

GraphCare: Enhancing Healthcare Predictions With Personalized Knowledge Graphs

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Overview

- Motivation
- Challenges
- Method
- Experiments

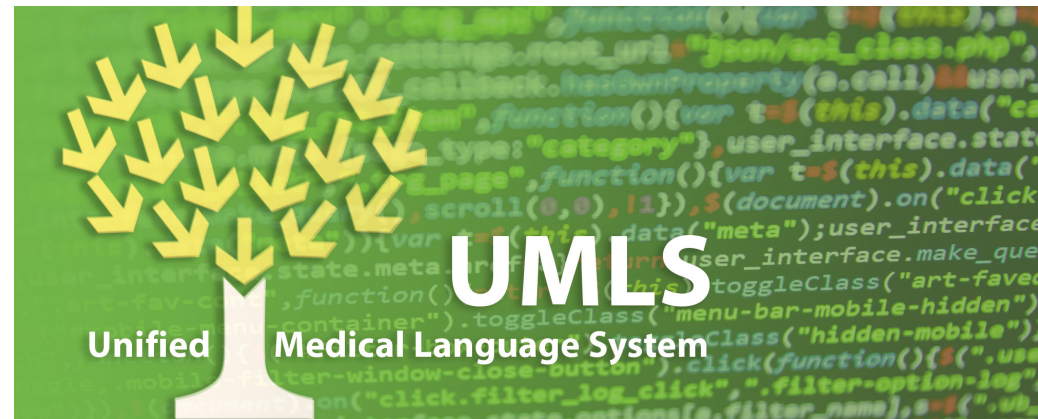
Motivation

1. Personalized KG is needed for personalized treatment.
2. Existing works mainly focus on
 - 1) simple hierarchical relations
 - 2) the inner graphical structure of EHRwhile there are many external biomedical KGs available

