

Recent approaches to Graph Neural Network

Andrei Nicolicioiu Iulia Duță {anicolicioiu,iduta}@bitdefender.com

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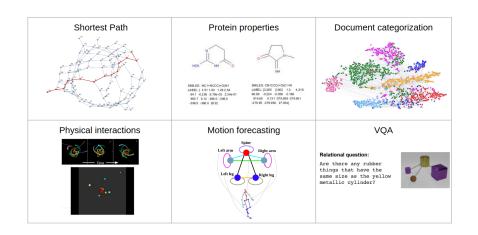
Introduction





Motivation





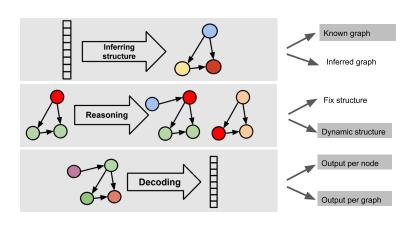
Motivation



- any data that can be structured in form of a graph has information that we want to use
- we want to design models that implicitly take advantage of known biases in the data
 - graphs gives us locality assumption: entities in a neighbourhood interacts more than distant ones
- we want models with properties similar with CNN
 - locality: interactions between neighbouring nodes
 - stationarity : all the interactions are the same at every position in graph

Graph approaches

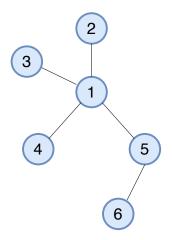




Capture from "Graph Networks: relational inductive biases for deep learning" talk, R. Pascanu



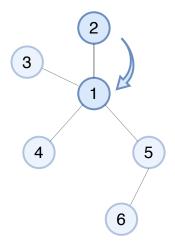
Process a graph in 3 steps





Process a graph in 3 steps

1. send messages from each node to its neighbours

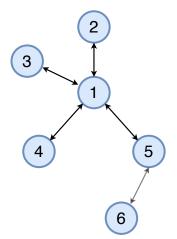


$$M_t(h_v^t, h_w^t, e_{vw}) \tag{1}$$



Process a graph in 3 steps

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$$m_v^{t+1} = \sum_{w \in N(v)} M_t(h_v^t, h_w^t, e_{vw})$$
 (2)



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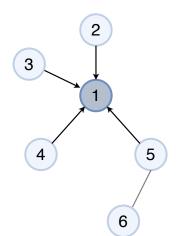
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2. with the received information update each node

$$h_{v}^{t+1} = U_{t}(h_{v}^{t}, m_{v}^{t+1})$$
 (3)





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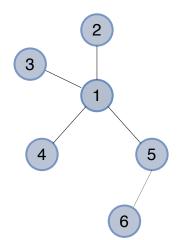
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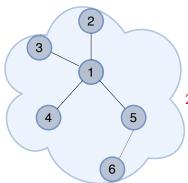
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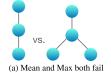
3. compute features for the whole graph with readout function *R*

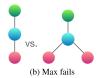


Neighborhood aggregation



- aggregation must be a permutation invariant function
- aggregation methods include: sum, avg, max-pooling, LSTM (Hamilton et al. [2017]), attention mechanisms
- ► Xu et al. [2019]



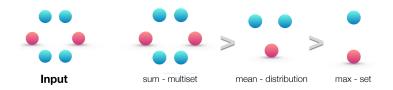




(c) Mean and Max both fair

Neighborhood aggregation





- sum: captures the full multiset
- ▶ mean: captures the distribution of elements of a given type
- ▶ max: captures the underlying set of a multiset

Convolutional networks on graphs for learning molecular fingerprints



Duvenaud, D. K., Maclaurin, D., Iparraguirre, J., Bombarell, R., Hirzel, T., Aspuru-Guzik, A., and Adams, R. P. Convolutional networks on graphs for learning molecular fingerprints. In Advances in neural information processing systems. NIPS 2015. Duvenaud et al. [2015]

Learning Molecular Fingerprints



- ▶ node v and edge vw vectorial representation: h_v , e_{vw}
- ▶ from each node *v* send to its neighbour *w* a linear projection of its node and edge features

$$M_t(h_v^t, h_w^t, e_{vw}) = W_t^{deg(v)}[h_v^t; e_{vw}]$$
 (5)

- no correlations between node and edge features
- different parameters for nodes with different degree
- update the node features by applying non-linearity on the sum of all messages

$$U_t(h_v^t, m_v^{t+1}) = \sigma(m_v^{t+1})$$
 (6)

