



Triplet Interaction Improves Graph Transformers

Accurate Molecular Graph Learning with Triplet Graph Transformers

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and

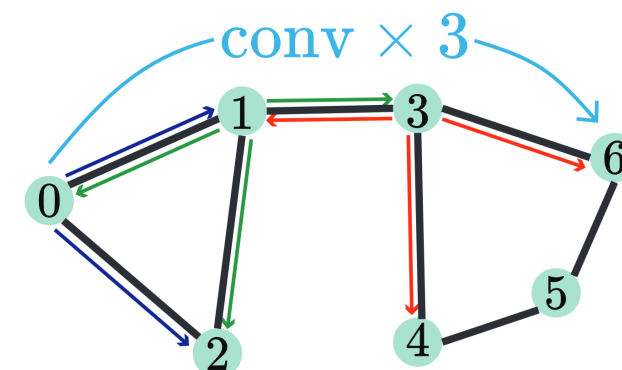
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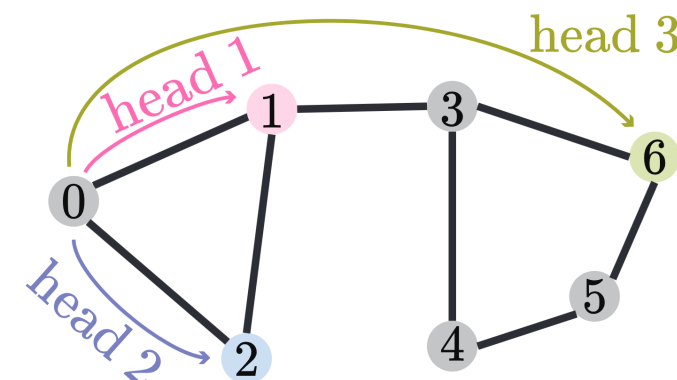


Graph Transformers vs. GCNs

- Long-distance, dynamic interaction
 - Not limited to neighbors
 - Attention weights are determined by the network
- Limitations of hand-crafted encodings/features
 - Positional-encoding based GTs (e.g., Graphormer)
 - Structural understanding \approx as good as the used positional encoding
 - Geometric GTs (e.g., Equiformer)
 - Geometric understanding \approx as good as the used geometric features
- **Goal: let the network form its own geometry / structure**



