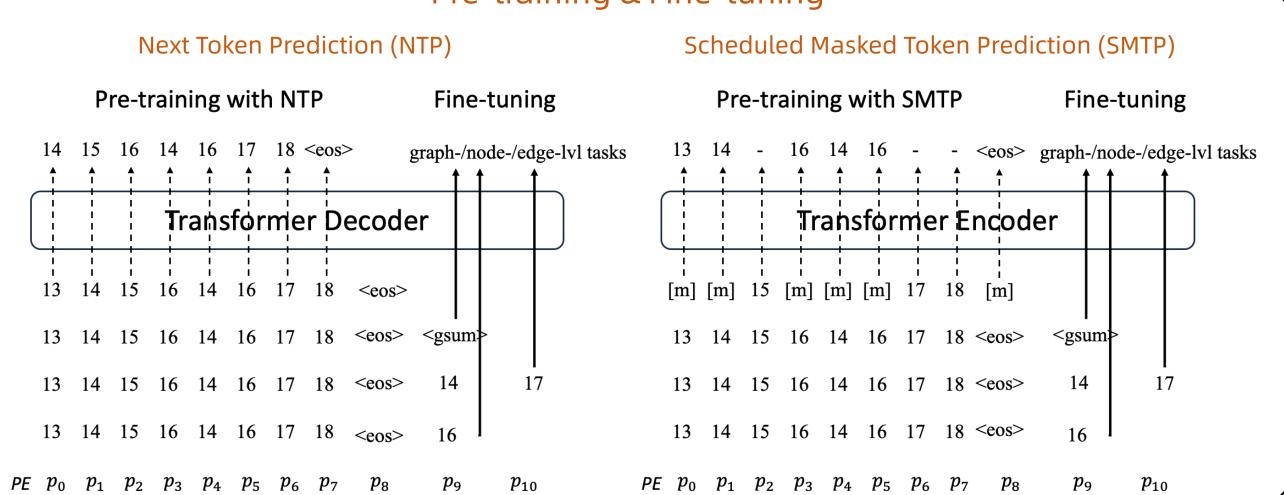
## Motivation

- Graph has not benefited from the transformer architecture, like NLP/CV/Audio
- Unifying graph with other modalities is problematic due to inconsistent architecture
- Graph has not benefited from scaling up model sizes

## Method



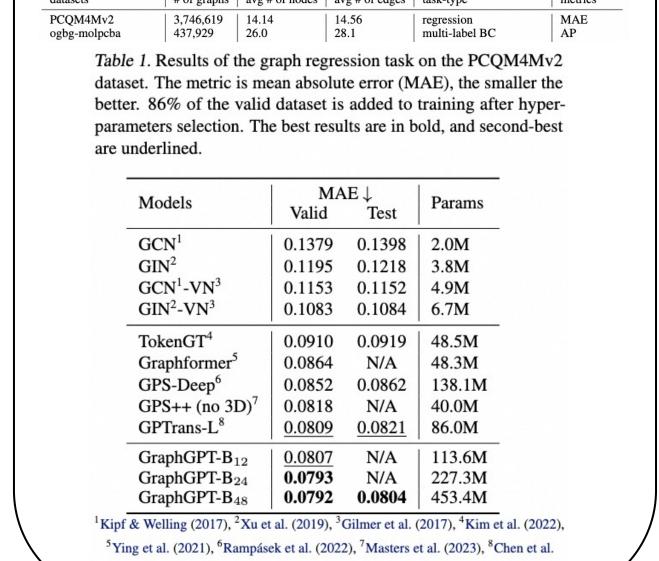
#### Pre-training & Fine-tuning



## Graph-level Task

- PCQM4Mv2 contains > 3.7 million organic molecules from PubChemQC (Nakata & Shimazaki, 2017). Nodes represent atoms (9D attributes: atomic number, chirality, etc.), and edges denote chemical bonds (3D attributes: bond type, stereochemistry, conjugation).
- ogbg-molpcba is a smaller molecular dataset (Wu et al., 2017) with the same node/edge attributes.