Learning Molecular Fingerprints



- ▶ in order to get information from neighbours of order T, they use T message and update steps
- the final representation of the graph is the sum of projections of all nodes representations at every time step

$$R = \sum_{v \in G, t \in [1, T]} softmax(V_t h_v^t)$$
 (7)

Efficient computations



- ► the message from one node to another is just a linear projection of its features *Wh*
- all the projections are then summed
- these two operations could be written as matrix multiplications

$$H_{t+1} = AH_tW (8)$$

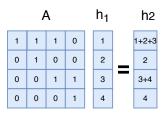
- $ightharpoonup H_tW$ projects all the features
- A is the adjacency matrix, and AH sums the neighbours of every node
- a model with T steps could just be made by stacking T such multiplications:

$$H_t = \sigma(A\sigma(...(AH_0W_0))W_t) \tag{9}$$

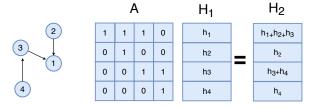
▶ this is also done in Kipf and Welling [2017]





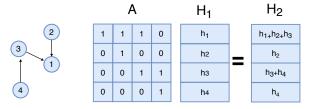




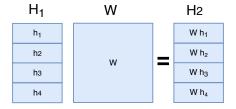


► AH: for every node sums all the neighbours features



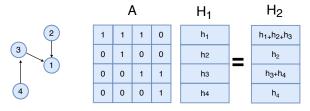


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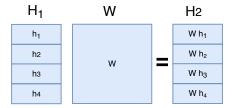


 $ightharpoonup H_1W$: applies linearity W to every row





► AH: for every node sums all the neighbours features



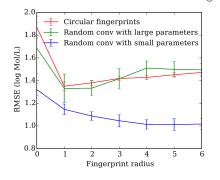
- $ightharpoonup H_1W$: applies linearity W to every row
- \triangleright AHW: projects all node features h_v and sums by A

Experiments



.NH₂

vertices represent individual atoms and edges represent bonds



- convolutional graph is used to compute features that are used as input to a classifier
- even with random weights the graph convolutional method is better than existing methods

Results



Dataset Units	Solubility [4] log Mol/L	Drug efficacy [5] EC ₅₀ in nM	Photovoltaic efficiency [8] percent
Predict mean Circular FPs + linear layer Circular FPs + neural net Neural FPs + linear layer	$ \begin{vmatrix} 4.29 \pm 0.40 \\ 1.71 \pm 0.13 \\ 1.40 \pm 0.13 \\ 0.77 \pm 0.11 \end{vmatrix} $	1.47 ± 0.07 1.13 ± 0.03 1.36 ± 0.10 1.15 ± 0.02	6.40 ± 0.09 2.63 ± 0.09 2.00 ± 0.09 2.58 ± 0.18
Neural FPs + neural net	$\textbf{0.52} \pm \textbf{0.07}$	1.16 ± 0.03	1.43 ± 0.09

Interaction Networks for Learning about Objects, Relations and Physics



Battaglia, P., Pascanu, R., Lai, M., and Rezende, D. J. (2016). Interaction networks for learning about objects, relations and physics. In Advances in neural information processing systems. NIPS 2016
Battaglia et al. [2016]

Interaction Networks



- \blacktriangleright the message from v to w is an MLP that takes into account:
 - source object representation
 - target object representation
 - relation attributes

$$M_t(h_v^t, h_w^t, e_{vw}) = \mathsf{MLP}([h_v^t; \boldsymbol{h}_w^t; e_{vw}]) \tag{10}$$

- update function is also an MLP that takes into account:
 - current node representation
 - sum of all received messages
 - external effects on current node

$$U_t(h_v^t, m_v^t, x_v) = \mathsf{MLP}([\boldsymbol{h_v^t}; m_v^t; \boldsymbol{x_v}])$$
 (11)

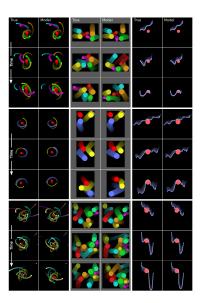
the aggregation step is used only when a global representation is needed:

$$R = \mathbf{MLP}(\sum_{v \in G} (h_v^T)) \tag{12}$$

 the global representation use intermediate representation from every layer

Experiments

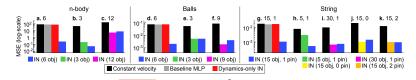




- generalized well to systems with fewer or greater number of objects (due to shared parameters across nodes)
- network trained on single-step predictions can be used to simulate thousands of steps