(a) Convolution



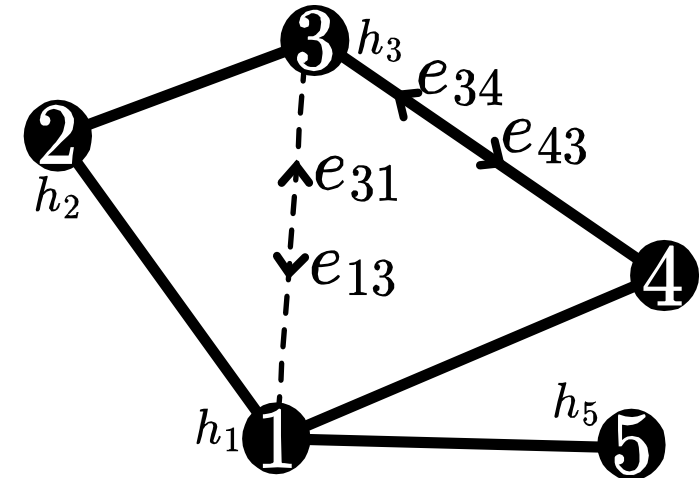
(b) Self-attention





Graph Structure and Pair Representations

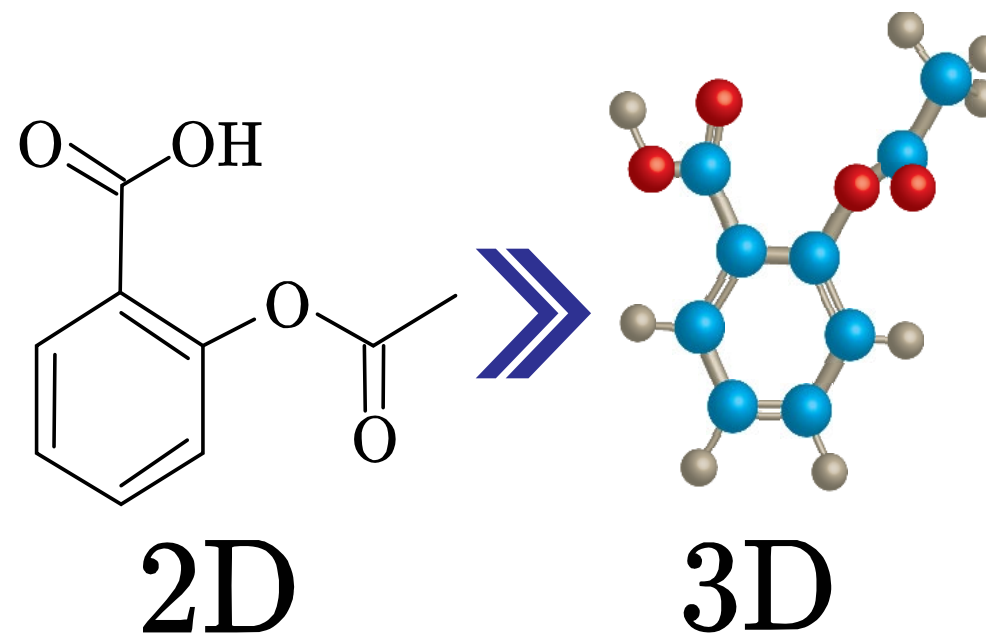
- Pairs \rightarrow directed existing/non-existing edges
 - e.g., $3 \rightarrow 4$, $4 \rightarrow 3$, $3 \rightarrow 1$, $1 \rightarrow 3$
- For graphs pair representations (e_{ij}) can be as important as node representations (h_i)
 - Allow the structure of the graph to evolve over layers
 - Refine structure/topology internally in case of inaccuracies
 - Directly perform pair related task
 - Link prediction
 - Edge classification
 - Distance Prediction
- EGT (Edge-augmented Graph Transformer)
 - Make pairs (2-tuple) first class citizen, just like nodes
 - Break free of the input graph topology
 - **Limitation: only 2nd-order interaction**





3D Molecular Geometry

- 2D
 - Bonds + Atoms (i.e., chemical formula)
- 3D
 - Coordinates
 - Often interatomic distances is enough
- 3D shape directly dictates molecular property
 - But costly to compute (QM simulation required)
- **Train a model: 2D→3D**
 - i.e., predict the molecular geometry



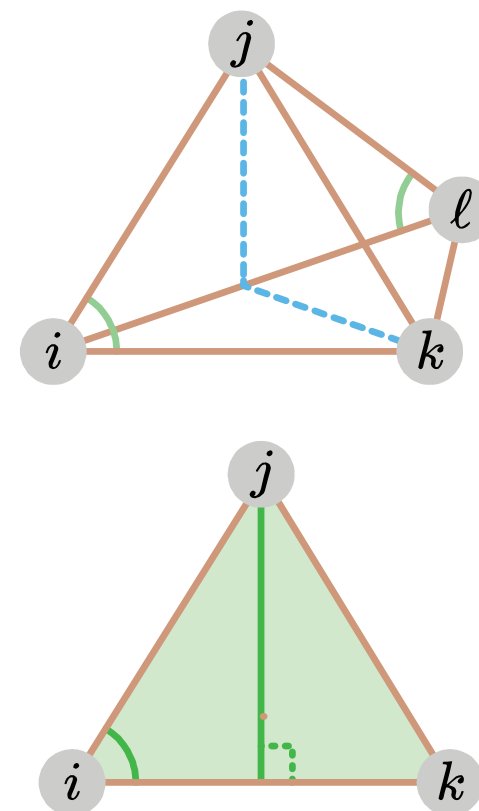


K-order interaction vs. K-order features

- **2nd Order (i, j) : Pairwise distances**
- **3rd Order (i, j, k) : Angles, area of triangles, etc.**
- **4th Order (i, j, k, ℓ) : Dihedral angles, volume of tetrahedrons, etc.**

We need either higher order interactions or higher order features for full geometric understanding

- **Crucial for 3D geometry prediction**
- **Our contribution: Third order interaction**
 - Pairs (i, j) , (j, k) , (i, k) within the 3-tuple (i, j, k)





Why higher order interactions?

