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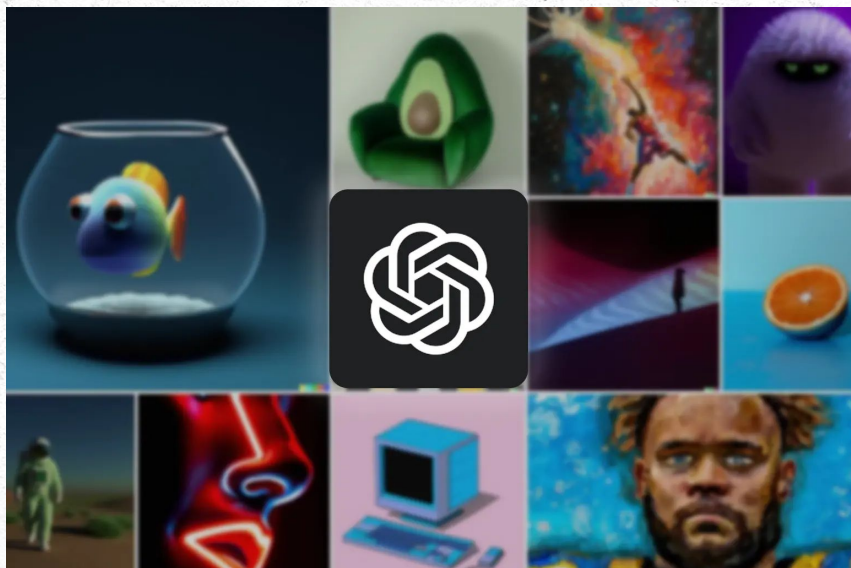
# Latent 3D Graph Diffusion

Yuning You<sup>1</sup>, Ruida Zhou<sup>2</sup>, Jiwoong Park<sup>1</sup>, Haotian Xu<sup>1</sup>,  
Chao Tian<sup>1</sup>, Zhangyang Wang<sup>3</sup>, Yang Shen<sup>1</sup>

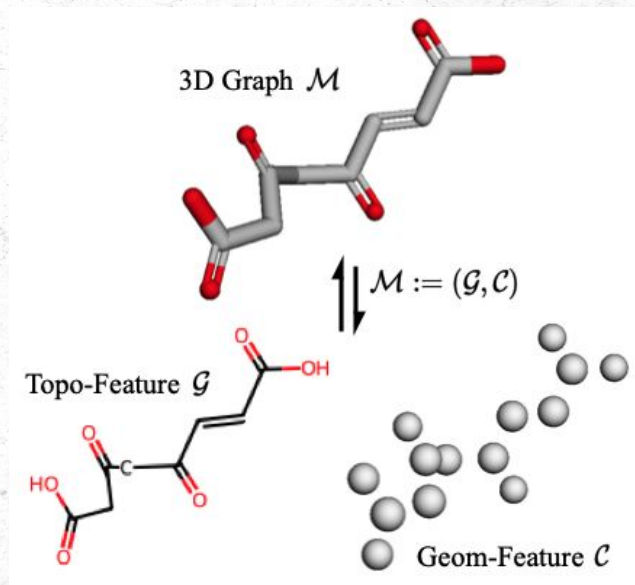
<sup>1</sup>Texas A&M University, <sup>2</sup>University of California, Los Angeles, <sup>3</sup>University of Texas at Austin

