

DeepMind

Everything is Connected

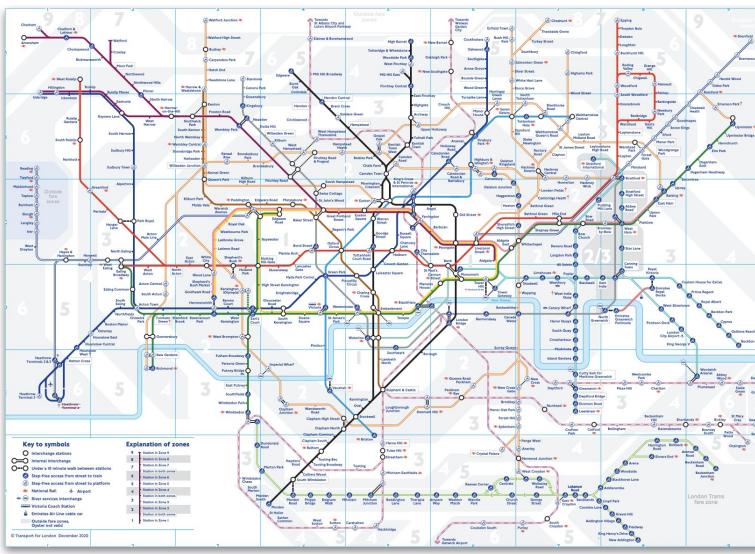
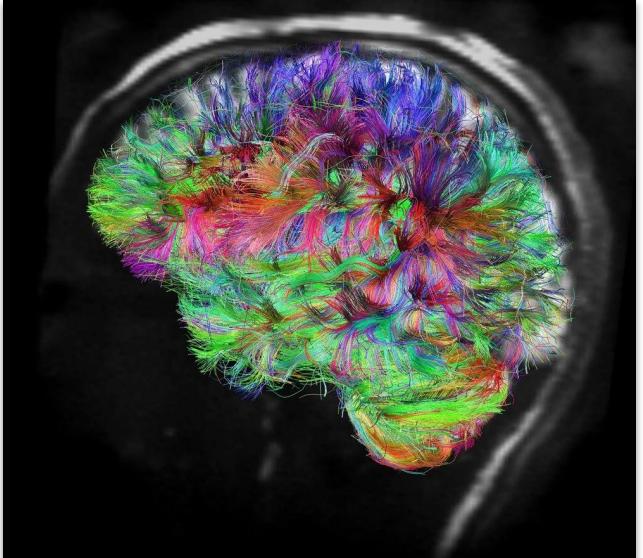
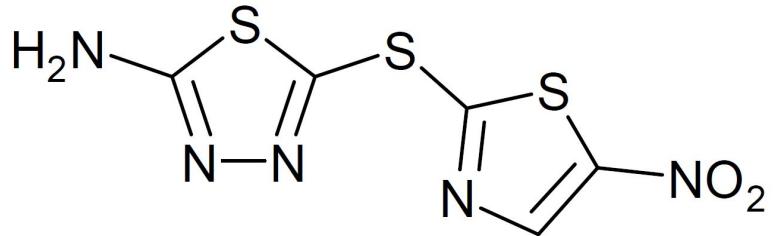
Deep Learning on Graphs

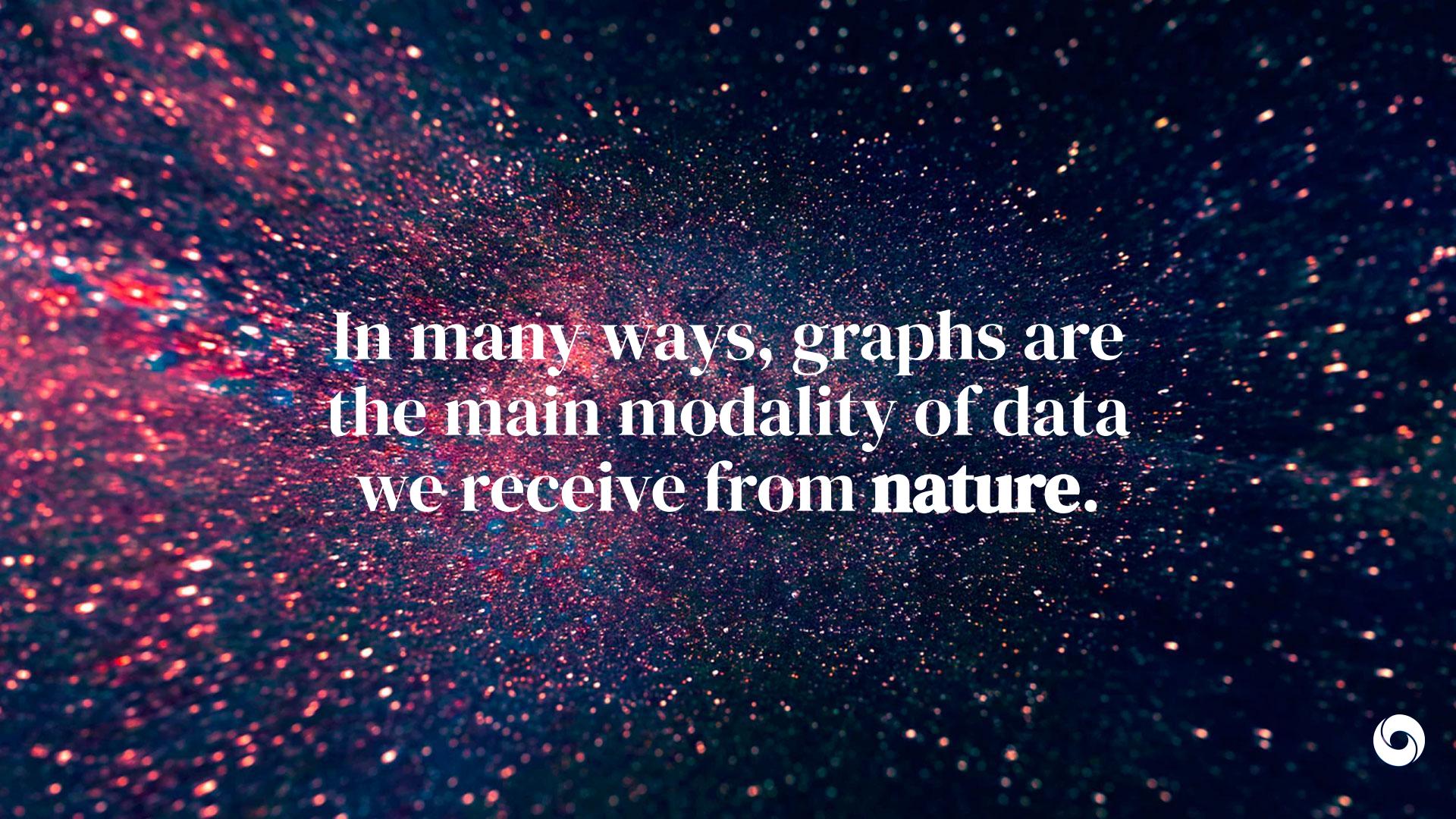
Petar Veličković

MLinPL x CUAI Cambridge Pre-meeting
30 October 2021



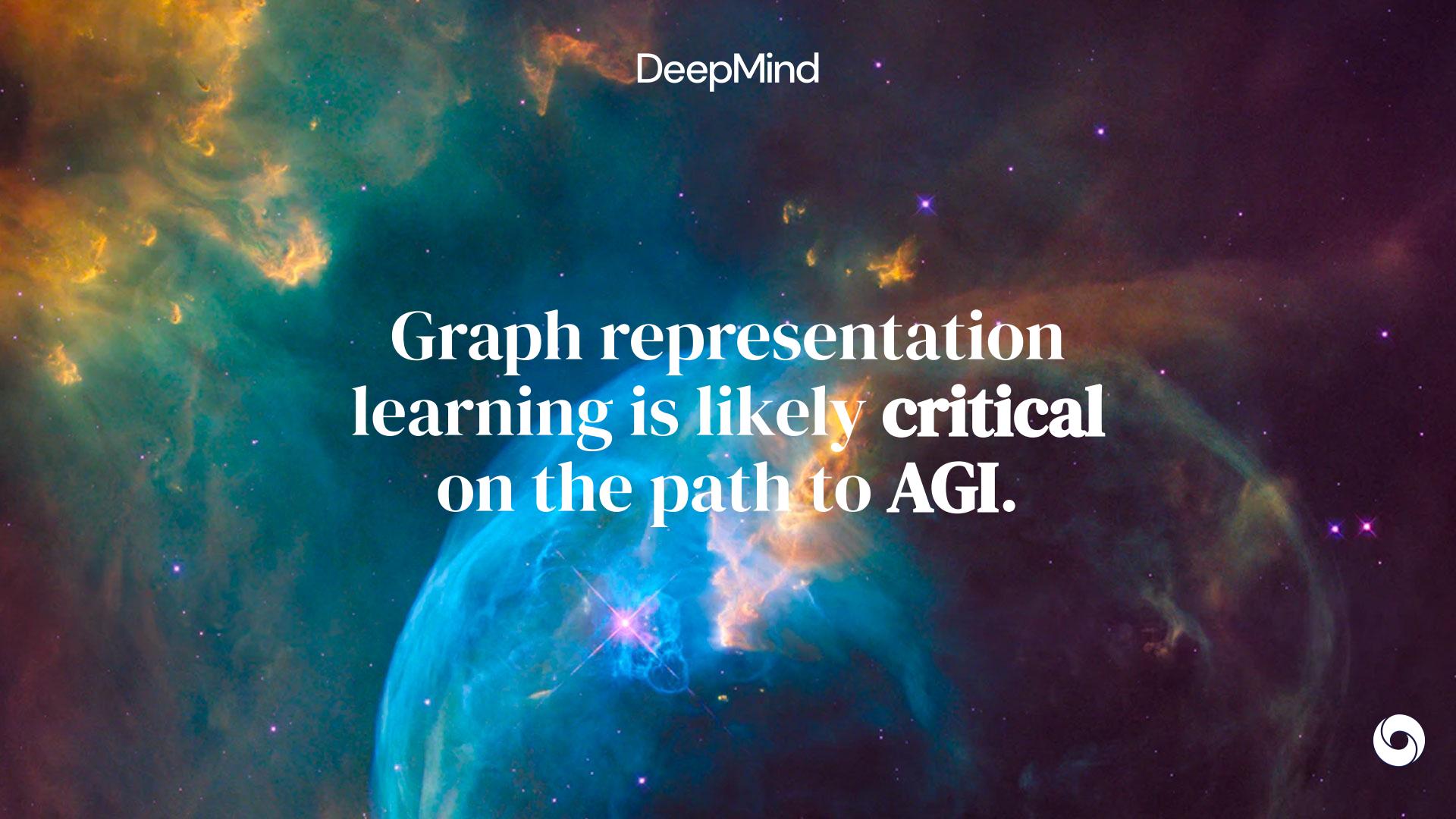
Graphs are everywhere!





In many ways, graphs are
the main modality of data
we receive from nature.