LLM as Knowledge Base



UMLS-KG

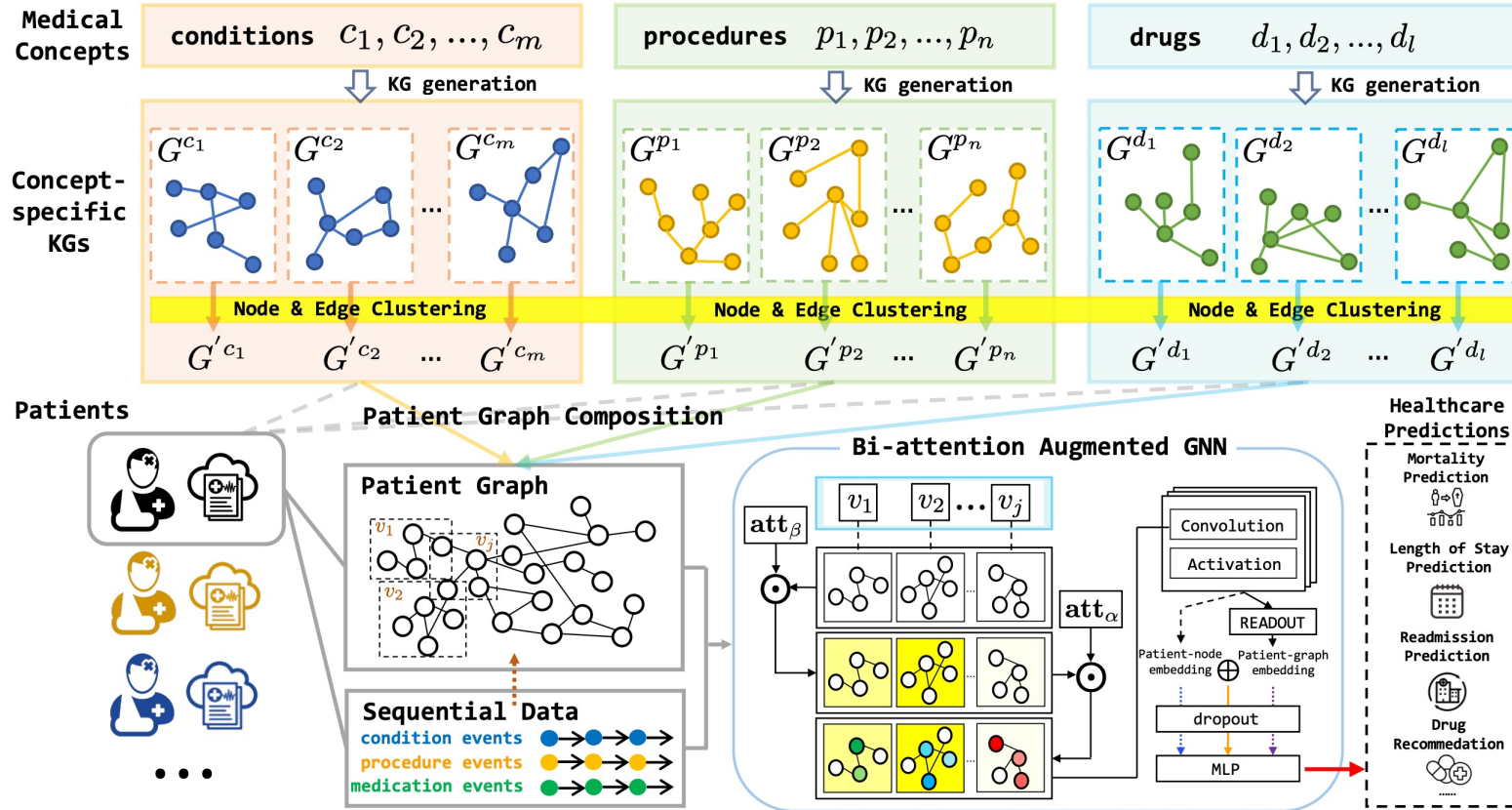


Challenges

1. How to construct personalized KGs utilizing external knowledge bases?
 - We construct medical concept-specific KGs
2. How to improve time-series clinical (EHR) predictions with those KGs?
 - We treat personalized knowledge graphs as patient representations

Method

Our proposed method - GraphCare



Step 1: Generate concept-specific KGs for every medical concept using LLM prompts and by subsampling from existing KGs. Perform clustering on nodes and edges across these KGs.

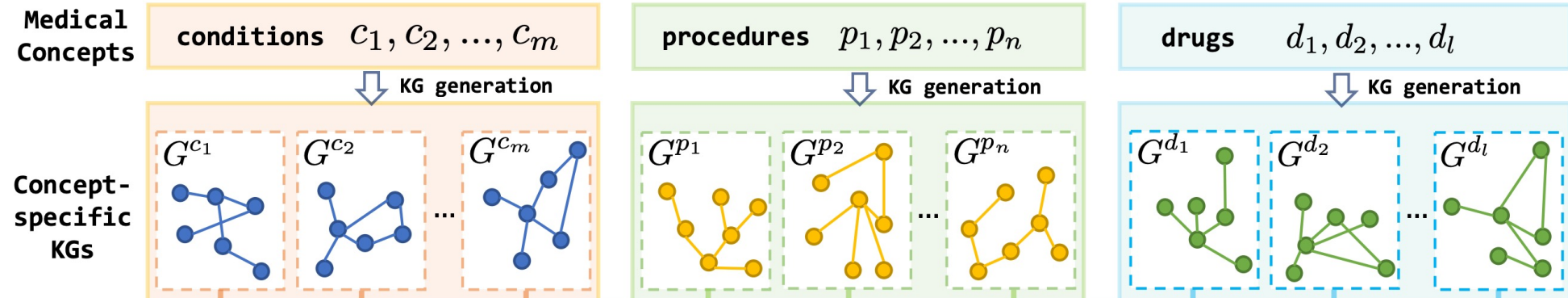
Step 2: For each patient, merge relevant concept-specific KGs to form a personalized KG.

Step 3: Employ Bi-attention Augmented (BAT) Graph Neural Network (GNN) to predict based on the personalized KGs.

Method

GraphCare – Concept-specific KG Generation

Step 1: Generate concept-specific KGs for every medical concept using LLM prompts and by subsampling from existing KGs. Perform clustering on nodes and edges across these KGs.



1.1 Knowledge extraction from LLM through prompting:

Given a prompt (a medical condition/procedure/drug), extrapolate as many relationships as possible of it and provide a list of updates.

The relationships should be helpful for healthcare prediction (e.g., drug recommendation, mortality prediction, readmission prediction ...)

Each update should be exactly in format of [ENTITY 1, RELATIONSHIP, ENTITY 2]. The relationship is directed, so the order matters.

Both ENTITY 1 and ENTITY 2 should be noun.

Any element in [ENTITY 1, RELATIONSHIP, ENTITY 2] should be conclusive, make it as short as possible.