# Background

## Generative AI on Images



## Generative AI on 3D Graphs



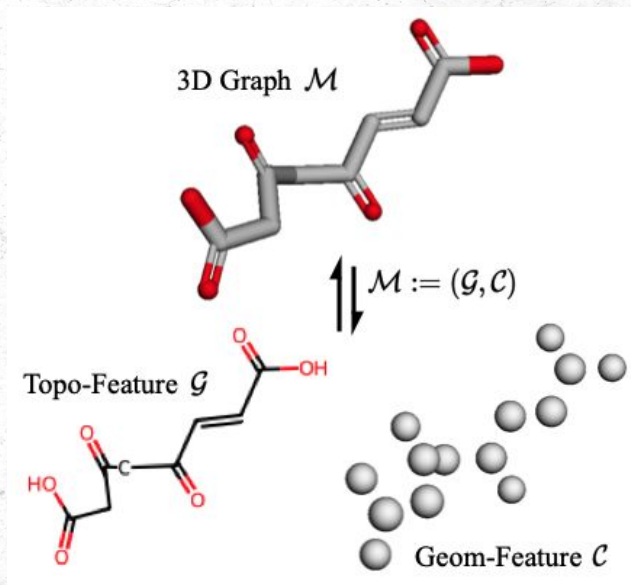


# Central Question

In what (latent) space should we learn the distribution of graphs with neural networks?

- Original space: Topology $\otimes$ Geometry;
- Symmetry constraint: Data identities are invariant to certain transformations;
- High dimensionality: Data dimensionality depends on graph size;
- Low-dimensional manifold: Data are supposed to distribute in the LD manifold.

Generative AI on 3D Graphs



# Answer: What justifies a “good” space for graph generative models?

- We focus on diffusion generative model.
- Assuming there are forward and reverse mappings between the original 3D graph space and a latent space;
- The diffusion model is trained on the latent space to capture distribution, then:

$$\boxed{\text{3D Graph Diffusion Performance}} \leq \boxed{\text{Latent Space Reconstruction Quality}} + \boxed{\text{Symmetry Preservation}} \times \boxed{\text{Data Dimensionality}}.$$

**Proposition 2.** (3D graph diffusion could benefit from the **lower-dimensional latent space if appropriately constructed**. See proof in Append. A.2) Assume there existing mappings  $\vec{h} : \mathbb{R}^{D'} \rightarrow \mathbb{R}^{D''}$ ,  $\overleftarrow{h} : \mathbb{R}^{D''} \rightarrow \mathbb{R}^{D'}$  that  $D'' < D'$  and  $\overleftarrow{h}$  is injective. Assume DGM now is trained in  $\mathbb{R}^{D''}$  to model  $\vec{p}_{\text{data}}(\mathbf{z}) = \Pr\{\mathbf{x}_M : \vec{h}(\mathbf{x}_M) = \mathbf{z}, \mathbf{x}_M \sim p_{\text{data}}\}$  with  $p_{\theta}(\mathbf{z})$ , and it is evaluated in  $\mathbb{R}^{D'}$  on  $\overleftarrow{p}_{\theta}([\mathbf{x}_M]_{\Pi, \Omega}) = \Pr\{\mathbf{z} : \overleftarrow{h}(\mathbf{z}) \in [\mathbf{x}_M]_{\Pi, \Omega}, \mathbf{z} \sim p_{\theta}\}$  (as in Propos. 1), and the assumptions in Propos. 1 retain for the score estimator  $f_{\theta}$  and mapping distribution. Then, it holds:

$$\boxed{\text{TV}(\overleftarrow{p}_{\theta}, \tilde{p}_{\text{data}})} \lesssim \boxed{\text{TV}(\overleftarrow{p}_{\text{data}}, \tilde{p}_{\text{data}})} + \boxed{\bar{\alpha}(p_{\theta}, \vec{h}, \overleftarrow{h}, \Pi, \Omega) \left( \sqrt{\text{KL}(\vec{p}_{\text{data}} \| \mathcal{N}_{D''})} e^{-T} + (L\sqrt{D''} + Lm + \varepsilon_{\text{score}})\sqrt{T} \right)}, \quad (3)$$

where  $\overleftarrow{p}_{\text{data}}([\mathbf{x}_M]_{\Pi, \Omega}) = \Pr\{\mathbf{x}'_M : \overleftarrow{h}(\vec{h}(\mathbf{x}'_M)) \in [\mathbf{x}_M]_{\Pi, \Omega}, \mathbf{x}'_M \sim p_{\text{data}}\}$ , and  $\bar{\alpha}(\cdot)$  depends on both the latent diffusion architecture that  $\bar{\alpha}(p_{\theta}, \vec{h}, \overleftarrow{h}, \Pi, \Omega) = \alpha(\overleftarrow{p}_{\theta}, \Pi, \Omega)$  if  $\overleftarrow{p}_{\text{data}} = p_{\text{data}}$ .  $\square$

