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Graph neural networks and an application to molecule generation

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OUTLINE

- ▶ Graph neural networks (GNNs)
- ▶ Applications of GNNs
- ▶ A method for molecule generation
- ▶ Some experimental results



Graph Neural Networks (GNNs)

Graphs

- ▶ they allow to represent
 - ▶ patterns (nodes with attached vector features)
 - ▶ **their relationships (edges with attached vector features)**

GNNs

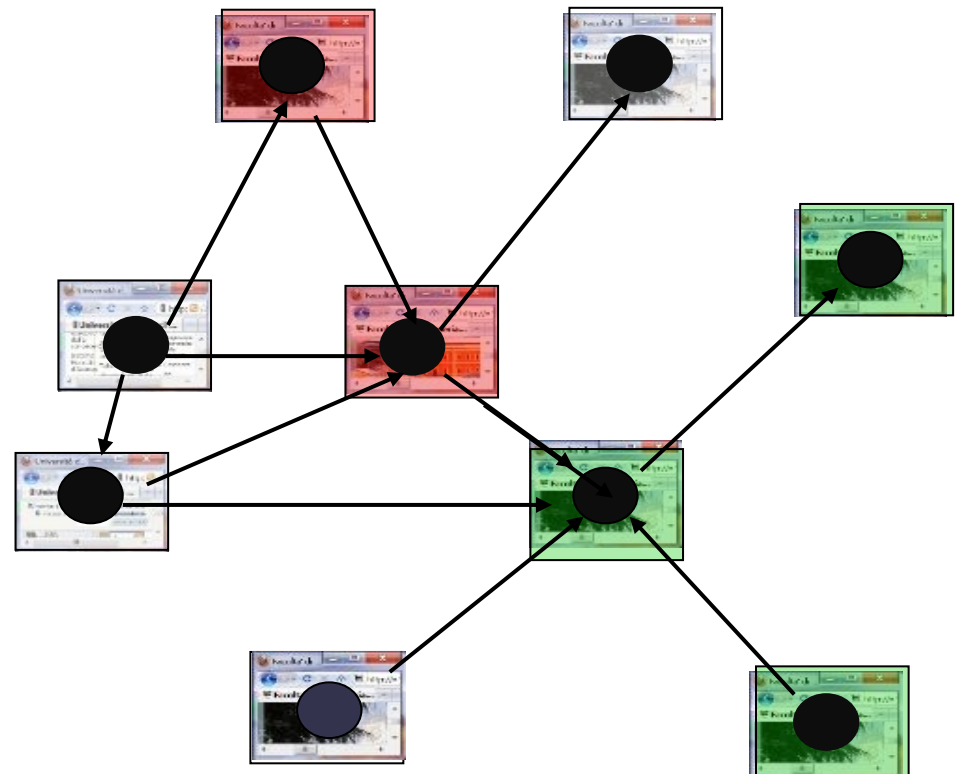
- ▶ A class of machine learning models for graph processing
- ▶ They take in input a graph and return an output at each node/edge



An example: web spam detection

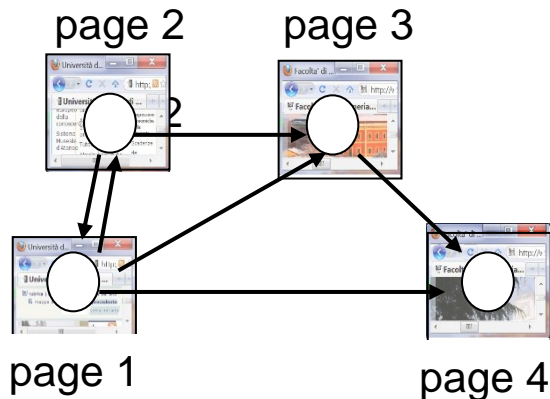
- ▶ Goal: learning by examples to detect spam pages using
 - Web connectivity
 - Page content

Red=spam
Green=not spam
Black=Unkown

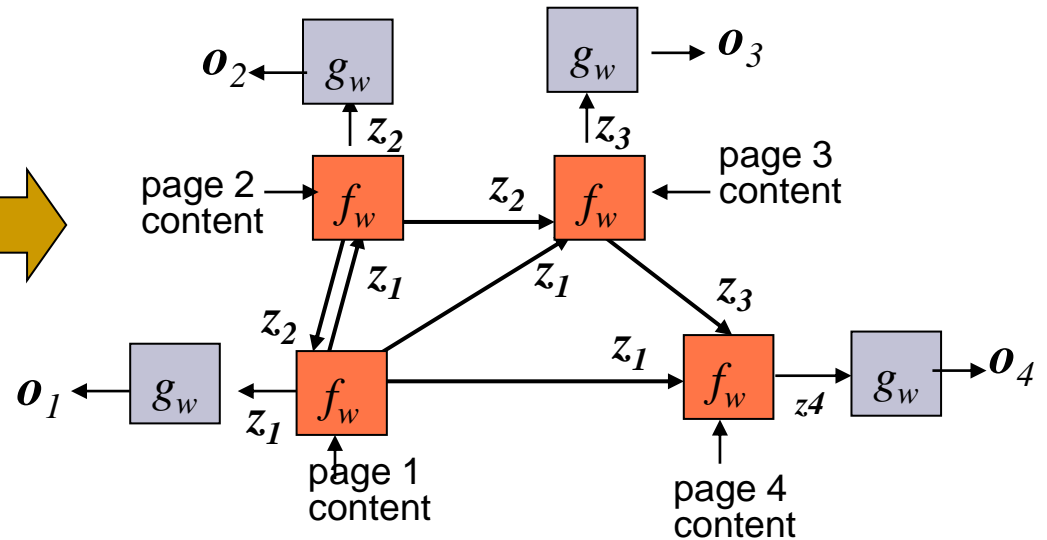
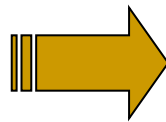


The GNN model (the first version)