Do this in both breadth and depth. Expand [ENTITY 1, RELATIONSHIP, ENTITY 2] until the size reaches 100.

{example}

prompt: {term}

updates:

```
if category == "condition":
```

```
example = \
```

```
"""Example:
```

```
prompt: systemic lupus erythematosus
```

```
updates: [[systemic lupus erythematosus, is an, autoimmune condition], [systemic lupus erythematosus, may cause, nephritis], [anti-nuclear antigen, is a test for, systemic lupus erythematosus], [systemic lupus erythematosus, is treated with, steroids], [methylprednisolone, is a, steroid]]
```

```
"""
```

```
elif category == "procedure":
```

```
example = \
```

```
"""Example:
```

```
prompt: endoscopy
```

```
updates: [[endoscopy, is a, medical procedure], [endoscopy, used for, diagnosis], [endoscopic biopsy, is a type of, endoscopy], [endoscopic biopsy, can detect, ulcers]]
```

```
"""
```

```
elif category == "drug":
```

```
example = \
```

```
"""Example:
```

```
prompt: iobenzamic acid
```

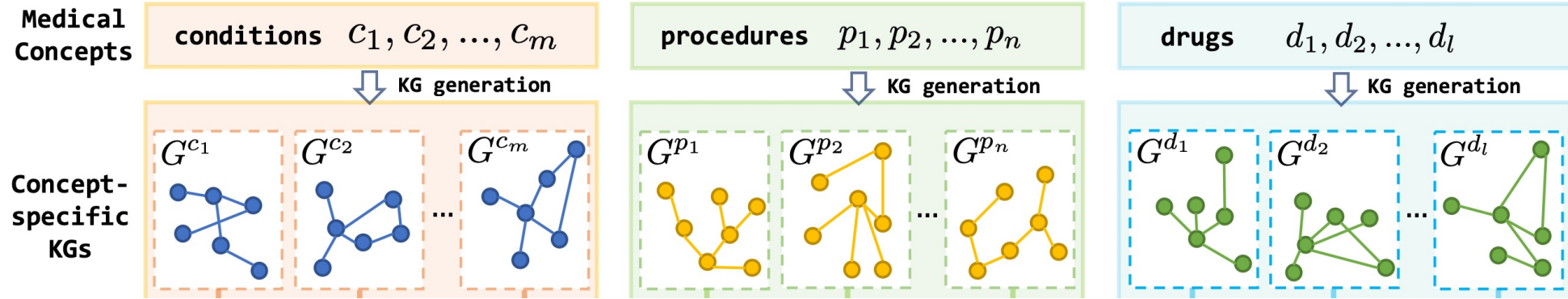
```
updates: [[iobenzamic acid, is a, drug], [iobenzamic acid, may have, side effects], [side effects, can include, nausea], [iobenzamic acid, used as, X-ray contrast agent], [iobenzamic acid, formula, C16H13I3N2O3]]
```

```
"""
```

Method

GraphCare – Concept-specific KG Generation

Step 1: Generate concept-specific KGs for every medical concept using LLM prompts and by subsampling from existing KGs. Perform clustering on nodes and edges across these KGs.



