

# Optimizing over trained GNNs via symmetry breaking

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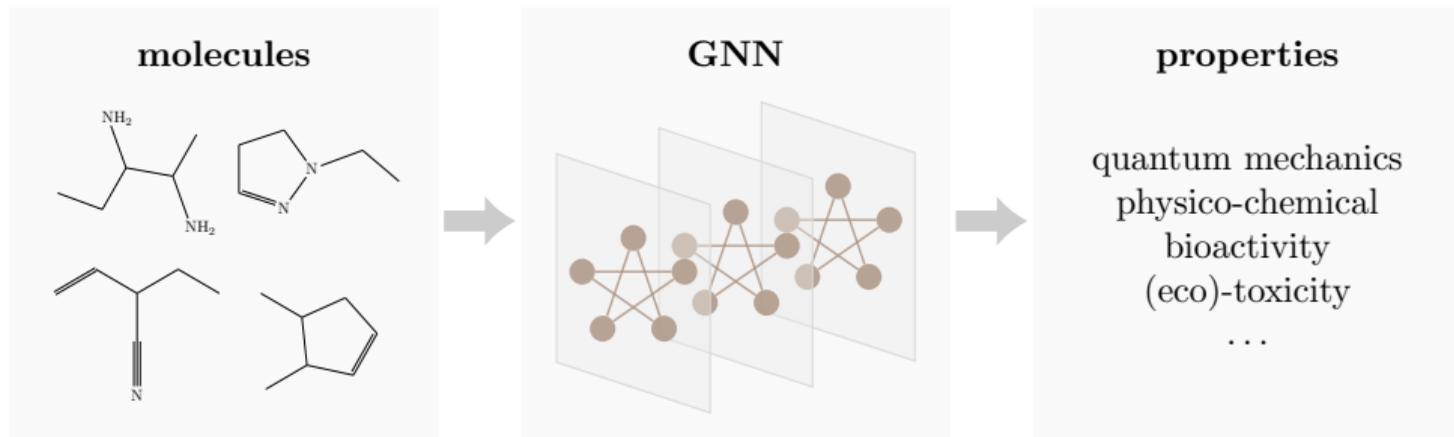
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# Motivation: Forward & Backward problems over GNNs

Prediction (Forward): What are the properties for a given molecule?



Optimization (Backward): What is the optimal molecule with desired properties?

## Problem definition

Given a trained GNN, we aim to find the input with optimal property <sup>1</sup>:

$$\begin{aligned}(X^*, A^*) &= \arg \min_{(X, A)} GNN(X, A) \\ &s.t. \quad f_j(X, A) \leq 0, j \in \mathcal{J} \\ &\quad \quad g_k(X, A) = 0, k \in \mathcal{K}\end{aligned}$$

where  $X$  denotes features,  $A$  is the adjacency matrix of input graph,  $f_j, g_k$  are problem-specific constraints, and  $\mathcal{J}, \mathcal{K}$  are index sets.

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<sup>1</sup>Optimality is defined on this given GNN instead of true properties.

# Symmetry issue

## Observation

GNN is permutation invariant<sup>2</sup>: isomorphic graphs have the same output.

## Good for training

Different indexing of a graph data will not influence its output.

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<sup>2</sup>In this work, we only consider GNNs that are permutation invariant.

# Symmetry issue

## Observation

GNN is permutation invariant<sup>2</sup>: isomorphic graphs have the same output.

## Good for training

Different indexing of a graph data will not influence its output.

## Bad for optimization

Each graph indexing corresponds to a solution, which significantly enlarges the searching space.

For example, there are  $4! = 24$  different indexing for this molecule:



<sup>2</sup>In this work, we only consider GNNs that are permutation invariant.

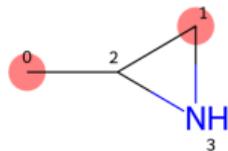
## Symmetry-breaking constraints I<sup>3</sup>

Each node (except 0) should be linked with a node with smaller index:

$$\forall v \in [N] \setminus \{0\}, \exists u < v, \text{ s.t. } A_{u,v} = 1 \quad (\text{S1})$$

i.e., the subgraph induced by nodes  $\{0, 1, \dots, v\}$  is connected.