Experiments





Domain	Constant velocity	Baseline	Dynamics-only IN	IN
n-body	82	79	76	0.25
Balls	0.074	0.072	0.074	0.0020
String	0.018	0.016	0.017	0.0011

- Constant velocity: output the input velocity
- ▶ Baseline: MLP over a flattened vector data
- ► Dynamics-only IN: Interaction model without message from neighbours
- ▶ IN: Interaction Network model

Graph Attention Networks

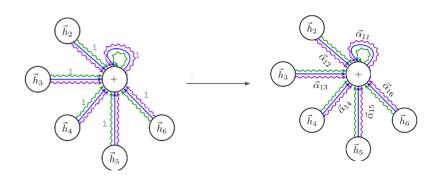


P. Velikovi, G. Cucurull, A. Casanova, A. Romero, P. Li, and Y. Bengio. Graph attention networks. In International Conference on Learning Representations, ICLR 2018 Velikovi et al. [2018]

Graph Attention Networks



▶ introduce an aggregation module for received messages:



$$\alpha_{ij} = \frac{\exp(ReLU(a^T[Wh_i; Wh_j]))}{\sum_{k \in N_i} \exp(ReLU(a^T[Wh_i; Wh_k]))}$$
(13)

Graph Attention Networks



- replace the summation used to aggregate neighbours with a weighted sum based on an attention module
- offer a better visualization of how the network use the structure
- can be used with unordered, variable number of neighbours

Experiments



Two kinds of tasks:

- transductive: Cora, Citeseer, Pubmed document categorization
 - the entire dataset (train + test) forms a single graph known apriori
 - ▶ 1 graph, 20 nodes per class with known labels, 3-7 classes
- inductive: PPI multilabeling on proteins:
 - train and test splits have different graphs
 - ▶ 20 graphs, 2372 nodes in average, 50 feats per node, 121 classes

Experiments



► Transductive:

Transductive				
Method	Cora	Citeseer	Pubmed	
MLP	55.1%	46.5%	71.4%	
ManiReg (Belkin et al., 2006)	59.5%	60.1%	70.7%	
SemiEmb (Weston et al., 2012)	59.0%	59.6%	71.7%	
LP (Zhu et al., 2003)	68.0%	45.3%	63.0%	
DeepWalk (Perozzi et al., 2014)	67.2%	43.2%	65.3%	
ICA (Lu & Getoor, 2003)	75.1%	69.1%	73.9%	
Planetoid (Yang et al., 2016)	75.7%	64.7%	77.2%	
Chebyshev (Defferrard et al., 2016)	81.2%	69.8%	74.4%	
GCN (Kipf & Welling 2017)	81.5%	70.3%	79.0%	
MoNet (Monti et al., 2016)	$81.7\pm0.5\%$	_	$78.8\pm0.3\%$	
GCN-64*	$81.4 \pm 0.5\%$	$70.9 \pm 0.5\%$	79.0 ± 0.3%	
GAT (ours)	$83.0 \pm 0.7\%$	72.5 \pm 0.7%	79.0 ± 0.3%	

Inductive:

Inductive

Method	PPI
Random	0.396
MLP	0.422
GraphSAGE-GCN (Hamilton et al., 2017)	0.500
GraphSAGE-mean (Hamilton et al., 2017)	0.598
GraphSAGE-LSTM (Hamilton et al., 2017)	0.612
GraphSAGE-pool (Hamilton et al., 2017)	0.600
GraphSAGE*	0.768
Const-GAT (ours)	0.934 ± 0.006
GAT (ours)	0.973 ± 0.002

Gated graph sequence neural networks



Li, Yujia, Tarlow, Daniel, Brockschmidt, Marc, and Zemel, Richard. Gated graph sequence neural networks. ICLR, 2016. Li et al. [2016]

Gated graph sequence neural networks



- used for tasks with sequential output
- intuition: use recurrent network with graph operations at every step
- from each node send a linear projection (dependent on the type of edge) of its node features

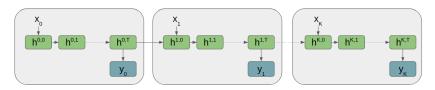
$$M_t(h_v^t, h_w^t, e_{vw}) = W_{e_{vw}} h_v^t + b$$
 (14)

ightharpoonup compute m_{V} as the sum of all messages and update the state of each node by using recurrent GRU cell rules

$$U_t(h_v^t, m_v^{t+1}) = GRU(h_v^t, m_v^{t+1})$$
 (15)