1.2 Subgraph sampling from existing KG:

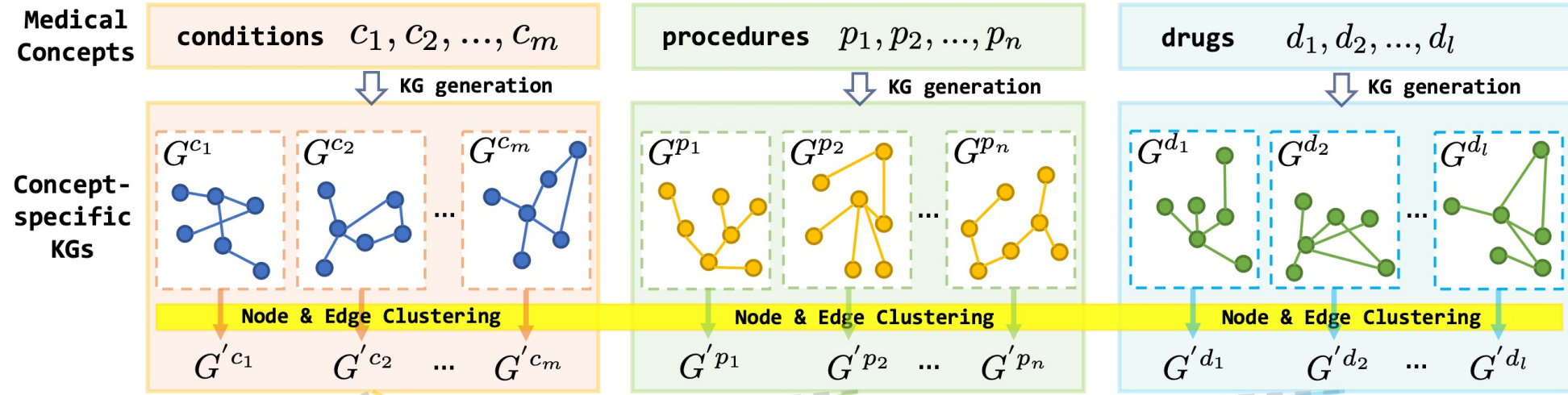
```
1: procedure SUBGRAPHSAMPLING(medical concept  $e$ , KG  $\mathcal{G}$ , hop limit  $\kappa$ , window size  $\epsilon$ )
2:   Initialize an empty list  $Q$  and an empty graph  $G_{\text{sub}(\kappa)}^e = (\mathcal{V}_{\text{sub}(\kappa)}^e, \mathcal{E}_{\text{sub}(\kappa)}^e)$ 
3:   Add  $e$  to  $Q$ 
4:    $\mathcal{V}_{\text{sub}(\kappa)}^e \leftarrow \mathcal{V}_{\text{sub}(\kappa)}^e \cup \{e\}$ 
5:   for  $i = 1$  to  $\kappa$  do
6:     Initialize an empty list  $Q_{\text{next}}$ 
7:     for all  $\text{ent} \in Q$  do
8:       if  $i = 1$  then
9:         Retrieve all triples  $(\text{ent}, \text{rel}, \text{ent}')$  or  $(\text{ent}', \text{rel}, \text{ent})$  from  $\mathcal{G}$ 
10:      else
11:        Randomly retrieve  $\epsilon$  triples  $(\text{ent}, \text{rel}, \text{ent}')$  or  $(\text{ent}', \text{rel}, \text{ent})$  from  $\mathcal{G}$ 
12:      end if
13:      Add retrieved triples to  $\mathcal{E}_{\text{sub}(\kappa)}^e$ 
14:       $\mathcal{V}_{\text{sub}(\kappa)}^e \leftarrow \mathcal{V}_{\text{sub}(\kappa)}^e \cup \{\text{ent}'\}$ 
15:      Add  $\text{ent}'$  to  $Q_{\text{next}}$ 
16:    end for
17:     $Q \leftarrow Q_{\text{next}}$ 
18:  end for
19:  return  $G_{\text{sub}(\kappa)}^e$ 
```

(For the first hop, we retrieve all the triples containing the concept. For the second (and higher) hop, we randomly retrieve triples of window size ϵ)

Method

GraphCare – Concept-specific KG Generation

Step 1: Generate concept-specific KGs for every medical concept using LLM prompts and by subsampling from existing KGs. Perform clustering on nodes and edges across these KGs.



1.3 Node & Edge Clustering

Based on the word embedding of nodes and edges, we apply agglomerative clustering to get two mappings:

$$\begin{aligned} C_{\mathcal{V}} : \mathcal{V} &\rightarrow \mathcal{V}' \\ C_{\mathcal{E}} : \mathcal{E} &\rightarrow \mathcal{E}' \end{aligned}$$

New global graph (node clusters as new nodes; edge clusters as new edges):