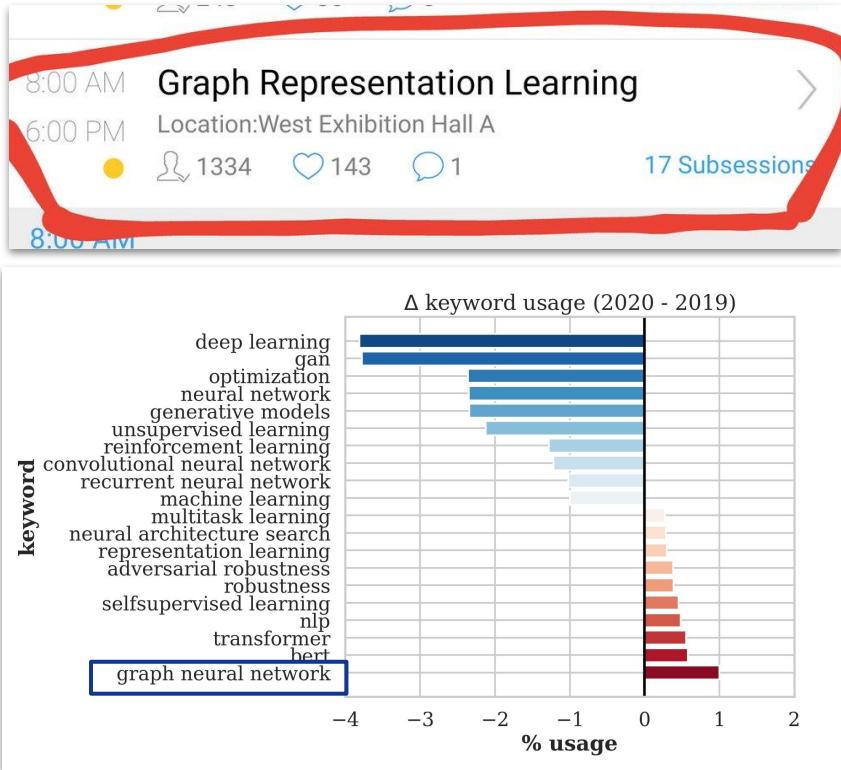
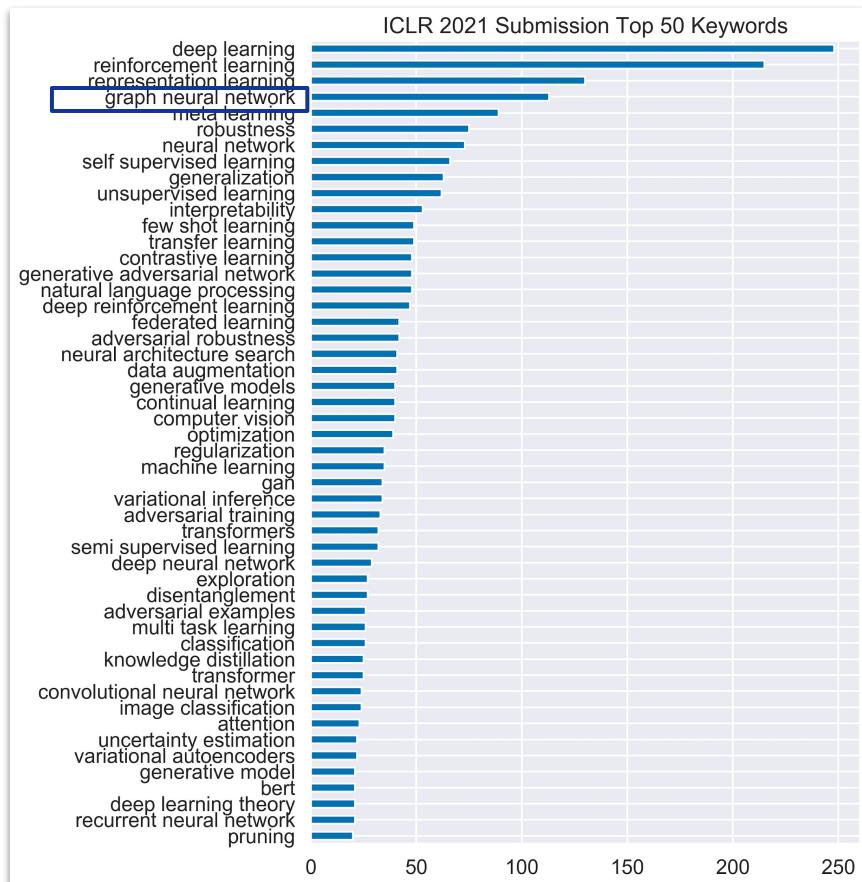


DeepMind

Graph representation
learning is likely **critical**
on the path to AGI.



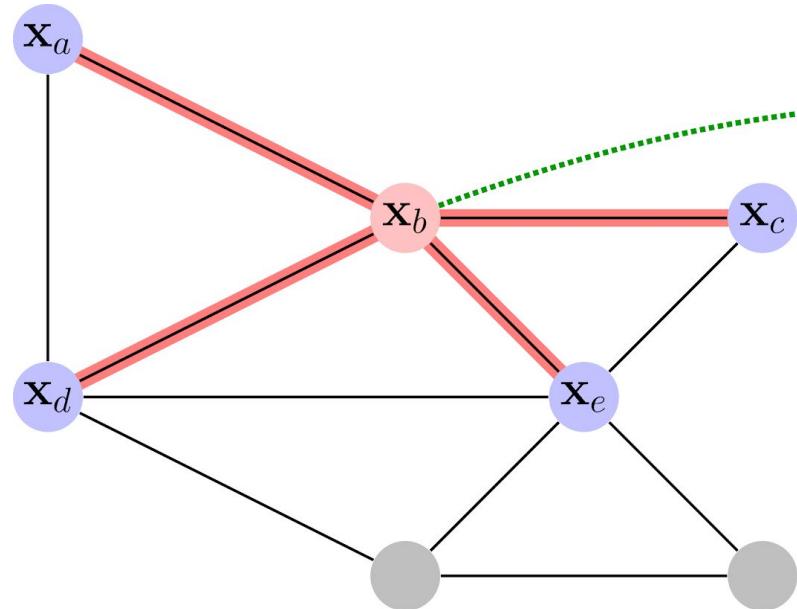
A very hot research topic



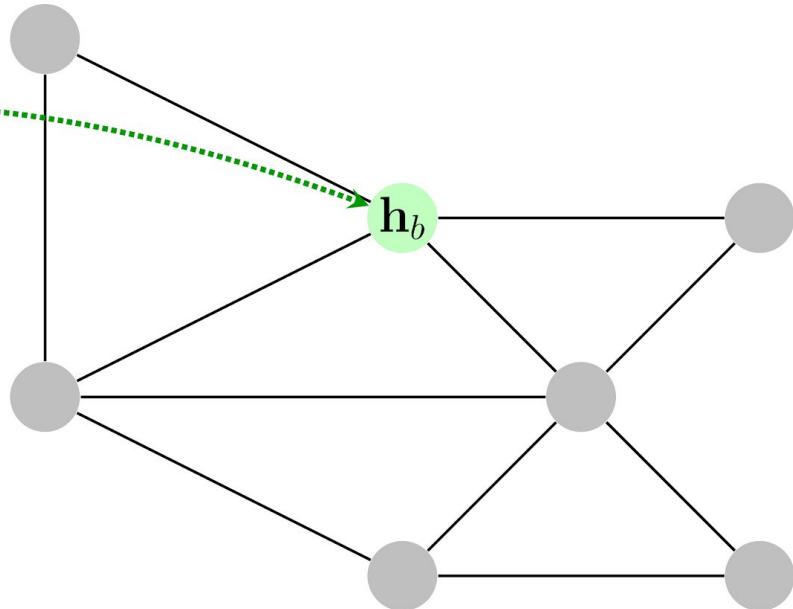
GRL is currently experiencing
its "ImageNet" moment



Graph neural networks (GNNs)



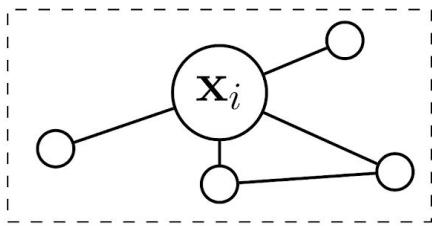
$$g(\mathbf{x}_b, \mathbf{X}_{\mathcal{N}_b})$$



$$\mathbf{X}_{\mathcal{N}_b} = \{\{\mathbf{x}_a, \mathbf{x}_b, \mathbf{x}_c, \mathbf{x}_d, \mathbf{x}_e\}\}$$



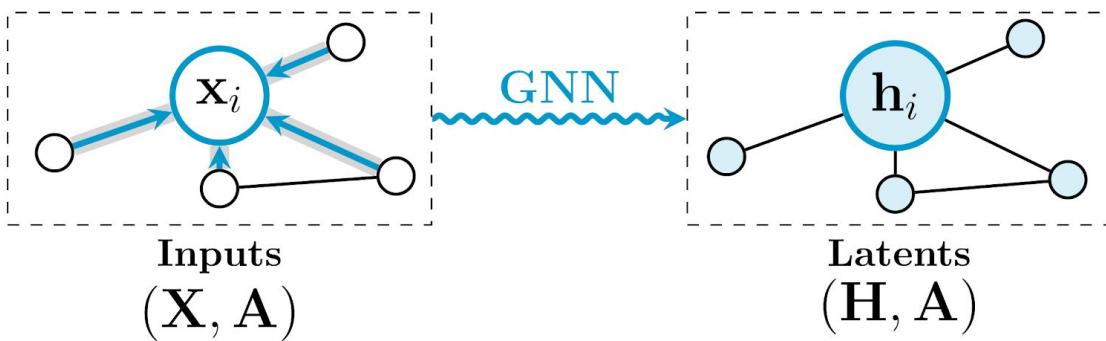
General blueprint for learning on graphs



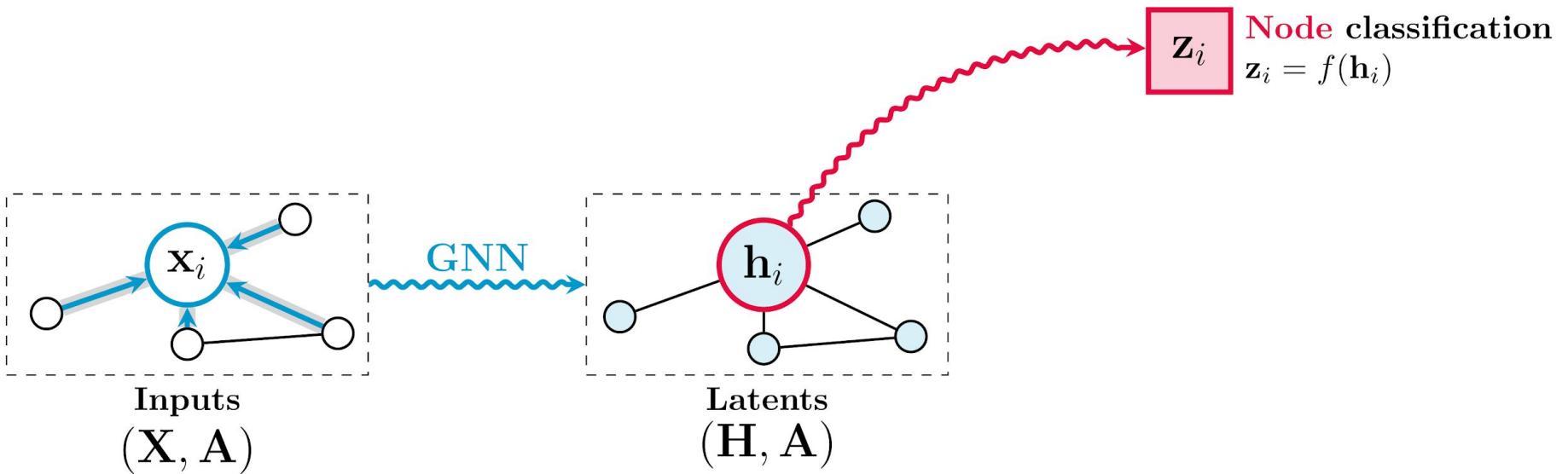
Inputs
 (\mathbf{X}, \mathbf{A})



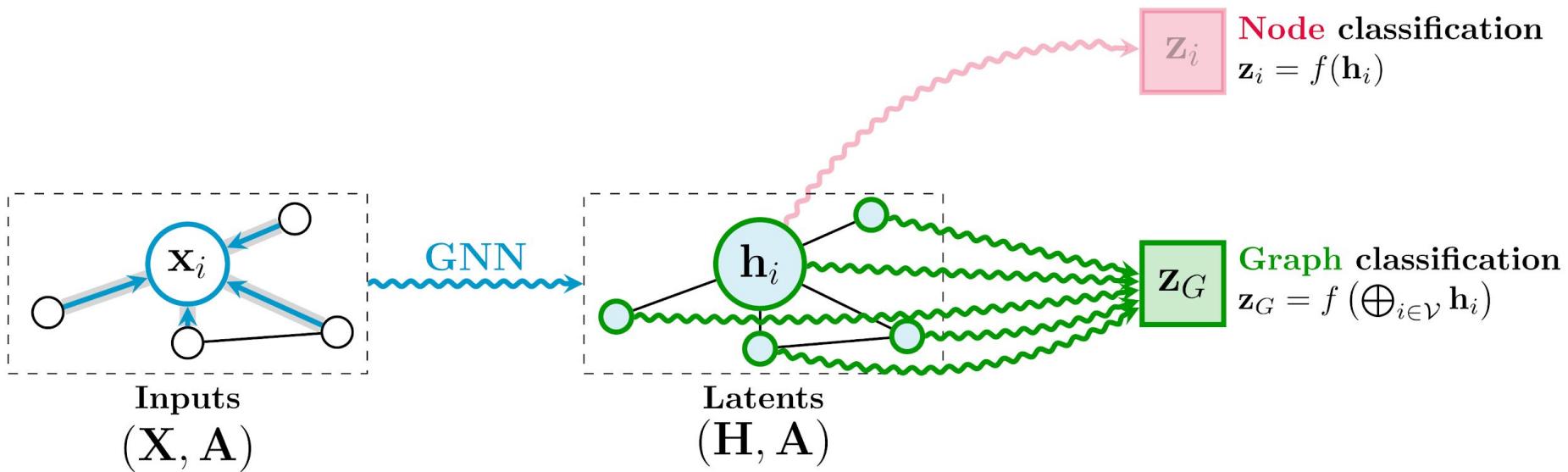
General blueprint for learning on graphs