#### PCQM4M-v2: 3.7M molecules

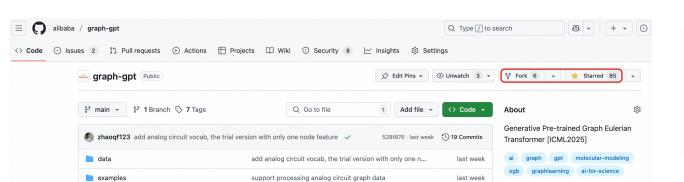


#### OGBG-MOLPCBA: 438K molecules

Table 2. Results of the graph classification task on the ogbgmolpcba dataset. All the baseline results are from the OGB leaderboard or the corresponding papers. † indicates the model is pretrained on PCQM4M-v2 dataset.

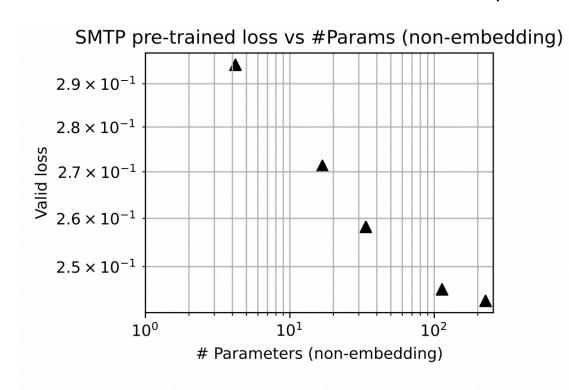
Models	Average Pred Test	cision (%) ↑ Valid	Params
$GCN^1$	$20.20_{\pm 0.24}$	$20.59_{\pm 0.33}$	0.57M
$GIN^2$	$22.66_{\pm0.28}$	$23.05 {\scriptstyle \pm 0.27}$	1.92M
$GINE^3$ - $VN^4$	$29.17_{\pm 0.15}$	$30.65_{\pm0.30}$	6.1M
NGIN <sup>5</sup> -VN <sup>4</sup>	$30.07_{\pm 0.37}$	$30.59_{\pm 0.56}$	44.19M
$PDF^6$	$30.31_{\pm 0.26}$	$31.15 {\scriptstyle \pm 0.20}$	3.84M
Graphormer-L <sup>†7</sup>	$31.40_{\pm 0.32}$	$32.27_{\pm0.24}$	119.5M
EGT-Larger <sup>†8</sup>	$29.61_{\pm0.24}$	N/A	110.8M
GRPE-Large <sup>†9</sup>	$31.50_{\pm0.10}$	N/A	118.3M
GPTrans-L <sup>†10</sup>	$ullet{32.43_{\pm 0.22}}$	N/A	86.0M
GraphGPT-M <sup>†</sup>	$30.13_{\pm 0.25}$	$31.62_{\pm0.24}$	37.7M
GraphGPT- $\mathrm{B}_{12}^{\dagger}$	$31.28_{\pm0.23}$	$32.27_{\pm 0.15}$	113.6M
GraphGPT- $B_{24}^{\dagger}$	$31.81_{\pm0.1}$	$\overline{32.54_{\pm 0.2}}$	227.3M
1	2		1

- <sup>1</sup>Kipf & Welling (2017), <sup>2</sup>Xu et al. (2019), <sup>3</sup>Brossard et al. (2020), <sup>4</sup>Gilmer et al. (2017), <sup>5</sup>Zhang & Li (2021), <sup>6</sup>Yang et al. (2023), <sup>7</sup>Ying et al. (2021), <sup>8</sup>Hussain et al. (2022), <sup>9</sup>Park et al. (2022), <sup>10</sup>Chen et al. (2023b)
- *SOTA*: On PCQM4Mv2, GraphGPT achieves a test MAE of **0.0804**, significantly outperforming the previous SOTA (*0.0821*, GPTrans)
- *vs GTs*: GTs like TokenGT, Graphformer, GPS, and GPTrans requires handcrafted features or intricate architectures to encode structural information, while GraphGPT attains superior performance without manual feature engineering.
- vs GNNs: GraphGPT surpasses GNNs by a substantial margin.
- Parameter Efficiency. GraphGPT's larger parameter count may reflect its capacity to implicitly learn features that other GTs encode manually. Generative pre-training also allocates model capacity to generation tasks, potentially limiting discriminative performance of models at smaller scales.