- ▶ Two f and g neural modules (an instance for each node)
 - ▶ f compute a state z_v for each node
 - ▶ g compute an output o_v for each node
 - ▶ f modules are **locally connected** so as the graph
- ▶ All the modules share the same parameters



**Input graph
i.e., the web**



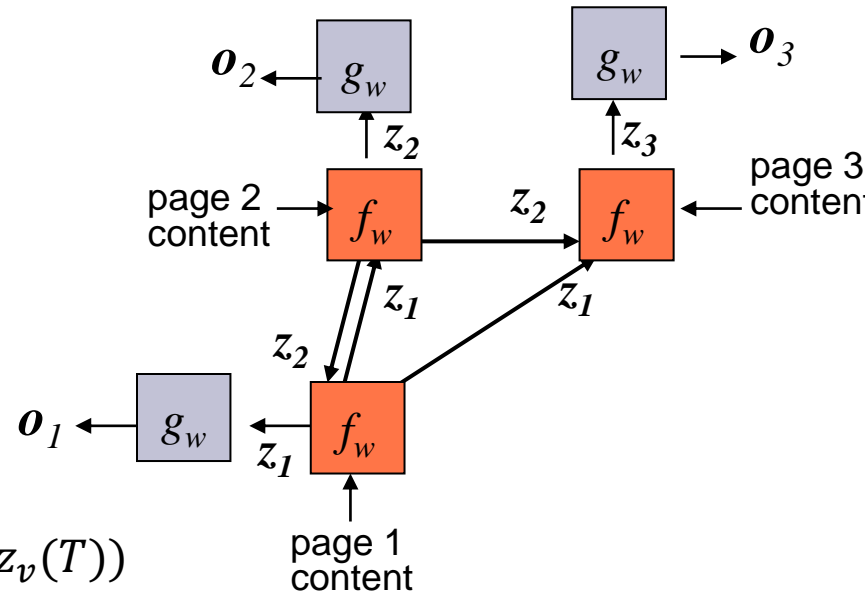
Encoding network



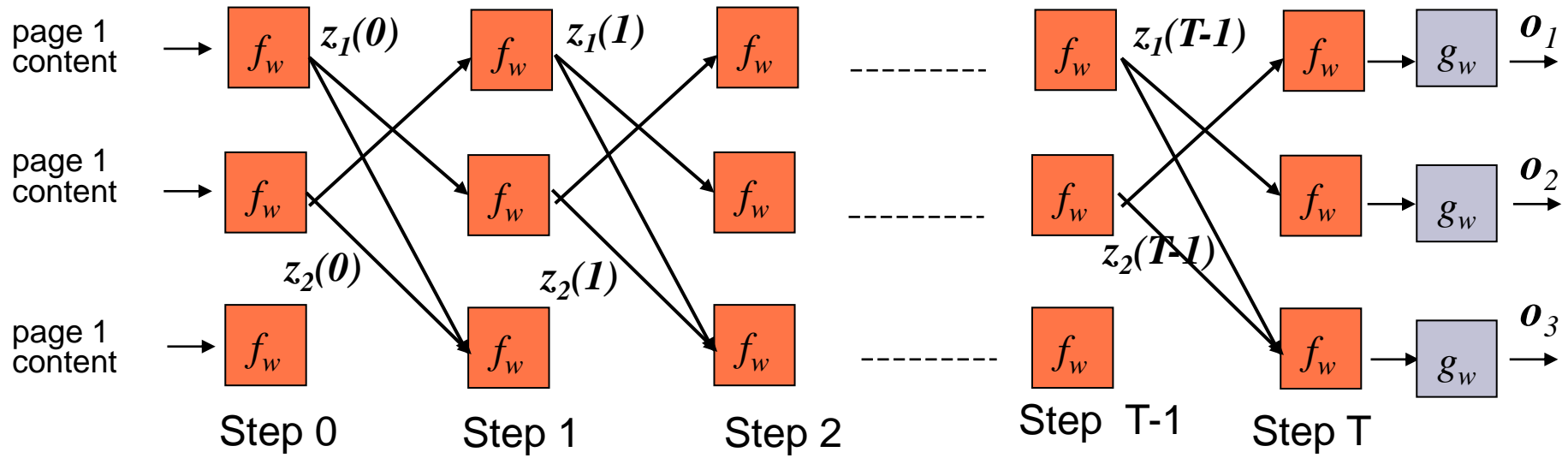
Output computation

- ▶ The output is computed by iterating the computation of the state until the state converges