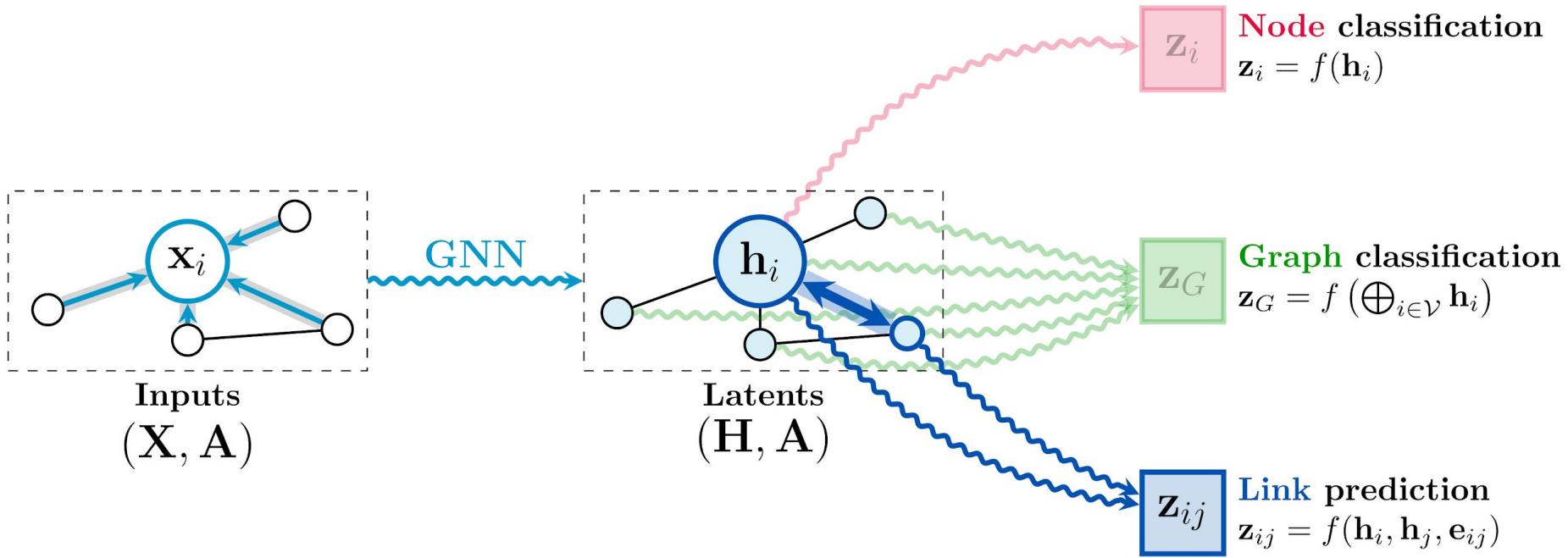
General blueprint for learning on graphs



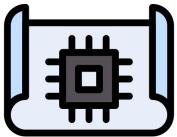
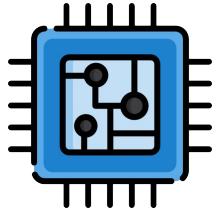
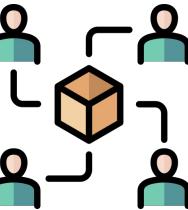
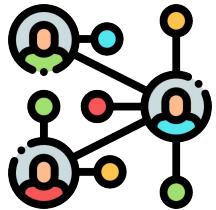
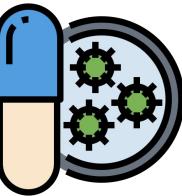
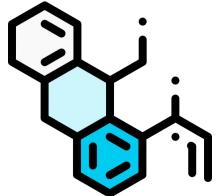
General blueprint for learning on graphs



General blueprint for learning on graphs

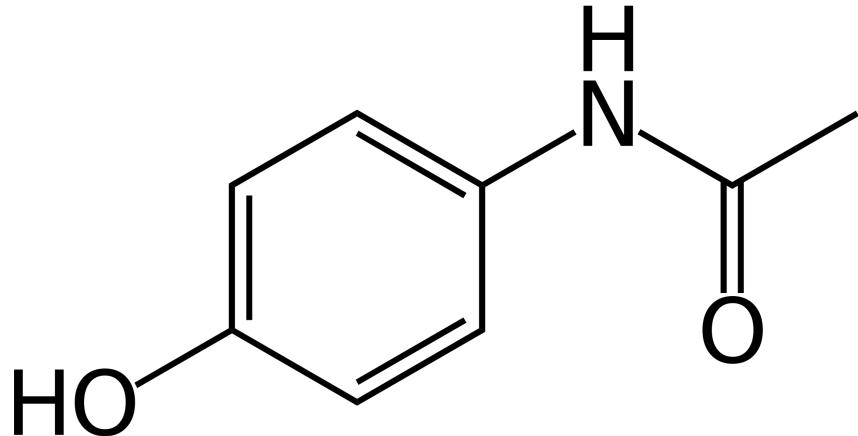
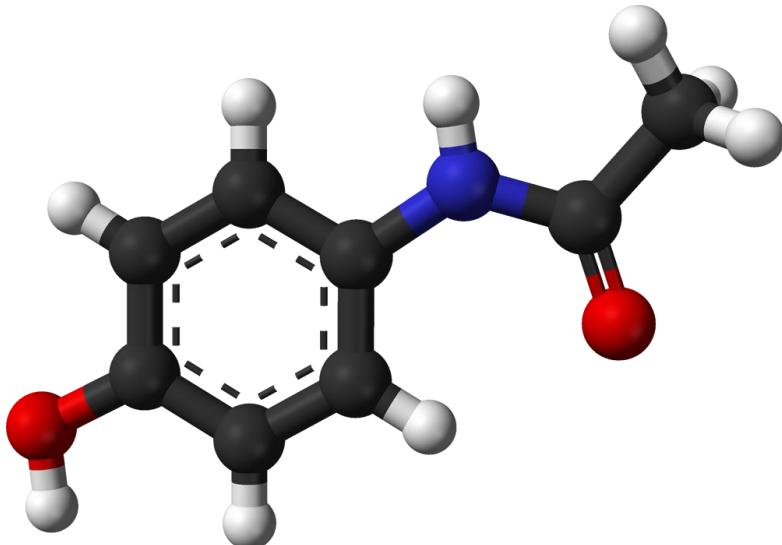


Impactful applications in science and industry



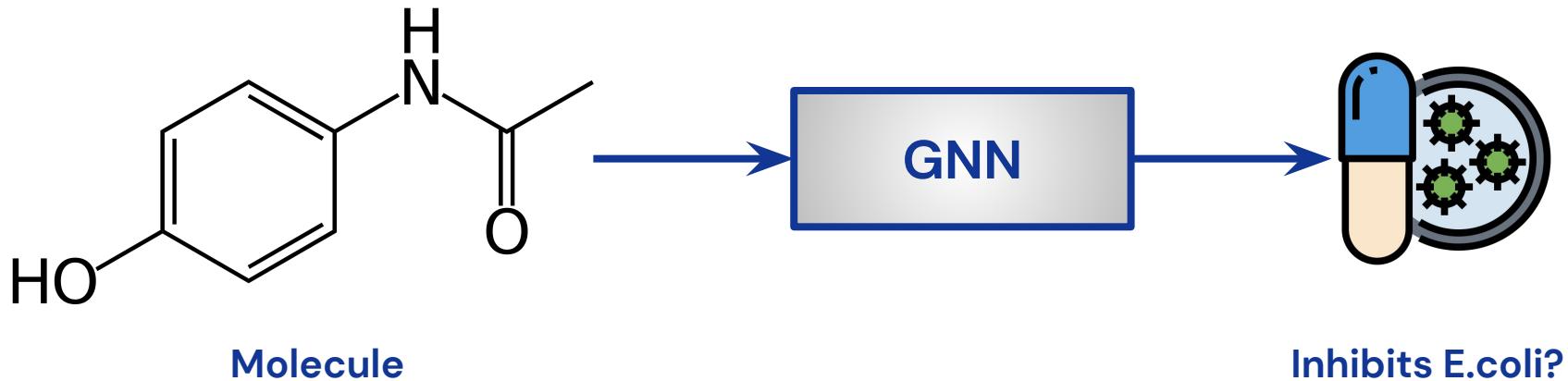
Molecules are graphs!

- A very natural way to represent molecules is as a **graph**
 - **Atoms** as nodes, **bonds** as edges
 - Features such as **atom type, charge, bond type...**



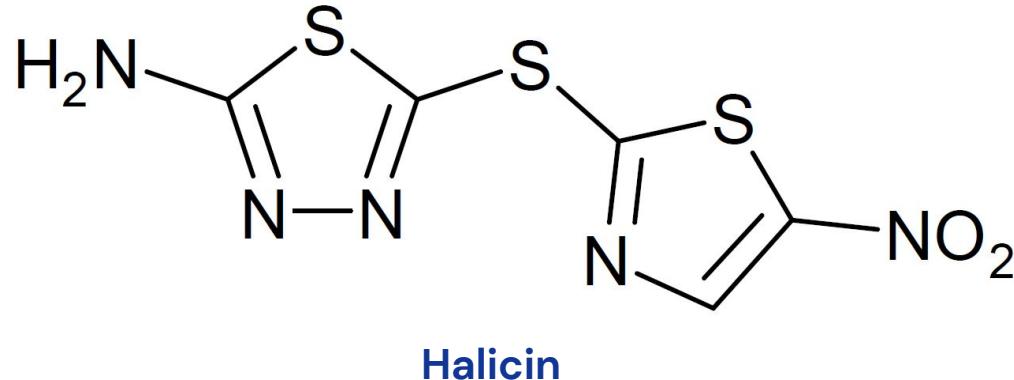
GNNs for molecule classification

- Interesting task to predict is, for example, whether the molecule is a potent **drug**.
 - Can do *binary classification* on whether the drug will inhibit certain bacteria. (*E.coli*)
 - Train on a **curated dataset** for compounds where response is known.



Follow-up study

- Once trained, the model can be applied to *any* molecule.
 - Execute on a large dataset of known candidate molecules.
 - Select the ~top-100 candidates from your GNN model.
 - Have chemists thoroughly investigate those (after some additional filtering).
- Discover a previously overlooked compound that is a **highly potent** antibiotic!



...Achieve wide acclaim!

Arguably the most popularised **success story** of graph neural networks to date!

Cell

A Deep Learning Approach to Antibiotic Discovery

Graphical Abstract

The graphical abstract illustrates a deep learning pipeline for antibiotic discovery. It starts with a 'Directed message passing neural network' (DMPNN) processing two molecules. The output of the DMPNN is used to predict antibiotic activity ('Antibiotic predictions (upper limit $10^{11} +$)'). This prediction is compared against a 'Training set (10^4 molecules)' to validate the model ('Model validation'). The validation process involves 'Growth [antibiotic]' curves. The validated model is then used to predict activity for a large dataset ('Chemical space') represented by a funnel. The final output is 'New Antibiotics'. A legend indicates: 1. Training set; 2. Model validation; 3. Antibiotic predictions; 4. DMPNN; 5. Chemical space.

Authors
Jonathan M. Stokes, Kevin Yang, Kyle Swanson, ..., Tommi S. Jaakkola, Regina Barzilay, James J. Collins

Correspondence
regina@csail.mit.edu (R.B.), jimjc@mit.edu (J.J.C.)

