

FACET: A Fragment-Aware Conformer Ensemble Transformer

Duy M. H. Nguyen^{1,2,3}, Trung Q. Nguyen³, Ha T. H. Le³, Mai Thanh Nhat Truong³, TrungTin Nguyen^{4,5}, Nhat Ho⁶, Khoa D Doan⁷

Duy Duong-Tran⁸, Li Shen⁸, Daniel Sonntag^{3,9}, James Zou¹⁰, Mathias Niepert^{1, 2}, Hyojin Kim¹¹, Jonathan E Allen¹¹

¹Max Planck Research School for Intelligent Systems (IMPRS-IS) ²University of Stuttgart ³German Research Center for Artificial Intelligence (DFKI)

⁴ARC Centre of Excellence for the Mathematical Analysis of Cellular Systems ⁵School of Mathematical Sciences, Queensland University of Technology

⁶University of Texas at Austin ⁷VinUniversity ⁸University of Pennsylvania ⁹Oldenburg University ¹⁰Stanford University ¹¹Lawrence Livermore National Labs

Correspondence to: Trung Q. Nguyen <trung.nguyen@dfki.de>, Duy M. H. Nguyen <hong01@dfki.de>



Motivation and Challenges

Motivation

- 2D graphs lack geometry
- single 3D conformer misses flexibility
- ensemble captures thermodynamic reality

Challenges

- naive aggregation ignores geometry
- FGW alignment is accurate but expensive
- poor scalability to large datasets



Our Contributions

- replace costly FGW alignment with a trainable Graph Transformer
- introduce fragment-level structural priors
- achieve over 6x speedup compared to FGW baselines and SOTA results



Fragment-Enhanced 2D Representation

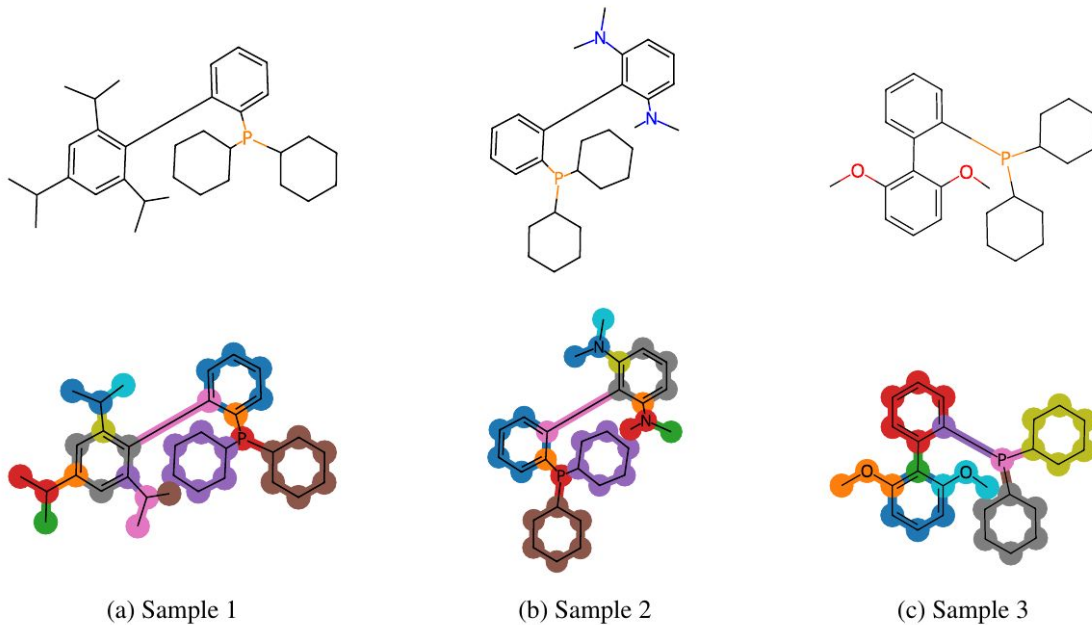
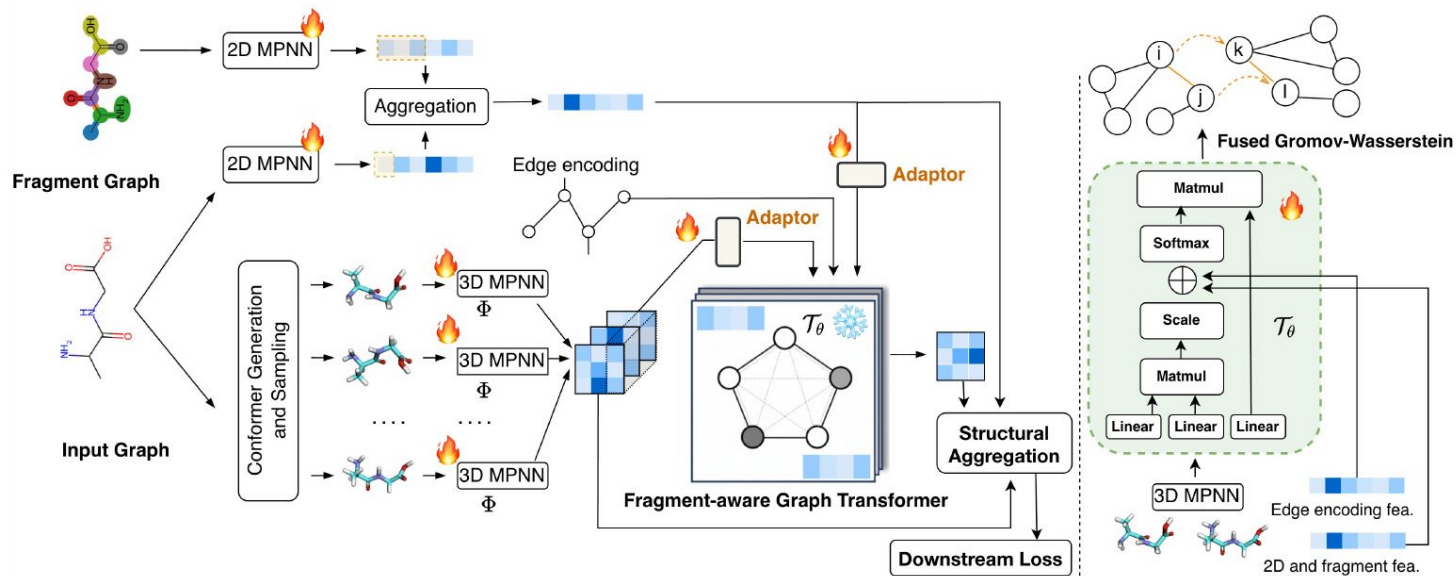


