

Graph Neural Processes

Towards Bayesian Graph Neural Networks

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Introduction

Problem

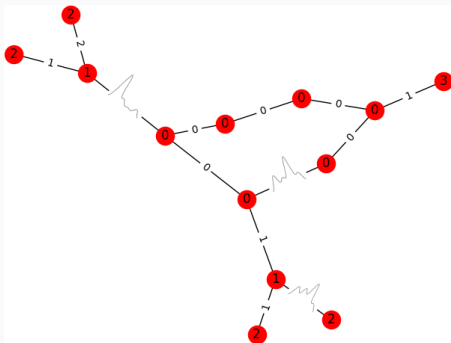
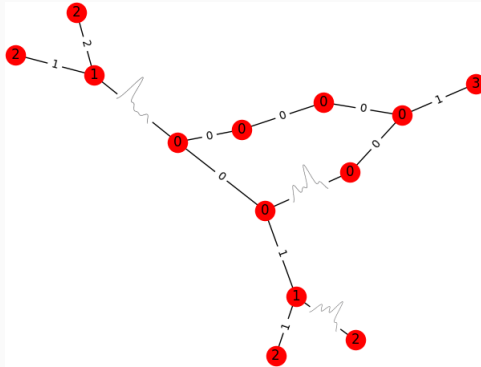


Figure 1: Graph with imputed edge distributions

Problem



$$G = (a, V, E)$$

$$V = \{v_i\}_{i=1:N^v}$$

$$E = \{e_k, v_k, u_k\}_{k=1:N^e}$$

Motivation

- Visual scene understanding
- Few-shot learning
- Learning dynamics of physical systems
- Traffic prediction
- Multi-agent systems
- Natural language processing
- semi-supervised text classification

Background

- Operate on Graph Structured Input

- Operate on Graph Structured Data
- Neural Operations on Graphs (e.g., spectral convolution)

Use Eigeninformation from traditional Graph Laplacian to perform convolution

$$L = D - A$$

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- Fixed sized input

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- Neural Operations on Graphs (e.g., spectral convolution)
- Fixed sized input
- Difficult to scale
- No measure of uncertainty

$$P(Y_T|X_T, X_C, Y_C) \iff P(Y_T|X_T, r_C)$$

Conditional Neural Processes

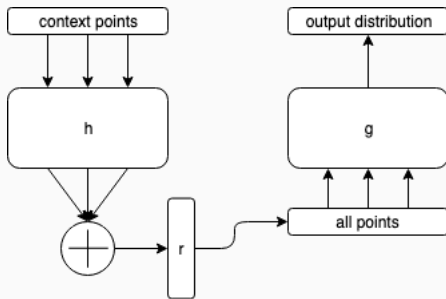
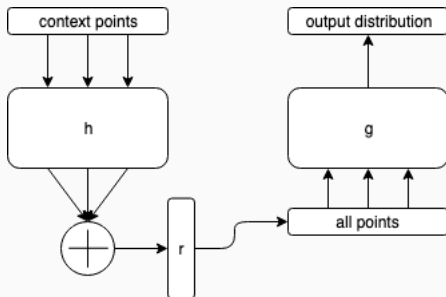


Figure 2: Conditional Neural Process Architecture

Conditional Neural Processes



$$r_i = h_{\theta}(\vec{x}_i) \quad \forall \vec{x}_i \in X_C \quad (1)$$

$$r_C = r_1 \oplus r_2 \oplus r_3 \oplus \dots \oplus r_n \quad (2)$$

$$z_i = g_{\phi}(\vec{y}_i | r_C) \quad \forall \vec{y}_i \in X_T \quad (3)$$

- Flexible

- Flexible
- Scalable

- Flexible
- Scalable
- Measures uncertainty

Our Method

- Conditional Neural Process on Graphs

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- Edges are context points

- Conditional Neural Process on Graphs
- Edges are context points
- Spectral Features