Generative Pre-trained Graph Eulerian Transformer by Q Zhao · 2023 · Cited by 16— We introduceGraphGPT, a novel self-supervised generative pretrained model for graph learning based on the Graph Eulerian Transformer (GET).

#### Scaling up model sizes consistent improvement up to 200M parameters



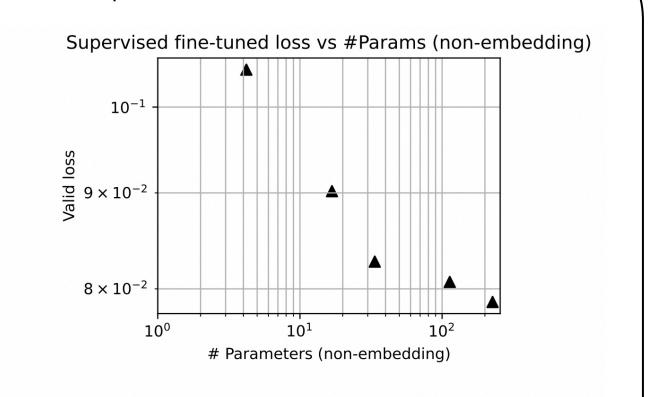


Figure 3. Log-log plot of pre-training loss and supervised fine-tuning loss versus the number of non-embedding parameters for the Mini/Small/Medium/Base/Base24 model configurations (see Table 11) on the PCQM4M-v2 dataset.

#### Graph Structure Understanding (GSU)

ataset stats: totall	y 45000 graph	ns		
	train	valid	test-small	test-large
indices	[0, 30000)	[30000, 35000)	[35000, 40000)	[40000, 45000)
connected graphs	63.13%	62.78%	62.76%	13.78%
avg # node	15.58	15.54%	15.61	63.08
min # node	4	4	4	26
max # node	25	25	25	100

- Pre-training (PT) is highly beneficial: 32% → 92%
- PT on other datasets also improves GSU on this dataset, sometimes even better: a vs b/c/d
  - This holds true even when PT includes node and
- edge attribute prediction: a vs e/f/g• PT on real graphs outperforms random graphs: c vs d, (a+b+c) vs (a+b+d)
- More data & diverse PT enhance generalization: a vs

(a+b)/(a+c)/(b+c) vs (a+b+c)/(a+b+d)