$$\longrightarrow G' = (\mathcal{V}', \mathcal{E}')$$

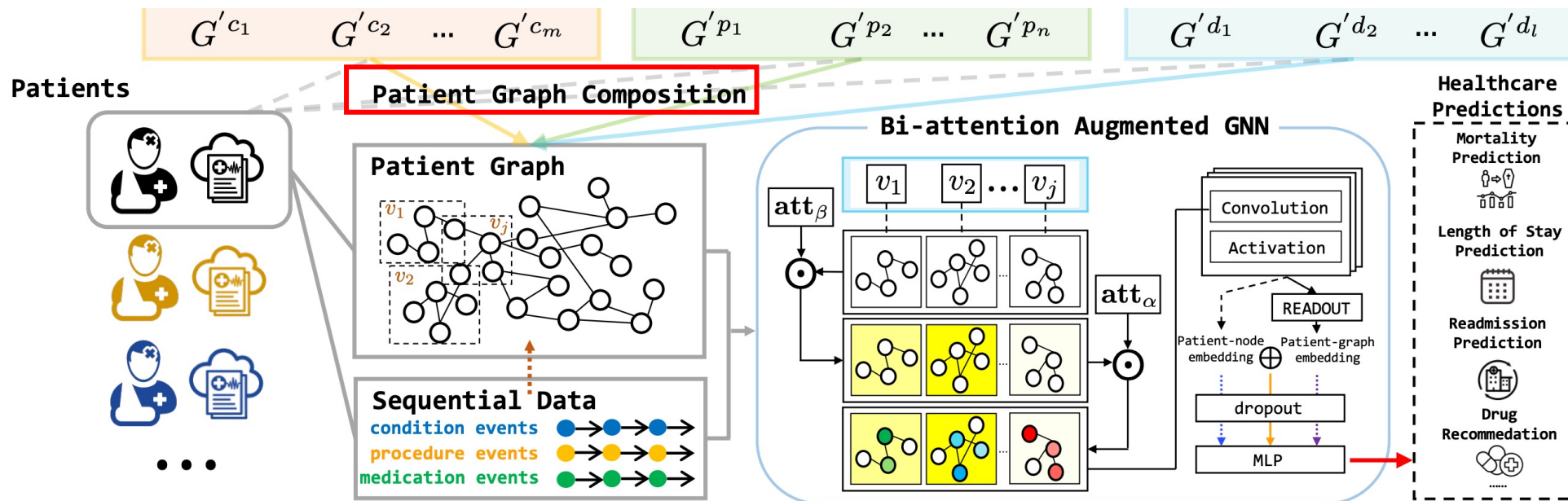
New concept-specific graph:

$$\longrightarrow G'^e = (\mathcal{V}'^e, \mathcal{E}'^e) \subset G'$$

Method

GraphCare – Personalized KG Composition

Step 2: For each patient, merge relevant concept-specific KGs to form a personalized KG.

