

Figure 1. SGT on ZINC: Test MAE v.s. Batch Size (BS). # Training epochs are adjusted per batch-size for the same total update steps: $400 * BS/32$. The first 10% epochs are in the warmup stage.

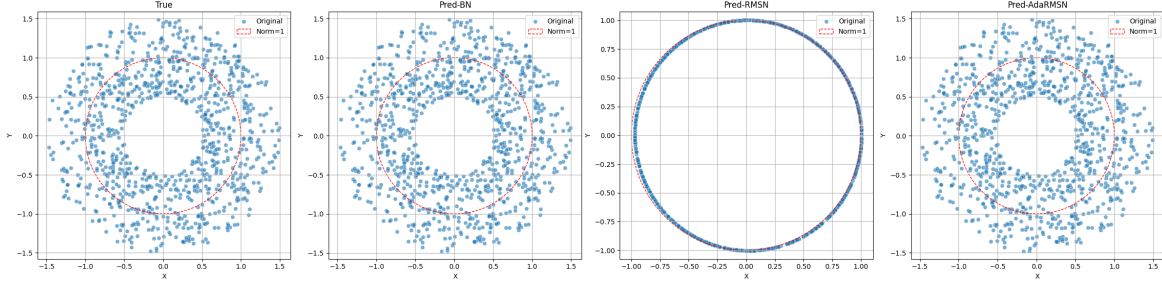


Figure 2. (Case Study of AdaRMSN) Visualization of Input and Pred data points. Overfitting test on Auto-encoders of 2-dim ($Linear \rightarrow BN/RMSN/AdaRMSN \rightarrow Linear$): each model is trained 5000 epochs via AdamW without regularization. (together with Figure. 3)

Table 1. Performance on ZINC. GPS+s L_2 : integrating s L_2 attention into GPS without changing other parts. (run 3 trials)

ZINC	GPS	GPS+s L_2	SGT
MAE (\downarrow)	0.070 ± 0.004	0.0693 ± 0.0023	0.0566 ± 0.002

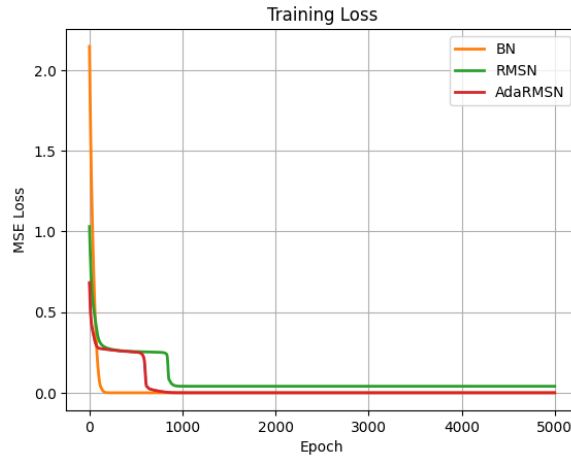


Figure 3. (Case Study of AdaRMSN) Training curves of overfitting test. (together with Figure. 2)

Table 2. Comparison of peak GPU memory usage and per-epoch training time for GRIT and SGT. Dataset: Peptides-Structure (15K graphs); Model config.: 5 transformer layers, 96 channels, batch size 32. Hardware: a single Nvidia V100 GPU with 32GB memory, supported by 80 Intel Xeon Gold 6140 CPUs running at 2.30GHz

Model	GPU Memory (GB)	Training Time (Sec/Epoch)
GRIT	29.16	141.60
SGT	25.07	100.68
Improv.	~14.03%	~28.9%

Table 3. Performance on PCQM4Mv2 (over 3.7M graphs). The eval. pipeline follows Rampásek et al. (2022); no 3D-info included.

PCQM4Mv2	Val MAE (\downarrow)	# Param.
Graphormer	0.0864	48.3M
GPS	0.0858	19.4M
GRIT	0.0859	16.6M
SGT	0.0856	17.6M

Table 4. Performance comparison across different models on various datasets. Best results are highlighted in bold. * indicates the difference to the best is not statistically significant (by two-tail T-test)

Model	ZINC MAE (\downarrow)	SP-CIFAR Acc. (\uparrow)	SP-MNIST Acc. (\uparrow)	PATTERN W.Acc. (\uparrow)	CLUSTER W.Acc. (\uparrow)	Peptides-Struct MAE (\downarrow)	Peptides-Func AP (\uparrow)
Exphormer	-	74.69 \pm 0.125	98.55 \pm 0.037	86.74 \pm 0.015	78.07 \pm 0.037	0.2481 \pm 0.0007	0.6527 \pm 0.0043
GEAET	-	76.634 \pm 0.427	98.513 \pm 0.086	86.993 \pm 0.026	-	0.2445\pm0.0013	-
GEANet	0.193 \pm 0.001	73.857 \pm 0.306	98.315 \pm 0.097	85.607 \pm 0.038	77.013 \pm 0.224	0.2512 \pm 0.0003	0.6722 \pm 0.0065
SGT	0.0566\pm0.002	78.560\pm0.700	98.614\pm0.096	89.752\pm0.030	80.027\pm0.114	0.2450 \pm 0.0017*	0.6961\pm0.0062

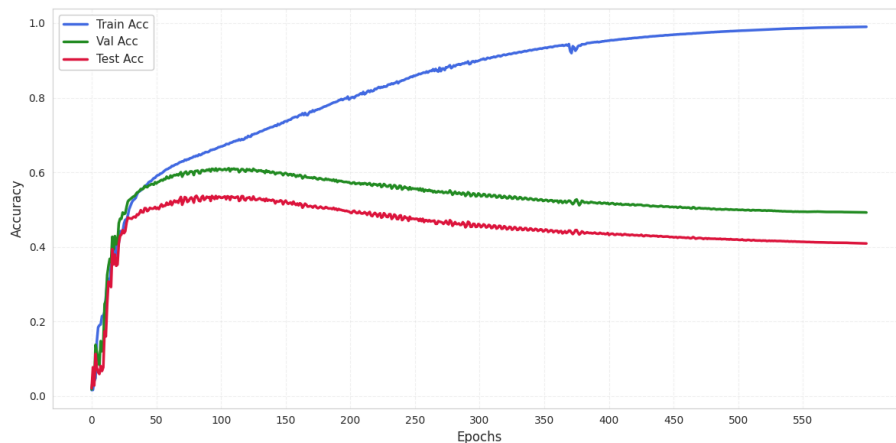


Figure 4. Sanity check of Exp-SGT on a large-scale graph in OGBN-ArXiv (169,343 nodes). Use the same configuration as Exphormer and remove all regularizations to validate the trainability via an overfitting test.