In Brief
A trained deep neural network predicts antibiotic activity in molecules that are structurally different from known antibiotics, among which Halicin exhibits efficacy against broad-spectrum bacterial infections in mice.

Drug Repurposing Hub HALICIN
Halicin is shown interacting with a bacterial cell, causing a change in pH (ΔpH) and leading to bacterial cell death. The chemical structure of Halicin is shown: CN1=CSC(=S)c2cc([N+]([O-])=O)cc21.

ZINC15 Database
The ZINC15 Database contains various chemical compounds. Two specific molecules are highlighted:
1. **Rapidly bactericidal Broad-spectrum**: O=[N+]([O-])c1cc(Br)c(O)n2c(N)nc(O)c2n1
2. **Low MIC Broad-spectrum**: CC1(C)C2=C1C(=O)N3C=C4C=C3C=C4C=C2N3Cc5ccc(O)cc5

(Stokes et al., Cell'20)



...Achieve wide acclaim!

Arguably the most popularised **success story** of graph neural networks to date!

The screenshot shows a news article from the **nature** website. At the top, there are logos for **Cell** and **ARTICLE**, and a blue "Subscribe" button. Below the header, the word **ARTICLE** is faintly visible. The main title of the article is **Powerful antibiotics discovered using AI**. The subtitle reads: "Machine learning spots molecules that work even against 'untreatable' strains of bacteria." The author's name, **(Stokes et al., Cell'20)**, is mentioned at the bottom left. A small graphic at the bottom features a mouse icon, the names *Acinetobacter baumannii* and *Clostridioides difficile*, a green checkmark, and a chemical structure labeled "Broad-spectrum". The date **20 FEBRUARY 2020** is also present.

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Powerful antibiotics discovered using AI

Machine learning spots molecules that work even against 'untreatable' strains of bacteria.

(Stokes et al., Cell'20)

Acinetobacter baumannii
Clostridioides difficile

Broad-spectrum

20 FEBRUARY 2020



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Arguably the most popular

nature

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Powerful and

Machine learning spots
bacteria.

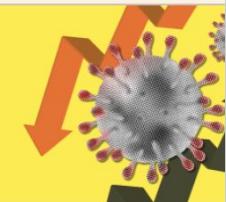
(Stokes et al., Cell'20)

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AI discovers antibiotics to treat drug-resistant diseases

Machine learning uncovers potent new drug able to kill 35 powerful bacteria



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Our new guide for getting ahead

Scientists discover powerful antibiotic using AI

Machine learning uncovers potent new drug able to kill 35 powerful bacteria

(Stokes et al., Cell'20)

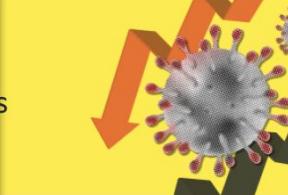
Machine learning uncovers potent new drug able to kill 35 powerful bacteria

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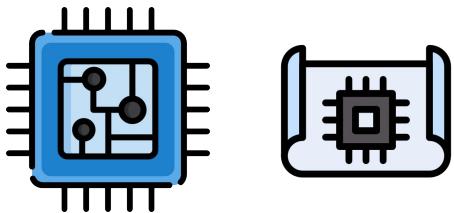
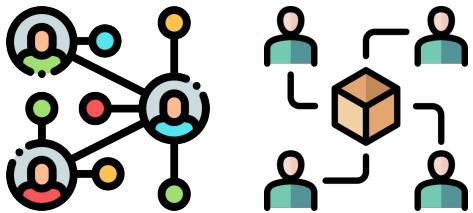
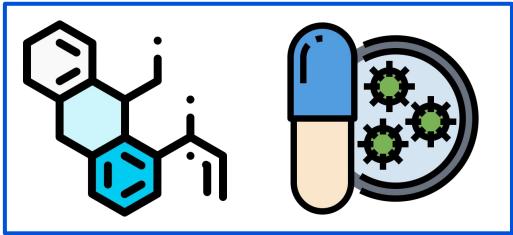
Anti-social robots harr increase social distanc

TIMES

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Impactful applications in science and industry



Cell
A Deep Learning Approach to Antibiotic Discovery

Graphical Abstract

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In Brief

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AI discovers antibiotics to treat drug-resistant diseases

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Scientists discover powerful antibiotic using AI

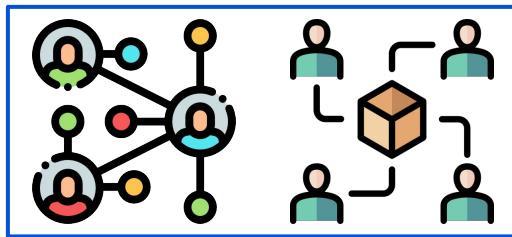
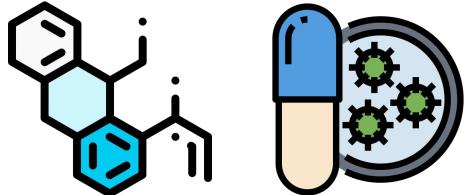
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This block displays a screenshot of a scientific article from the journal *Cell* titled "A Deep Learning Approach to Antibiotic Discovery". It includes a graphical abstract showing a directed message passing neural network and a chemical space plot. Below it is a BBC News page featuring an AI guide for worklife. To the right is a snippet from the Financial Times about AI discovering antibiotics. At the bottom is a news article from the BBC about scientists discovering a powerful antibiotic using AI.

Virtual drug screening



Impactful applications in science and industry



PinSage: A new graph convolutional neural network for web-scale recommender systems



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PUBLICATION

P-Companion: A principled framework for diversified complementary product recommendation

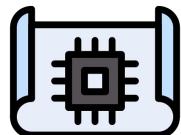
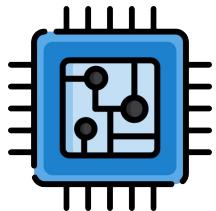
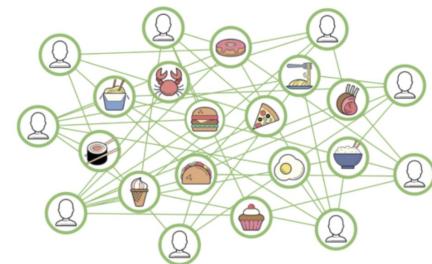
By Junheng Hao, Tong Zhao, Jin Li, Xin Luna Dong, Christos Faloutsos, Yizhou Sun, Wei Wang

2020

Food Discovery with Uber Eats: Using Graph Learning to Power Recommendations

Ankit Jain, Isaac Liu, Ankur Sarda, and Piero Molino

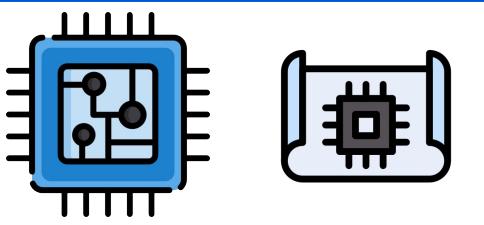
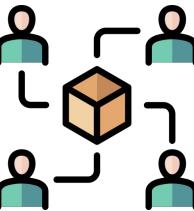
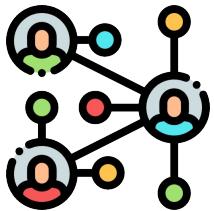
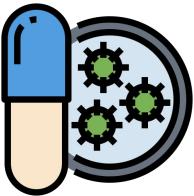
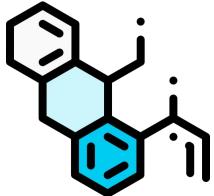
December 4, 2019



Recommender systems



Impactful applications in science and industry



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Article | Published: 09 June 2021

A graph placement methodology for fast chip design

Azalia Mirhoseini, Anna Goldie, Mustafa Yazgan, Joe Wenjie Jiang, Ebrahim Songhori, Shen Wang, Young-Joon Lee, Eric Johnson, Omkar Pathak, Azade Nazi, Jiwoo Pak, Andy Tong, Kavya Srinivasa, William Hang, Emre Tuncer, Quoc V. Le, James Laudon, Richard Ho, Roger Carpenter & Jeff Dean

GOOGLE \ TECH \ ARTIFICIAL INTELLIGENCE

Google is using AI to design its next generation of AI chips more quickly than humans can

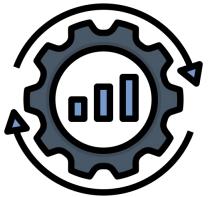
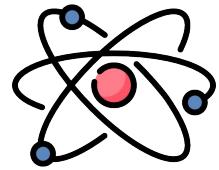
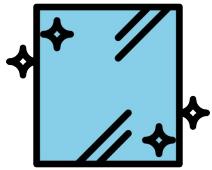
Designs that take humans months can be matched or beaten by AI in six hours

By James Vincent | Jun 10, 2021, 9:13am EDT

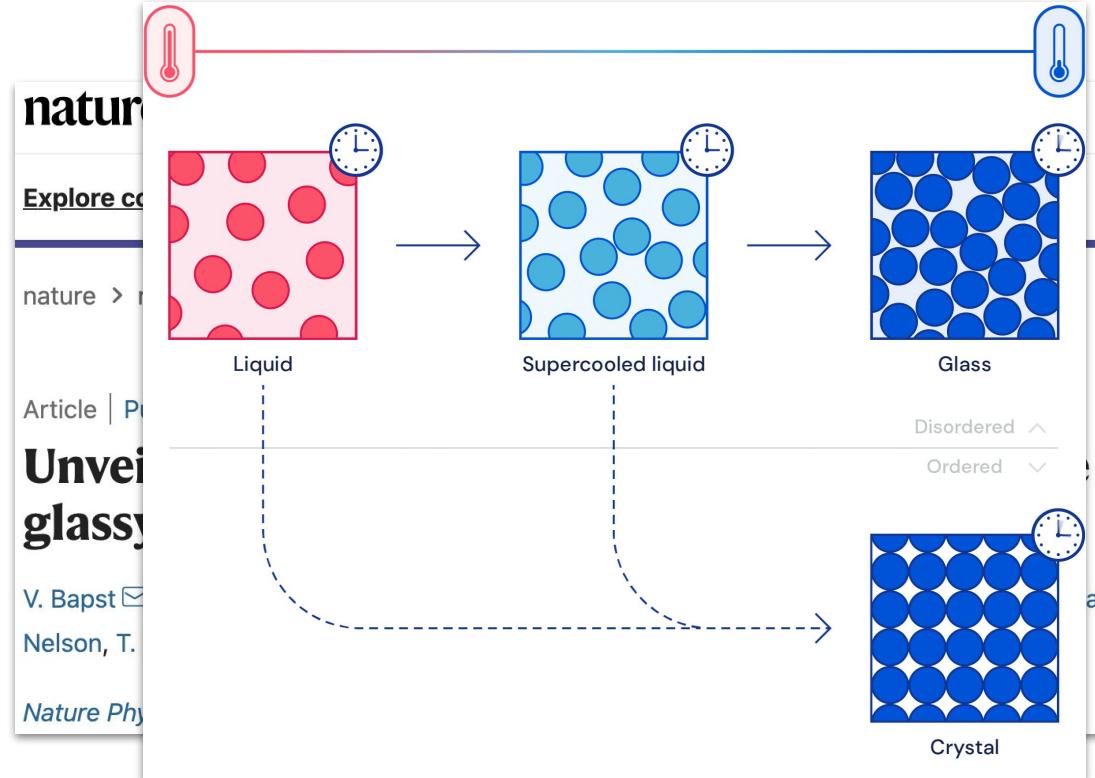
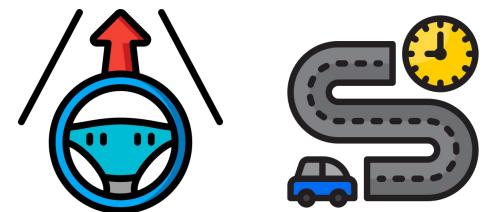
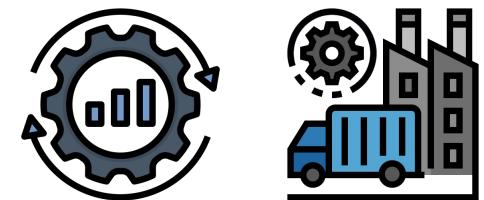
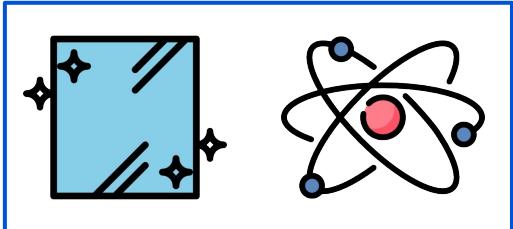
Chip design (TPUv5)