#### Graph triangle counting: 1~10 (10-class classification)

Models	Accuracy (%) ↑ T-small T-large		Params
	1-Sman	T-large	
$GIN^1$	$71.53_{\pm 0.94}$	$33.54_{\pm0.30}$	0.15M
Transformer <sup>2</sup>	$12.08_{\pm0.31}$	$10.01_{\pm 0.04}$	0.2M
Transformer-LapPE <sup>3</sup>	$78.29_{\pm0.25}$	$10.64_{\pm 2.94}$	0.2M
Transformer-RWSE <sup>3</sup>	$99.40_{\pm0.10}$	$54.76_{\pm 7.24}$	0.2M
Graphormer <sup>4</sup>	$99.09_{\pm0.31}$	$\overline{42.34_{\pm 6.48}}$	0.2M
GET-B	32.60 <sub>±1.86</sub>	$13.99_{\pm 1.78}$	113.5M
GraphGPT-B <sup>a</sup>	$92.16_{\pm0.28}$	$26.51_{\pm 1.01}$	113.5M
GraphGPT-B <sup>b</sup>	$81.38_{\pm0.27}$	$37.68_{\pm0.99}$	113.5M
GraphGPT-B <sup>c</sup>	$99.08_{\pm0.14}$	$38.80_{\pm 3.60}$	113.5M
GraphGPT-B <sup>d</sup>	$90.93_{\pm 0.51}$	$40.79_{\pm 1.40}$	113.5M
GraphGPT-B <sup>e</sup>	$64.28_{\pm0.33}$	$17.38_{\pm0.61}$	113.5M
GraphGPT-B <sup>f</sup>	$86.14_{\pm 7.38}$	$26.94_{\pm 4.80}$	113.5M
GraphGPT-B <sup>g</sup>	$86.57_{\pm 2.74}$	$23.45_{\pm 1.44}$	113.5M
GraphGPT-B <sup>a+b</sup> ✓	84.83 <sub>±0.81</sub>	$39.62_{\pm 1.84}$	113.5M
GraphGPT-B <sup>a+c</sup>	$98.68_{\pm0.18}$	$50.07_{\pm 3.28}$	113.5M
GraphGPT-B <sup>b+c</sup>	$98.26_{\pm0.30}$	$52.33_{\pm 2.61}$	113.5M
GraphGPT-B <sup>a+b+d</sup> <	$89.98_{\pm0.54}$	$33.45_{\pm 2.51}$	113.5M
GraphGPT-Ma+b+c	$95.07_{\pm 0.67}$	$51.72_{\pm 1.12}$	33.7M
GraphGPT-B <sup>a+b+c</sup> <	$98.63_{\pm 0.18}$	$58.96_{\pm 1.90}$	113.5M

Pre-trained with: <sup>a</sup>Triangles (45K), <sup>b</sup>Reddit-threads (0.22M), <sup>c</sup>Internal dataset (3.1M), <sup>d</sup>Random graphs (3.1M), <sup>e</sup>PCQM4M-v2 (3.7M), <sup>f</sup>ogbl-ppa (1), gogbn-proteins (1).

The Hits@100 score on the test and validation sets. The higher, the better

## Edge-level Task

Leaderboard for ogbl-ppa

Package: >=1.1.1

datasets	# of graphs	avg # of nodes	avg # of edges	task-type	metrics
PCQM4Mv2	3,746,619	14.14	14.56	regression	MAE
ogbg-molpcba	437,929	26.0	28.1	multi-label BC	AP
reddit-threads	203,088	23.9	24.9	BC	ROC-AUC
Triangles	45,000	20.9	32.7	multi-class classification	ACC
Internal dataset	3,100,000	24.8	54.7	N/A	N/A
Random Graph <sub><math>p=0.05</math></sub>	3,100,000	67.0	124.7	N/A	N/A
Random Graph $_{p=0.03}$	3,100,000	67.1	74.8	N/A	N/A
Random Graph $_{p=0.01}$	3,100,000	67.1	25.0	N/A	N/A
ogbl-ppa	1	576,289	30,326,273	BC	HR@100
ogbl-citation2	1	2,927,963	30,561,187	BC	MRR
ogbn-proteins	1	132,534	39,561,252	multi-label BC	ROC-AUC
ogbn-arxiv	1	169,343	1,166,243	multi-class classification	ACC

