

# INVERSE PROBLEMS OVER TRAINED GRAPH NEURAL NETWORKS

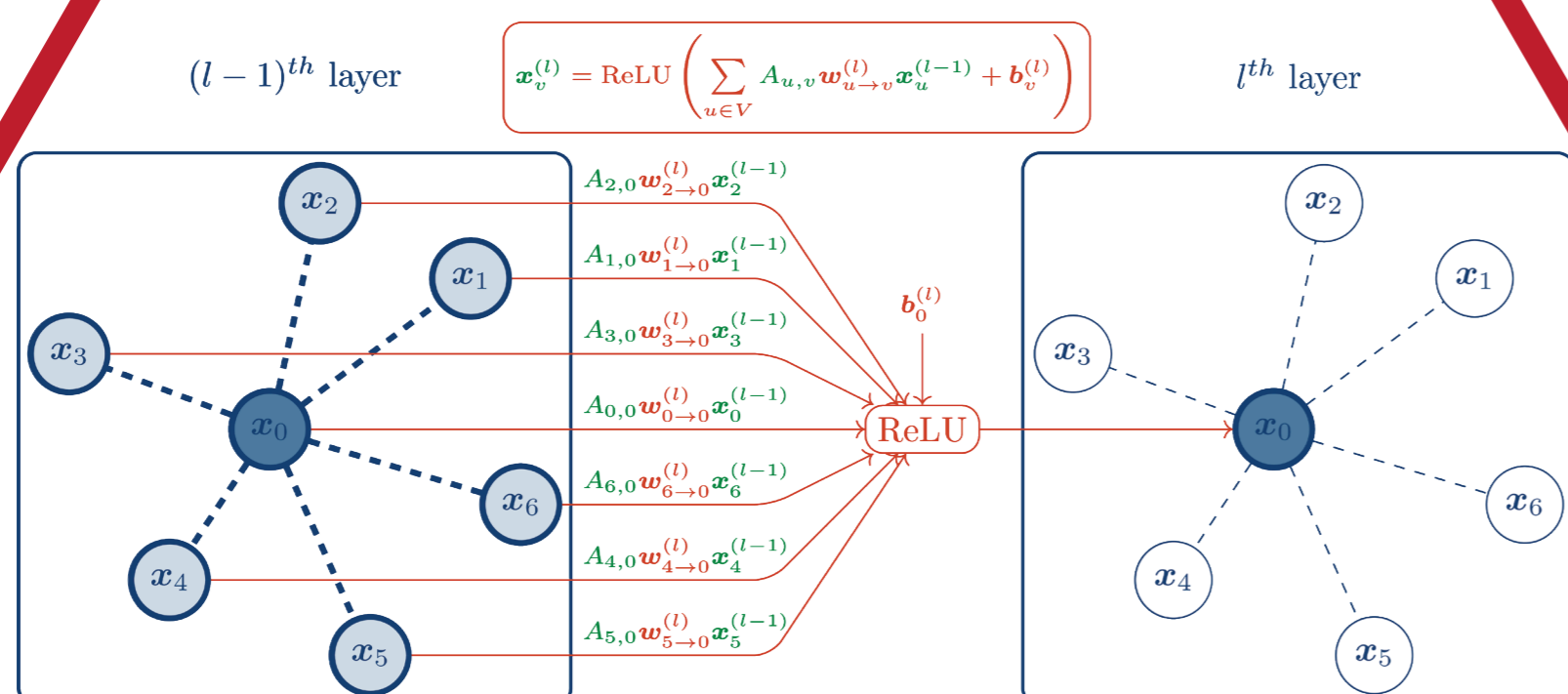
## OPTIMIZATION, SYMMETRY-BREAKING, AND APPLICATIONS

Shiqiang Zhang<sup>1\*</sup>, Juan S. Campos<sup>1</sup>, Christian W. Feldmann<sup>2</sup>,  
Darren Budd<sup>3</sup>, Frederik Sandfort<sup>2</sup>, Miriam Mathea<sup>2</sup>, Ruth Misener<sup>1</sup>  
\*s.zhang21@imperial.ac.uk

<sup>1</sup>Imperial College London, London, SW7 2AZ, UK  
<sup>2</sup>BASF SE, Ludwigshafen am Rhein, 67056, Germany  
<sup>3</sup>BASF plc, Stockport, SK1 3GG, UK

### Optimization

A GNN layer with unknown graph:



How to encode a GNN to an optimization problem?  
(without doing math and programming)



OMLT automatically encodes a trained GNN to its corresponding optimization formulation

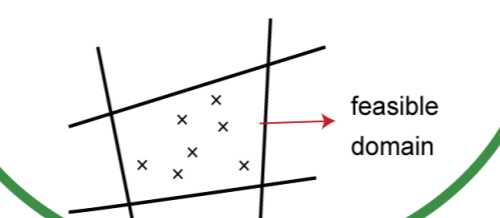
#### Requirements

- size of molecules
- types of atoms
- composition of molecules
- substructure inclusion
- substructure exclusion
- .....