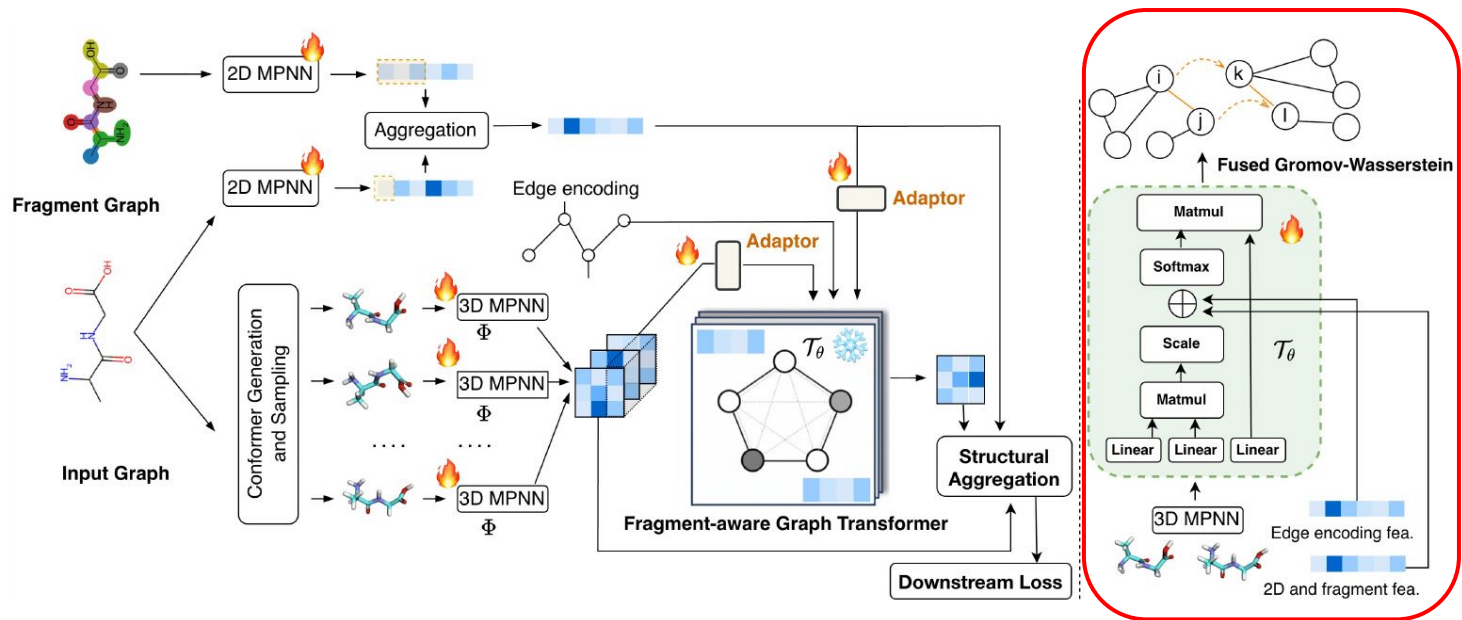
Figure 6: RingsPaths decomposition on three samples of the **Kraken** dataset. Top: 2D molecules; bottom: corresponding RingsPaths decomposition results.