Models	ogbl-ppa HR@100 (%)↑	ogbl-citation2 MRR (%) ↑
Common Neighbor Adamic Adar	$27.65_{\pm 0.00}$ $32.45_{\pm 0.00}$	$\begin{array}{ c c c c c c }\hline 51.47_{\pm 0.00} \\ 51.89_{\pm 0.00} \\ \hline \end{array}$
Resource Allocation <sup>1</sup>	$49.33_{\pm 0.00}$	$51.89\pm0.00$ $51.98\pm0.00$
Node2Vec <sup>2</sup>	$22.26_{\pm0.83}$	$61.41_{\pm 0.11}$
Matrix Factorization <sup>3</sup>	$32.29_{\pm 0.94}$	$51.86_{\pm 4.43}$
GCN <sup>4</sup>	$18.67_{\pm 1.32}$	84.74 <sub>±0.21</sub>
GraphSAGE <sup>5</sup>	$16.55_{\pm 2.40}$	$82.60_{\pm0.36}$
SEAL <sup>6</sup>	$48.80_{\pm 3.16}$	$87.67_{\pm0.32}$
AGDN <sup>7</sup>	$41.23_{\pm 1.59}$	$85.49_{\pm0.29}$
SIEG <sup>8</sup>	$63.22_{\pm 1.74}$	$90.18_{\pm0.15}$
MPLP <sup>9</sup>	$65.24_{\pm 1.50}$	$90.72_{\pm0.12}$
RefinedGAE <sup>10</sup>	$73.74_{\pm 0.92}$	$84.55_{\pm0.15}$
GraphGPT-M	$65.44_{\pm0.43}$	$92.82_{\pm 0.27}$
GraphGPT-B	$68.76_{\pm0.67}$	$\overline{93.05_{\pm 0.20}}$
GraphGPT-XXL	$\bf 76.55_{\pm 0.67}$	N/A

Table 4. Results of the link prediction task on the ogbl-ppa and

ogbl-citation2 datasets.

Leaderboard for ogbl-citation2 The MRR score on the test and validation sets. The higher, the better. Package: >=1.2.4 <sup>1</sup>Zhou et al. (2009), <sup>2</sup>Grover & Leskovec (2016), <sup>3</sup>Mnih & Salakhutdinov (2008). <sup>4</sup>Kipf & Welling (2017), <sup>5</sup>Hamilton et al. (2017), <sup>6</sup>Zhang et al. (2021), <sup>7</sup>Sun et al (2020), <sup>8</sup>Shi et al. (2024), <sup>9</sup>Dong et al. (2023), <sup>10</sup>Ma et al. (2024)

- Performance Superiority: GraphGPT significantly outperforms all baseline methods, including GNNs, heuristic models, and latent-factor approaches, across both datasets.
- Scalability. GraphGPT scales seamlessly to 2 billion parameters, achieving sustained performance gains with increasing model size.
- Transformer Efficacy. To our knowledge, GraphGPT is the first transformer-based model to achieve SOTA results on ogbl-ppa and ogbl-citation2, demonstrating the viability of sequence-driven architectures for large-scale edge-level tasks.

## Node-level Task

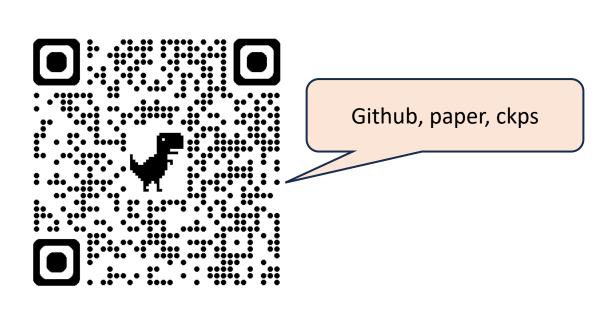
#### Table 5. Results of the node classification task on the ogbn-proteins and ogbn-arxiv datasets.