- Using higher order features such as angles implies
 - An initial estimate of geometry is required
 - Features are only as accurate as the estimate
 - Specialized for geometric graphs only
- Using higher order interactions implies
 - No estimate of geometry is required, a simple graph is enough
 - The network can form representations that are more refined than the initial estimate
 - Applicable to both geometric and non-geometric graphs
- **Our work improves geometric expressivity without losing generality**
- **We break free from the inaccuracies of the initial geometry (when given)**





Our Contribution

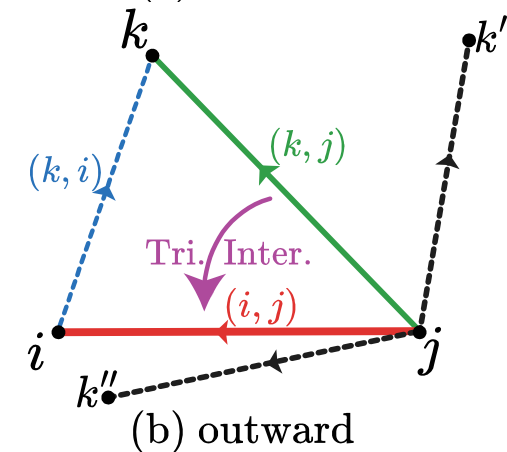
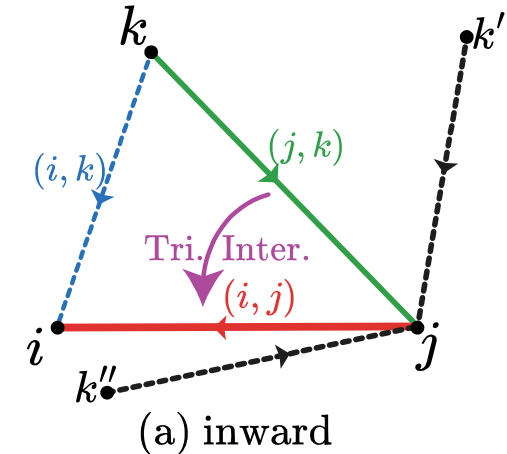
Triplet Graph Transformer





Triplet Interaction within a 3-tuple (i, j, k)

- Participants (inward): $(i, j), (j, k), (i, k)$
- (i, j) gathers information from (j, k)
- Without triplet interaction
 - $(j, k) \rightarrow j \rightarrow (i, j)$
 - Bottleneck at node j
- With triplet interaction
 - $(j, k) \xrightarrow{(i, k)} (i, j)$
- (i, k) also participates in this process
- Similarly: outward update





Triplet Interaction within a 3-tuple (i, j, k)

- Two mechanisms
 - Triplet Attention (TGT-At)

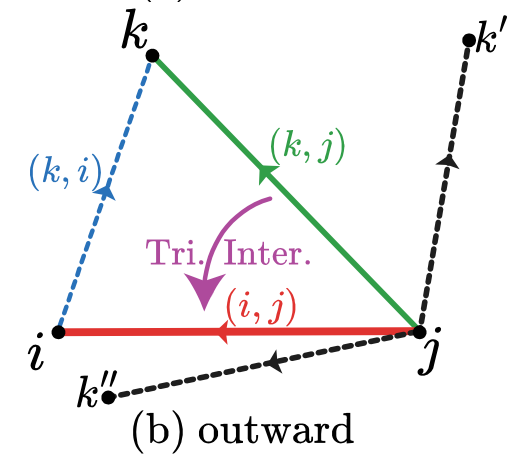
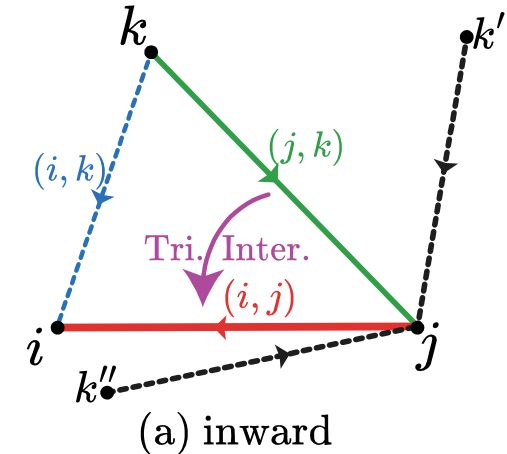
$$\mathbf{o}_{ij}^{\text{in}} = \sum_{k=1}^N a_{ijk}^{\text{in}} \mathbf{v}_{jk}^{\text{in}} ; \quad a_{ijk}^{\text{in}} = \text{softmax}_k \left(\frac{1}{\sqrt{d}} \mathbf{q}_{ij}^{\text{in}} \cdot \mathbf{p}_{jk}^{\text{in}} + b_{ik}^{\text{in}} \right) \times \sigma(g_{ik}^{\text{in}})$$