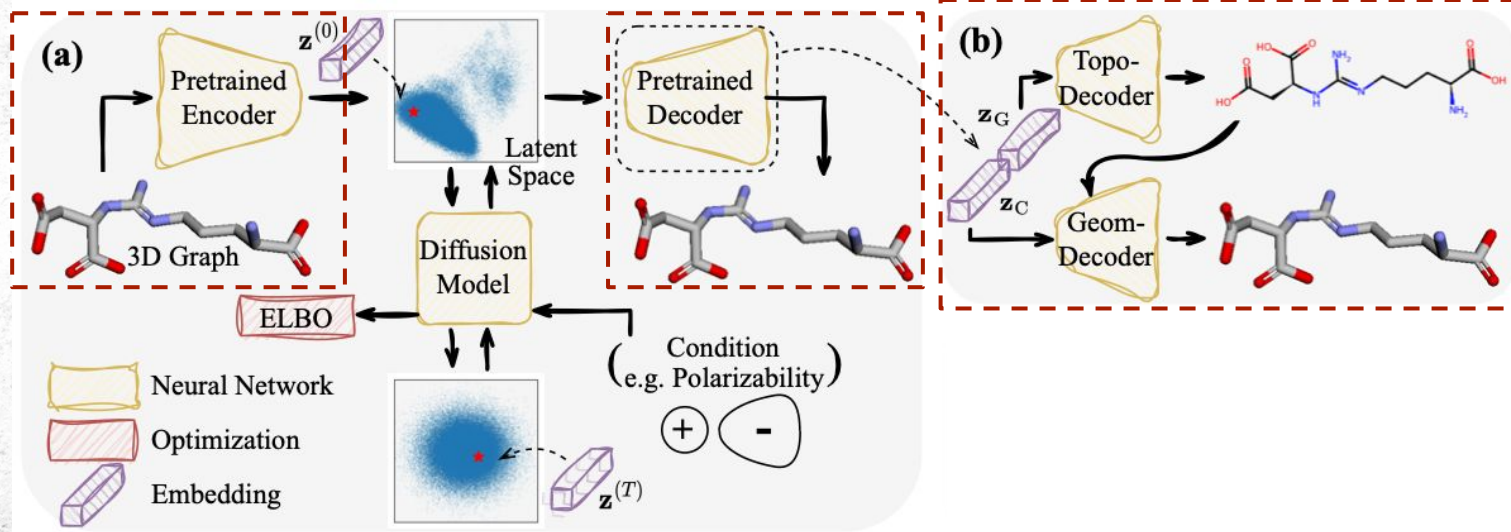


# Answer: How to construct a qualified latent space for graph diffusion?

- We develop the framework termed latent 3D graph diffusion.
- Cascaded auto-encoder for 3D graphs:

