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A Theoretical and Empirical Investigation into the Equivalence of Graph Neural Networks and the Weisfeiler-Leman Algorithm

From the Faculty of Mathematics, Physics, and Computer Science approved for the purpose of obtaining the academic degree of Bachelor of Sciences.

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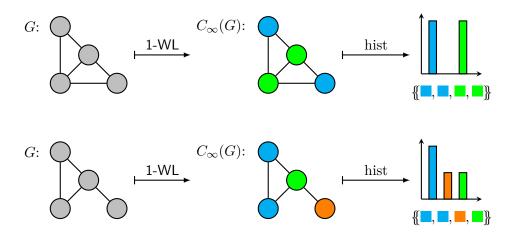
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Table 1: Overview of the mean absolute error and the standard deviation (logMAE) on large-scale (multi-target) molecular regression tasks. We highlighted the lowest error for the 1-WL+NN and GNN models for each dataset.

	Dataset						
Method	ALCHEMY	Alchemy (10k)	ZINC	ZINC (10K)			
1-WL+NN	$0.600\ \pm0.004\ -0.625\ \pm0.032$	$0.305 \pm 0.001 - 1.740 \pm 0.042$	$0.229 ~\pm 0.003$	$0.465~\pm 0.009$			
GNN	$0.523 \pm 0.016 - 0.705 \pm 0.035$	$0.282\ \pm0.002\ -1.890\ \pm0.031$	0.104 ± 0.005	0.298 ± 0.034			

Table 2: Accuracy and standard deviation in percent achieved by the best-performing 1-WL+NN, 1-WL:GNN, and GNN model on each classification dataset.

26.11	Dataset						
Model	Enzymes	IMDB-BINARY	Mutag	Nci1	PROTEINS	REDDIT-BINARY	
1-WL+NN	$48.3{\scriptstyle~\pm 8.1}$	$72.4{\scriptstyle~\pm4.1}$	$85.1~\pm 8.6$	83.6 ± 2.2	75.2 ± 3.9	78.4 ± 2.7	
GNN	34.4 ± 7.0	$74.7 \pm\! 3.8$	84.6 ±8.7	79.9 ± 2.2	74.3 ± 5.1	86.9 ±3.2	



test Morris et al. [2019]

Bibliography

[1] C. Morris, M. Ritzert, M. Fey, W. L. Hamilton, J. E. Lenssen, G. Rattan, and M. Grohe. Weisfeiler and leman go neural: Higher-order graph neural networks. In *Proceedings of the AAAI conference on artificial intelligence*, volume 33, pages 4602–4609, 2019.