Impactful applications from DeepMind



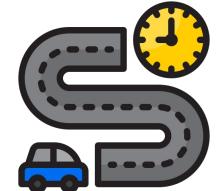
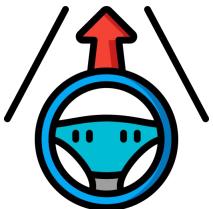
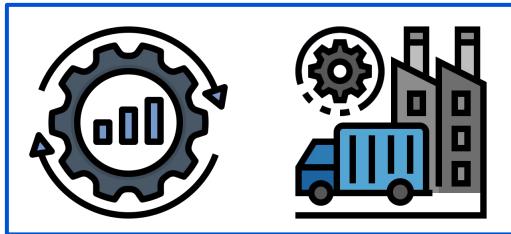
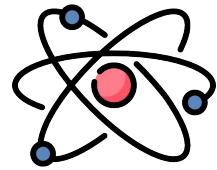
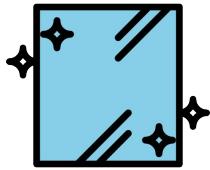
Impactful applications from DeepMind



Glassy dynamics



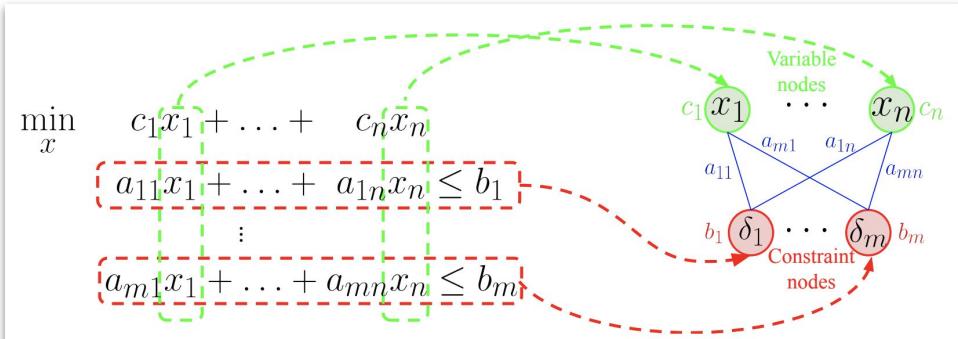
Impactful applications from DeepMind



Solving Mixed Integer Programs Using Neural Networks

Vinod Nair^{*†1}, Sergey Bartunov^{*1}, Felix Gimeno^{*1}, Ingrid von Glehn^{*1}, Paweł Lichocki^{*2}, Ivan Lobov^{*1}, Brendan O'Donoghue^{*1}, Nicolas Sonnerat^{*1}, Christian Tjandraatmadja^{*2}, Pengming Wang^{*1}, Ravichandra Addanki¹, Tharindi Hapuarachchi¹, Thomas Keck¹, James Keeling¹, Pushmeet Kohli¹, Ira Ktena¹, Yujia Li¹, Oriol Vinyals¹, Yori Zwols¹

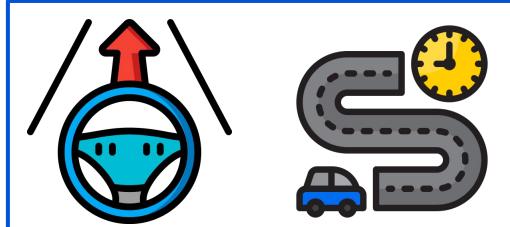
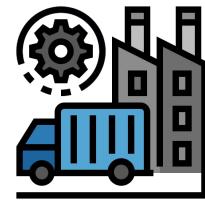
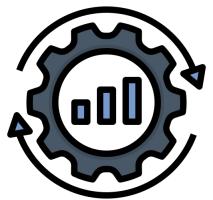
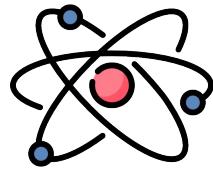
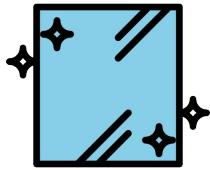
¹DeepMind, ²Google Research



Combinatorial optimisation



Impactful applications from DeepMind



ETA Prediction with Graph Neural Networks in Google Maps

Austin Derrow-Pinion¹, Jennifer She¹, David Wong^{2*}, Oliver Lange³, Todd Hester^{4*}, Luis Perez^{5*}, Marc Nunkesser³, Seongjae Lee³, Xueying Guo³, Brett Wiltshire¹, Peter W. Battaglia¹, Vishal Gupta¹, Ang Li¹, Zhongwen Xu^{6*}, Alvaro Sanchez-Gonzalez¹, Yujia Li¹ and Petar Veličković¹

¹DeepMind ²Waymo ³Google ⁴Amazon ⁵Facebook AI ⁶Sea AI Lab *work done while at DeepMind

{derrowap,jenshe,wongda,petarv}@google.com

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The Machine
Making sense of AI

DeepMind claims its AI improved Google Maps travel time estimates by up to 50%

Kyle Wiggers @Kyle_L_Wiggers September 3, 2020 7:00 AM

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Travel-time Prediction in Google Maps



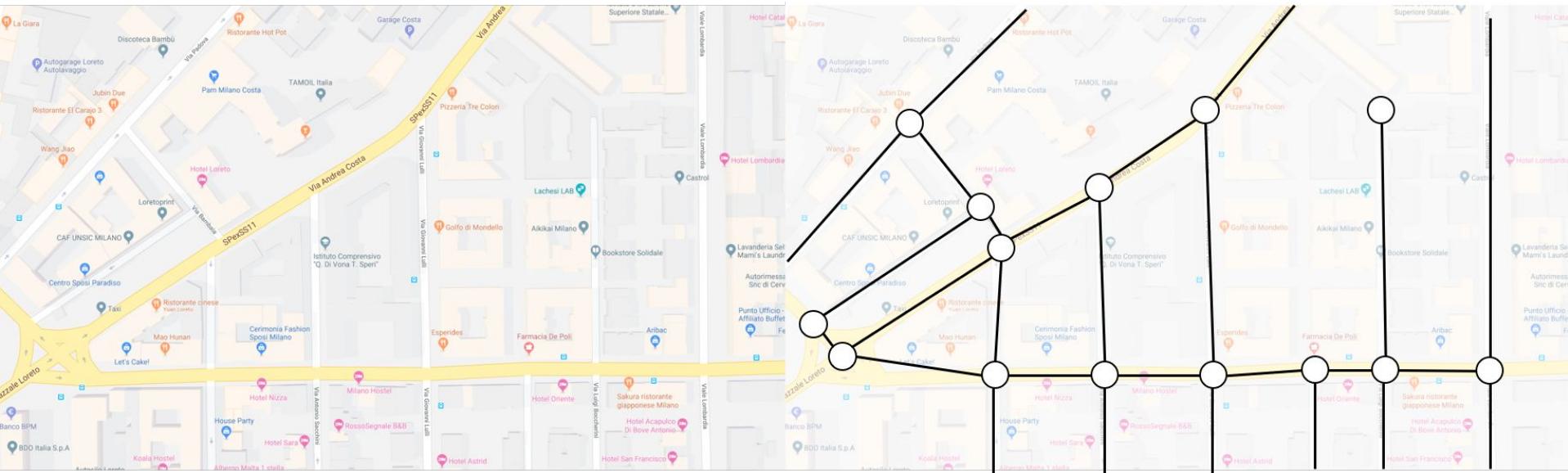


Enabling scalable traffic predictions with GNNs in Google Maps



Traffic maps are graphs!

Transportation maps (e.g. the ones found on Google Maps) naturally modelled as **graphs**.



Nodes could be **intersections**, and edges could be **roads**.



Estimated Time of Arrival (ETA) Prediction

- A critical service provided by Google Maps is **ETA prediction**.
 - Given a start-point and end-point, what is the expected travel time?
 - Important for both **users** and **ride-sharing/delivery** companies (using the Maps API).
- Relevant **node features**: road *length*, *current speeds*, *historical speeds*
- Use anonymised, crowd-sourced real-time / historical traffic data.
 - Not as reliable as e.g. physical speed sensors
 - Traffic conditions change dynamically and unpredictably
 - Most trips between [10min, 1h], requiring **near-future predictions**



DeepMind's approach: Graph Nets on Supersegments

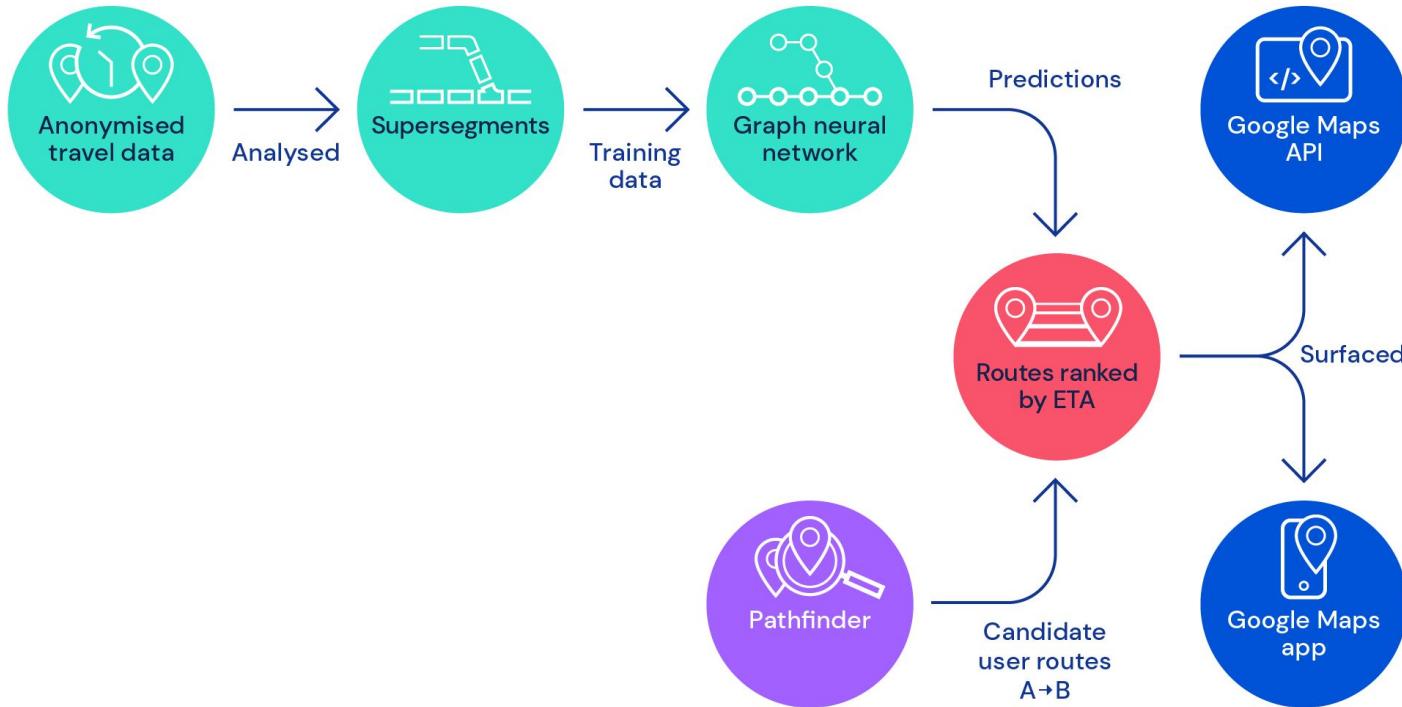
Partition candidate route into **supersegments**, sampled proportionally to (est.) **traffic density**.

Run a GNN over **supersegment** graph to estimate ETA (*graph regression*).