$$z_v(t) = f_w(l_v, l_{ne[v]}, x_{ne[v]}(t-1)) \quad o_v = g_w(l_v, z_v(T))$$



The unfolding network





Modern GNNs

Modern message passage models

$$z_v(t) = \text{COMBINE}_w(x_v(t-1), \text{AGGRREGATE}_w(x_{ne[v]}(t-1)))$$

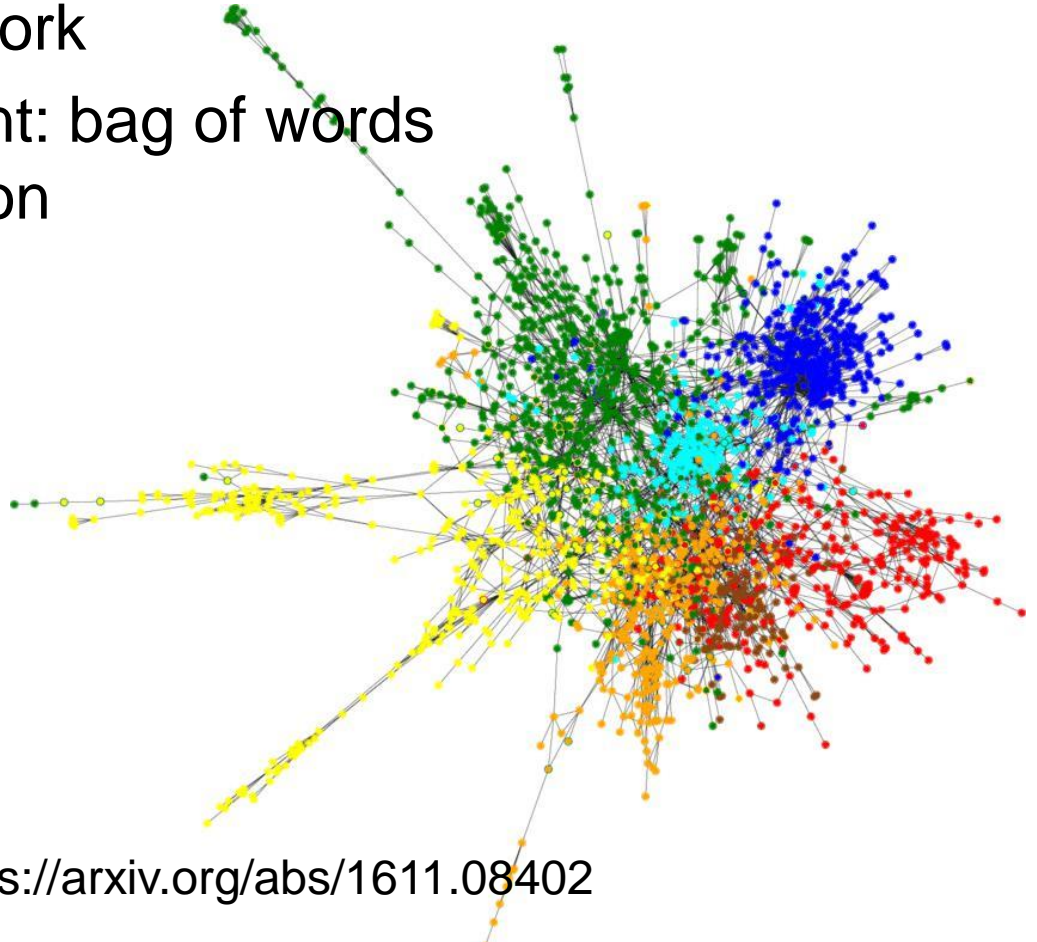
- ▶ Different parameters (and state dimension) at each step
- ▶ Special types of modules: pooling, focus of attention, explanation, ...
- ▶ Different types combination and aggregation functions
- ▶ GNNs for dynamic data
- ▶ ...



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Applications: document citation networks

- ▶ To classify research papers using
 - ▶ citation network
 - ▶ paper content: bag of words representation



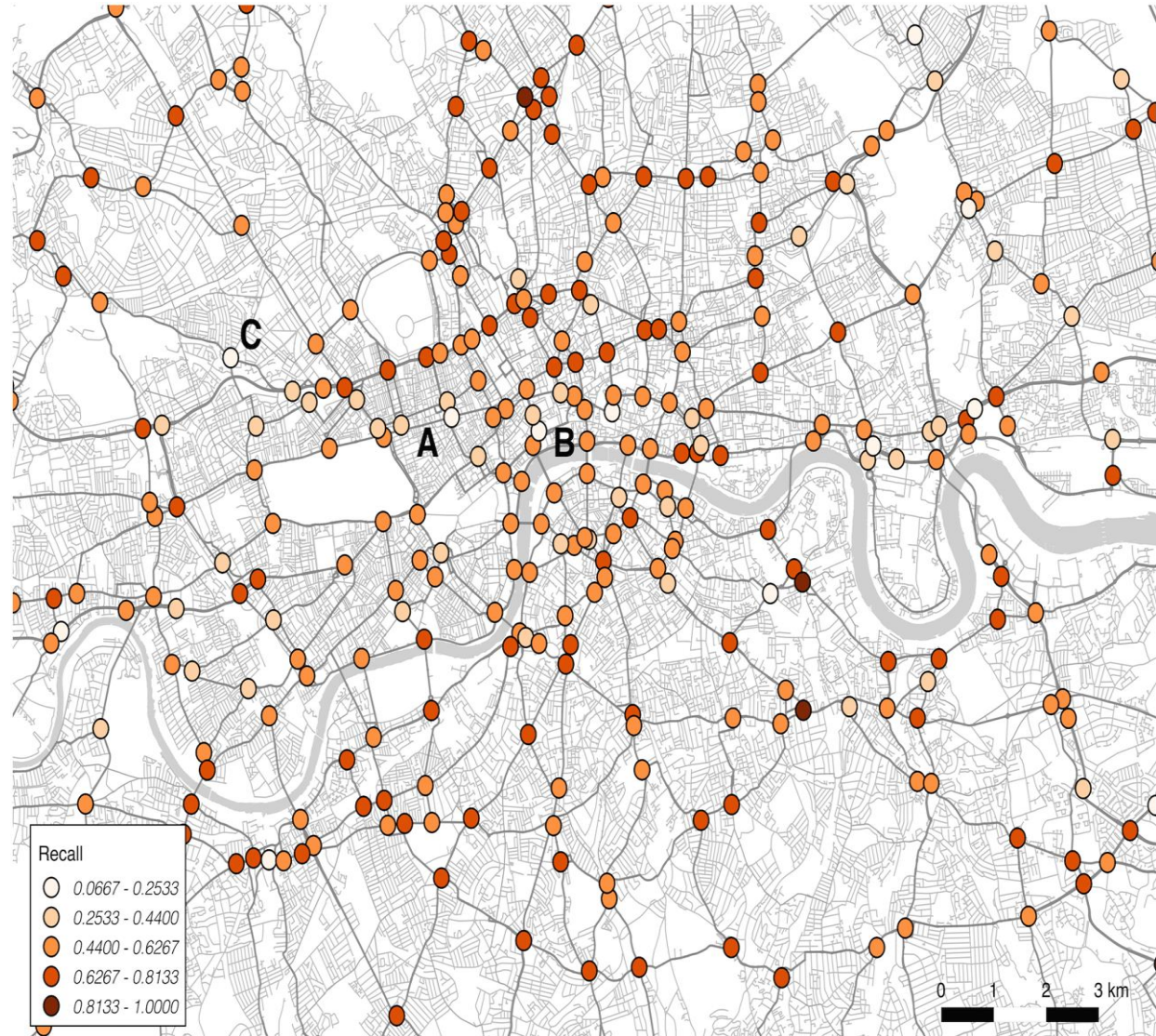
Source <https://arxiv.org/abs/1611.08402>



