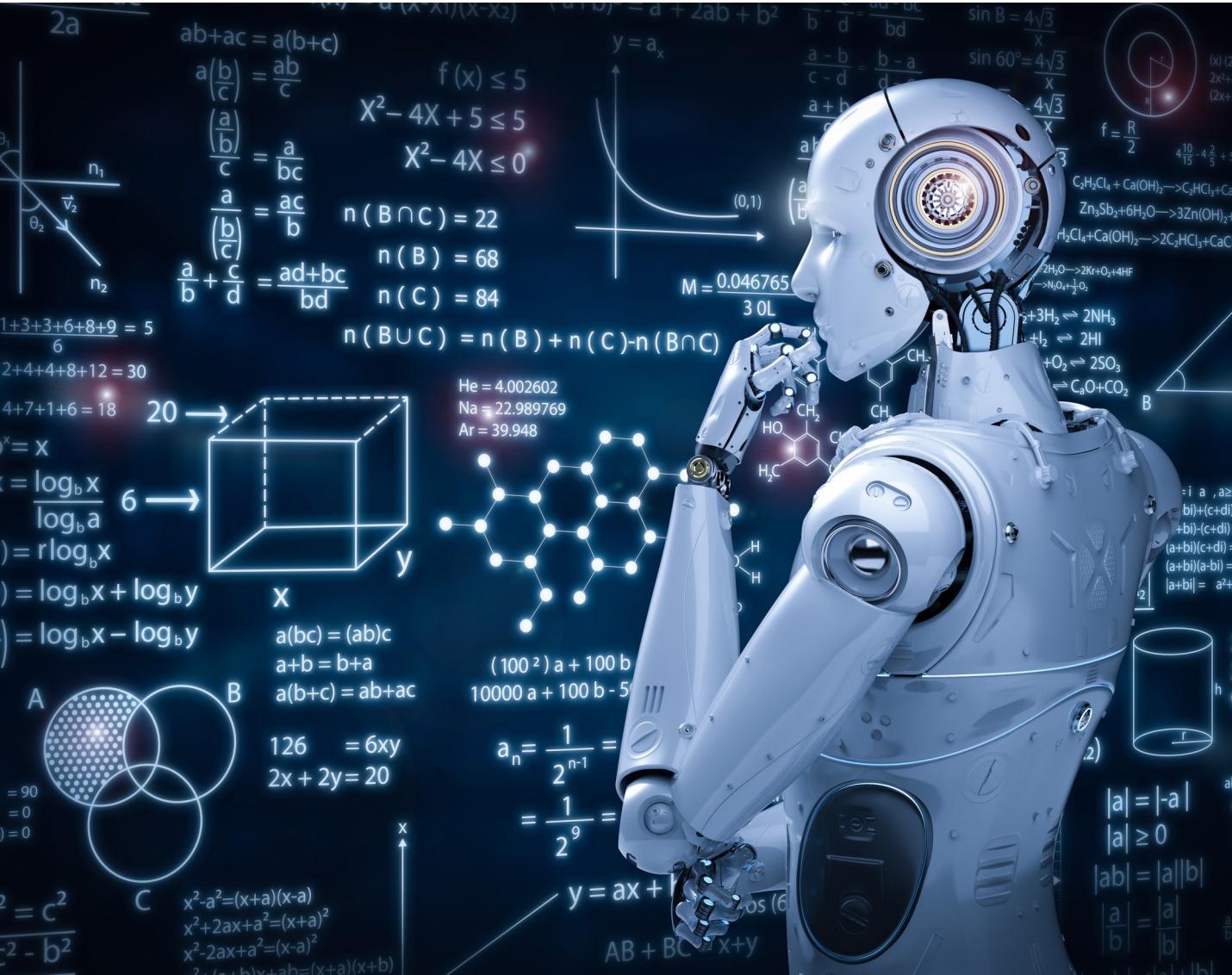




Weisfeiler and Leman Go Loopy

Raffaele Paolino, Sohir Maskey,
 Pascal Welke & Gitta Kutyniok

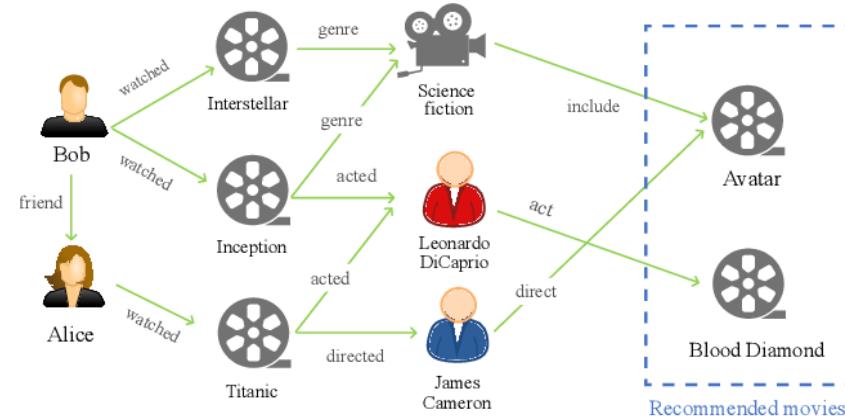


The Current Impact of Graph Neural Networks

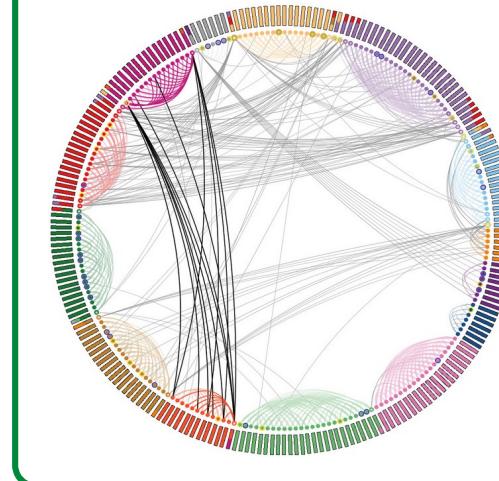
Social Networks



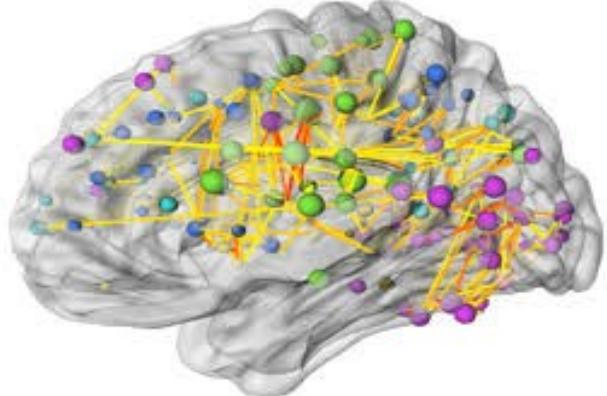
Recommender Systems



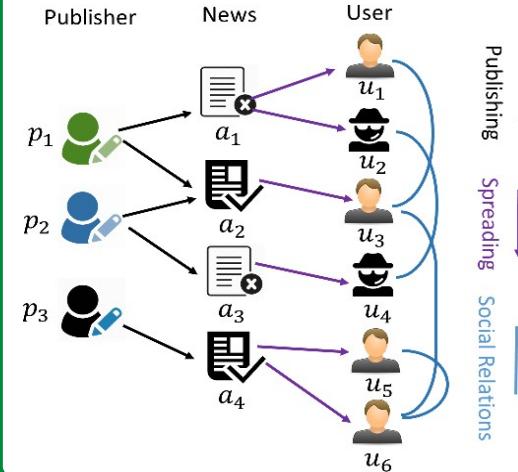
Drug Discovery



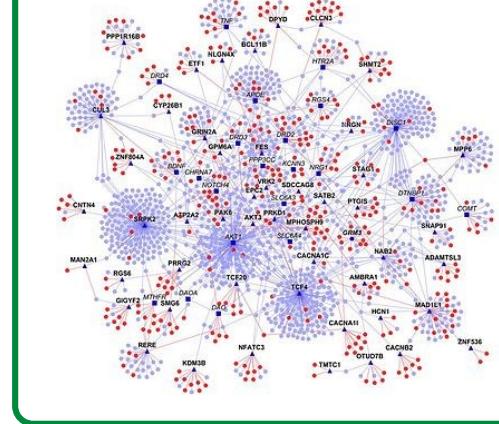
Functional Brain Networks



Fake News Detection



Drug Design



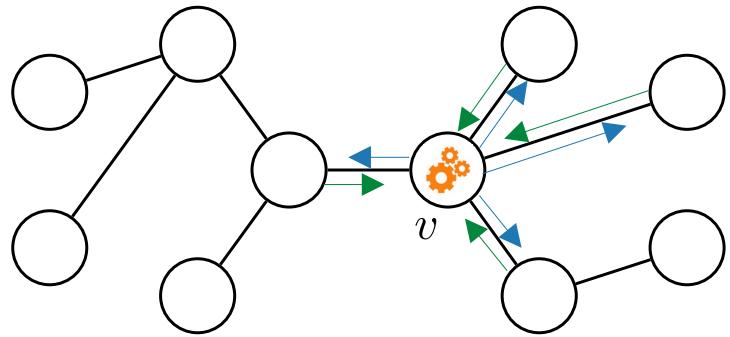
Gilmer, J., et al. (2017). *Neural message passing for quantum chemistry*.

Wang, J., et al. (2018). *Billion-scale commodity embedding for e-commerce recommendation in alibaba*.

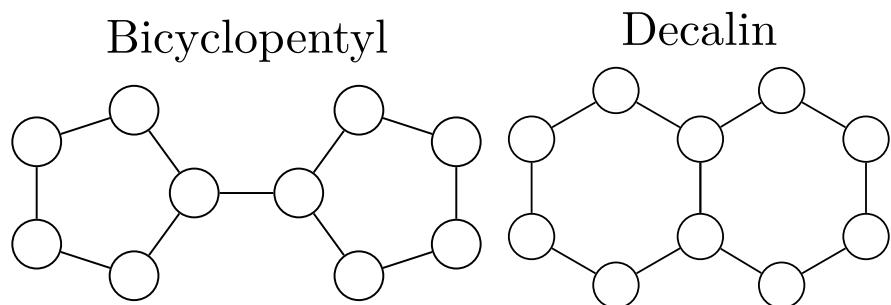
Monti, F., et al. (2019). *Fake news detection on social media using geometric deep learning*.