- More expressive, $O(N^3)$

- Triplet Aggregation (TGT-Ag)

$$\mathbf{o}_{ij}^{\text{in}} = \sum_{k=1}^N a_{ik}^{\text{in}} \mathbf{v}_{jk}^{\text{in}} ; \quad a_{ik}^{\text{in}} = \text{softmax}_k (b_{ik}^{\text{in}}) \times \sigma(g_{ik}^{\text{in}})$$

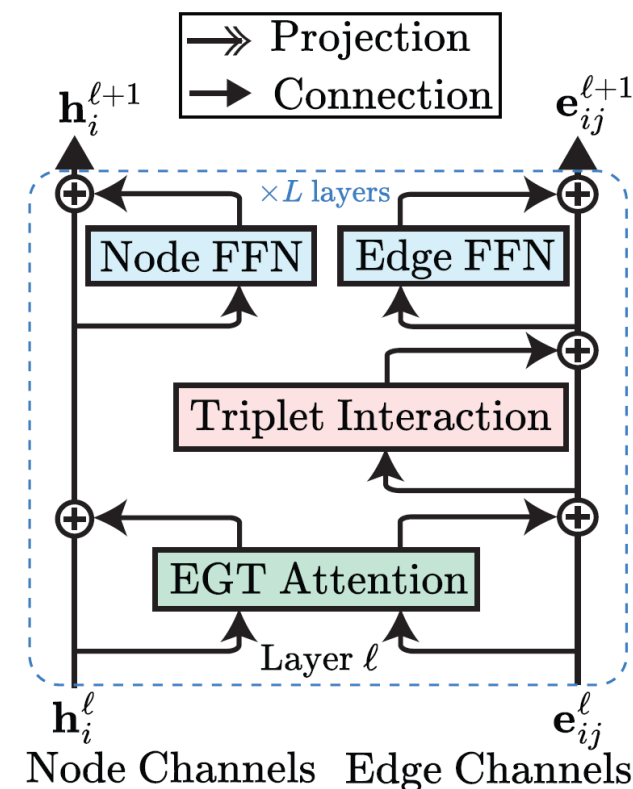
- More efficient, $O(N^{2.37})$





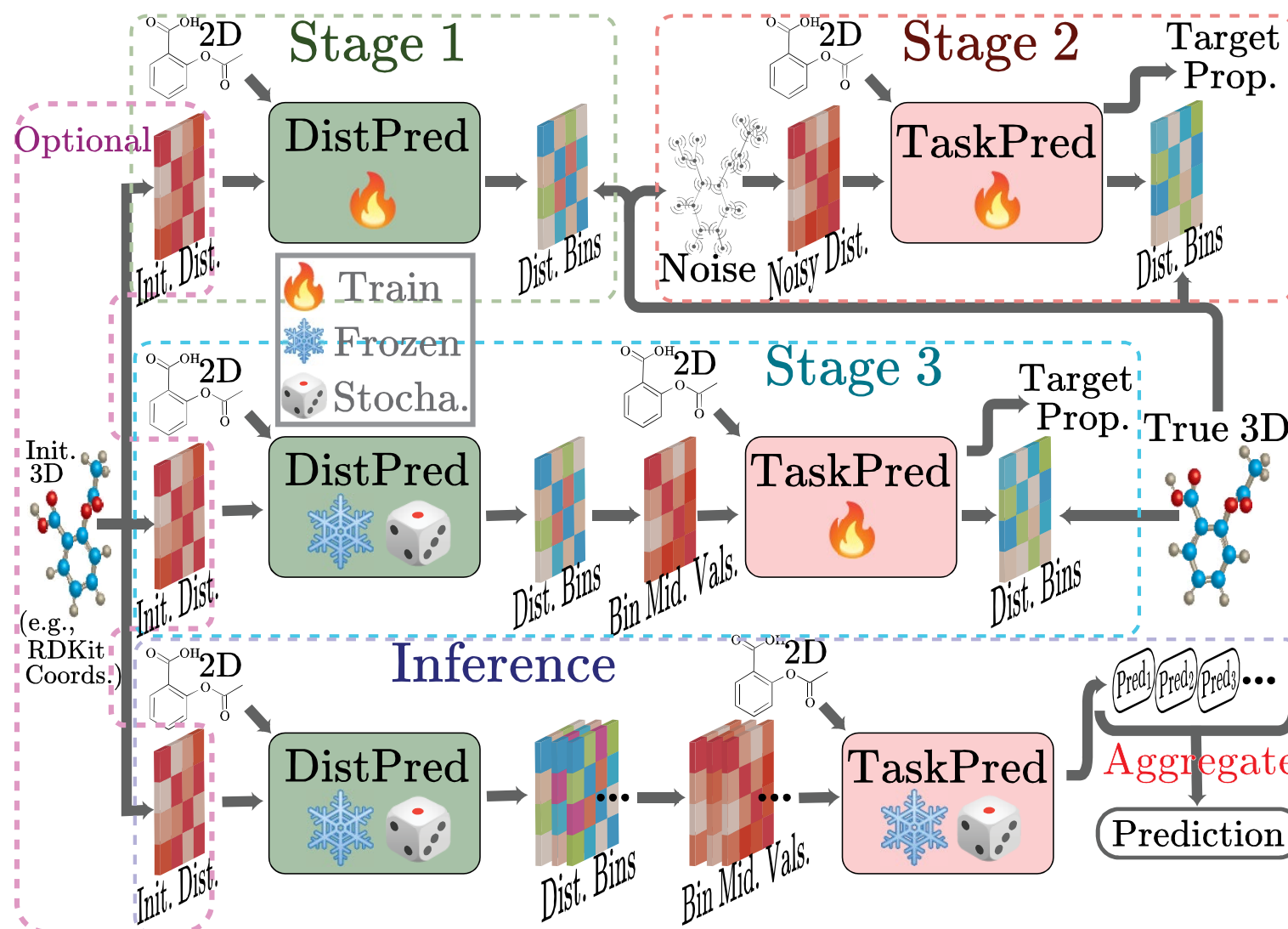
Network Architecture

- EGT (our previous work)
 - Node representations \rightarrow Node channels
 - Pair representations \rightarrow Edge channels
 - Only 2nd order interactions
- Triplet interaction (TGT)
 - Update pair representations based on each other
 - 3rd order interactions
- Pair embeddings are directly used for predicting (binned) pairwise atomic distances
- Useful for other geometric tasks as well (e.g., Traveling Salesman Problem)