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Road networks

- To predict traffic learning for previous examples of traffic loads

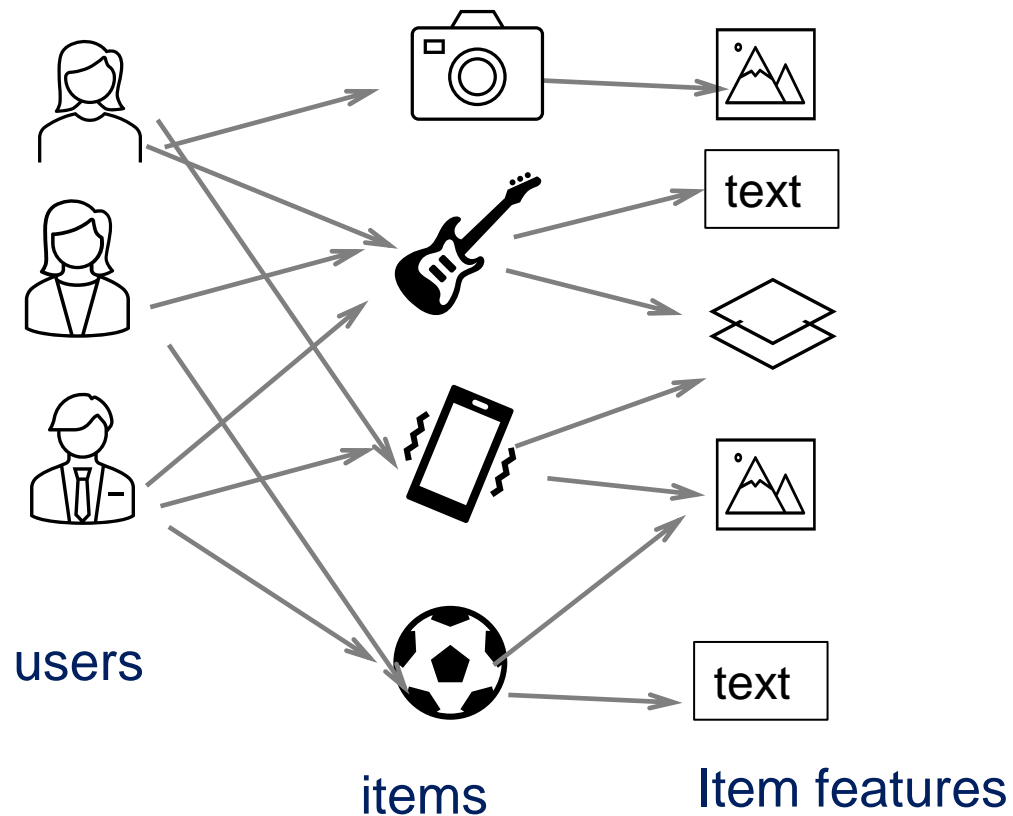


Source: <https://journals.plos.org/plosone/article/figure?id=10.1371/journal.pone.0127095.g004>



Large scale recommendation system

- ▶ To recommend items to users using
 - ▶ User-item network
 - ▶ Item features, eg images, text, ...





Other examples of applications

- ▶ Microsoft have experimented GNNs for program understanding
- ▶ Deep mind have used GNNs for traffic prediction, protein function prediction,
- ▶ Facebook has used GNNs to encode wikipedia graph
- ▶ Researchers at Large Hadron Collider at CERN will use GNNs to analyse data
- ▶

A future general case?

Previous applications regard homogenous graphs for single tasks

A company/organization aggregates into a single data warehouse all its information

- ▶ The data look as a single big graph

A lot of nice new problems to play with

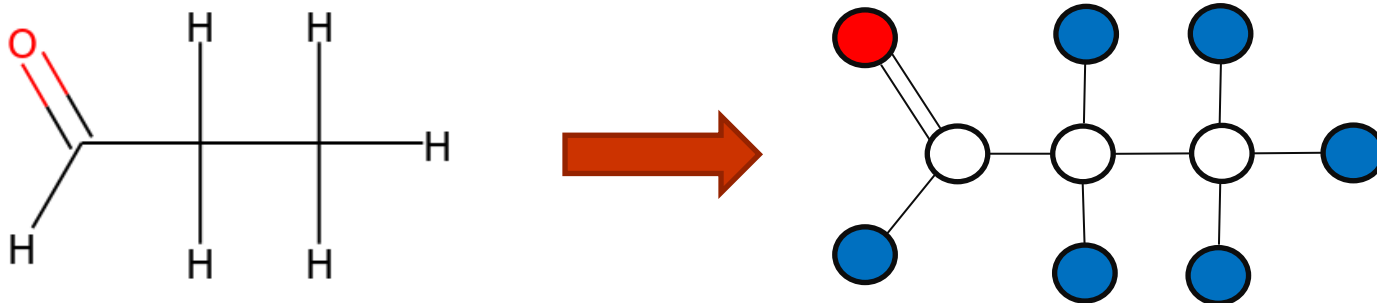
- ▶ Heterogenous patterns (nodes/edges)
- ▶ Different tasks to be solved (in sequence, contemporaneously)
- ▶ Different types of learning involved, e.g. inductive and transductive learning
- ▶ ...