10 out of 24 solutions violate (S1), for example:



Node 1 is not linked with node 0.

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<sup>3</sup>N: number of nodes.

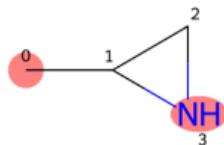
## Symmetry-breaking constraints II<sup>4</sup>

Node 0 has the minimal function value under a designed hierarchical function  $h : \mathbb{R}^F \rightarrow \mathbb{R}$  defined over features:

$$h(X_0) \leq h(X_v), \forall v \in [N] \setminus \{0\} \quad (\text{S2})$$

i.e., node 0 has the most "special" features under the action of  $h$ .

11 out of 14 solutions violate (S2), for example:



Construct  $h$  such that  $h(N) < h(C)$ , then the nitrogen atom should be indexed 0.

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<sup>4</sup> $F$ : number of features.

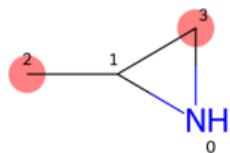
## Symmetry-breaking constraints III<sup>5</sup>

The neighbor set of a node with smaller index has smaller lexicographical order:

$$LO(\mathcal{N}(v) \setminus \{v+1\}) \leq LO(\mathcal{N}(v+1) \setminus \{v\}), \forall v \in [N-1] \setminus \{0\} \quad (\text{S3})$$

i.e., node  $v$  has "stronger" neighbors comparing to node  $v+1$ .

2 out of 3 solutions violate (S3), for example:



$$\mathcal{N}(2) = \{1\}, \mathcal{N}(3) = \{0, 1\}$$

$$LO(\mathcal{N}(2) \setminus \{3\}) > LO(\mathcal{N}(3) \setminus \{2\})$$

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<sup>5</sup> $\mathcal{N}(\cdot)$ : neighbor set.  $LO(\cdot)$ : lexicographical order.

# Do these constraints reduce the diversity of the feasible set?

Algorithm 1 yields at least one feasible indexing for any graph (see proofs in the paper).

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## Algorithm 1 Indexing algorithm

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**Input:**  $G = (V, E)$  with node set  $V = \{v_0, v_1, \dots, v_{N-1}\}$  ( $N := |V|$ ). Denote the neighbor set of node  $v$  as  $\mathcal{N}(v), \forall v \in V$ .

$\mathcal{I}(v_0) \leftarrow 0$  ▷ Assume that  $v_0$  is indexed with 0  
 $s \leftarrow 1$  ▷ Index for next node  
 $V_1^1 \leftarrow \{v_0\}$  ▷ Initialize set of indexed nodes

**while**  $s < N$  **do**

$V_2^s \leftarrow V \setminus V_1^s$  ▷ Set of unindexed nodes  
 $\mathcal{N}^s(v) \leftarrow \{\mathcal{I}(u) \mid u \in \mathcal{N}(v) \cap V_1^s\}, \forall v \in V_2^s$  ▷ Obtain all indexed neighbors  
 $rank^s(v) \leftarrow |\{LO(\mathcal{N}^s(u)) < LO(\mathcal{N}^s(v)) \mid \forall u \in V_2^s\}|, \forall v \in V_2^s$  ▷ Assign a rank to each unindexed node

$\mathcal{I}^s(v) \leftarrow \begin{cases} \mathcal{I}(v), & \forall v \in V_1^s \\ rank^s(v) + s, & \forall v \in V_2^s \end{cases}$  ▷ Assign temporary indexes

$\mathcal{N}_t^s(v) \leftarrow \{\mathcal{I}^s(u) \mid u \in \mathcal{N}(v)\}, \forall v \in V_2^s$  ▷ Define temporary neighbor sets based on  $\mathcal{I}^s$   
 $v^s \leftarrow \arg \min_{v \in V_2^s} LO(\mathcal{N}_t^s(v))$  ▷ Neighbors of  $v^s$  has minimal order

▷ If multiple nodes share the same minimal order, arbitrarily choose one  
▷ Index  $s$  to node  $v^s$