Fan, W., et al. (2019). *Graph neural networks for social recommendation*.

The Current Weaknesses of Graph Neural Networks



$$x_v^{(k+1)} = \text{MLP}^{(k+1)} \left((1 + \varepsilon^{(k+1)}) x_v^{(k)} + \sum_{j \in \mathcal{N}(v)} x_j^{(k)} \right)$$



- The paradigm of GNN is **message passing**.
 - **Messages** are collected from neighboring nodes.
 - Messages are used to **update** the node features.
 - The new message is **sent** to neighboring nodes.

- Message-passing cannot separate all non-isomorphic graphs.
 - They cannot approximate all functions on graphs.
 - No universal approximation theorem.

Which are the graphs that Message Passing Neural Networks can distinguish?

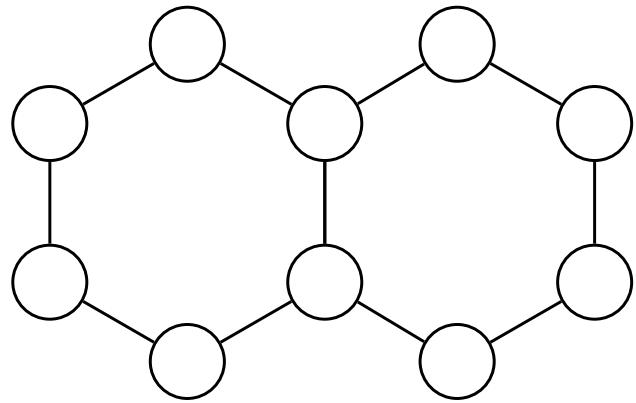
Weisfeiler-Leman Graph Isomorphism Test



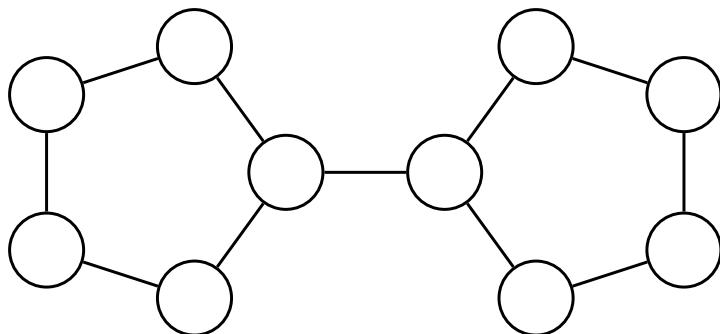
Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



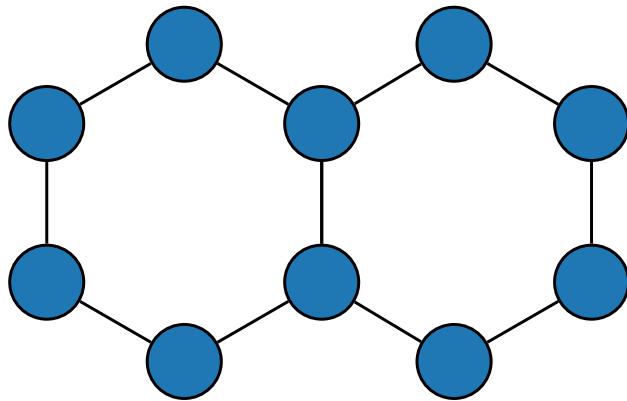
Bicyclopentyl



Weisfeiler-Leman Graph Isomorphism Test

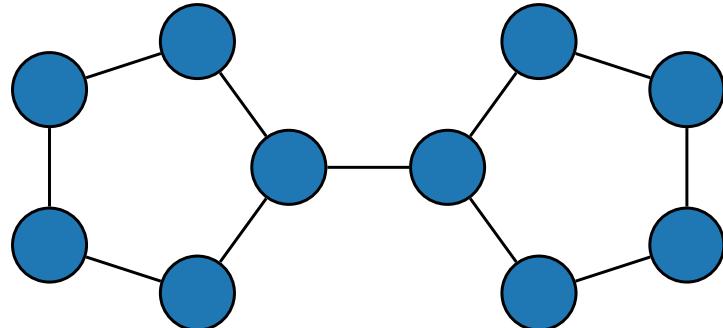
Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



1. Assign to each node the same label •.

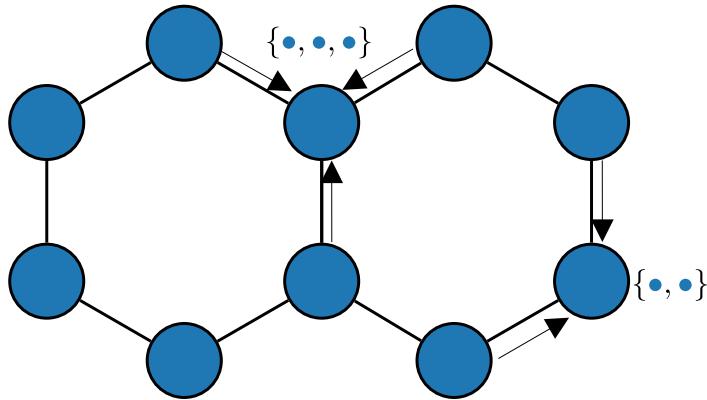
Bicyclopentyl



Weisfeiler-Leman Graph Isomorphism Test

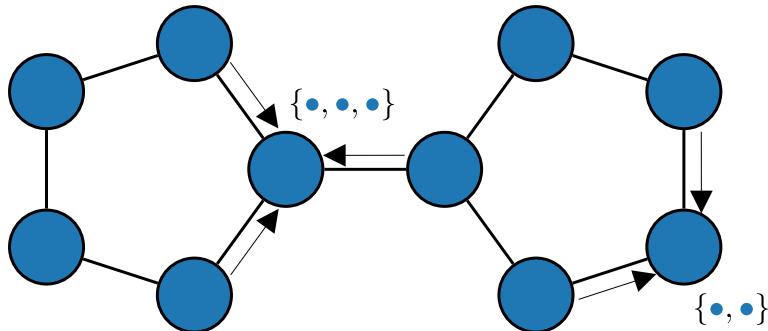
Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



1. Assign to each node the same label \bullet .
2. Collect all labels from direct neighbors.

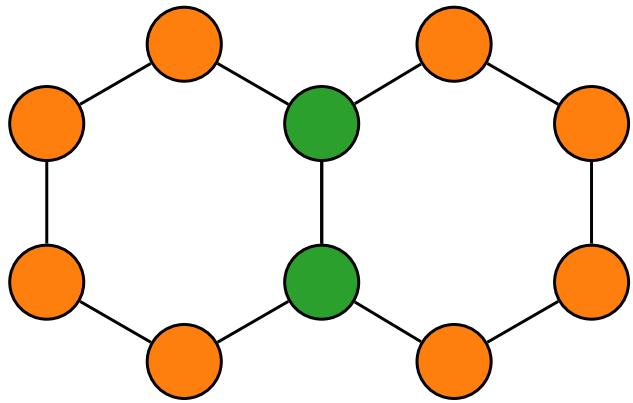
Bicyclopentyl



Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

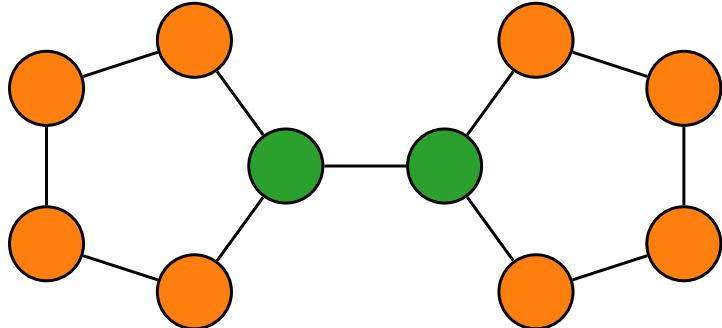
Decalin



1. Assign to each node the same label \bullet .
2. Collect all labels from direct neighbors.
3. Compress to get a new label

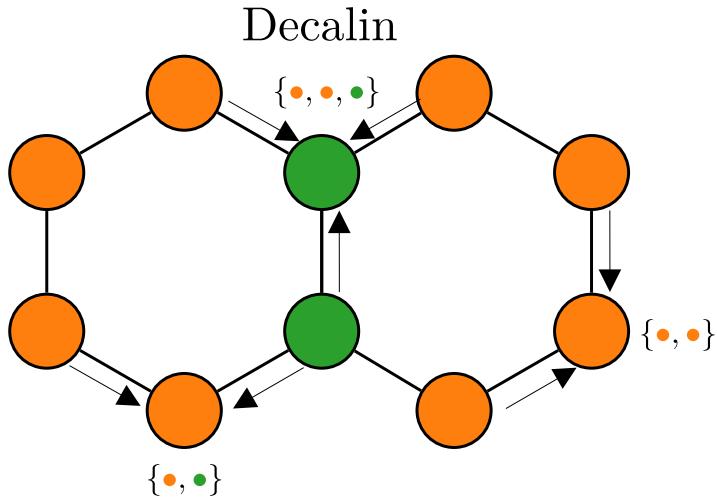
$$\{\bullet, \bullet\} \mapsto \bullet \quad \{\bullet, \bullet, \bullet\} \mapsto \bullet$$

Bicyclopentyl



Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic



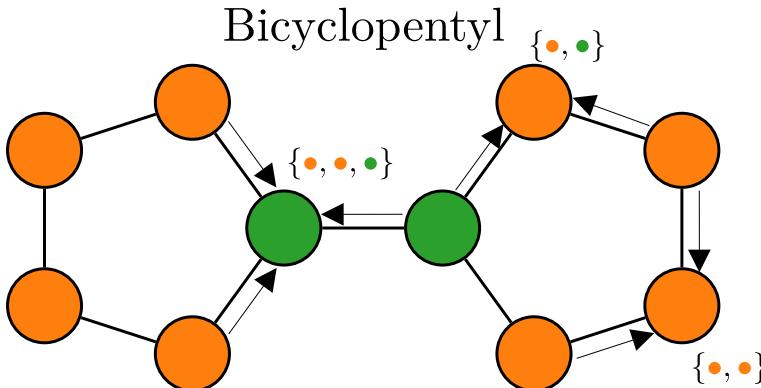
1. Assign to each node the same label \bullet .