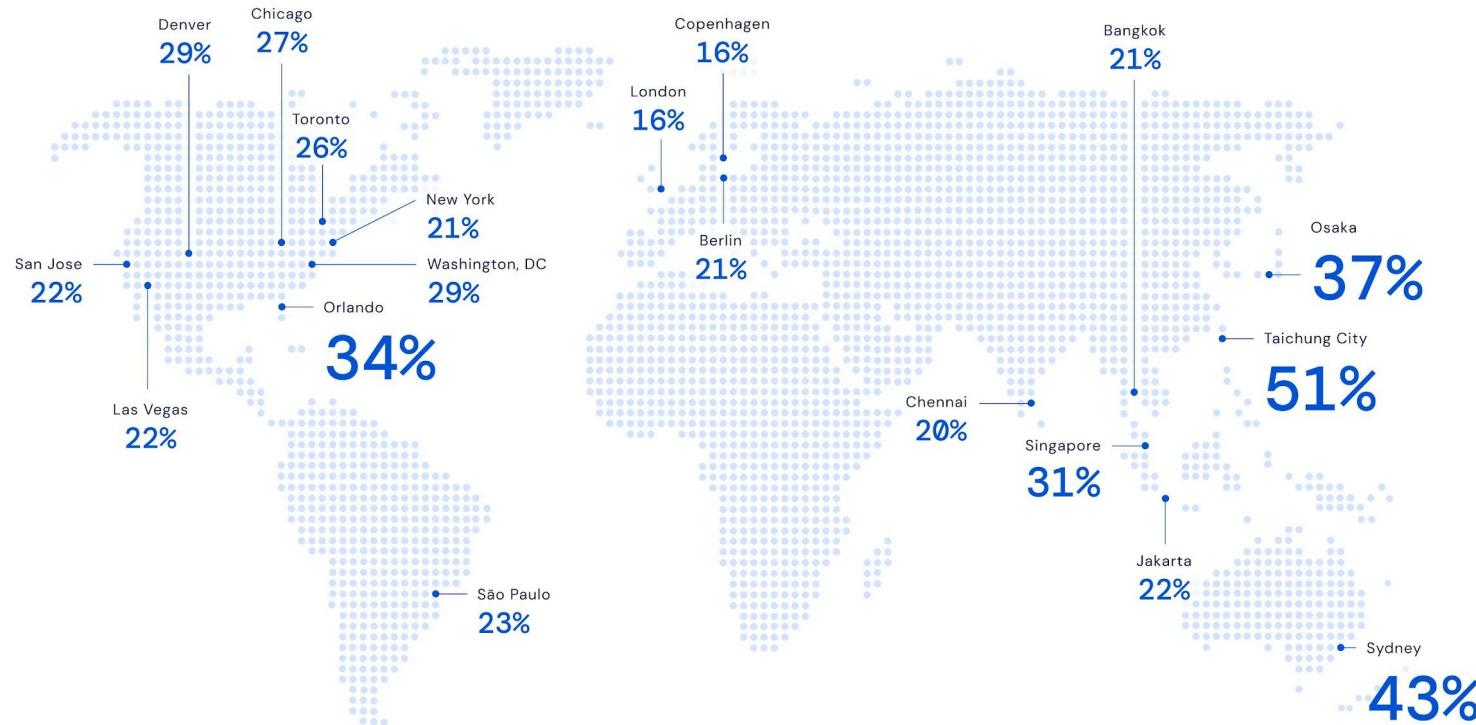
Overall pipeline

Rank candidate routes by predicted ETA, surface information to Google Maps.



Returns

Already **deployed** worldwide, significantly reducing negative ETA outcomes!



DeepMind

I

Getting in on
the action



Rich ecosystem of libraries



PyTorch
geometric

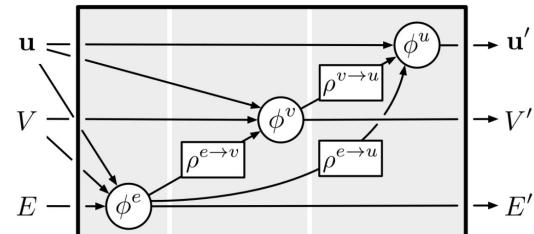
github.com/rusty1s/pytorch_geometric

DGL
dgl.ai



Spektral

graphneural.network



github.com/deepmind/graph_nets



github.com/deepmind/jraph

Rich ecosystem of datasets



ogb.stanford.edu



PyTorch
geometric

<https://pytorch-geometric.readthedocs.io/en/latest/modules/datasets.html>



graphlearning.io

Benchmarking Graph Neural Networks

github.com/graphdeeplearning/benchmarking-gnns



Getting into it!

- I recently compiled a list of many useful GNN resources in a **Twitter thread**
 - https://twitter.com/PetarV_93/status/1306689702020382720
- When you feel ready, I **highly** recommend Alekса Gordić's GitHub repository on GATs:
 - <https://github.com/gordicaleksa/pytorch-GAT>
 - Arguably the most *gentle* introduction to GNN implementations



III

Graph Isomorphism Testing



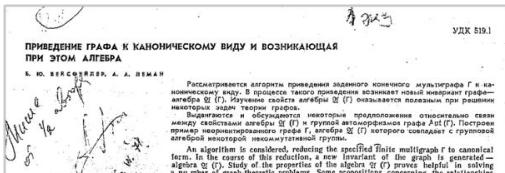
How *powerful* are Graph Neural Networks?

- GNNs are a powerful tool for processing real-world graph data
 - But they won't solve **any** task specified on a graph accurately!
- Canonical example: deciding *graph isomorphism*
 - Am I able to use my GNN to **distinguish** two *non-isomorphic* graphs? ($\mathbf{h}_{G1} \neq \mathbf{h}_{G2}$)
 - If I can't, any kind of task discriminating them is *hopeless*
- We will assess the **power** of GNNs by which graphs they are able to **distinguish**.



Weisfeiler-Lehman Test

- Simple but powerful way of distinguishing: pass **random hashes of sums** along the edges
 - Iterate until hashes don't change.
 - "Possibly isomorphic"* if hash histograms are the same.



1. Рассмотрим произвольный конечный граф Γ и его матрицу смежности $A(\Gamma)(a_i)$; здесь $a_i = \{x_1, x_2, \dots, x_n\}$ — множество вершин графа Γ и $a_i = \{j_1, j_2, \dots, j_n\}$. В случае необходимости мы будем называть матрицу смежности $A(\Gamma)$ метрикой Γ , а ее элементы — вершинами Γ . Видим, что мы будем называть матрицу смежности при каноническом виде алгеброй $\mathfrak{A}(\Gamma)$, а ее элементы — вершинами алгебры $\mathfrak{A}(\Gamma)$.

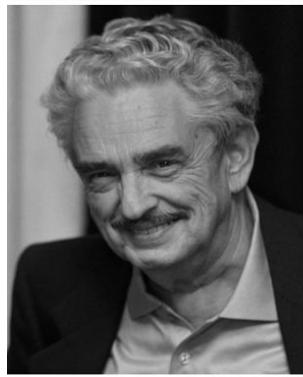
An algorithm is considered, reducing the specified finite multigraph Γ to canonical form. In this case the resulting, so called, "algebra" $\mathfrak{A}(\Gamma)$ is generated by a number of graph-theoretic problems. Some problems concerning the relationships between vertices of Γ are also considered. The main problem of this article is the construction of the automorphism group $Aut(\Gamma)$. An example of non-oriented graph $\tilde{\Gamma}$ is constructed whose group $\mathfrak{A}(\tilde{\Gamma})$ coincides with the group $\mathfrak{A}(G)$ of a non-commutative group.

В п. 1 указан процесс приведения графа Γ к каноническому виду, состоящий в согласованном перенесении строк и столбцов матрицы смежности $A(\Gamma)$ изображенного графа Γ в алгебру $\mathfrak{A}(\Gamma)$.

Рассмотрим для простоты канонический вид графа Γ без хроматической метки, единственным компонентом которого является единственный цикл. Допустим, что у вершин x_1, x_2, \dots, x_n есть хроматические метки a_1, a_2, \dots, a_n , так чтобы вершины с одинаковыми хроматическими метками были соединены ребрами. Тогда в результате этого упорядочения в соответствии с остаточными характеристиками вершин x_1, x_2, \dots, x_n получим, что в алгебре $\mathfrak{A}(\Gamma)$ вершины с одинаковыми остаточными характеристиками вершинами x_1, x_2, \dots, x_n будут соединены ребрами, а остальные вершины x_1, x_2, \dots, x_n не будут соединены ребрами.

В п. 2 описан процесс приведения канонического вида Γ к каноническому виду, использующий применение специальных операторов к матрице, полученной из $A(\Gamma)$ алгеброй $\mathfrak{A}(\Gamma)$. Алгебра $\mathfrak{A}(\Gamma)$ определяется в п. 1. Алгоритм приведения вершин x_1, x_2, \dots, x_n к каноническому виду, определяемому в п. 2, описан в п. 3.

Заметим, что описание алгоритма в п. 3 не содержит никаких методик, вложенных в [1] и [2].



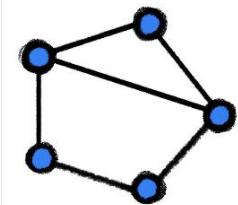
A. Lehman



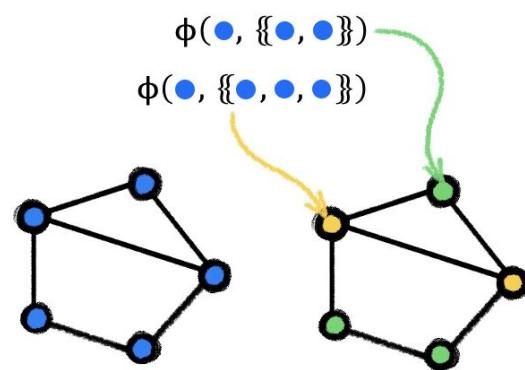
B. Weisfeiler



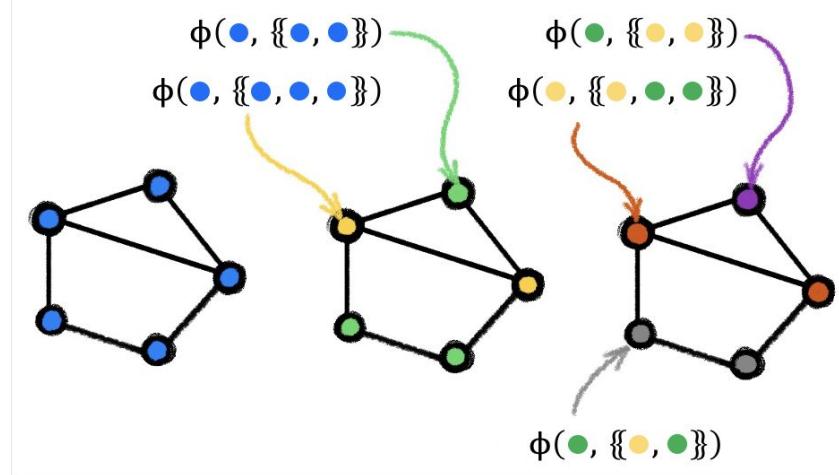
Let's run the WL Test!