Training and Inference



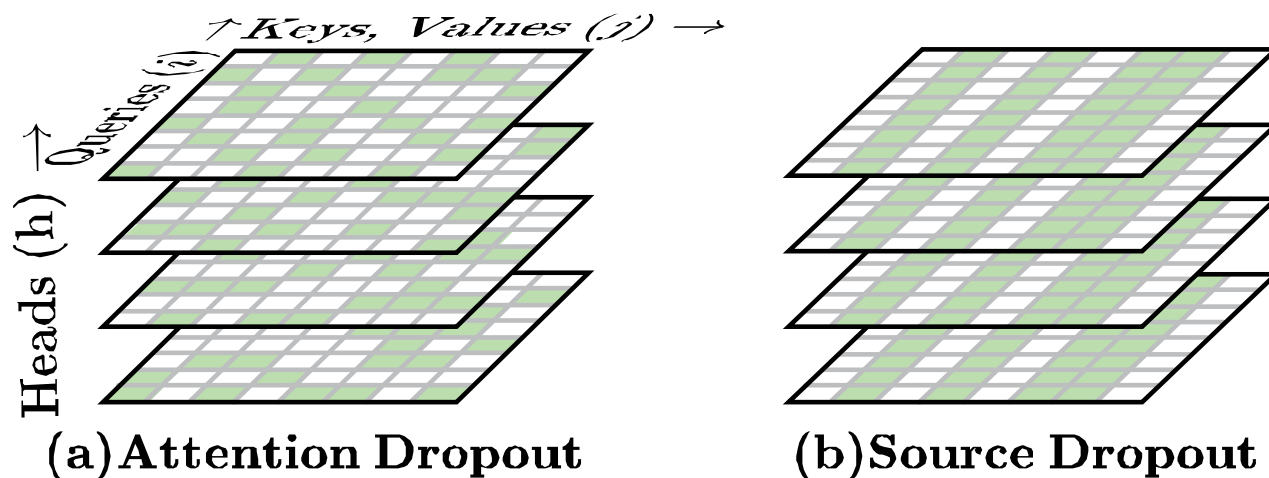


Other Contributions

- Locally smoothed 3D noise for pretraining the task predictor

$$\mathbf{r}'_i = \mathbf{r}_i + \sum_{j=1}^N e^{-\frac{\|\mathbf{r}_i - \mathbf{r}_j\|}{\nu}} \mathbf{u}_j; \text{ where } \mathbf{u}_j \sim \mathcal{N}(0, \sigma^2 \mathbf{I})$$

- Source dropout: Stronger regularization for graph transformers





Key Contributions

- Triplet Graph Transformer (TGT)
 - Novel triplet interaction mechanisms for direct pair-to-pair communication
 - Accurately models geometry in graphs
- Two-stage model
 - Separate distance and task predictors
 - Eliminates need for initial 3D coordinates
- Three-stage training procedure and stochastic inference
 - Significantly improves training efficiency and predictive performance
- TGT for graph learning
 - Traveling Salesman Problem





Quantum Chemistry (Large-scale)

- Predict in absence of ground truth 3D

(a) PCQM4Mv2 (Molecules)

Model	MAE↓ (meV)
EGT	86.2
Transformer-M	78.2
Uni-Mol+ (+RDKit)	70.5
<u>TGT-At</u>	69.8
<u>TGT-At (+RDKit)</u>	68.3

(b) OC20 IS2RE (Crystals)

Model	MAE↓ (meV)	EwT↑ (%)
SphereNet	618.8	3.32
EquiFormer	466	5.66
Uni-Mol+	414.3	8.23
<u>TGT-At</u>	414.7	8.3





Quantum Chemistry (Transfer Learning)

- Ground truth 3D is provided
- Fine-tuned from PCQM4Mv2

(a) QM9 (Molecules)

Method	μ	α	ϵ_H	ϵ_L	$\Delta\epsilon$	ZPVE	C_v
3D Infomax	0.034	0.075	29.8	25.7	48.8	1.67	0.033
SphereNet	0.025	0.053	22.8	18.9	31.1	1.12	0.024
Equiformer	0.011	0.046	15	14	30	1.26	0.023
Transformer-M	0.037	0.041	17.5	16.2	27.4	1.18	0.022
<u>TGT-Ag</u>	0.025	0.040	9.9	9.7	17.4	1.18	0.020





Molecular Property (Transfer Learning)

- Non-quantum property prediction and drug discovery
- Use frozen (not finetuned) distance predictor from PCQM4Mv2
 - Provides more accurate 3D information than RDKit

MOLPCBA (Property)

Model	AP↑ (%)
PHC-GNN	29.47
Graphormer	31.40
<u>TGT-Ag+RDKit</u>	31.44
<u>TGT-Ag+DP</u>	31.67