2. Collect all labels from direct neighbors.

3. Compress to get a new label

$$\{\bullet, \bullet\} \mapsto \bullet \quad \{\bullet, \bullet, \bullet\} \mapsto \bullet$$

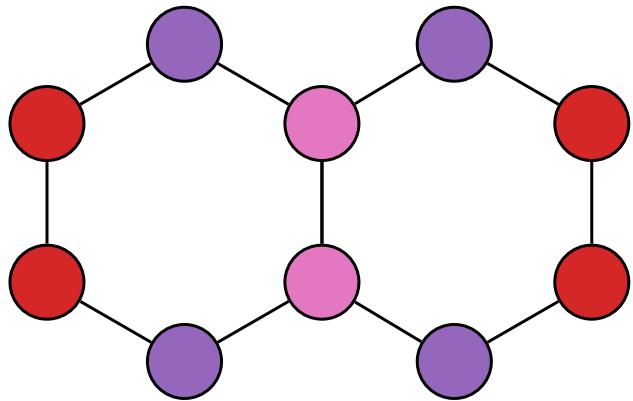
4. Repeat until there is no change in the partition of nodes with same label.



Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



1. Assign to each node the same label \bullet .

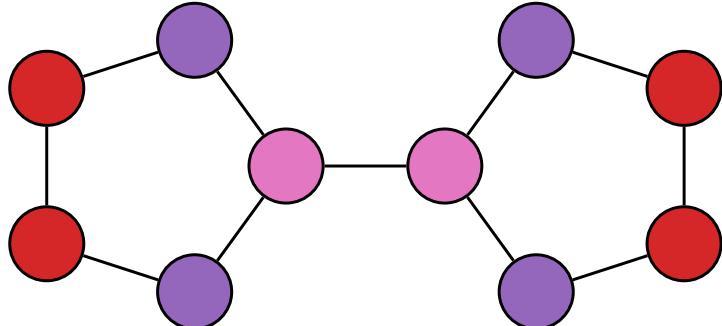
2. Collect all labels from direct neighbors.

3. Compress to get a new label

$$\{\bullet, \bullet\} \mapsto \bullet \quad \{\bullet, \text{green}\} \mapsto \bullet \quad \{\bullet, \bullet, \text{green}\} \mapsto \bullet$$

4. Repeat until there is no change in the partition of nodes with same label.

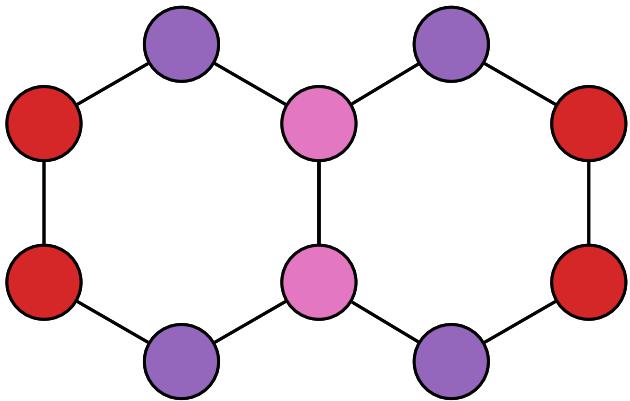
Bicyclopentyl



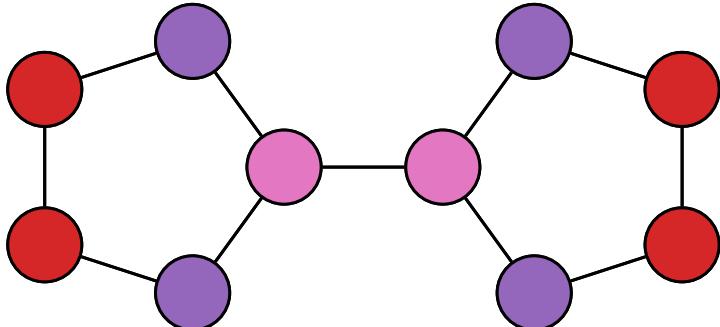
Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



Bicyclopentyl



1. Assign to each node the same label \bullet .

2. Collect all labels from direct neighbors.

3. Compress to get a new label

$$\{\bullet, \bullet\} \mapsto \bullet \quad \{\bullet, \circ\} \mapsto \circ \quad \{\bullet, \circ, \circ\} \mapsto \circ$$

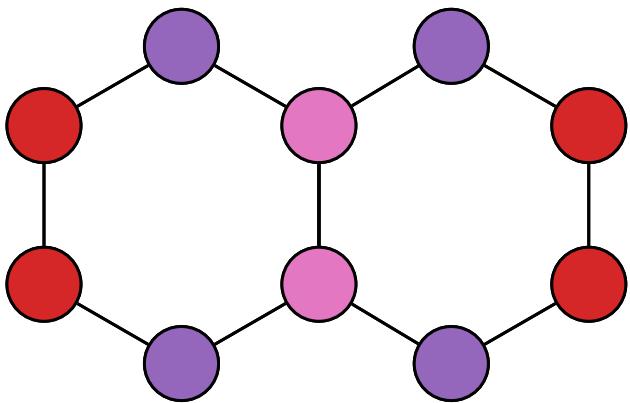
4. Repeat until there is no change in the partition of nodes with same label.

5. Compare the histogram of labels: if different, the graphs are definitely non-isomorphic.

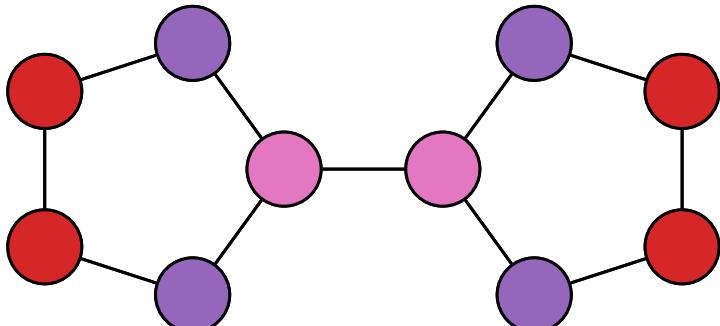
Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



Bicyclopentyl



1. Assign to each node the same label \bullet .

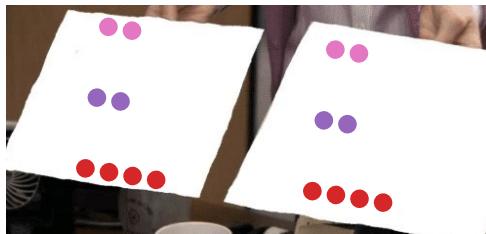
2. Collect all labels from direct neighbors.

3. Compress to get a new label

$$\{\bullet, \bullet\} \mapsto \bullet \quad \{\bullet, \circ\} \mapsto \circ \quad \{\bullet, \circ, \circ\} \mapsto \circ$$

4. Repeat until there is no change in the partition of nodes with same label.

5. Compare the histogram of labels: if different, the graphs are definitely non-isomorphic.

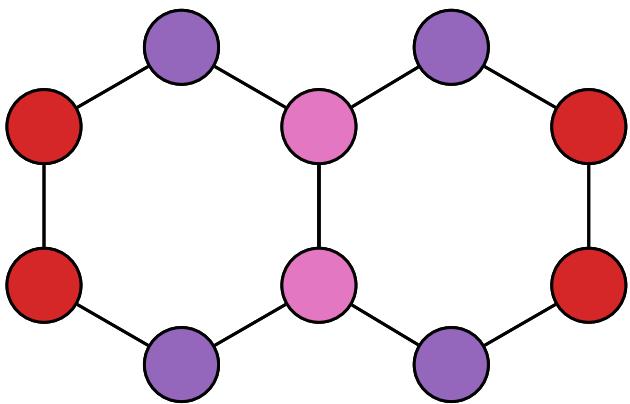


Corporate needs you to find the differences between these histograms

Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



1. Assign to each node the same label \bullet .

2. Collect all labels from direct neighbors.

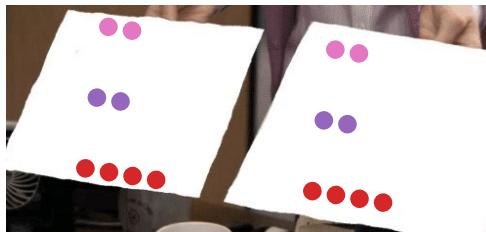
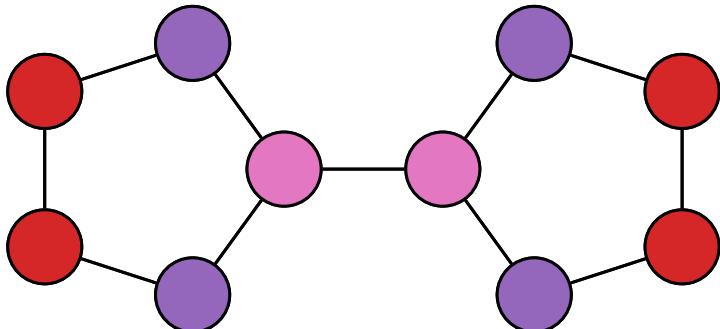
3. Compress to get a new label

$$\{\bullet, \bullet\} \mapsto \bullet \quad \{\bullet, \circ\} \mapsto \circ \quad \{\bullet, \circ, \circ\} \mapsto \circ$$

4. Repeat until there is no change in the partition of nodes with same label.

5. Compare the histogram of labels: if different, the graphs are definitely non-isomorphic.

Bicyclopentyl



Corporate needs you to find the differences between these histograms

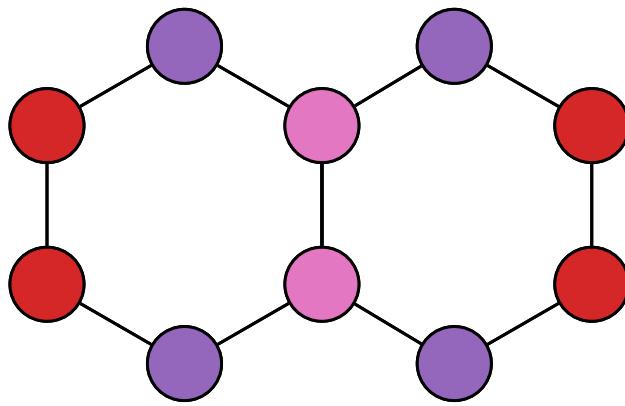


They are the same histograms!