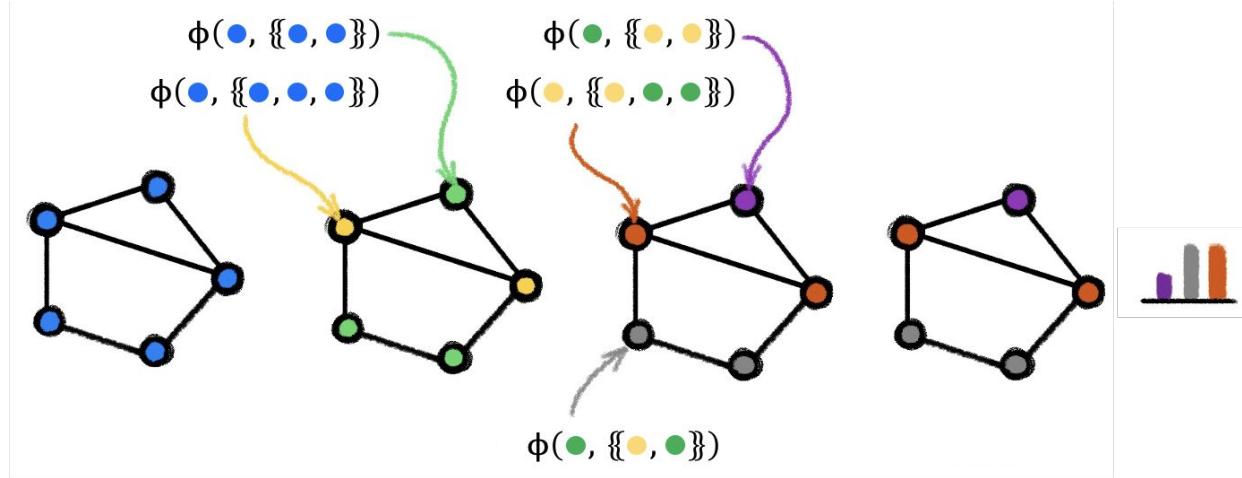
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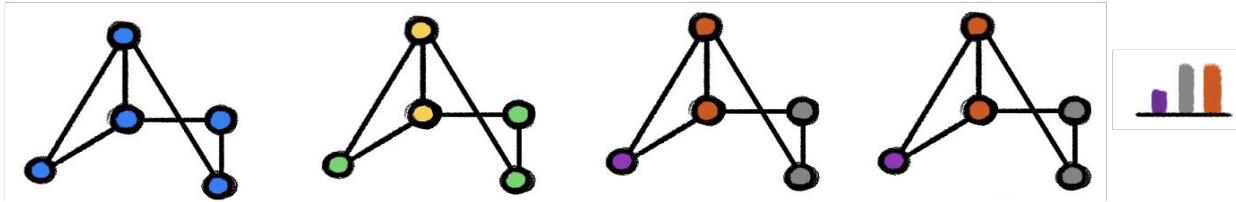
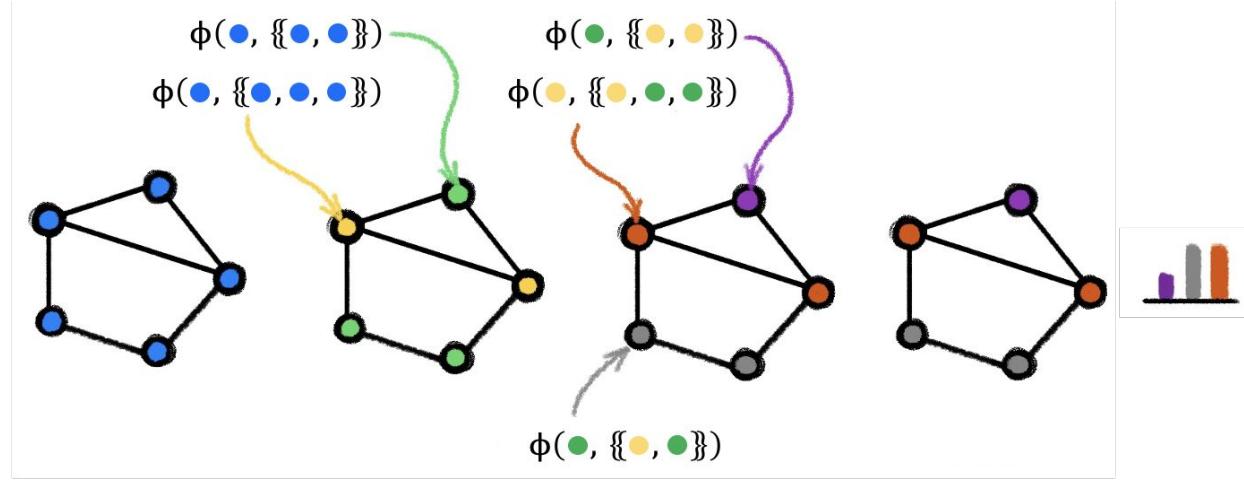
Let's run the WL Test!



Let's run the WL Test!

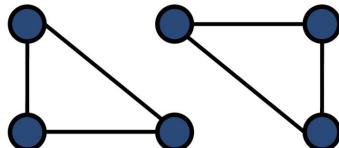
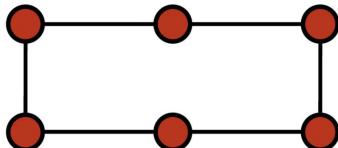


Let's run the WL Test!



Weisfeiler-Lehman Test

- Connection to conv-GNNs spotted very early; e.g. by GCN (Kipf & Welling, ICLR'17)
- **Untrained GNNs** can hence work very well!
 - Untrained ~ random hash
- The test does **fail** at times, however:



Algorithm 1: WL-1 algorithm (Weisfeiler & Lehmann, 1968)

Input: Initial node coloring $(h_1^{(0)}, h_2^{(0)}, \dots, h_N^{(0)})$

Output: Final node coloring $(h_1^{(T)}, h_2^{(T)}, \dots, h_N^{(T)})$

$t \leftarrow 0;$

repeat

for $v_i \in \mathcal{V}$ **do**

$h_i^{(t+1)} \leftarrow \text{hash} \left(\sum_{j \in \mathcal{N}_i} h_j^{(t)} \right);$

$t \leftarrow t + 1;$

until stable node coloring is reached;

$$\mathbf{h}_i = \phi \left(\mathbf{x}_i, \bigoplus_{j \in \mathcal{N}_i} c_{ij} \psi(\mathbf{x}_j) \right)$$



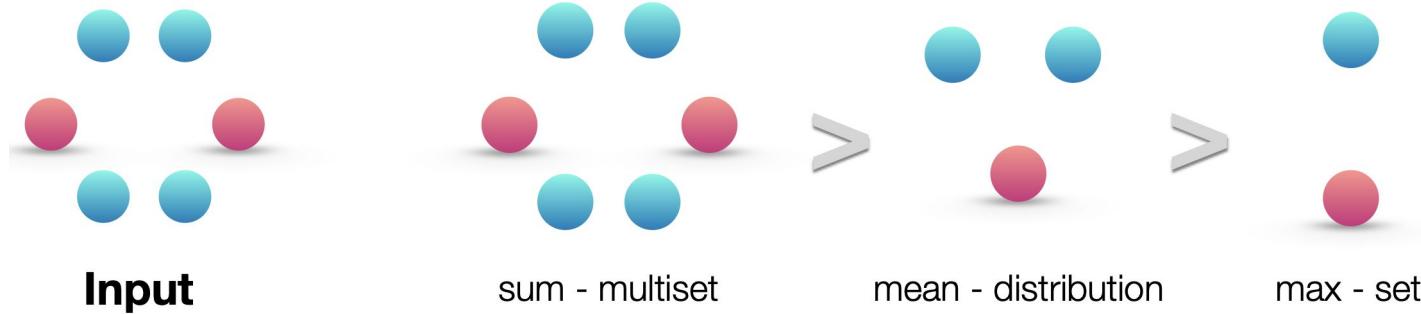
GNNs are no more powerful than 1-WL

- Over *discrete* features, GNNs can only be **as powerful** as the 1-WL test described before!



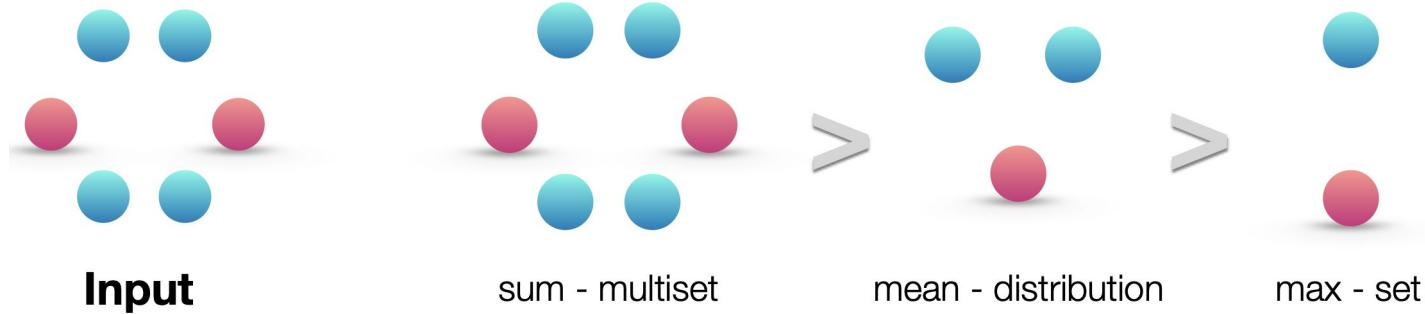
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- One important condition for maximal power is an *injective* aggregator (e.g. **sum**)



GNNs are no more powerful than 1-WL

- Over *discrete* features, GNNs can only be **as powerful** as the 1-WL test described before!
- One important condition for maximal power is an *injective* aggregator (e.g. **sum**)

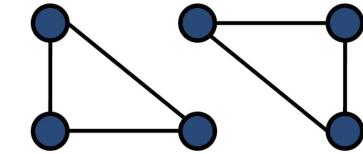
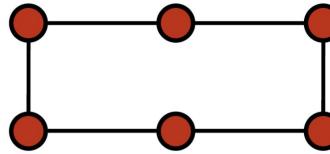


- Graph isomorphism network (**GIN**; Xu et al., ICLR'19) proposes a simple, maximally-expressive GNN, following this principle:

$$h_v^{(k)} = \text{MLP}^{(k)} \left(\left(1 + \epsilon^{(k)} \right) \cdot h_v^{(k-1)} + \sum_{u \in \mathcal{N}(v)} h_u^{(k-1)} \right)$$



Higher-order GNNs



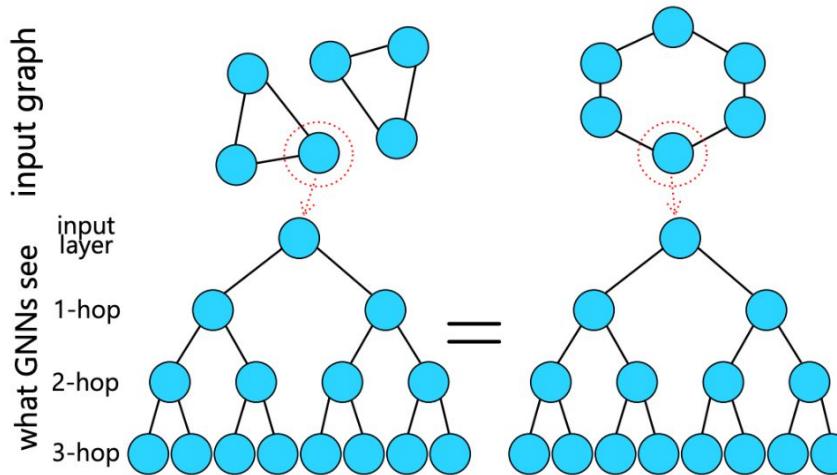
- We can make GNNs stronger by analysing **failure cases** of 1-WL!
 - Very active area, with many open problems!



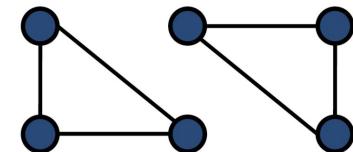
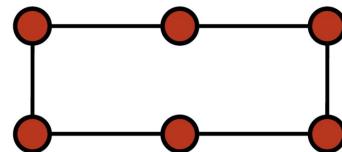
Higher-order GNNs



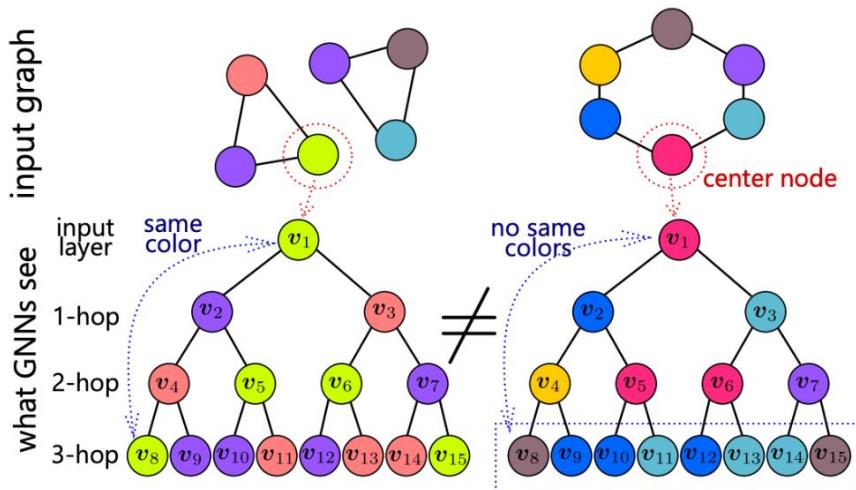
- We can make GNNs stronger by analysing **failure cases** of 1-WL!
- For example, just like 1-WL, GNNs cannot detect **closed triangles**
 - This is because, from a GNN's perspective, all nodes look the same!
 - Can you think of a simple fix?