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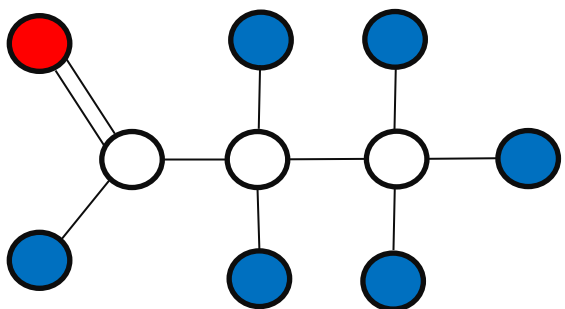
Molecular generative Graph Neural Networks for Drug Discovery (MG²N²) *with P. Bongini, M Bianchini*

Molecules are usually represented as undirected graphs: atoms correspond to nodes and bonds correspond to edges.

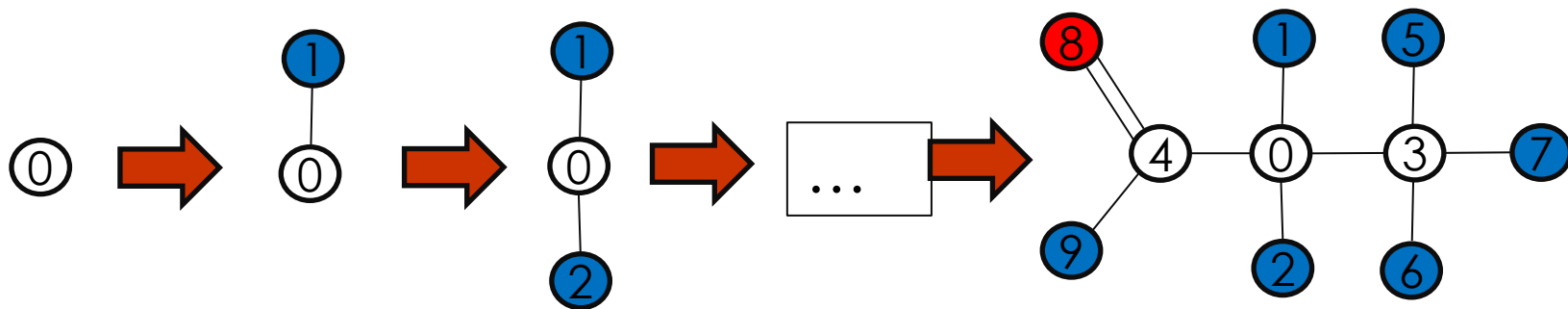




Molecular generative Graph Neural Networks for Drug Discovery (MG²N²)



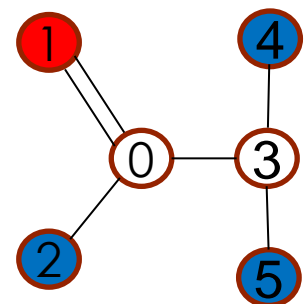
Generation starts from a graph composed of a single atom (sampled from the training set distribution). Each step is a standalone problem.



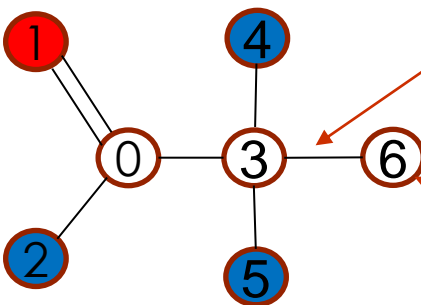


Molecular generative Graph Neural Networks for Drug Discovery (MG²N²)

Each step is split into three subproblems



P1: Do I generate another neighbour for node 3? Of which type?



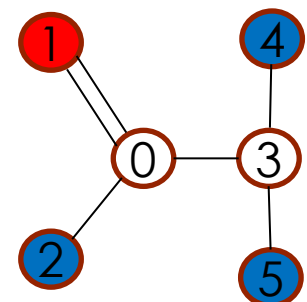
P2: Which is the type of bond (3,6)?

P3: Do I generate any extra bond: (0,6), (1,6), (2,6), (4,6), (5,6)?

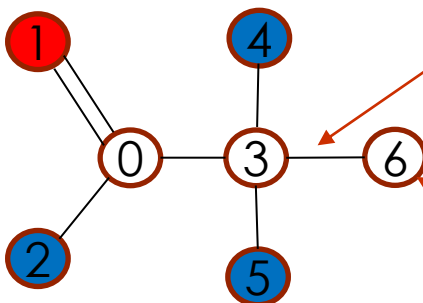


GNN Modules

Each step is split into three subproblems



P1: A node-classifier GNN is asked to predict the next neighbour of node 3: (C, H, N, O, F, Stop). If the answer is «Stop», jump to node 4.

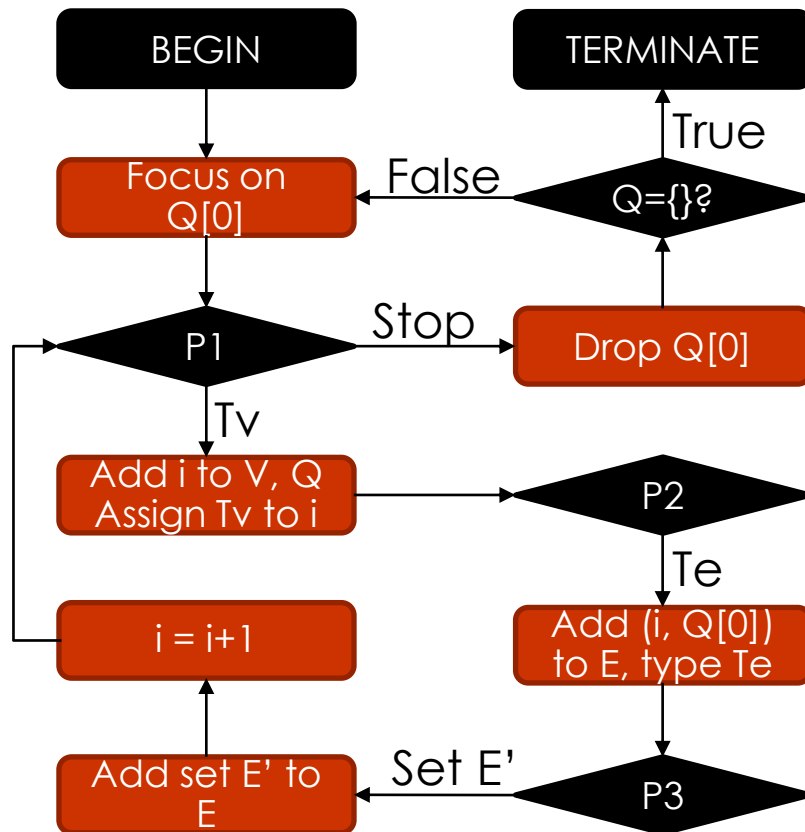


P2: Otherwise, an edge-classifier GNN is asked to predict the type of bond (3,6): (I, II, III).