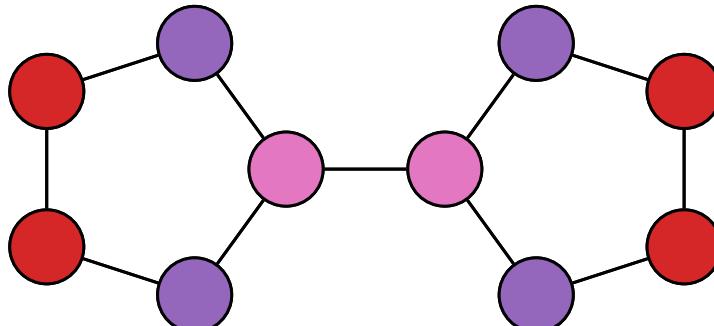
Weisfeiler-Leman Graph Isomorphism Test

Simplest algorithm to check whether two graphs are non-isomorphic

Decalin



Bicyclopentyl



Advantage:

- Locality: only local computations needs to be performed.

Disadvantage:

- Locality: unable to capture higher-order interactions.



Proposition (Which graphs can WL distinguish?). *The Weisfeiler-Leman algorithm can distinguish trees and forests (disjoint union of trees).*

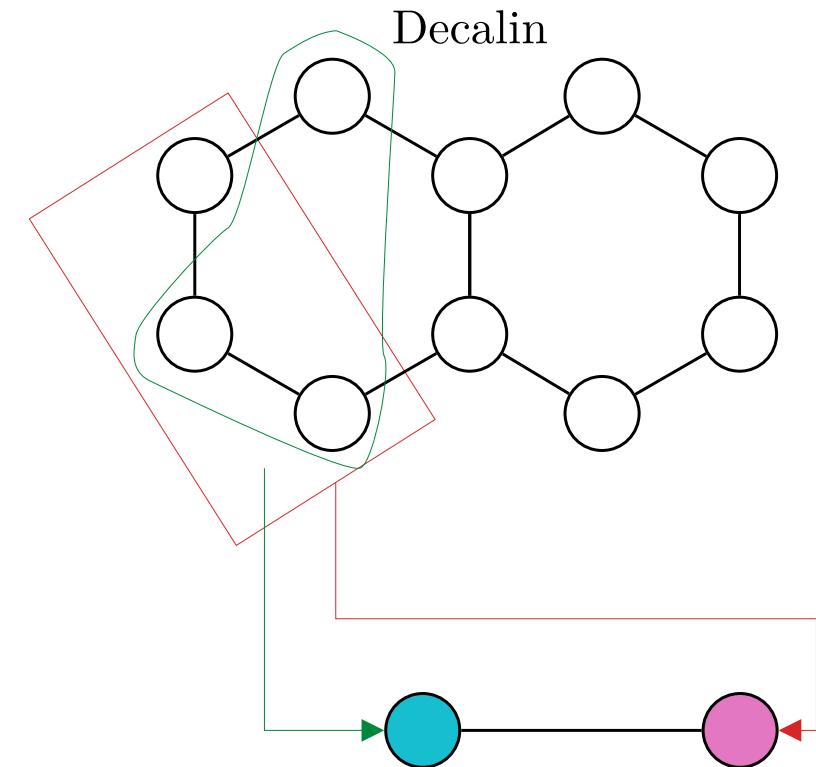
Proposition (Expressive Power of GIN). *GIN is as powerful as the Weisfeiler-Leman test.*

Higher-Order Weisfeiler-Leman Tests



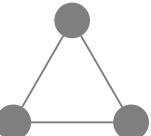
Higher-Order Weisfeiler-Leman Tests

Simplest improvement of Weisfeiler-Leman Graph Isomorphism Test



1. From the original graph G with nodes $V(G)$ and edges $E(G)$, create an auxiliary graph F .
 - $V(F) := \left\{ \mathbf{g} = \{g_i\}_{i=1}^k \mid g_i \in V(G) \right\} = V(G)^k$;
 - $E(F) := \left\{ (\mathbf{g}_1, \mathbf{g}_2) \mid d_H(\mathbf{g}_1, \mathbf{g}_2) = 1, \mathbf{g}_1, \mathbf{g}_2 \in V(G)^k \right\}$.
2. Give each k -tuple a label corresponding to the isomorphic type of the subgraph induced in the original graph.
3. Apply WL algorithm to the auxiliary graphs.

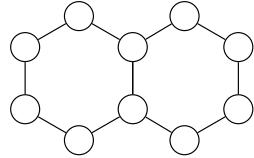
Isomorphic
Types:



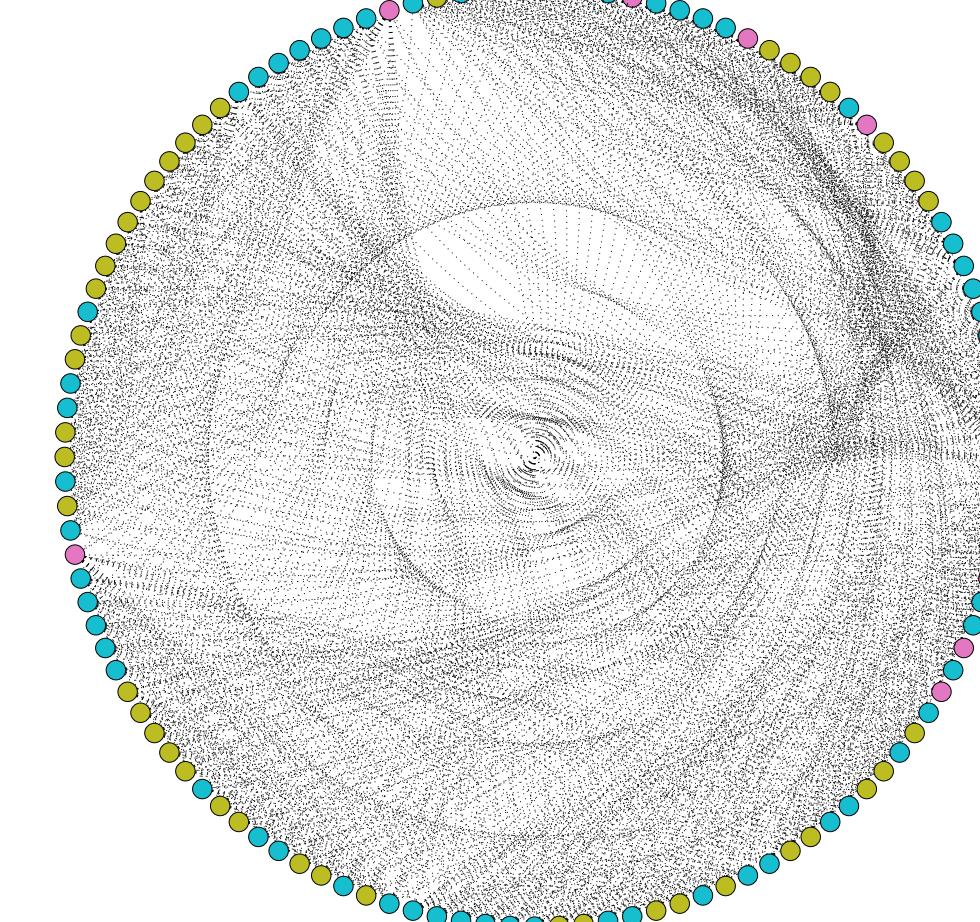
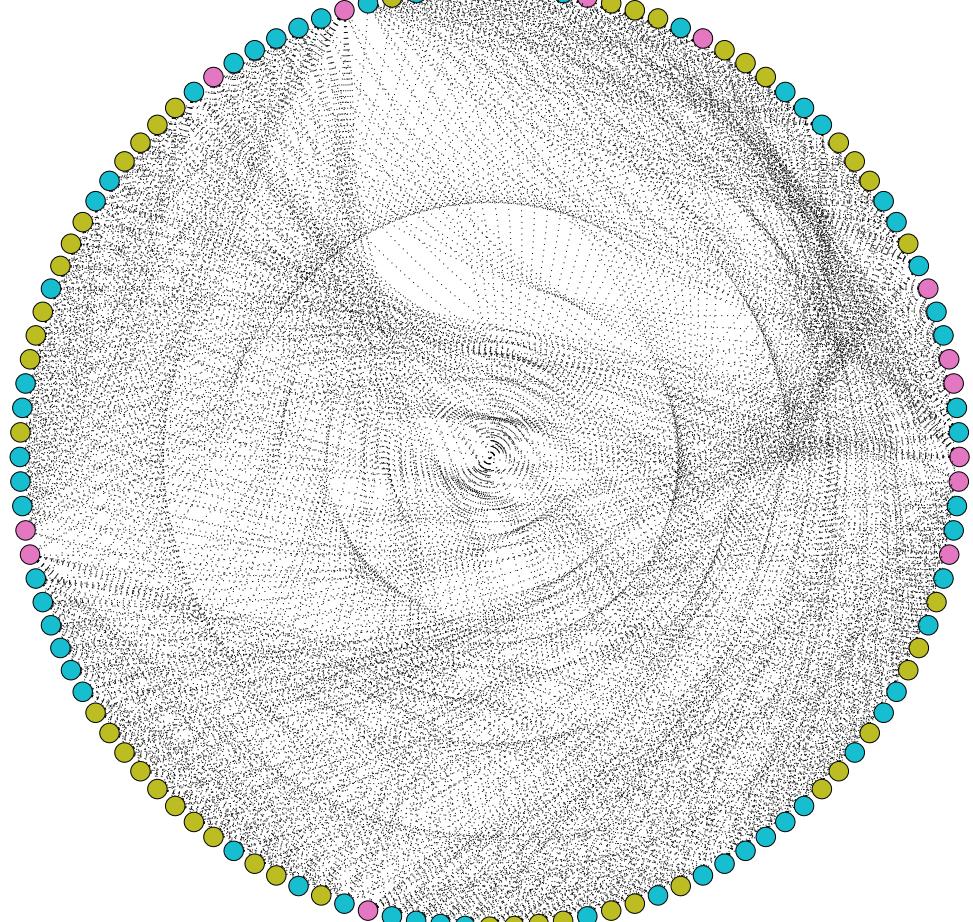
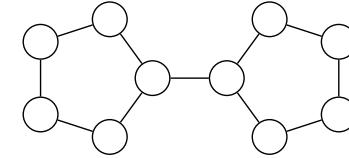
Higher-Order Weisfeiler-Leman Tests