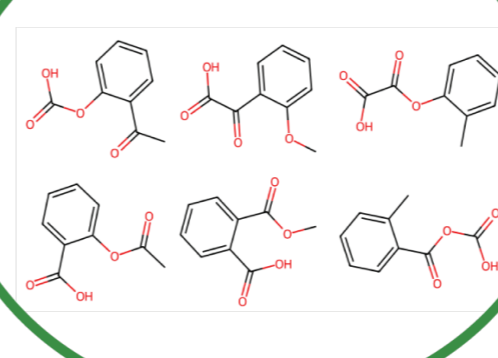


#### Optimization

- Variables (binary): features
- Constraints (linear): requirements



#### Molecules



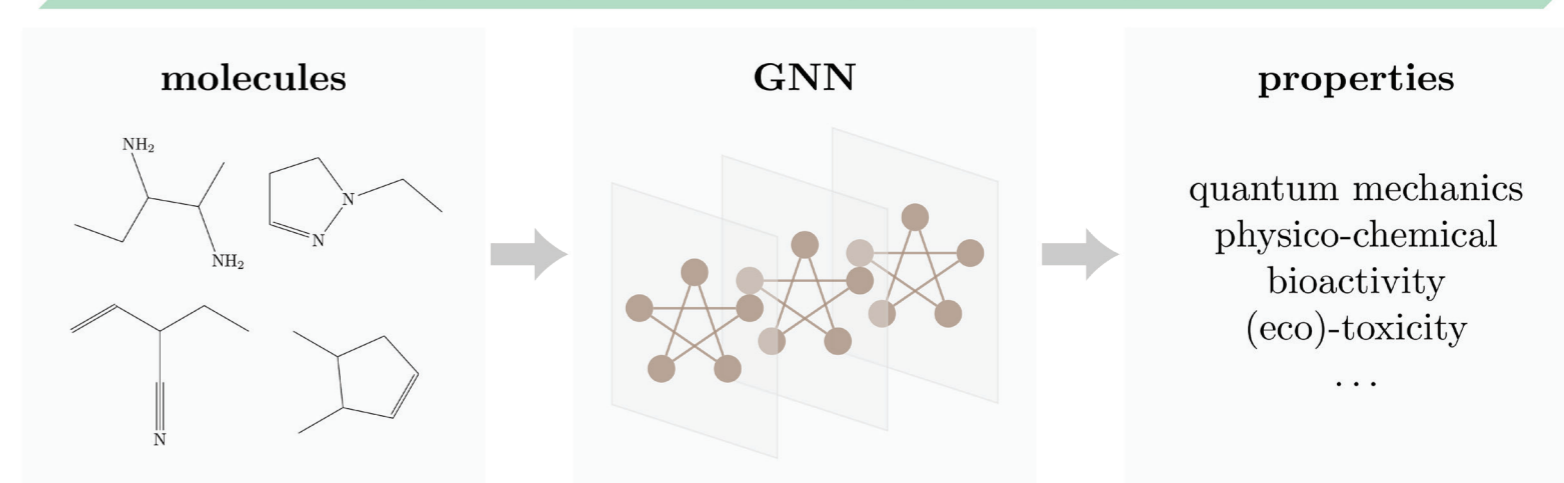
### Molecular Design

How to find desired molecules?