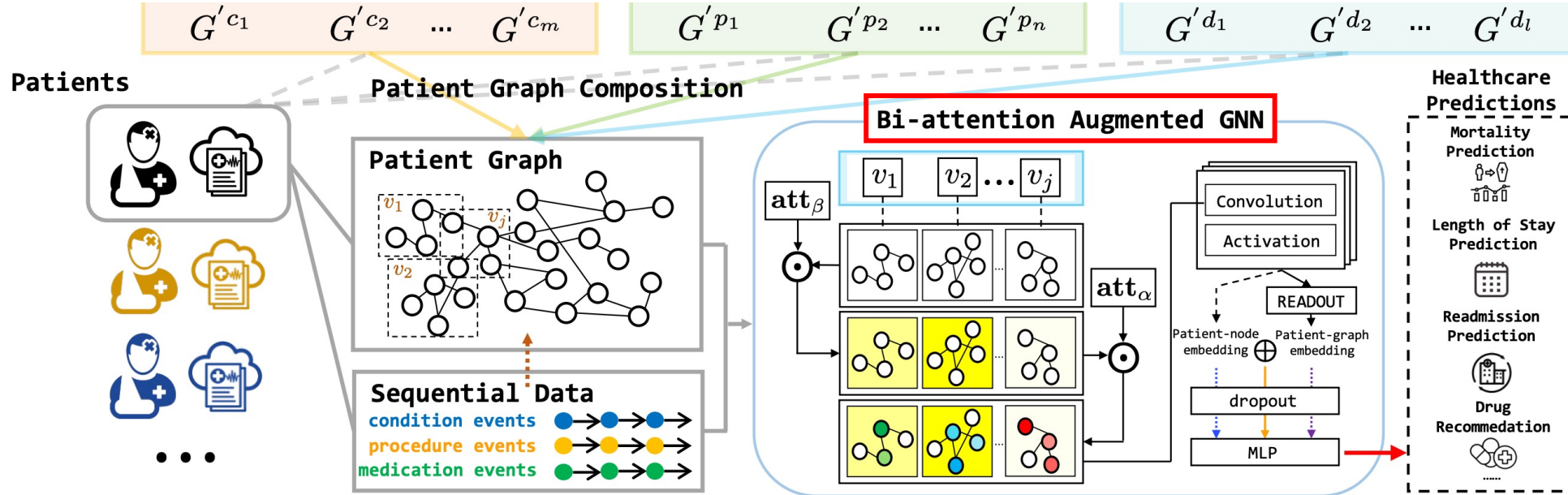


For each patient, we compose their personalized KG by merging the clustered KGs of their medical concepts. We create a patient node (\mathcal{P}) and connect it to its direct EHR nodes in the graph. The personalized KG for a patient can be represented as $G_{\text{pat}} = (\mathcal{V}_{\text{pat}}, \mathcal{E}_{\text{pat}})$, where $\mathcal{V}_{\text{pat}} = \mathcal{P} \cup \{\mathcal{V}^{e_1}, \mathcal{V}^{e_2}, \dots, \mathcal{V}^{e_\omega}\}$ and $\mathcal{E}_{\text{pat}} = \epsilon \cup \{\mathcal{E}^{e_1}, \mathcal{E}^{e_2}, \dots, \mathcal{E}^{e_\omega}\}$, with $\{e_1, e_2, \dots, e_\omega\}$ being the medical concepts directly associated with the patient, ω being the number of concepts, and ϵ being the edge connecting \mathcal{P} and $\{e_1, e_2, \dots, e_\omega\}$. Further, as a patient is represented as a sequence of J visits (Choi et al., 2016a), the *visit-subgraphs* for patient i can be represented as $G_{\text{pat}(i)} = \{G_{i,1}, G_{i,2}, \dots, G_{i,J}\} = \{(\mathcal{V}_{i,1}, \mathcal{E}_{i,1}), (\mathcal{V}_{i,2}, \mathcal{E}_{i,2}), \dots, (\mathcal{V}_{i,J}, \mathcal{E}_{i,J})\}$ for visits $\{x_1, x_2, \dots, x_J\}$ where $\mathcal{V}_{i,j} \subseteq \mathcal{V}_{\text{pat}(i)}$ and $\mathcal{E}_{i,j} \subseteq \mathcal{E}_{\text{pat}(i)}$ for $1 \leq j \leq J$.

Method

GraphCare – Bi-attention Augmented GNN

Step 3: Employ Bi-attention Augmented (BAT) Graph Neural Network (GNN) to predict based on the personalized KGs.



$$\alpha_{i,j,1}, \dots, \alpha_{i,j,M} = \text{Softmax}(\mathbf{W}_\alpha \mathbf{g}_{i,j} + \mathbf{b}_\alpha),$$

$$\beta_{i,1}, \dots, \beta_{i,N} = \lambda^\top \text{Tanh}(\mathbf{w}_\beta^\top \mathbf{G}_i + \mathbf{b}_\beta), \quad \text{where } \lambda = [\lambda_1, \dots, \lambda_N],$$

$$\mathbf{h}_{i,j,k}^{(l+1)} = \sigma \left(\mathbf{W}^{(l)} \sum_{j' \in J, k' \in \mathcal{N}(k) \cup \{k\}} \left(\underbrace{\alpha_{i,j',k'}^{(l)} \beta_{i,j'}^{(l)} \mathbf{h}_{i,j',k'}^{(l)}}_{\text{Node aggregation term}} + \underbrace{\mathbf{w}_{\mathcal{R}(k,k')}^{(l)} \mathbf{h}_{(i,j,k) \leftrightarrow (i,j',k')}}_{\text{Edge aggregation term}} \right) + \mathbf{b}^{(l)} \right)$$

Method

GraphCare – Bi-attention Augmented GNN

Step 3: Employ Bi-attention Augmented (BAT) Graph Neural Network (GNN) to predict based on the personalized KGs.

Attention Initialization

Table 4: Keyword candidates we attempted for attention initialization. We **highlight** the keywords we finally used in the experiments.