Simplest improvement of Weisfeiler-Leman Graph Isomorphism Test

Decalin



Bicyclopentyl

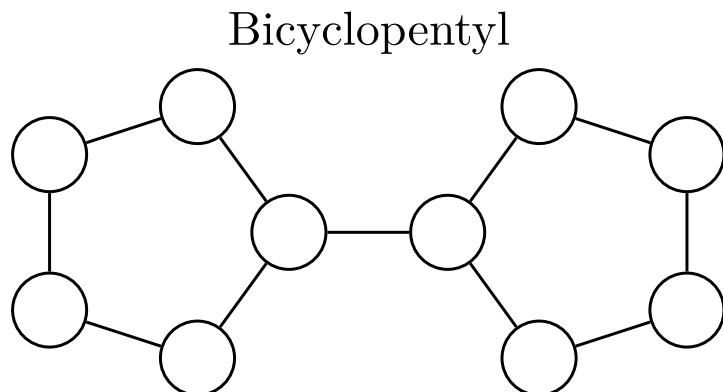
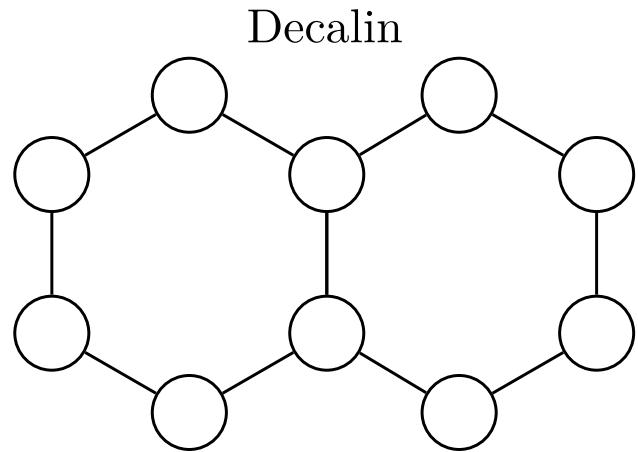


Loopy Weisfeiler-Leman



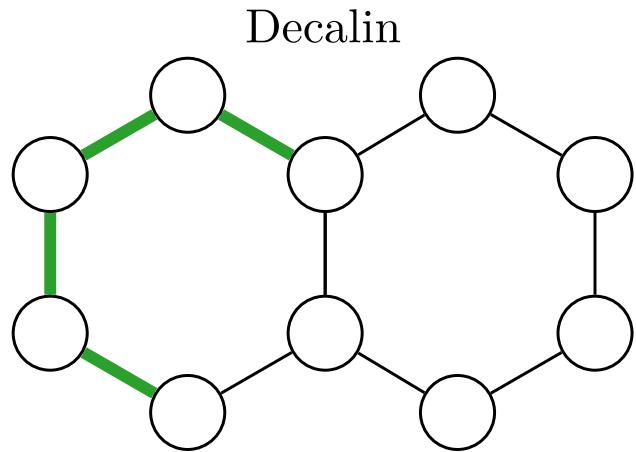
Loopy Weisfeiler Leman

Our approach to account for loops



Loopy Weisfeiler Leman

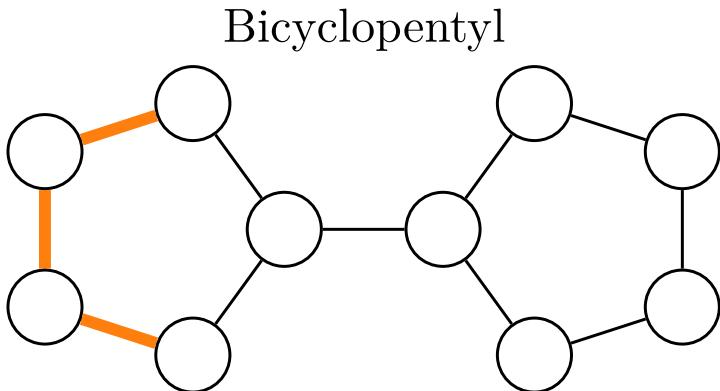
Our approach to account for loops



Definition (Simple Path). *Given a graph G , a simple path of length r is a collection $\mathbf{p} = \{p_i\}_{i=1}^{r+1}$ of $r + 1$ nodes such that consecutive nodes are adjacent, i.e.,*

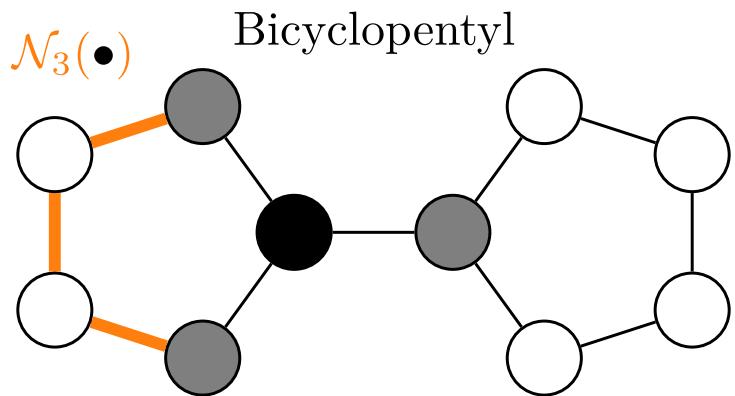
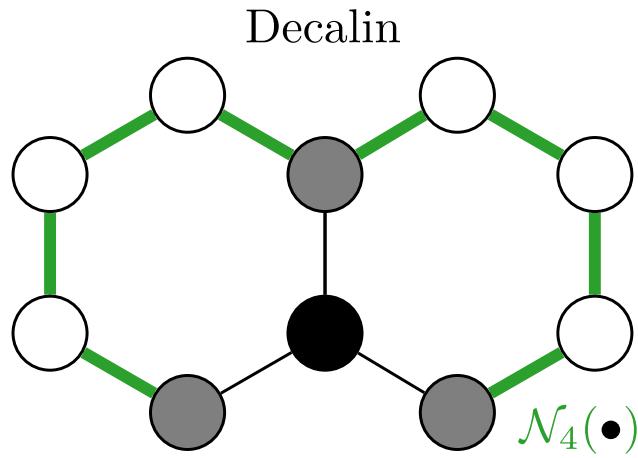
$$\{p_i, p_{i+1}\} \in E(G), \forall i \in \{1, \dots, r\},$$

and there are no repeated nodes, i.e., $i \neq j \implies p_i \neq p_j$.



Loopy Weisfeiler Leman

Our approach to account for loops



Definition (Simple Path). *Given a graph G , a simple path of length r is a collection $\mathbf{p} = \{p_i\}_{i=1}^{r+1}$ of $r + 1$ nodes such that consecutive nodes are adjacent, i.e.,*

$$\{p_i, p_{i+1}\} \in E(G), \forall i \in \{1, \dots, r\},$$

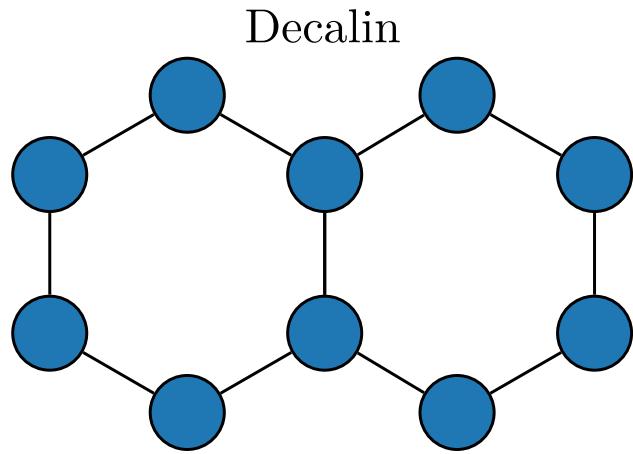
and there are no repeated nodes, i.e., $i \neq j \implies p_i \neq p_j$.