$\mathcal{I}(v^s) = s$  ▷ Add  $v^s$  to set of indexed nodes  
 $V_1^{s+1} \leftarrow V_1^s \cup \{v^s\}$  ▷ Next index is  $s + 1$   
 $s \leftarrow s + 1$

**end while**

**Output:**  $\mathcal{I}(v), v \in V$  ▷ Result indexing

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## Mixed-integer formulation for GNNs

Since the input graph structure is not fixed, all elements in the adjacency matrix are variables:

$$\mathbf{x}_v^{(l)} = \sigma \left( \sum_{u \in V} e_{u \rightarrow v} \mathbf{w}_{u \rightarrow v}^{(l)} \mathbf{x}_u^{(l-1)} + \mathbf{b}_v^{(l)} \right)$$

where

- $\mathbf{x}_v^{(l)}$ : (continuous or discrete) **variables**, the features of node  $v$  in  $l$ -th layer.
- $e_{u \rightarrow v}$ : binary **variable**, the existence of edge  $u \rightarrow v$ .
- $\mathbf{w}_{u \rightarrow v}^{(l)}, \mathbf{b}_v^{(l)}$ : **constants**, weights and biases of  $l$ -th layer.

Bilinear terms  $e_{u \rightarrow v} \mathbf{x}_u^{(l-1)}$  result in a mixed-integer quadratically constrained optimization problem (MIQCP), which can be handled by state-of-the-art solvers such as Gurobi.

Alternatively, they can be reformulated in a linear way using big-M formulation.

# Numerical results

Optimal molecular design:

- atom  $\rightarrow$  node, bond  $\rightarrow$  edge
- atom type, #neighbors, ...  $\rightarrow$  features
- chemical requirements  $\rightarrow$  constraints

Our numerical results show that:

- Symmetry-breaking constraints significantly reduce the searching space.
- After breaking symmetry, the solving time is largely decreased.

Table 1: Numbers of feasible solutions for QM7.

$N$	(S1)	(S1) - (S2)	(S1) - (S3)
4	3,323	726	416
5	67,020	11,747	3,003
6	> 2,500,000	<b>443,757</b>	<b>50,951</b>
7	> 2,500,000	> 2,500,000	504,952

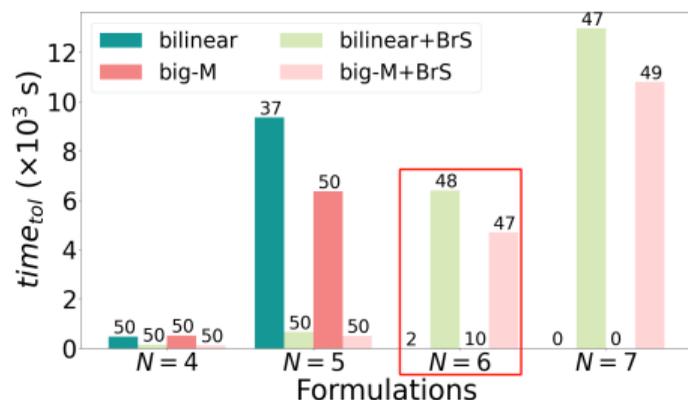


Figure 1: Average solving time over 50 runs.

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*Thanks for your attention!*

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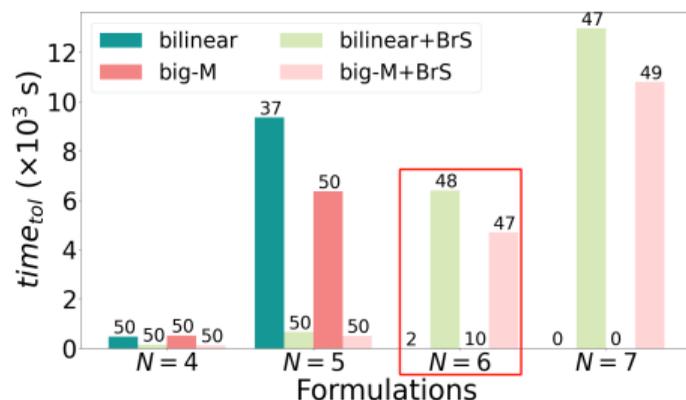


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