Models	ogbn-proteins ROC-AUC (%) ↑	ogbn-arxiv Accuracy (%) ↑
GCN <sup>1,2</sup>	$77.29_{\pm0.46}$	$73.53_{\pm0.12}$
GraphSAGE <sup>1,3</sup>	$82.21_{\pm 0.32}$	$73.00_{\pm0.28}$
GAT <sup>1,4</sup>	$85.01_{\pm 0.46}$	$73.30_{\pm0.18}$
DRGAT <sup>5</sup>	N/A	$74.16 _{\pm 0.07}$
AGDN <sup>6</sup>	$88.65_{\pm0.13}$	$73.41_{\pm 0.25}$
DeeperGCN <sup>7</sup>	$85.80_{\pm0.17}$	$71.92_{\pm 0.16}$
GraphGPS <sup>1,8</sup>	$77.15_{\pm 0.64}$	$71.23_{\pm 0.59}$
NAGphormer <sup>1,9</sup>	$72.17 {\scriptstyle \pm 0.45}$	$70.88_{\pm0.24}$
Exphormer <sup>1,10</sup>	$77.62_{\pm0.33}$	$72.32_{\pm0.36}$
GOAT <sup>1,11</sup>	$79.31_{\pm 0.42}$	$72.76_{\pm0.29}$
NodeFormer <sup>1,12</sup>	$77.86_{\pm0.84}$	$67.78_{\pm0.28}$
SGFormer <sup>1,13</sup>	$79.92_{\pm0.48}$	$72.76_{\pm0.33}$
Polynormer <sup>1,14</sup>	$79.53_{\pm 0.67}$	$73.40_{\pm0.22}$
GraphGPT-S	$83.4_{\pm 0.00}$	$71.2_{\pm 0.00}$
GraphGPT-M	$84.3_{\pm 0.00}$	$71.8_{\pm 0.00}$
GraphGPT-B	$85.5_{\pm 0.00}$	$72.2_{\pm 0.00}$

(2017), <sup>5</sup>Zhang et al. (2023), <sup>6</sup>Sun et al. (2020), <sup>7</sup>Li et al. (2020), <sup>8</sup>Rampásek et al. (2022), 9Chen et al. (2023a), 10Shirzad et al. (2023), 11Kong et al. (2023), 12Wu et al. (2022), <sup>13</sup>Wu et al. (2024), <sup>14</sup>Deng et al. (2024)

#### • Ogbn-proteins: Undirected, weighted graph of 132,534 proteins (nodes) with 8D edge attributes encoding association strengths.

• Ogbn-arxiv: Citation network of 169,343 papers; tasks involve predicting 40 subject categories.



- GraphGPT outperforms/matches classic GNNs. But still lags behind some customized GNN variants.
- It significantly improves or equals GTs.

#### Various Model Sizes

Model-size	Hidden-size	# of layers	# of heads	Params (excluding e
Mini	256	4	4	4.2M
S (Small)	512	4	8	16.8M
M (Medium)	512	8	8	33.6M
B / B <sub>12</sub> (Base)	768	12	12	113.2M
B <sub>24</sub> (Base24)	768	24	12	226.5M
B <sub>48</sub> (Base48)	768	48	12	453.0M
L (Large)	1024	24	16	402.7M
XXL (XXLarge)	1600	48	25	2.0B

Statistics of GraphGPT models of different sizes. The GraphGPT-Base is of the same scale as Bert-Base (Devlin et al., 2019).

## Ablation

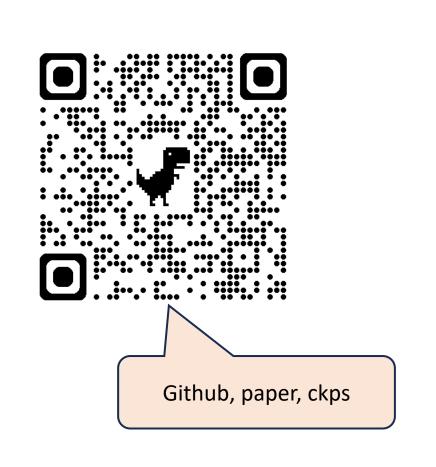
#### Pre-training

Table 6. Ablation study of pre-training on the datasets of various types of tasks. \* means both molpcba and PCQM4Mv2 datasets are used for SMTP pre-training, and † indicates that the model is further trained using PCQM4M-v2's regression task. For the PCQM4Mv2 dataset, the metric is MAE, the lower the better.