Definition (r -Neighborhood). *Given a graph G and an integer $r \geq 1$, we define the r -neighborhood $\mathcal{N}_r(v)$ of $v \in V(G)$ as the set of all simple paths of length r between distinct direct neighbors of v which do not contain v , i.e.,*

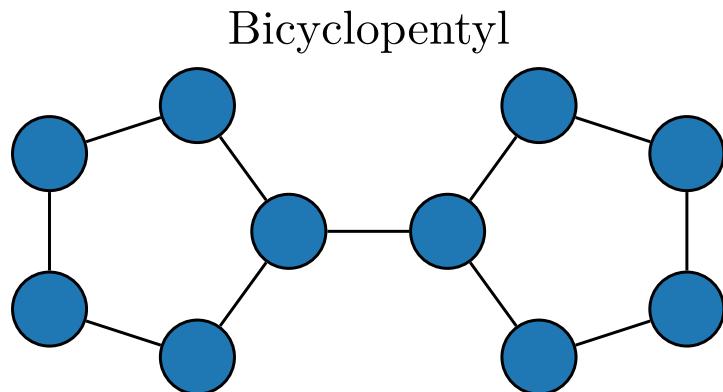
$$\mathcal{N}_r(v) := \{\mathbf{p} \mid \mathbf{p} \text{ } r\text{-path, } p_1, p_{r+1} \in \mathcal{N}(v), v \notin \mathbf{p}\}.$$

Loopy Weisfeiler Leman

Our approach to account for loops

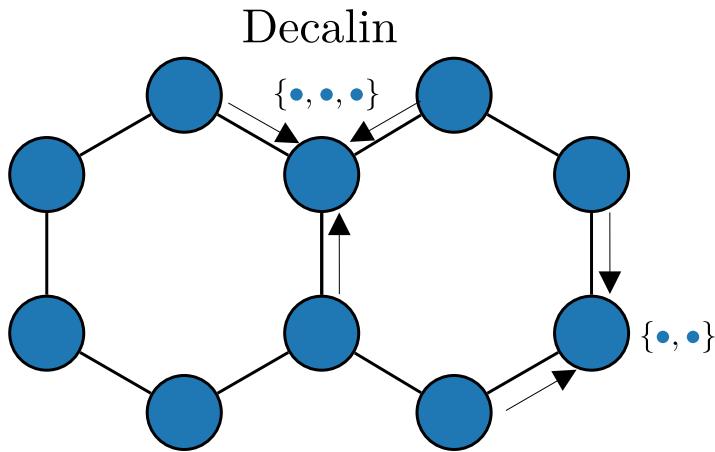


1. Assign to each node the same label •.

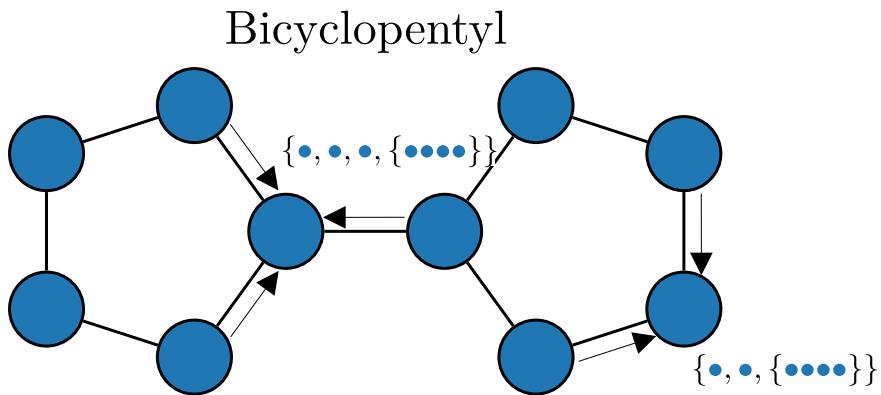


Loopy Weisfeiler Leman

Our approach to account for loops

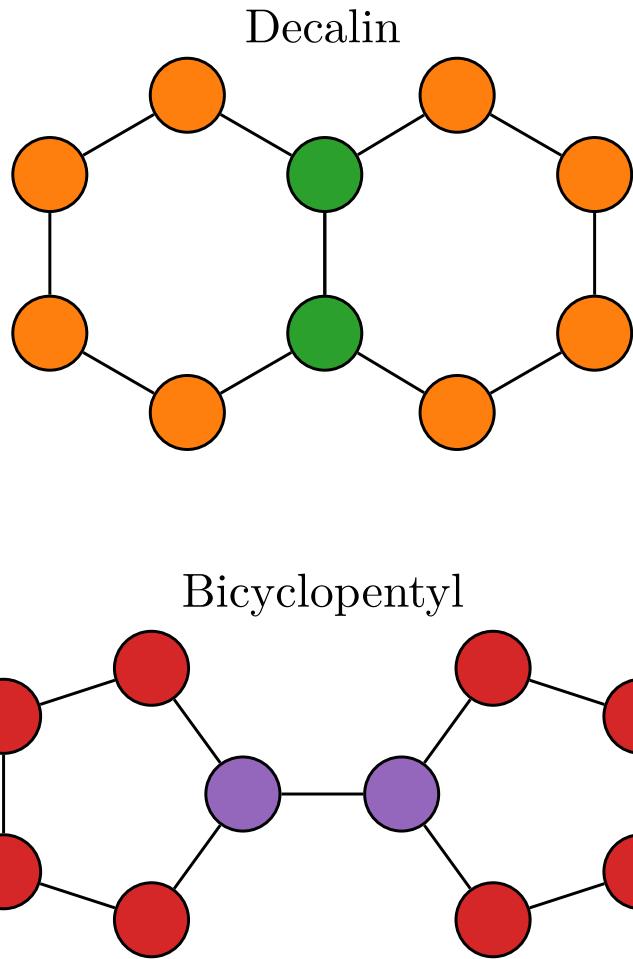


1. Assign to each node the same label \bullet .
2. Collect all labels from direct neighbors and r -neighbors.



Loopy Weisfeiler Leman

Our approach to account for loops



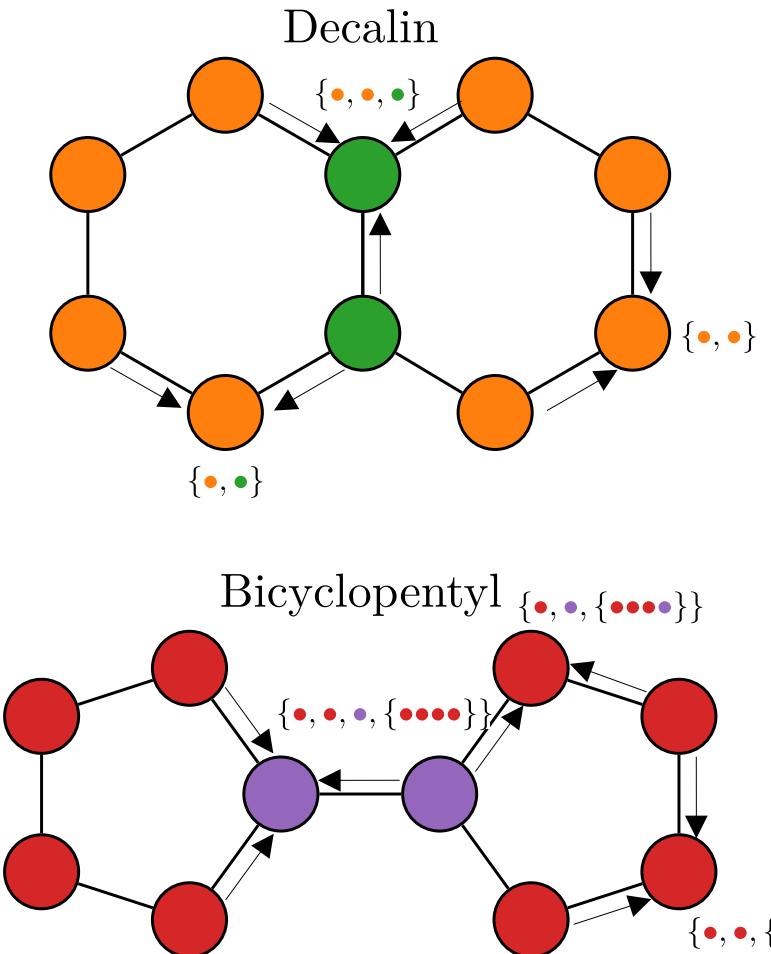
1. Assign to each node the same label \bullet .
2. Collect all labels from direct neighbors and r -neighbors.
3. Compress to get a new label

$$\begin{aligned}\{\bullet, \bullet\} &\mapsto \bullet \\ \{\bullet, \bullet, \bullet\} &\mapsto \bullet \\ \{\bullet, \bullet, \{\bullet, \bullet, \bullet\}\} &\mapsto \bullet \\ \{\bullet, \bullet, \bullet, \{\bullet, \bullet, \bullet\}\} &\mapsto \bullet\end{aligned}$$

The colours are already different, so we can stop the algorithm.

Loopy Weisfeiler Leman

Our approach to account for loops



1. Assign to each node the same label \bullet .
2. Collect all labels from direct neighbors and r -neighbors.
3. Compress to get a new label

$$\begin{aligned}\{\bullet, \bullet\} &\mapsto \bullet \\ \{\bullet, \bullet, \bullet\} &\mapsto \bullet \\ \{\bullet, \bullet, \{\bullet, \bullet, \bullet\}\} &\mapsto \bullet \\ \{\bullet, \bullet, \bullet, \{\bullet, \bullet, \bullet\}\} &\mapsto \bullet\end{aligned}$$

4. Repeat until there is no change in the partition of nodes with same label.

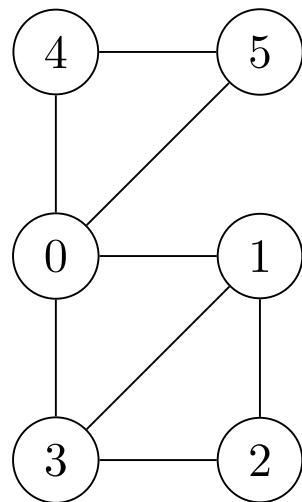
The colours are already different, so we can stop the algorithm.