Gated graph sequence neural networks





readout function

$$R_k = tanh(\sum_{v \in G} \sigma(i(h_v^{k,T}, x_v)) \odot tanh(j(h_v^{k,T}, x_v)))$$
 (16)

- depending on the task the model could
 - receive input x_{k+1} after every output y_k
 - continue processing without any input

Experiments



Artificial tasks:

- ▶ *bAbl task 4*: (Two Arguments) to recognize subjects and objects in text
- ▶ *bAbl task 15*: (Deduction) reasoning from general statements to examples
- ► bAbl task 16: (Induction) reasoning from examples to statements
- ▶ bAbl task 18: (Size) reasoning about size of an object
- ► bAbl task 19*: (Path Finding) find the path between 2 locations
- ► Shortest Path*: find the shortest path between 2 locations (unique)
- ► Eulerian Circuit*: find eulerian circuit between 2 locations.

^{*}Sequencial tasks

Experiments



► Single output:

Task	RNN	LSTM	GG-NN
bAbI Task 4	97.3±1.9 (250)	97.4±2.0 (250)	100.0±0.0 (50)
bAbI Task 15	48.6 ± 1.9 (950)	50.3 ± 1.3 (950)	100.0 ± 0.0 (50)
bAbI Task 16	33.0 ± 1.9 (950)	37.5 ± 0.9 (950)	100.0 ± 0.0 (50)
bAbI Task 18	88.9±0.9 (950)	88.9±0.8 (950)	100.0 ± 0.0 (50)

► Sequencial output:

Task	RNN	LSTM		GGS-NNs	
bAbI Task 19 Shortest Path Eulerian Circuit	24.7±2.7 (950) 9.7±1.7 (950) 0.3±0.2 (950)	28.2±1.3 (950) 10.5±1.2 (950) 0.1±0.2 (950)	100.0± 0.0 (50)	92.5±5.9 (100)	99.0±1.1 (250)

Overview



Fingerprints:

$$M_{t}(h_{v}^{t}, h_{w}^{t}, e_{vw}) = W_{t}^{deg(v)}[h_{v}^{t}; e_{vw}]$$
 $U_{t}(h_{v}^{t}, m_{v}^{t+1}) = \sigma(m_{v}^{t+1})$
 $R = \sum_{v \in G, t \in [1, T]} softmax(V_{t}h_{v}^{t})$

Attention:

$$M_{t}(h_{v}^{t}, h_{w}^{t}) = W_{t}h_{v}^{t}$$

$$U_{t}(h_{v}^{t}, m_{v}^{t+1}) = \sum_{w \in N(v)} \alpha_{i}M_{t}(h_{v}^{t}, h_{w}^{t})$$

$$- - -$$

Interactions:

$$\begin{aligned} M_{t}(h_{v}^{t}, h_{w}^{t}, e_{vw}) &= MLP([h_{v}^{t}; h_{w}^{t}; e_{vw}]) \\ U_{t}(h_{v}^{t}, m_{v}^{t}, x_{v}) &= MLP([h_{v}^{t}; m_{v}^{t}; x_{v}]) \\ R &= MLP(\sum_{v \in G} (h_{v}^{T})) \end{aligned}$$

Gated:

$$\begin{aligned} M_t(h_v^t, h_w^t, \mathbf{e}_{vw}) &= W_{\mathbf{e}_{vw}} h_v^t + b \\ U_t(h_v^t, m_v^{t+1}) &= GRU(h_v^t, m_v^{t+1}) \\ R_k &= tanh(\sum_{v \in G} \sigma(i(h_v^{k,T}, x_v)) \odot \\ \odot tanh(j(h_v^{k,T}, x_v))) \end{aligned}$$

Questions?



Thank you!

References I



- P. Battaglia, R. Pascanu, M. Lai, D. J. Rezende, et al. Interaction networks for learning about objects, relations and physics. In *Advances in neural information processing systems*, pages 4502–4510, 2016.
- D. K. Duvenaud, D. Maclaurin, J. Iparraguirre, R. Bombarell, T. Hirzel, A. Aspuru-Guzik, and R. P. Adams. Convolutional networks on graphs for learning molecular fingerprints. In Advances in neural information processing systems, pages 2224–2232, 2015.
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- K. Xu, W. Hu, J. Leskovec, and S. Jegelka. How powerful are graph neural networks? In *International Conference on Learning Representations*, 2019. URL https://openreview.net/forum?id=ryGs6iA5Km.