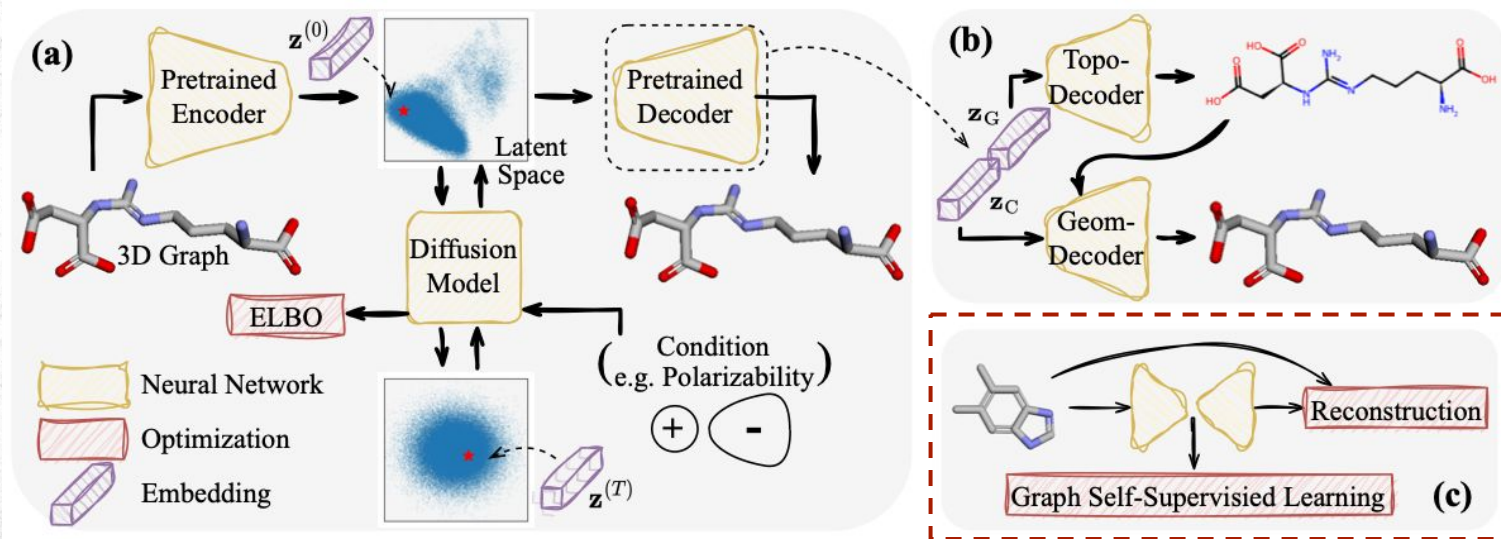
$$\text{Encoding: } \mathbf{z}_G = \vec{h}_{\phi_{1,G}}(\mathcal{G}), \quad \mathbf{z}_C = \vec{h}_{\phi_{1,C}}(\mathcal{C}), \quad \mathbf{z} = [\mathbf{z}_G; \mathbf{z}_C],$$

$$\text{Decoding: } \bar{\mathcal{G}} = \overleftarrow{h}_{\phi_{2,G}}(\mathbf{z}_G), \quad \bar{\mathcal{C}} = \overleftarrow{h}_{\phi_{2,C}}(\bar{\mathcal{G}}, \mathbf{z}_C),$$



# Answer: How to regularize the latent space to introduce domain prior?

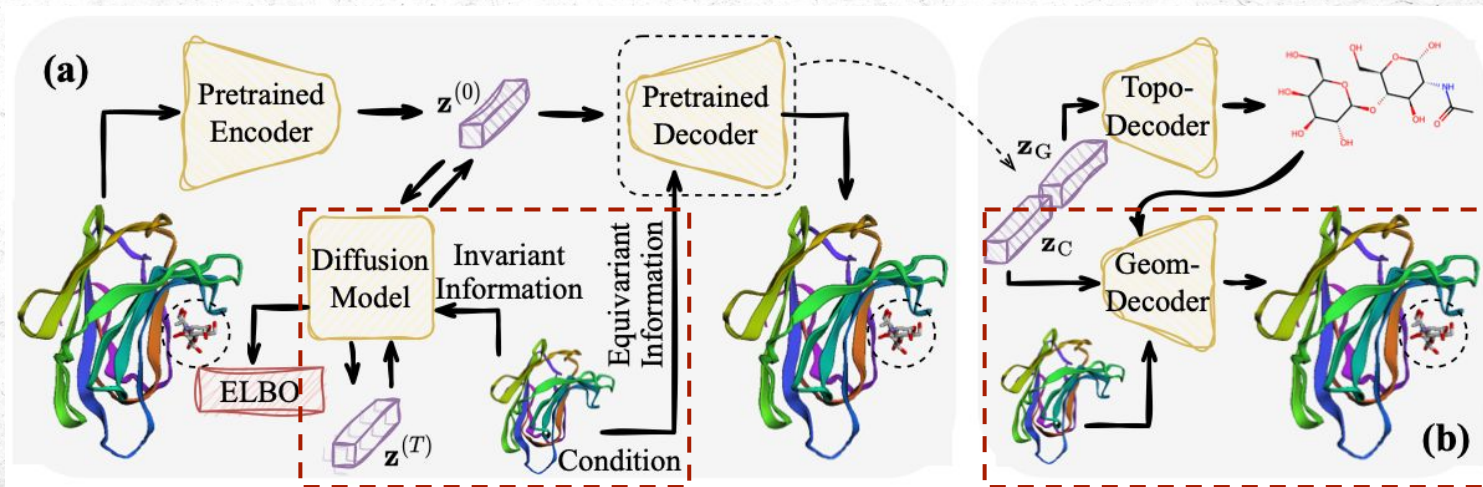
- We develop the framework termed latent 3D graph diffusion.
- Graph self-supervised learning regularized auto-encoding:
  - Graph contrastive learning (GraphCL, NeurIPS'20).