P3: An edge predictor GNN decides if any extra bonds should be generated for the new node (and of which type).



Algorithm Chart



Generation of a graph G:

$G=(V,E)$

V: Vertex Set

E: Edge Set

Q: Expansion Queue

S: Starting Distribution

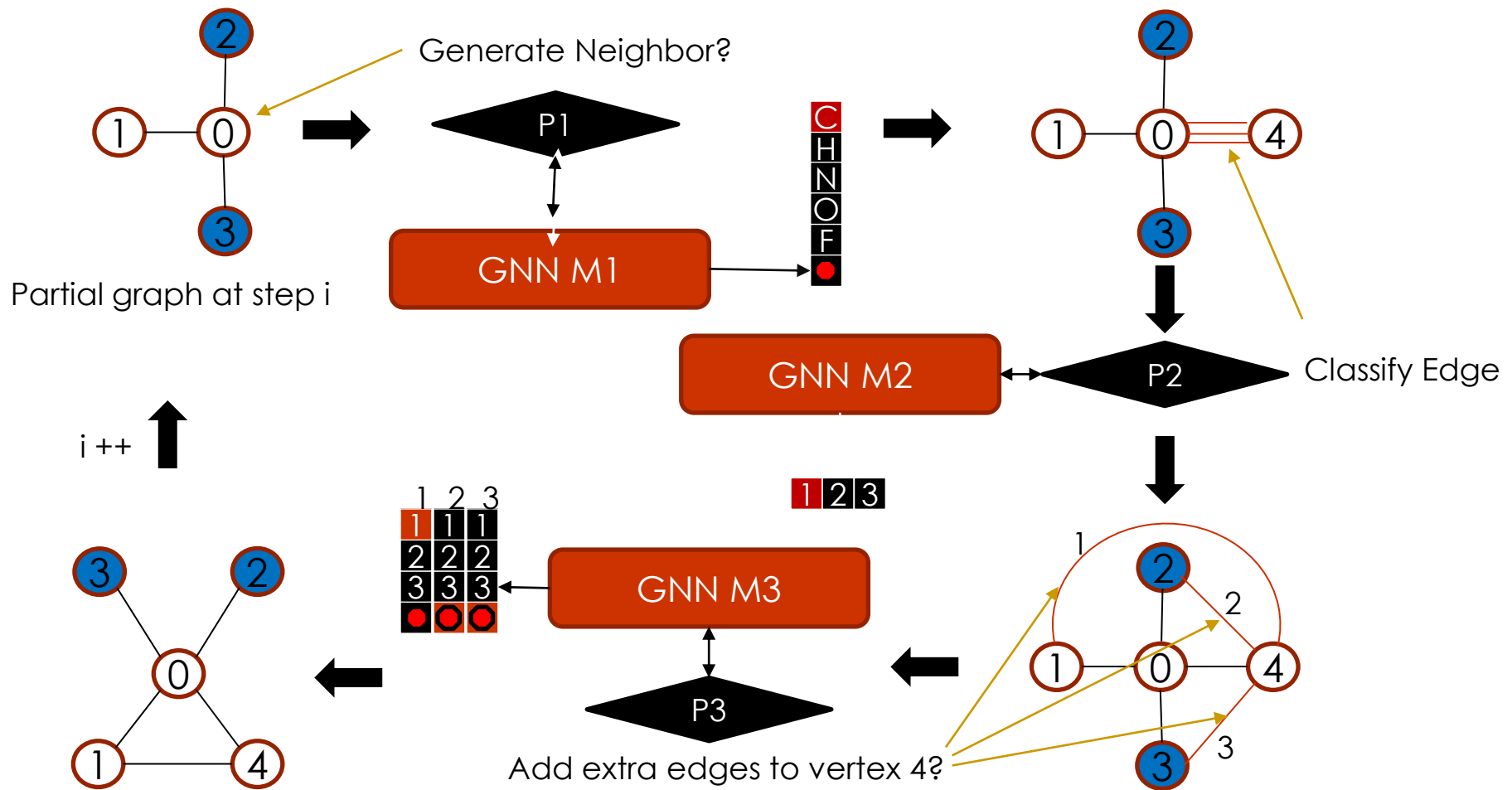
P1: Choose if to generate neighbor, and its type.

P2: Choose edge type

P3: Choose which edges to generate, and their types.