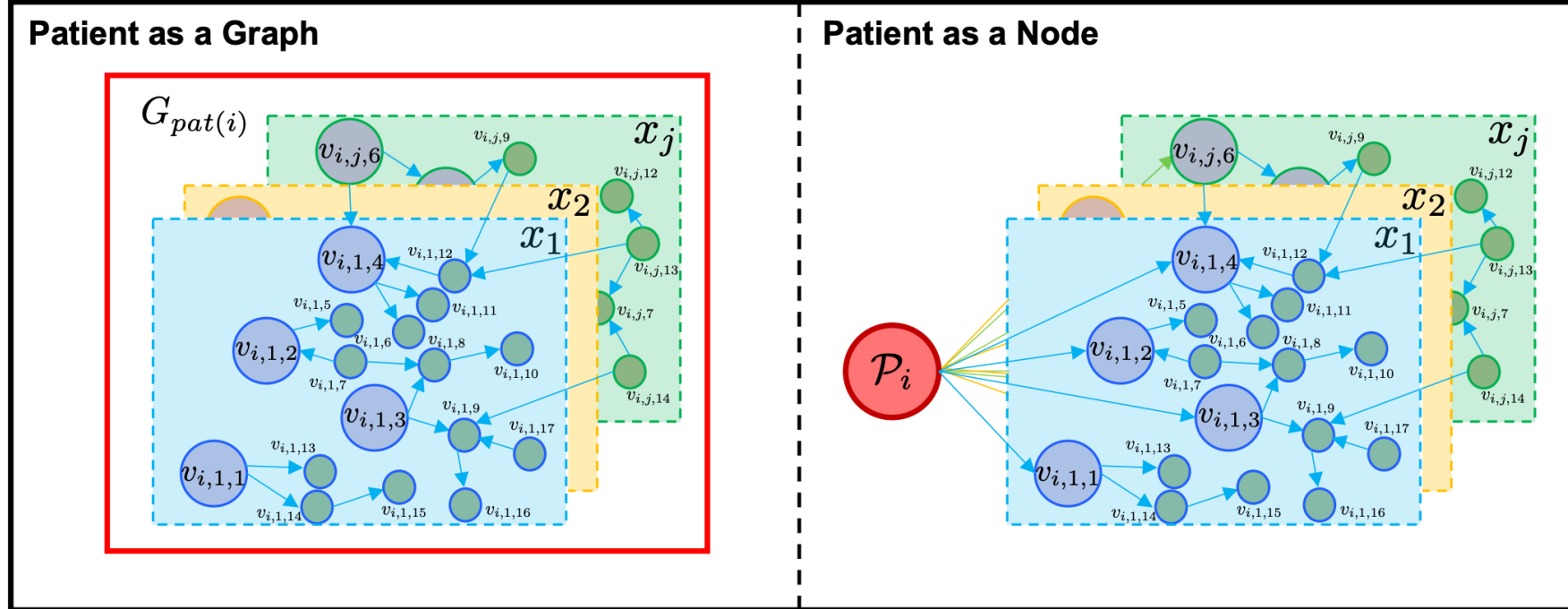
Task	Conditions	Procedures	Drugs
MT.	<i>terminal condition,</i> <i>critical diagnosis,</i> <i>end-stage,</i> <i>life-threatening</i>	<i>critical interventions,</i> <i>life-saving measures,</i> <i>resuscitation,</i> <i>emergency procedure</i>	<i>palliative medication,</i> <i>end-of-life drugs,</i> <i>life support drugs,</i> <i>emergency meds</i>
RA.	<i>chronic ailment,</i> <i>postoperative complication,</i> <i>recurrent,</i> <i>readmission-prone</i>	<i>follow-up procedure,</i> <i>secondary intervention,</i> <i>post-treatment,</i> <i>treatment review</i>	<i>maintenance medication,</i> <i>postoperative drugs,</i> <i>treatment continuation,</i> <i>follow-up meds</i>
LOS	<i>acute condition,</i> <i>severe diagnosis,</i> <i>long-term ailment,</i> <i>extended-care diagnosis</i>	<i>major surgery,</i> <i>intensive procedure,</i> <i>long recovery intervention,</i> <i>extended hospitalization</i>	-
Drug.	<i>chronic disease,</i> <i>acute diagnosis,</i> <i>symptomatic,</i> <i>treatable condition</i>	<i>diagnostic procedure,</i> <i>treatment procedure,</i> <i>medical intervention,</i> <i>drug-indicative procedure</i>	-

Method

GraphCare – Bi-attention Augmented GNN

Step 3: Employ Bi-attention Augmented (BAT) Graph Neural Network (GNN) to predict based on the personalized KGs.