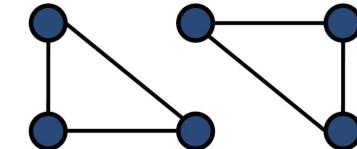
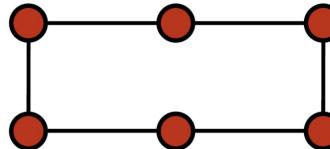
Higher-order GNNs



- We can make GNNs stronger by analysing **failure cases** of 1-WL!
- For example, just like 1-WL, GNNs cannot detect **closed triangles**
 - Augment nodes with **randomised features** (Sato et al., SDM'21)
 - Now a node can “see itself” k hops away!



Higher-order GNNs

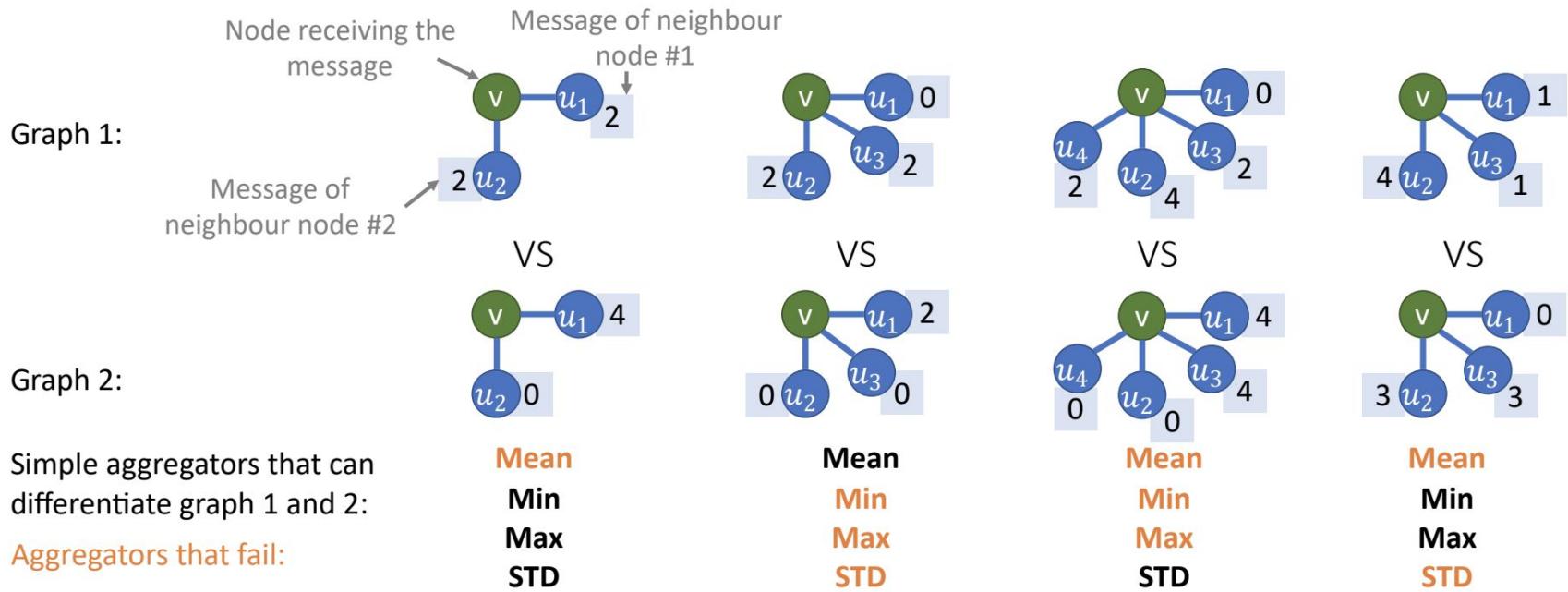


- We can make GNNs stronger by analysing **failure cases** of 1-WL!
- For example, just like 1-WL, GNNs cannot detect **closed triangles**
 - Augment nodes with randomised/positional features (Sato et al., SDM'21)
 - Explored by RP-GNN (Murphy et al., ICML'19) and P-GNN (You et al., ICML'19)
 - Can also literally **count** interesting subgraphs (Bouritsas et al., 2020)
- Fixing “failure cases” of 1-WL yields many classes of *higher-order GNNs*
- They can broadly be categorised into three groups:
 - Modifying **features** (as above)
 - Modifying the **message passing rule**; e.g. DGN (Beaini, Passaro et al. (2020))
 - Modifying the **graph structure**; e.g. 1-2-3-GNNs (Morris et al., AAAI'19)



Going beyond discrete features

- What happens when features are **continuous**? (real-world apps / latent GNN states)
 - ... the proof for injectivity of sum (hence GINs' expressivity) **falls apart**



Which is best? Neither.

- There doesn't seem to be a clear single "winner" aggregator here...
- In fact, we prove in the PNA paper (Corso, Cavalleri et al., NeurIPS'20) that **there isn't one!**

Theorem 1 (Number of aggregators needed). *In order to discriminate between multisets of size n whose underlying set is \mathbb{R} , at least n aggregators are needed.*

- The proof is (in my opinion) **really cool!** (relies on **Borsuk–Ulam theorem**)
- PNA proposes empirically powerful **combination** of aggregators for general-purpose GNNs:

$$\bigoplus = \underbrace{\begin{bmatrix} I \\ S(D, \alpha = 1) \\ S(D, \alpha = -1) \end{bmatrix}}_{\text{scalers}} \otimes \underbrace{\begin{bmatrix} \mu \\ \sigma \\ \max \\ \min \end{bmatrix}}_{\text{aggregators}}$$



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III

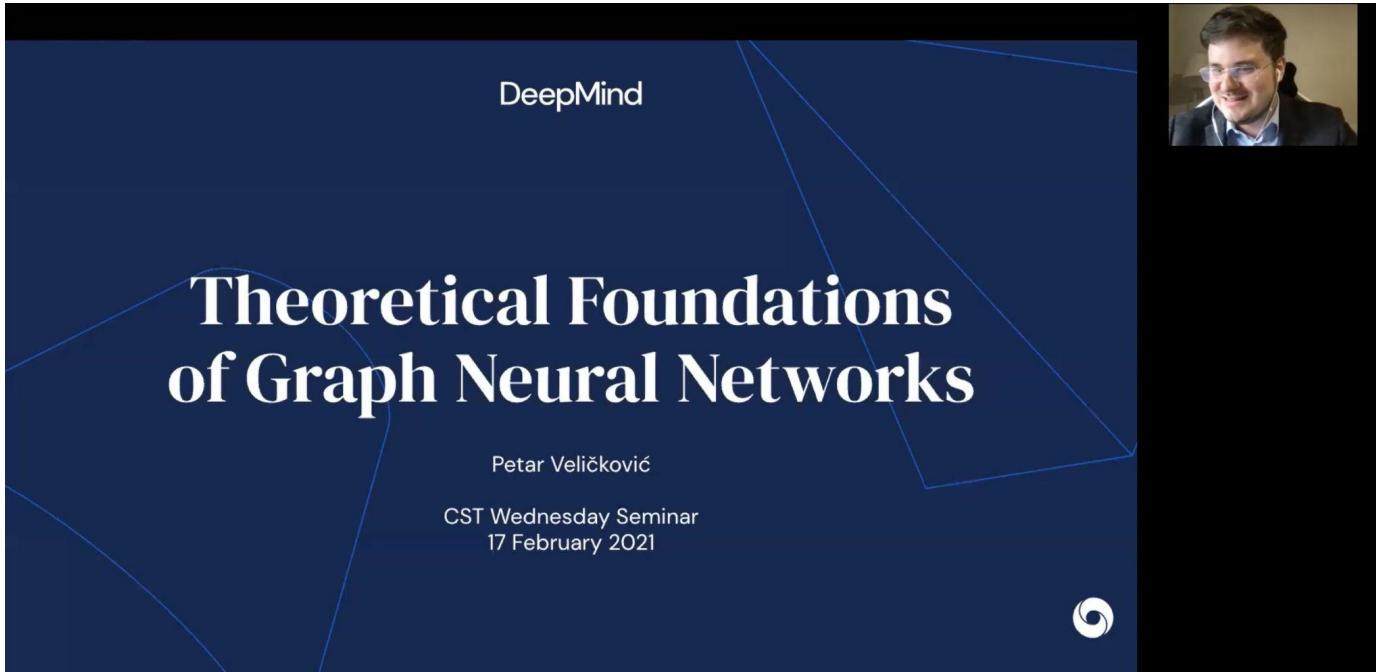
Further
resources



Further insight: graph representation learning

If GNNs are new(ish) to you, I recently gave a useful talk on **theoretical GNN foundations**:

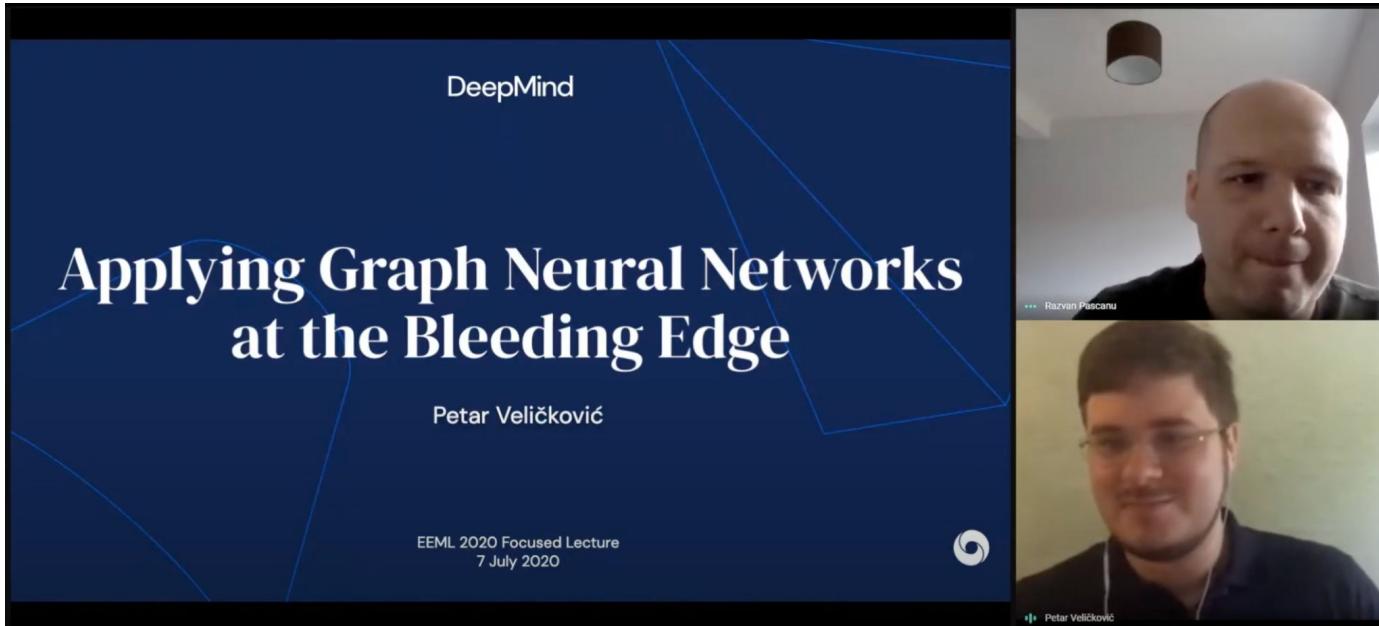
<https://www.youtube.com/watch?v=uF53xsT7mjc>



Further insight: bleeding-edge applications

For an in-depth view of bleeding edge applications of GNNs, check out my **EEML 2020 talk**:

<https://www.youtube.com/watch?v=fpb3j33RfTc>



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Thank you!

Questions?

petarv@deepmind.com | <https://petar-v.com>