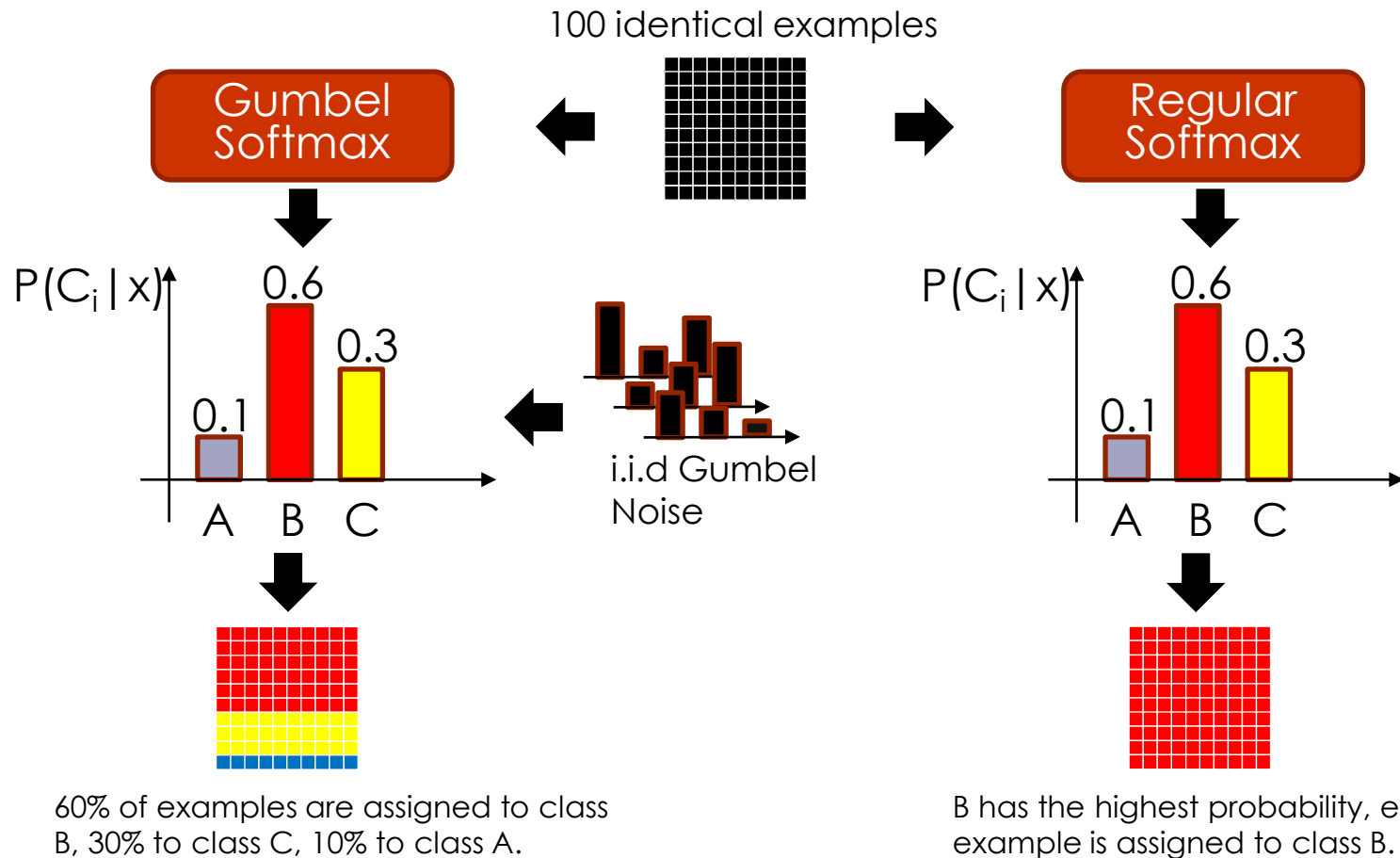
Initial conditions: $E= \{\}$, $V=\{0\}$, type of vertex 0 sampled from training set, $i=1$, $Q=\{0\}$.

Generation Step Example





Gumbel Softmax Classifier



About MG²N²

- ▶ GNN can use the whole graph information to decide
- ▶ Generation by iterative methods is "more explainable"
- ▶ GNN modules are trained separately
 - ▶ Make training/retraining much more flexible
 - ▶ It is based on an assumption their independence



QM9 Dataset

- ▶ Dataset of 134k molecules
- ▶ 5 atom types (CHNOF)
- ▶ 3 bond types (Single, double, triple)
- ▶ The objective is to generate new molecules (not found in QM9) which are chemically valid
- ▶ New molecules should have similar chemical properties with respect to a held-out test set (proof of generalization)

Experiments

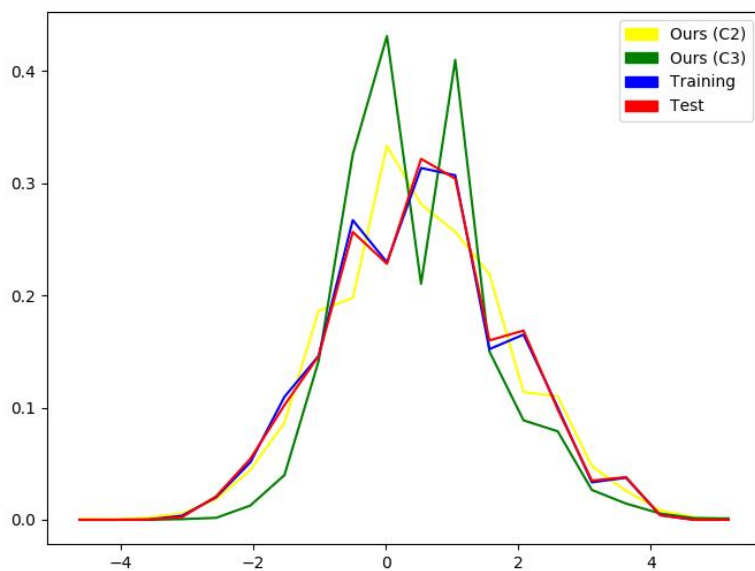
- ▶ Each experiment consists in generating 10K molecule graphs
- ▶ A held-out test set of 10K graphs will be used to compare their chemical properties
- ▶ Chemical Validity, Novelty and Uniqueness are assessed with the RdKit package
- ▶ Molecular Mass, logP and QED are measured with RdKit as well