GNN emulating Loopy Weisfeiler-Leman

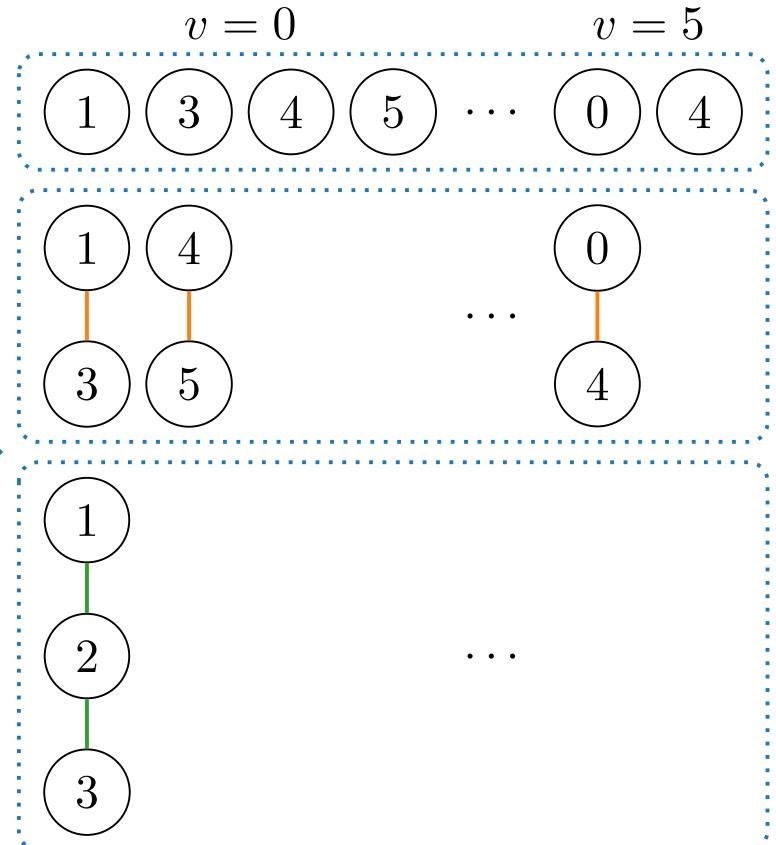


Loopy Graph Neural Network

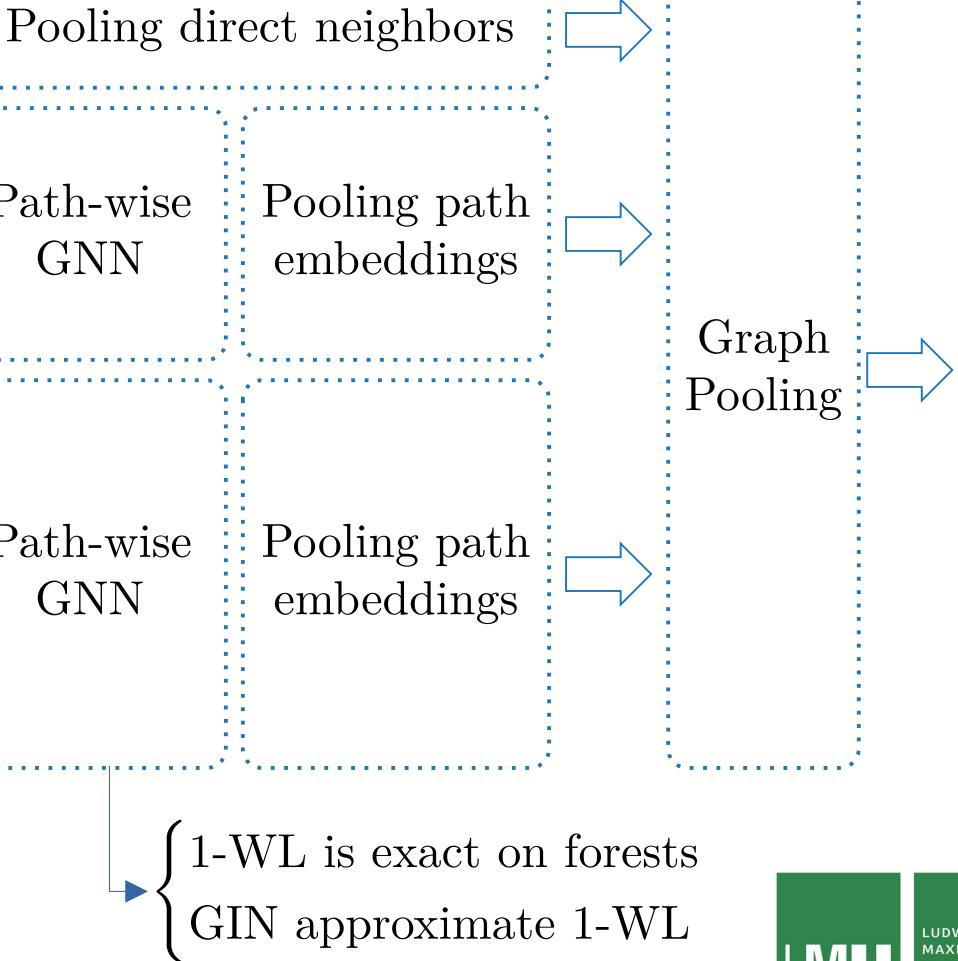
Raw graph



Preprocessing: extracting $\mathcal{N}_r(v)$



Training: paths-to-graph embedding



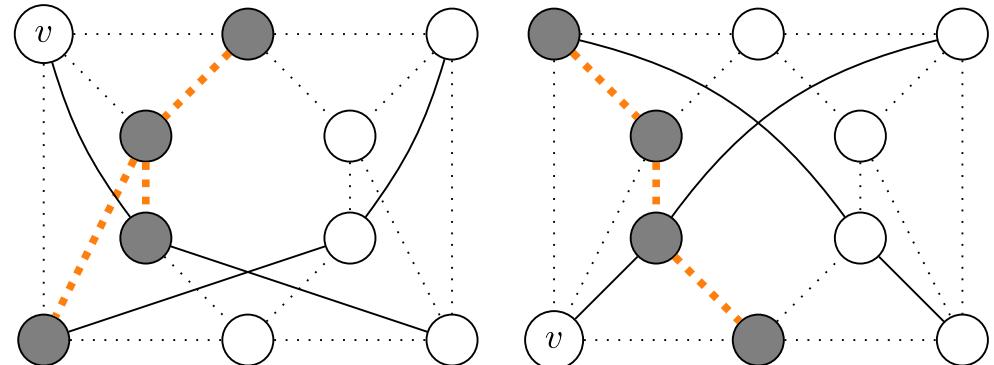
Expressivity of Loopy Weisfeiler-Leman



Isomorphism Expressivity

Standard Expressivity Measure

F



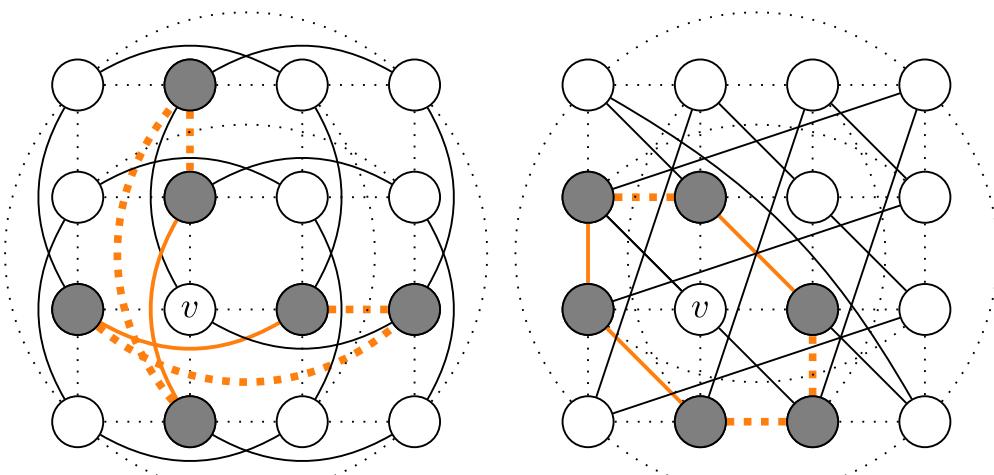
1-WL equivalent

G

→ Comparison with k -WL hierarchy.

Proposition (Hierarchy of r - ℓ WL). *Let $r > 0$.*

- i) r - ℓ WL is strictly more powerful than 1-WL;
- i) r - ℓ WL is strictly more powerful than $(r - 1)$ - ℓ WL;



4×4 rook graph

Shrikhande graph

3-WL equivalent

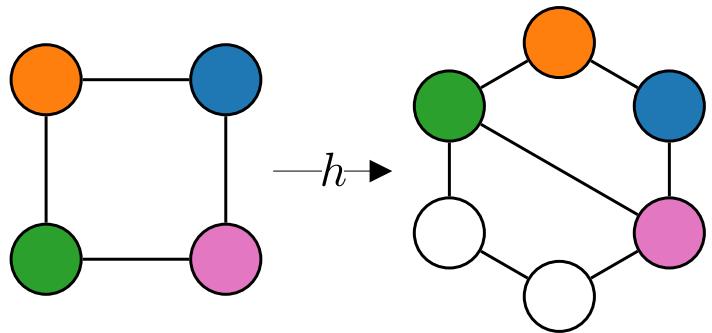
Advantages:

- Easy to analyze.

Disadvantages:

- Only qualitative measure.

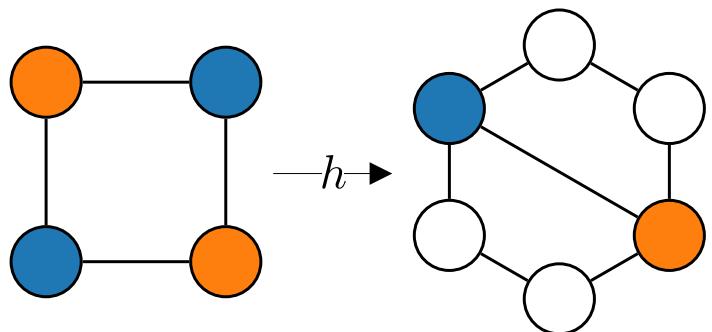
Subgraph Expressivity



Definition (Subgraph Isomorphism). A subgraph isomorphism is an injective homomorphism. The set of subgraph isomorphisms from F to G is denoted by $\text{Sub}(F, G)$, and its cardinality by $\text{sub}(F, G) = |\text{Sub}(F, G)|$.

Homomorphism Expressivity

Quantitative Expressivity Measure: Counting Patterns



Definition (Homomorphism). Let F and G be two graphs. A homomorphism from F to G is a map $h : V(F) \rightarrow V(G)$ such that

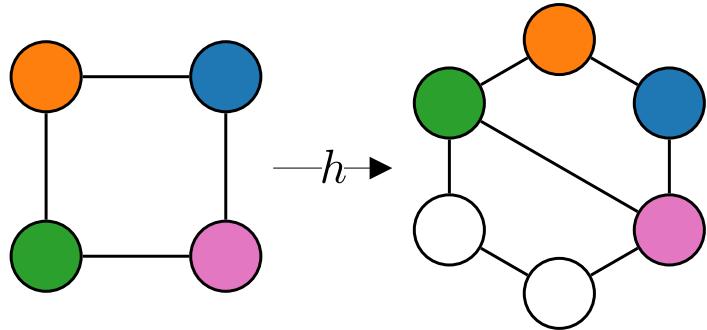
$$\{u, v\} \in E(F) \implies \{h(u), h(v)\} \in E(G).$$

The set of homomorphisms from F to G is denoted by $\text{Hom}(F, G)$, and its cardinality by $\text{hom}(F, G) := |\text{Hom}(F, G)|$.