Normalized Symmetric Graph Laplacian

$$L = I_n - D^{-1/2}AD^{-1/2}$$

$$\Lambda = \sigma(L)$$

Take the eigenvectors of the graph laplacian and select the first k to obtain the local spectral eigen features

$$\Lambda|_k = (\Lambda_{kj})_{\substack{k \in r \\ 1 \leq j \leq m}}$$

Features

Or, more clearly

$$\Lambda|_k = (\Lambda_{kj})_{\substack{k \in r \\ 1 \leq j \leq m}}$$

```
def get_local_spectral_eigen_features(G, ind, k):  
    """  
    G - graph (networkx graph)  
    ind - location of the context point edge in the adjacency matrix  
    k - hyperparameter tuned based on the size of the input graphs in X_C  
    """  
  
    A = nx.adjacency_matrix(G).toarray()  
    N = A.shape[0]  
  
    diags = A.sum(axis=1)**(-1/2)  
    D = scipy.sparse.spdiags(diags.flatten(), [0], N, N, format='csr').toarray()  
  
    L = np.eye(N) - D.dot(A).dot(D) # calculate normalized graph laplacian  
    val, vec = np.linalg.eig(L)  
  
    return vec[ind][:k]
```

Additionally the value and degree of each node on the edge are used as features

How does it work?

Algorithm Walk Through

```
1: Let  $X$  input graphs
2: for  $t = 0, \dots, n_{\text{epochs}}$  do
3:   for  $x_i$  in  $X$  do
4:     Sample  $p \leftarrow \text{unif}(p_0, p_1)$ 
5:     Assign  $n_{\text{context points}} \leftarrow p \cdot |\text{Edges}(x_i)|$ 
6:     Sparsely Sample  $x_i^{cp} \leftarrow x_i|_{n_{\text{context points}}}$ 
7:     Compute degree and adj matrix  $D, A$  for graph  $x_i^{cp}$ 
8:     Compute  $L \leftarrow I_{n_{\text{context points}}} - D^{-\frac{1}{2}}AD^{-\frac{1}{2}}$ 
9:     Define  $F^{cp}$  as empty feature matrix for  $x_i^{cp}$ 
10:    Define  $F$  as empty feature matrix for full graph  $x_i$ 
11:    for edge  $k$  in  $x_i$  do
12:      Extract eigenfeatures  $\Lambda|_k$  from  $L$ 
13:      Concatenate  $[\Lambda|_k; v_k; u_k; d(v_k); d(u_k)]$  where  $v_k, u_k$  are the attribute values at the node, and  $d(v_k), d(u_k)$  the degree at the node.
14:      if edge  $k \in x_i^{cp}$  then
15:        Append features for context point to  $F^{cp}$ 
16:      end if
17:      Append features for all edges to  $F$ 
18:    end for
19:    Encode and aggregate  $r_C \leftarrow h_{\theta}(F^{cp})$ 
20:    Decode  $\tilde{x}_i \leftarrow g_{\phi}(F|r_C)$ 
21:    Calculate Loss  $l \leftarrow \mathcal{L}(\tilde{x}_i, x_i)$ 
22:    Step Optimizer
23:  end for
24: end for
```

Experiments

16 Datasets

Dataset	$ X $	$ \bar{N} $	$ \bar{E} $	$ \cup \{e_R\} $
AIDS	2000	15.69	16.20	3
BZR_MD	306	21.30	225.06	5
COX2_MD	303	26.28	335.12	5
DHFR_MD	393	23.87	283.01	5
ER_MD	446	21.33	234.85	5
Mutagenicity	4337	30.32	30.77	3
MUTAG	188	17.93	19.79	4
PTC_FM	349	14.11	14.48	4
PTC_FR	351	14.56	15.00	4
PTC_MM	336	13.97	14.32	4
Tox21_AHR	8169	18.09	18.50	4
Tox21_ARE	7167	16.28	16.52	4
Tox21_aromatase	7226	17.50	17.79	4
Tox21_ARLBD	8753	18.06	18.47	4
Tox21_ATAD5	9091	17.89	18.30	4
Tox21_ER	7697	17.58	17.94	4

Table 1: Features of the explored data sets

Results

Results

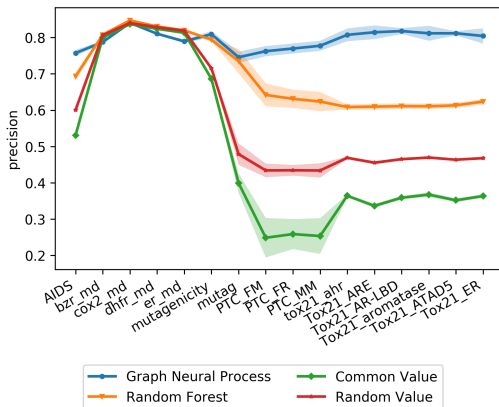


Figure 3: Experimental precision graph compared with baselines, we see our method performs achieves a $\sim .2$ higher precision on average.