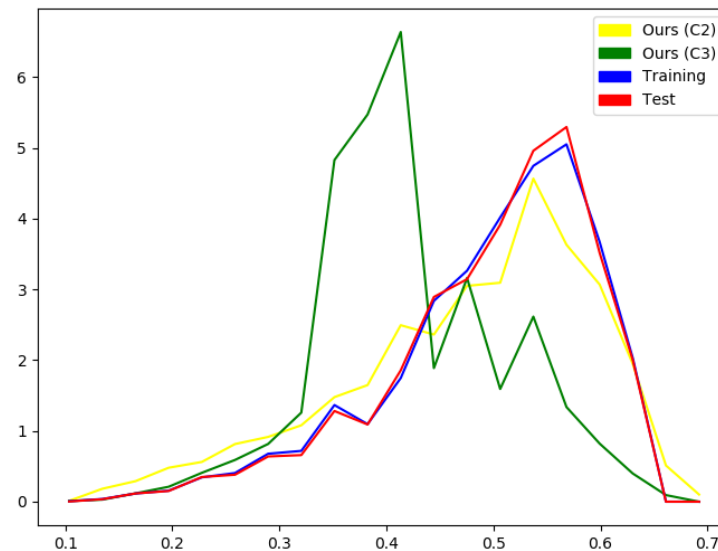
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Comparing chemical properties

logP score



QED score





Experimental Results

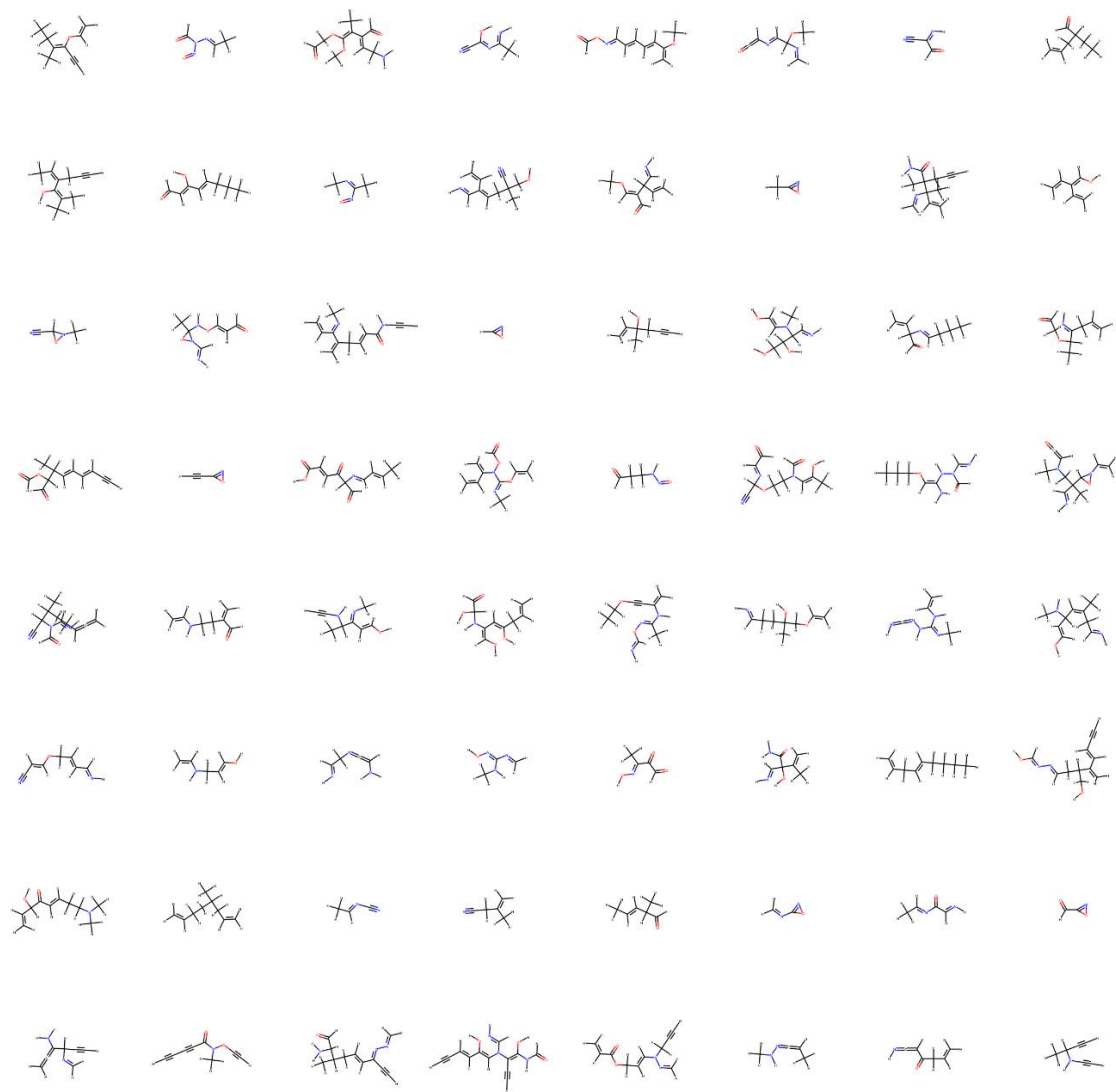
- ▶ A comparison was carried out with other models for the unconditioned generation of molecules.
- ▶ GrammarVAE is the best SMILES-based VAE on QM9
- ▶ MolGAN (RL-based) is the best sequential model on QM9
- ▶ GraphVAE is the best graph-based VAE (and state of the art) for unconditioned generation on QM9

Model	Valid	Unique	Novel	VUN	Avg. QED	Avg. logP	Avg. Mol. Wt.
GrammarVAE	0.602	0.093	0.809	0.045	-	-	-
GraphVAE	0.557	0.760	0.616	0.261	-	-	-
MolGAN	0.981	0.104	0.942	0.096	-	-	-
Ours(C2)	0.511	0.888	1.000	0.454	0.461 (0.116)	0.272 (1.336)	134.8 (45.7)
Ours(C3)	0.668	0.340	1.000	0.227	0.404 (0.088)	0.238 (1.093)	75.3 (52.8)
Test	-	-	-	-	0.482 (0.096)	0.270 (1.325)	127.3 (7.6)



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Generated Molecules





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Thank you for your attention!



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Reference

- ▶ Molecular generative Graph Neural Networks for Drug Discovery, P Bongini, M Bianchini, F Scarselli
Neurocomputing