LITPCBA (Drugs)

Model	ROC-AUC↑ (%)
GEM	78.4
GEM-2+RDKit	81.5
EGT+RDKit	81.2
<u>EGT+DP</u>	81.5





Traveling Salesman Problem(Graph Learning)

- Points on a 2D plane
- Edge classification

TSP

Model	F1↑ (%)
GatedGCN	83.8
ARGNP	85.5
EGT	85.3
<u>TGT-Agx4</u>	87.1

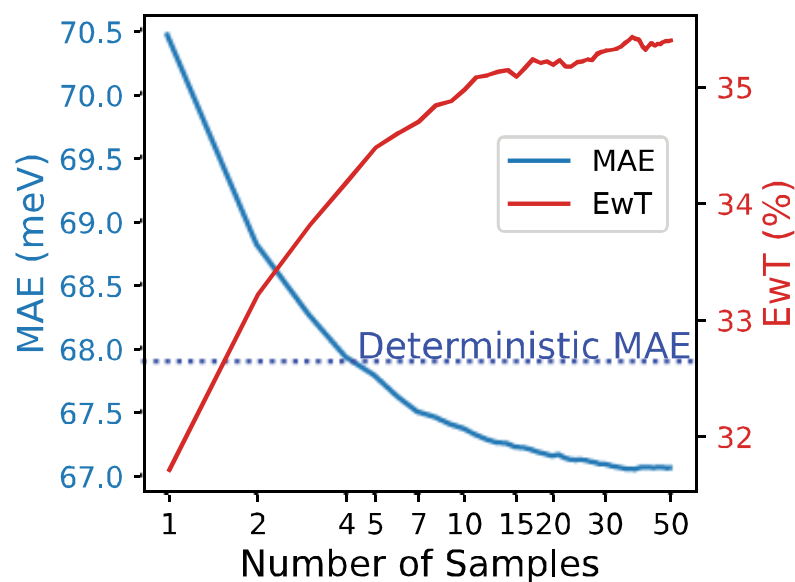


- Demonstrate the generality of our model

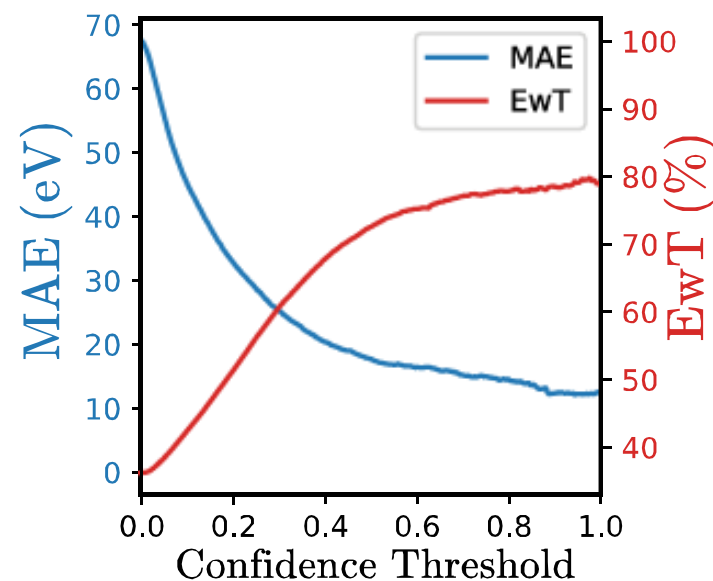


Merits of Stochastic Inference

- Outperforms deterministic inference with only ~4 samples
- Higher confidence implies higher accuracy



#Samples vs Performance



Confidence vs Performance





Future Work

- Explore use of triplet interaction for other graph learning tasks
 - Molecule and conformation generation
 - Link prediction
 - Combinatorial Problem
 - Self-supervised/semi-supervised/generative graph learning
- Improve compute and memory efficiency of triplet interaction
 - Sparsity
 - Linearity





Thank You

Please check out our paper for more details.

Paper: <https://arxiv.org/abs/2402.04538>

Implementation: <https://github.com/shamim-hussain/tgt>



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