DATASETS	PRE-TRAINING	TEST	VALID
	X	N/A	0.1086
PCQM4Mv2	NTP	N/A	0.0875
	SMTP	N/A	0.086
	×	12.8	13.31
	NTP	23.85	27.77
OGBG-MOLPCBA	SMTP	27.56	28.74
	SMTP*	27.2	28.49
	SMTP* + FT <sup>†</sup>	28.07	29.01
	×	41.28	40.14
OGBL-PPA	NTP	55.56	54.87
	SMTP	55.68	54.93
	×	57.52	61.19
OGBN-PROTEINS	NTP	75.61	80.47
	SMTP	83.02	86.41

## • Pre-training brings substantial improvements.

- SMTP > NTP in most cases.
- Strong in-domain transferability.



#### Node-reindex

Table 7. Ablation study of node re-indexing on the ogbg-molpcba dataset with two model sizes. PT means pre-training with NTP.

PARAMS	RE-INDEX	PT Loss	TEST	VALID
4.48M	×	0.0844	0.2310	0.2525
7.70171	/	0.0874	0.2385	0.2777
114.12M	X	0.0689	0.2270	0.2621
114.12IVI	/	0.0750	0.2517	0.2857

## Node-Identity Coding

Table 8. Ablation study of node identity encoding on the ogbl-ppa and ogbn-proteins datasets using NTP pre-training. NIE stands for Node identity encoding.

DATASETS	PARAMS	NIE	TEST	VALID
OGBL-PPA	14.75M	×	44.38 <b>55.56</b>	45.08 <b>54.87</b>
OGBN-PROTEINS	10.76M	×	60.22 <b>75.61</b>	65.66 <b>80.47</b>

## Limitations

- Model size is large, high computational resource is required
- A larger graph dataset is required to demonstrate superiority.
- Transferability: Pre-training is currently limited to same-domain datasets, making generalization to other graph data domains challenging.
- The transferability of graph structure understanding is evident.

## Outlook

- General Graph Structure Understanding [Graph] Foundation Model (GFM)
- Specialized Domain Understanding GFM (e.g., molecule)
- Combined with LLM, similar to Llava

## Synergy with diffusion LLM (dLLM)

- Speed and Performance: dLLM has shown superior generation speed and comparable performance compared to AR (autoregressive) LLM: Mercury and Gemini Diffusion.
- Same Pre-training: Masked dLLM like <u>LLaDa-8B</u>, <u>Dream-7B</u> share almost the same pre-training objectives as SMTP employed by GraphGPT. • SMTP > NTP: GraphGPT shows dLLM-like pre-training SMTP is much better than AR-like pre-training NTP in most graph datasets.
- Multi-modality: GraphGPT processes graph data in a way closely aligned with dLLM: using sequential tokens, a transformer encoder, and a masked token prediction objective. It implies graph data can be naturally incorporated in the dLLM.
- Al for Science: Some scientific data is naturally represented as graphs—for example, molecules and integrated circuits. Other scientific data, such as proteins and DNA/RNA, is represented as sequences. Unlike language, these data types lack the autoregressive (AR) inductive bias, making them better suited for modeling with dLLM.

### References

[1] Samar Khanna, Siddhant Kharbanda, Shufan Li, Harshit Varma, Eric Wang, Sawyer Birnbaum, Ziyang Luo, Yanis Miraoui, Akash Palrecha, Stefano Ermon, Aditya Grover, Volodymyr Kuleshov. Mercury: Ultra-Fast Language Models Based on Diffusion. arXiv preprint arXiv:2506.17298, 2025.

[2] Google DeepMind, Gemini Diffusion. url: <a href="https://deepmind.google/models/gemini-diffusion/">https://deepmind.google/models/gemini-diffusion/</a>.

[3] Weihua Hu, Matthias Fey, Marinka Zitnik, Yuxiao Dong, Hongyu Ren, Bowen Liu, Michele Catasta, Jure Leskovec. Open Graph Benchmark: Datasets for Machine Learning on Graphs. arXiv preprint arXiv: 2005.00687, 2020.

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[5] Jiacheng Ye, Zhihui Xie, Lin Zheng, Jiahui Gao, Zirui Wu, Xin Jiang, Zhenguo Li, and Lingpeng Kong. Dream 7B. url: https://hkunlp.github.io/blog/2025/dream/