FACET Architecture Overview



Approximate FGW with a Graph Transformer



FGW Approximation Quality

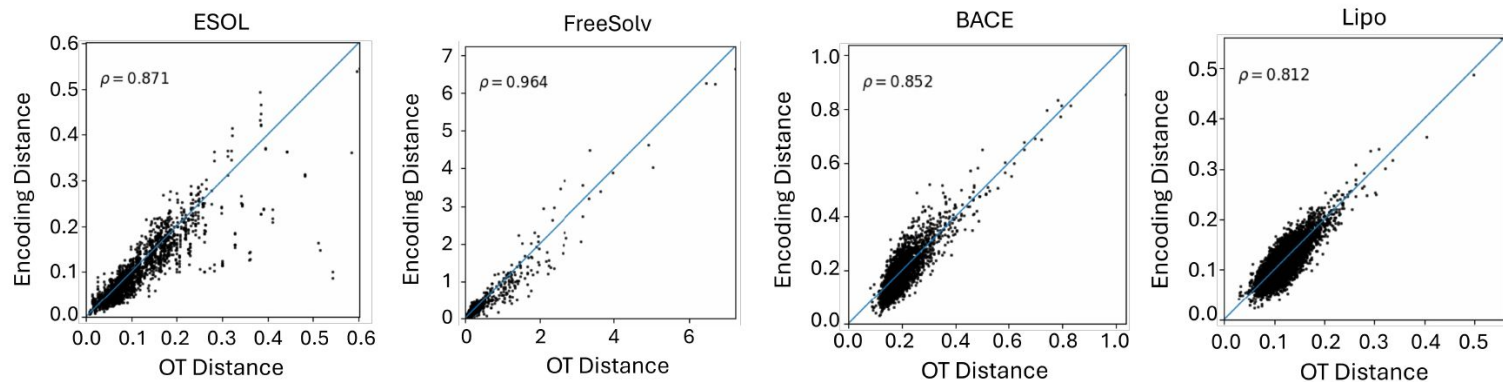


Figure 2: Correlations between FGW distance and trained GraphTransformer on four datasets in **MoleculeNet** benchmark. For each test molecule, we compute pairwise FGW distances between conformers and compare them with Euclidean distances between their Graph Transformer embeddings. The correlation ρ is reported, with the reference line $y = x$ shown in blue.



Scalability Results

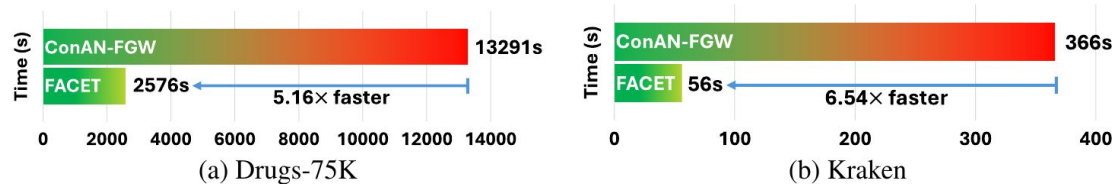


Figure 3: Comparison of the **one-epoch training time** of CONAN-FGW (Nguyen et al. 2024b) and the proposed FACET on the Drugs-75K and Kraken datasets from the MARCEL benchmark.

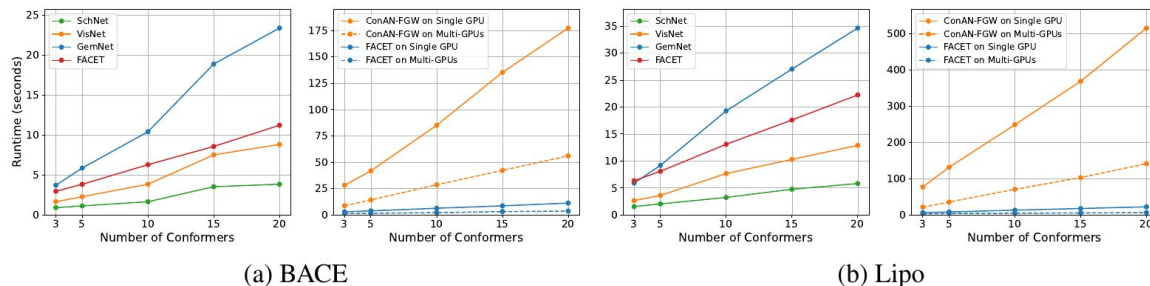


Figure 4: **Inference running time comparison** between FACET and other GNN-based methods on two datasets, BACE (a) and Lipo (b). Results are shown for both single-GPU and 4-GPU (multi-GPU) configurations. Reported runtimes represent the total time required to extract structural embeddings for all molecules in the test set of each dataset.