Results

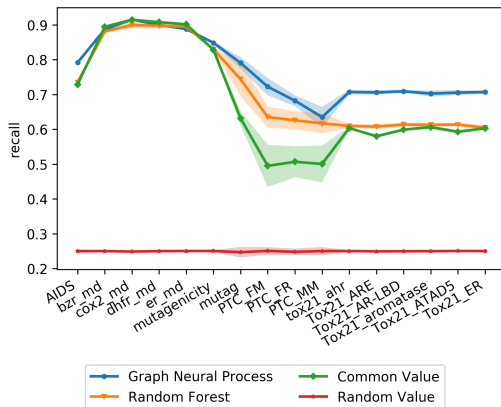


Figure 4: Experimental recall graph compared with baselines

Results

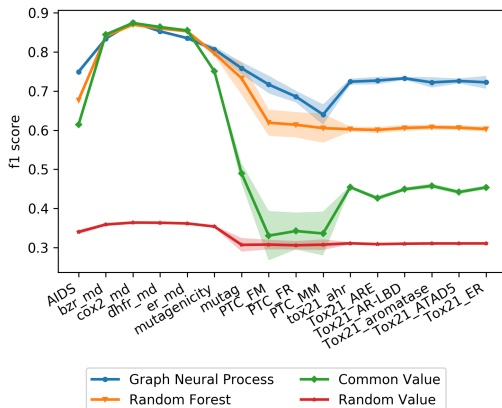


Figure 5: Experimental F1-score graph compared with baselines.

Results

Dataset	RF			RV			CV			GNP		
	P	R	F1	P	R	F1	P	R	F1	P	R	F1
AIDS	0.69	0.74	0.68	0.60	0.25	0.34	0.53	0.73	0.61	0.76	0.79	0.75
bzr_md	0.81	0.88	0.84	0.81	0.25	0.36	0.80	0.89	0.84	0.79	0.89	0.83
cox2_md	0.85	0.90	0.87	0.84	0.25	0.36	0.84	0.91	0.87	0.84	0.92	0.88
dhfr_md	0.83	0.90	0.86	0.83	0.25	0.36	0.82	0.91	0.86	0.81	0.90	0.85
er_md	0.82	0.90	0.85	0.82	0.25	0.36	0.81	0.90	0.86	0.79	0.89	0.84
mutagenicity	0.79	0.83	0.80	0.72	0.25	0.35	0.69	0.83	0.75	0.81	0.85	0.81
mutag	0.73	0.74	0.73	0.48	0.25	0.31	0.40	0.63	0.49	0.75	0.79	0.76
PTC_FM	0.64	0.64	0.62	0.43	0.25	0.31	0.25	0.50	0.33	0.76	0.72	0.72
PTC_FR	0.63	0.63	0.61	0.43	0.25	0.31	0.26	0.51	0.34	0.77	0.68	0.69
PTC_MM	0.62	0.62	0.61	0.43	0.25	0.31	0.25	0.50	0.34	0.78	0.64	0.64
tox21_ahr	0.61	0.61	0.60	0.47	0.25	0.31	0.36	0.60	0.45	0.81	0.71	0.72
Tox21_ARE	0.61	0.61	0.60	0.46	0.25	0.31	0.34	0.58	0.43	0.81	0.71	0.73
Tox21_AR-LBD	0.61	0.61	0.61	0.47	0.25	0.31	0.36	0.60	0.45	0.82	0.71	0.73
Tox21_aromatase	0.61	0.61	0.61	0.47	0.25	0.31	0.37	0.61	0.46	0.81	0.70	0.72
Tox21_ATAD5	0.61	0.61	0.61	0.46	0.25	0.31	0.35	0.59	0.44	0.81	0.71	0.73
Tox21_ER	0.62	0.60	0.60	0.47	0.25	0.31	0.36	0.60	0.45	0.80	0.71	0.72

Table 2: Experimental Results. RF=Random Forest; RV=Random edge label; CV=most common edge label; GNP=Graph Neural Process. P=precision; R=recall; F1=F1 score. Statistically significant bests are in bold with non-significant ties bolded across methods.

See <https://arxiv.org/abs/1902.10042> for complete list of references