→ Counting homomorphism is a complete isomorphic measure

$$G \cong H \iff \text{hom}(F, G) = \text{hom}(F, H) \quad \forall F$$

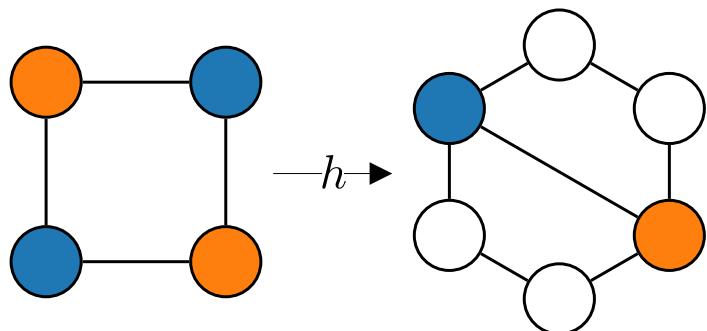
Subgraph Expressivity



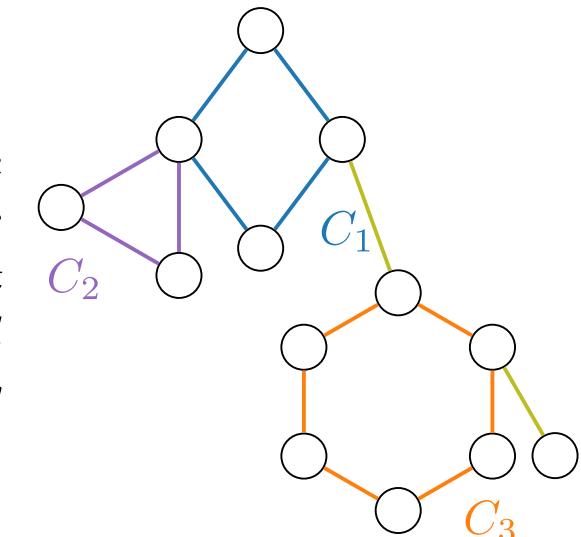
Proposition (Subgraph Counting Power of r - ℓ WL). *For any $r \geq 1$, r - ℓ WL can subgraph-count all cycles with at most $r + 2$ nodes.*

Homomorphism Expressivity

Quantitative Expressivity Measure: Counting Patterns



Definition (Cactus Graph). A cactus graph is a graph where every edge lies on at most one simple cycle. For $r \geq 2$, an r -cactus graph is a cactus where every simple cycle has at most r vertices. We denote by \mathcal{M} the set of all cactus graphs, and by \mathcal{M}^r the set comprising all q -cactus graphs for $q \leq r$.

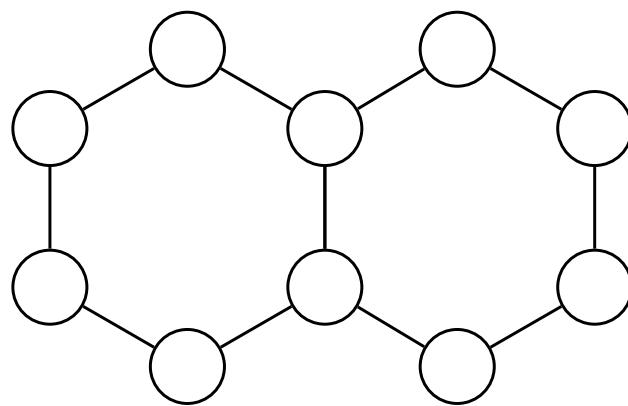


Proposition (Homomorphism Counting Power of r - ℓ WL). *Let $r \geq 0$. Then, r - ℓ WL can homomorphism-count \mathcal{M}^{r+2} .*

Subgraph Expressivity

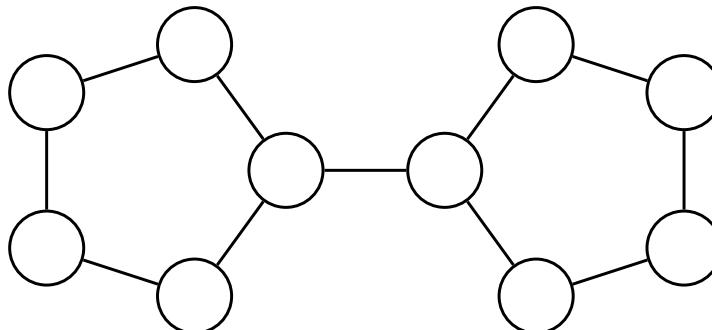
Homomorphism Expressivity

Decalin

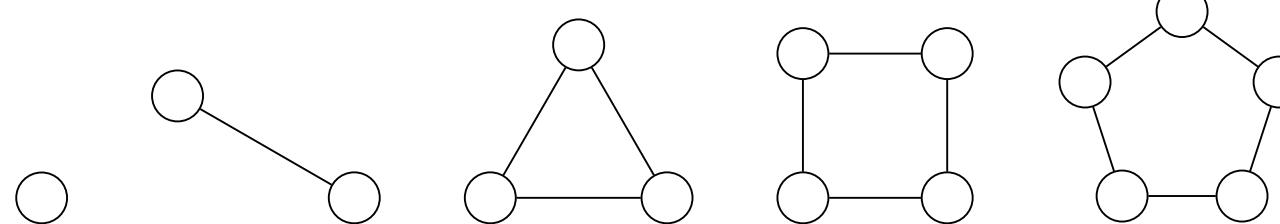


10	11	0	0	0
10	11	0	78	0

Bicyclopentyl



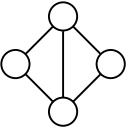
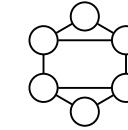
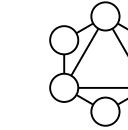
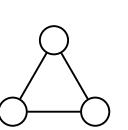
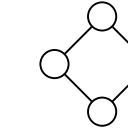
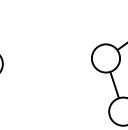
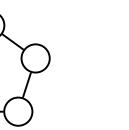
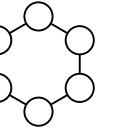
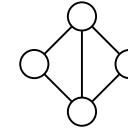
10	11	0	0	2
10	11	0	78	20



Patterns

Subgraph Expressivity

Homomorphism Expressivity

Model	hom(F, G)			sub(F, G)					
									
MPNN	0.300	0.233	0.254	0.358	0.208	0.188	0.146	0.261	0.205
Subgraph GNN	0.011	0.015	0.012	0.010	0.020	0.024	0.046	0.007	0.027
Local 2-GNN	0.008	0.008	0.010	0.008	0.011	0.017	0.034	0.007	0.016
Local 2-FGNN	0.003	0.005	0.004	0.003	0.004	0.010	0.020	0.003	0.010
$r\text{-}\ell\text{GIN}$	0.001 (r=2)	0.006 (r=3)	0.009 (r=3)	0.0005 (r=1)	0.0005 (r=2)	0.0003 (r=3)	0.0003 (r=4)	0.001 (r=2)	0.0004 (r=3)

Experiments on Molecular Datasets



Experiments on ZINC

Model	ZINC12K	ZINC250K
GIN	0.163 ± 0.004	0.088 ± 0.002
GCN	0.321 ± 0.009	-
GAT	0.384 ± 0.007	-
GSN	0.115 ± 0.012	-
CIN	0.079 ± 0.006	0.022 ± 0.002
NestedGNN	0.111 ± 0.003	0.029 ± 0.001
SUN	0.083 ± 0.003	-
GNNAK	0.080 ± 0.001	-
I2-GNN	0.083 ± 0.001	0.023 ± 0.001
DRFWL	0.077 ± 0.002	0.025 ± 0.003
SignNet	0.084 ± 0.004	0.024 ± 0.003
HIMP	0.151 ± 0.006	0.036 ± 0.002
PathNN	0.090 ± 0.004	-
5- ℓ GIN	0.072 ± 0.002	0.022 ± 0.001

Experiments on QM9

Target	Model							
	1-GNN	1-2-3-GNN	DTNN	DeepLRP	NestedGNN	I2-GNN	DRFFWL	5- ℓ GIN
μ	0.493	0.476	0.244	0.364	0.428	0.428	0.346	0.350 ± 0.011
α	0.78	0.27	0.95	0.298	0.290	0.230	0.222	0.217 ± 0.025
$\varepsilon_{\text{homo}}$	0.00321	0.00337	0.00388	0.00254	0.00265	0.00261	0.00226	0.00205 ± 0.00005
$\varepsilon_{\text{lumo}}$	0.00355	0.00351	0.00512	0.00277	0.00297	0.00267	0.00225	0.00216 ± 0.00004
$\Delta(\varepsilon)$	0.0049	0.0048	0.0112	0.00353	0.0038	0.0038	0.00324	0.00321 ± 0.00014
R^2	34.1	22.9	17.0	19.3	20.5	18.64	15.04	13.21 ± 0.19
ZVPE	0.00124	0.00019	0.00172	0.00055	0.0002	0.00014	0.00017	0.000127 ± 0.000003
U_0	2.32	0.0427	2.43	0.413	0.295	0.211	0.156	0.0418 ± 0.0520
U	2.08	0.111	2.43	0.413	0.361	0.206	0.153	0.023 ± 0.023
H	2.23	0.0419	2.43	0.413	0.305	0.269	0.145	0.0352 ± 0.0304
G	1.94	0.0469	2.43	0.413	0.489	0.261	0.156	0.0118 ± 0.0015
C_v	0.27	0.0944	2.43	0.129	0.174	0.0730	0.0901	0.0702 ± 0.0024

Thank you!