$$\min_{X,A} GNN(X,A), \text{ s.t. } f(X,A) \leq 0$$

features, graph

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



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

#### Part I: Optimal molecular design

data-driven, property-oriented  
structure-exploited, validity-ensured

#### Part II: Molecular generation

try Limeade!  
end-to-end, lightweight generator

### Symmetry Breaking

#### Symmetry issue!

One graph with N nodes has N! different indexing

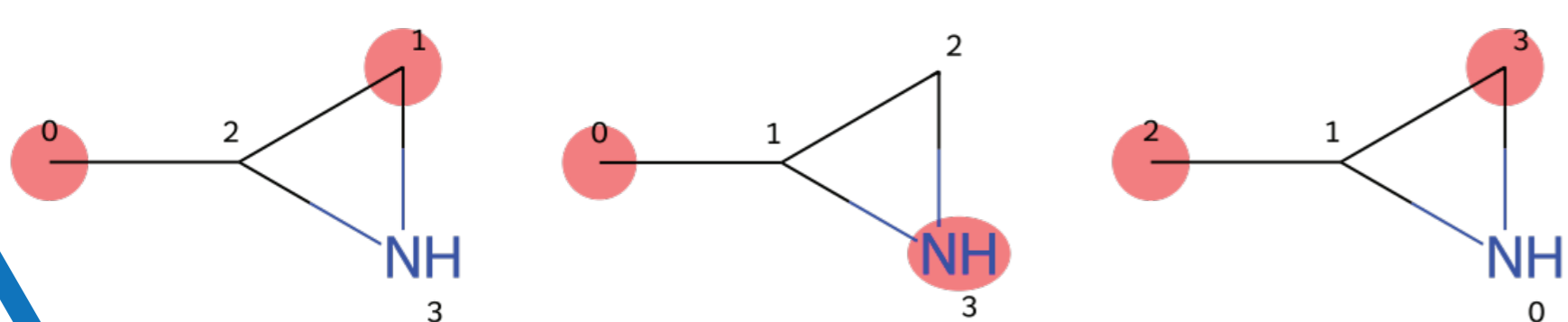
#### Symmetry-breaking constraints

(S1): The subgraph induced by nodes  $\{0,1,\dots,v\}$  is connected.

(S2): Node 0 has the most special features.

(S3): Node v has neighbors with smaller indexes compared to v+1.

#### Solutions removed by (S1) – (S3):



#### Generally applicable

suitable for any symmetry issue caused by graph isomorphism

#### Empirically powerful

remove >98% symmetries for molecular benchmarks

#### Theoretically guaranteed

at least one indexing is feasible given any graph

### AI Safety

#### GNNs are vulnerable!

Small input changes could lead to wrong predictions

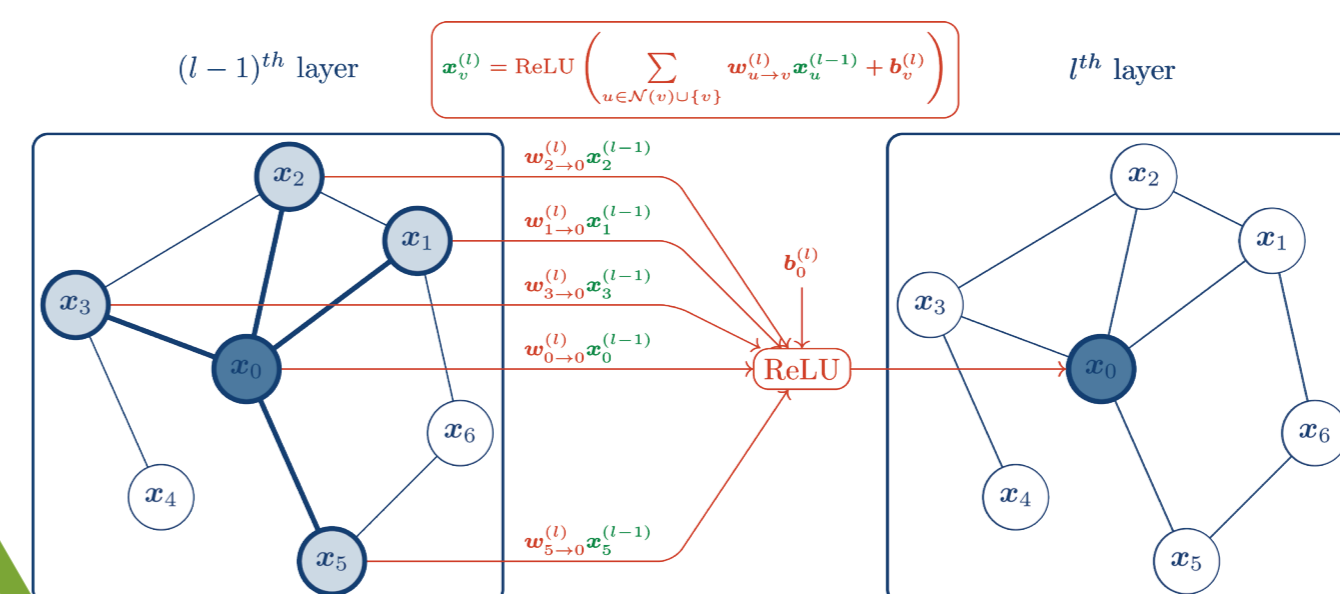
How to certify the robustness of a GNN?

$$m(c^*, c) := \min_{(X,A)} GNN_{c^*}(X,A) - GNN_c(X,A)$$

true label      correct prediction      wrong prediction

$$\text{s.t. } X \in \mathcal{P}(X^*), A \in \mathcal{P}(A^*)$$

robust      non-robust      perturb feature      perturb graph



Topology-based bounds tightening  
more attacks, tighter bounds, faster verification