# Answer: How to extend the framework to conditional generation?

- Equivariance constraint: When condition is a geometric object.
- 1. In/Equi-variant representations for condition inputs;
- 2. Invariant distribution modeling of latent embeddings;
- 3. Equivariant decoding to reconstruct 3D graphs.



# Experiments on Unconditional Generation

**Table 2:** Unconditional generation evaluation on validness of 3D molecules. Valid: proportion of (POF) chemically valid molecules; Valid&Uni: POF chemically valid and unique molecules; AtomSta: POF atoms with correct valency; MolSta: POF molecules without unstable atoms. Numbers(std) in **red** are the best results.

Methods	QM9				Drugs		Mean
	Valid	Valid&Uni	AtomSta	MolSta	Valid	AtomSta	
ENF	40.2	39.4	85.0	4.9	–	–	42.37
G-Schnet	85.5	80.3	95.7	68.1	–	–	82.40
GDM	–	–	97.0	63.2	90.8	75.0	81.50
GDM-Aug	90.4	89.5	97.6	71.6	91.8	77.7	86.43
EDM	91.9(0.5)	90.7(0.6)	98.7(0.1)	82.0(0.4)	92.6	81.3	89.53
EDM-Bridge	92.0	90.7	98.8(0.1)	84.6(0.3)	92.8	82.4(0.8)	90.21
GCDM	94.8(0.2)	93.3(0.0)	98.7(0.0)	85.7(0.4)	–	<b>89.0(0.8)</b>	92.30
MiDi	97.9	97.0	97.9	84.0	78.0	82.2	89.50
GraphLDM	83.6	82.7	<b>97.2</b>	70.5	97.2	76.2	84.56
GraphLDM-Aug	90.5	89.5	97.9	78.7	98.0	79.6	89.03
GeoLDM	93.8(0.4)	92.7(0.5)	<b>98.9(0.1)</b>	<b>89.4(0.5)</b>	99.3	84.4	93.08
Ours	<b>100.00(0.00)</b>	95.27(0.25)	97.57(0.02)	86.87(0.23)	<b>100.00(0.00)</b>	80.51(0.08)	<b>93.37</b>



# Experiments on Conditional Generation

**Table 6:** Conditional generation on protein binding targets evaluation. Assessment metrics QED/SA & Vina scores are calculated with RDKit (Landrum, 2013) & AutoDock (Huey et al., 2012), respectively.

Methods	QED $\uparrow$	SA $\uparrow$	HiAff $\uparrow$	Vina $\downarrow$	VDock $\downarrow$	Vina (Top-10%) $\downarrow$	Diversity $\uparrow$
LiGAN	0.39	0.59	21.1%	–	-6.33	–	0.66
GraphBP	0.43	0.49	14.2%	–	-4.80	-7.16	0.79
AR	0.51	0.63	37.9%	-5.75	-6.75	–	0.70
Pocket2Mol	0.56	<b>0.74</b>	48.4%	-5.14	-7.15	-8.71	0.69
TargetDiff	0.48	0.58	<b>58.1%</b>	-5.47	-7.80	-9.66	0.72
DiffSBDD	0.46	0.55	–	<b>-7.33</b>	–	-9.92	0.75
DecompDiff	0.45	0.61	64.4%	-5.67	<b>-8.39</b>	–	0.68
Ours	<b>0.60</b>	0.71	48.08%	-5.23	-6.85	<b>-12.34</b>	<b>0.80</b>



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# Thank You!

<https://yyou1996.github.io/>