SOTA Results — MoleculeNet

Table 3: Comparison of molecular property regression performance on the **MoleculeNet** benchmark (MSE ↓). The results of competing methods are adapted from [Nguyen et al. \(2024b\)](#). FACET uses a SchNet backbone.

Model	Lipo	ESOL	FreeSolv	BACE
2D-GAT	1.387 ± 0.206	2.288 ± 0.017	8.564 ± 1.345	1.844 ± 0.33
D-MPNN	0.534 ± 0.022	0.923 ± 0.045	4.213 ± 0.068	0.723 ± 0.021
Attentive FP	0.520 ± 0.001	0.771 ± 0.026	4.197 ± 0.193	-
PretrainGNN	0.545 ± 0.003	1.210 ± 0.005	6.392 ± 0.003	-
GROVER_large	0.676 ± 0.012	0.798 ± 0.018	5.162 ± 0.047	-
ChemBERTa-2*	0.639 ± 0.006	0.795 ± 0.033	-	1.858 ± 0.029
ChemRL-GEM	0.486 ± 0.008	0.706 ± 0.061	3.924 ± 0.436	-
MolFormer	0.492 ± 0.012	0.766 ± 0.026	5.485 ± 0.045	1.091 ± 0.021
ConfNet	1.360 ± 0.038	2.115 ± 0.484	-	1.329 ± 0.042
UniMol	0.374 ± 0.012	0.741 ± 0.014	2.867 ± 0.186	-
SchNet-scalar	0.704 ± 0.032	0.672 ± 0.027	1.608 ± 0.158	0.723 ± 0.100
SchNet-emb	0.589 ± 0.022	0.635 ± 0.057	1.587 ± 0.136	0.692 ± 0.028
ChemProp3D	0.602 ± 0.035	0.681 ± 0.023	2.014 ± 0.182	0.815 ± 0.170
CONAN	0.556 ± 0.013	0.571 ± 0.019	1.496 ± 0.158	0.635 ± 0.051
CONAN-FGW	0.422 ± 0.016	0.529 ± 0.022	1.068 ± 0.083	0.549 ± 0.016
FACET	0.424 ± 0.009	0.516 ± 0.044	0.967 ± 0.082	0.495 ± 0.034



SOTA Results — MARCEL

Table 4: Comparison of molecular property regression performance on the **MARCEL** benchmark (MAE ↓). The results of competing methods are adapted from [Zhu et al. \(2024a\)](#).

Category	Model	Drugs-75K			Kraken			
		IP	EA	χ	B ₅	L	BurB ₅	BurL
2D models	GIN	0.4354	0.4169	0.2260	0.3128	0.4003	0.1719	0.1200
	GIN+VN	0.4361	0.4169	0.2267	0.3567	0.4344	0.2422	0.1741
	ChemProp	0.4595	0.4417	0.2441	0.4850	0.5452	0.3002	0.1948
	GraphGPS	0.4351	0.4085	0.2212	0.3450	0.4363	0.2066	0.1500
3D models	SchNet	0.4394	0.4207	0.2243	0.3293	0.5458	0.2295	0.1861
	DimeNet++	0.4441	0.4233	0.2436	0.3510	0.4174	0.2097	0.1526
	GemNet	0.4069	0.3922	0.1970	0.2789	0.3754	0.1782	0.1635
	PaiNN	0.4505	0.4495	0.2324	0.3443	0.4471	0.2395	0.1673
	ClofNet	0.4393	0.4251	0.2378	0.4873	0.6417	0.2884	0.2529
	LEFTNet	0.4174	0.3964	0.2083	0.3072	0.4493	0.2176	0.1486
Ensemble Strategy with DeepSets	SchNet	0.4452	0.4232	0.2243	0.2704	0.4322	0.2024	0.1443
	DimeNet++	0.4126	0.3944	0.2267	0.2630	0.3468	0.1783	0.1185
	GemNet	0.4066	0.3910	0.2027	0.2313	0.3386	0.1589	0.0947
	PaiNN	0.4466	0.4269	0.2294	0.2225	0.3619	0.1693	0.1324
	ClofNet	0.4280	0.4033	0.2199	0.3228	0.4485	0.2178	0.1548
	LEFTNet	0.4149	0.3953	0.2069	0.2644	0.3643	0.2017	0.1386
FACET	SchNet	0.4235	0.3971	0.2155	0.2508	0.3982	0.1803	0.1245
	GemNet	0.3891	0.3852	0.1970	0.2225	0.3402	0.1503	0.0952



Thank you

Codes are available at <https://github.com/duyhominhnguyen/FACET>

Affiliations

