

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.

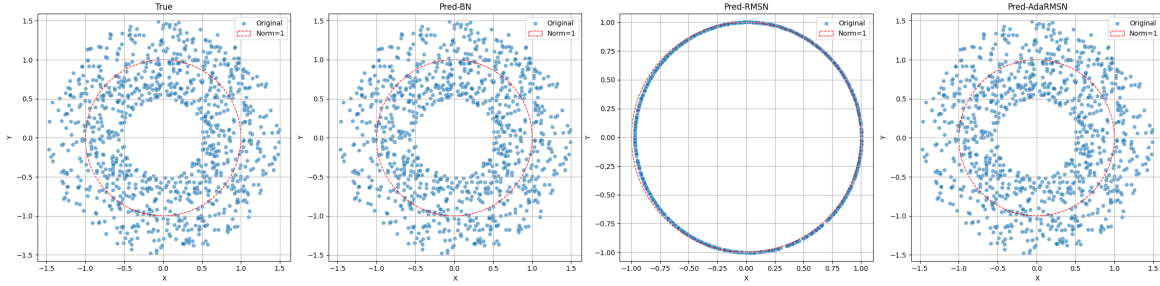


Figure 2. (Case Study of AdaRMSN) Visualization of Input and Pred data points. Overfitting test on Auto-encoders of 2-dim ( $Linear \rightarrow BN/RMSN/AdaRMSN \rightarrow Linear$ ): each model is trained 5000 epochs via AdamW without regularization. (together with Figure. 3)

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 $\pm$ 0.004	0.0693 $\pm$ 0.0023	0.0566 $\pm$ 0.002

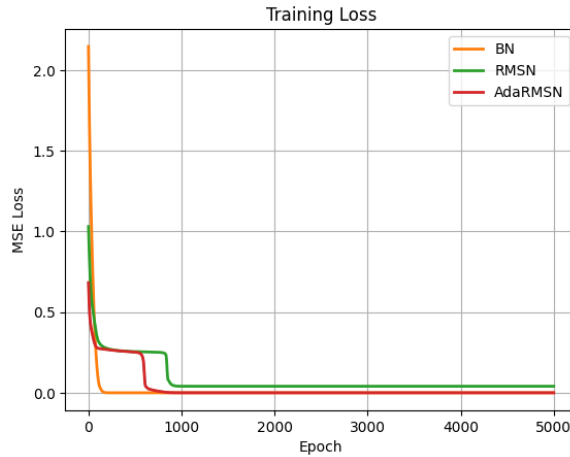


Figure 3. (Case Study of AdaRMSN) Training curves of overfitting test. (together with Figure. 2)

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
GAEAT	-	76.634 $\pm$ 0.427	98.513 $\pm$ 0.086	86.993 $\pm$ 0.026	-	<b>0.2445<math>\pm</math>0.0013</b>	-
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	<b>0.0566<math>\pm</math>0.002</b>	<b>78.560<math>\pm</math>0.700</b>	<b>98.614<math>\pm</math>0.096</b>	<b>89.752<math>\pm</math>0.030</b>	<b>80.027<math>\pm</math>0.114</b>	0.2450 $\pm$ 0.0017*	<b>0.6961<math>\pm</math>0.0062</b>

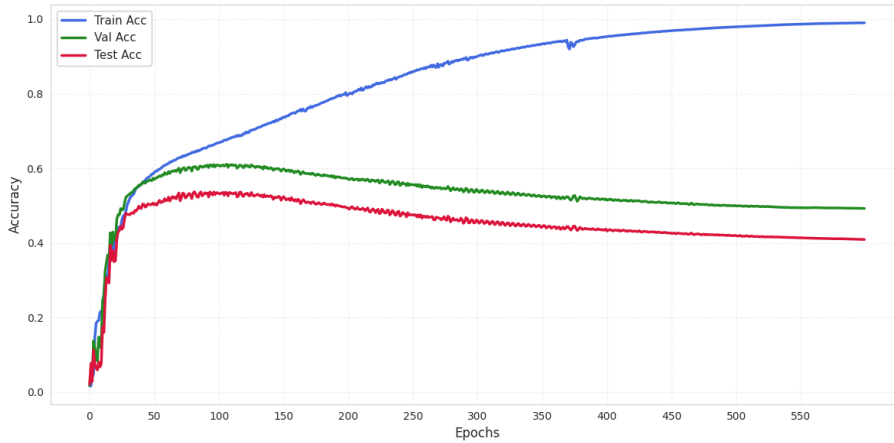


Figure 4. Sanity check of Exp-SGT on a large-scale graph in OGBN-ArXiv (169,343 nodes). Use the same configuration as Exphormer and remove all regularizations to validate the trainability via an overfitting test.