Patient Representation



$$\mathbf{h}_i^{G_{pat}} = \text{MEAN}\left(\sum_{j=1}^J \sum_{k=1}^{K_j} \mathbf{h}_{i,j,k}^{(L)}\right), \quad \mathbf{h}_i^{\mathcal{P}} = \text{MEAN}\left(\sum_{j=1}^J \sum_{k=1}^{K_j} \mathbb{1}_{i,j,k}^{\Delta} \mathbf{h}_{i,j,k}^{(L)}\right),$$

$$\mathbf{z}_i^{\text{graph}} = \text{MLP}(\mathbf{h}_i^{G_{pat}}), \quad \mathbf{z}_i^{\text{node}} = \text{MLP}(\mathbf{h}_i^{\mathcal{P}}), \quad \mathbf{z}_i^{\text{joint}} = \text{MLP}(\mathbf{h}_i^{G_{pat}} \oplus \mathbf{h}_i^{\mathcal{P}})$$

Experiments

Performance comparison of 4 prediction tasks on MIMIC-III and MIMIC-IV

Findings:

1. GraphCare consistently outperforms other methods on all tasks and datasets.
2. BAT outperforms other GNNs and graph transformers.
3. Performance gain on MIMIC-III is more obvious.

Model	Task 1: Mortality Prediction				Task 2: Readmission Prediction				
	MIMIC-III		MIMIC-IV		MIMIC-III		MIMIC-IV		
	AUPRC	AUROC	AUPRC	AUROC	AUPRC	AUROC	AUPRC	AUROC	
GRU	11.8 _(0.5)	61.3 _(0.9)	4.2 _(0.1)	69.0 _(0.8)	68.2 _(0.4)	65.4 _(0.8)	66.1 _(0.1)	66.2 _(0.1)	
Transformer	10.1 _(0.9)	57.2 _(1.3)	3.4 _(0.4)	65.1 _(1.2)	67.3 _(0.7)	63.9 _(1.1)	65.7 _(0.3)	65.3 _(0.4)	
RETAIN	9.6 _(0.6)	59.4 _(1.5)	3.8 _(0.4)	64.8 _(1.6)	65.1 _(1.0)	64.1 _(0.7)	66.2 _(0.3)	66.3 _(0.2)	
GRAM	11.4 _(0.7)	60.4 _(0.9)	4.4 _(0.3)	66.7 _(0.7)	67.2 _(0.8)	64.3 _(0.4)	66.1 _(0.2)	66.3 _(0.3)	
DeepPr	13.2 _(1.1)	60.8 _(0.4)	4.2 _(0.2)	68.9 _(0.9)	68.8 _(0.9)	66.5 _(0.4)	65.6 _(0.1)	65.4 _(0.2)	
AdaCare	11.1 _(0.4)	58.4 _(1.4)	4.6 _(0.3)	69.3 _(0.7)	68.6 _(0.6)	65.7 _(0.3)	65.9 _(0.0)	66.1 _(0.0)	
GRASP	9.9 _(1.1)	59.2 _(1.4)	4.7 _(0.1)	68.4 _(1.0)	69.2 _(0.4)	66.3 _(0.6)	66.3 _(0.3)	66.1 _(0.2)	
StageNet	12.4 _(0.3)	61.5 _(0.7)	4.2 _(0.3)	69.6 _(0.8)	69.3 _(0.6)	66.7 _(0.4)	66.1 _(0.1)	66.2 _(0.1)	
GRAPHCARE	w/ GAT	14.3 _(0.8)	67.8 _(1.1)	5.1 _(0.1)	71.8 _(1.0)	71.5 _(0.7)	68.1 _(0.6)	67.4 _(0.4)	67.3 _(0.4)
	w/ GINE	14.4 _(0.4)	67.3 _(1.3)	5.7 _(0.1)	72.0 _(1.1)	71.3 _(0.8)	68.0 _(0.4)	68.3 _(0.3)	67.5 _(0.4)
	w/ EGT	15.5 _(0.5)	69.1 _(1.0)	6.2 _(0.2)	71.3 _(0.7)	72.2 _(0.5)	68.8 _(0.4)	68.9 _(0.2)	67.6 _(0.3)
	w/ GPS	15.3 _(0.9)	68.8 _(0.8)	6.7 _(0.2)	72.7 _(0.9)	71.9 _(0.6)	68.5 _(0.6)	69.1 _(0.4)	67.9 _(0.4)
	w/ BAT	16.7 _(0.5)	70.3 _(0.5)	6.7 _(0.3)	73.1 _(0.5)	73.4 _(0.4)	69.7 _(0.5)	69.6 _(0.3)	68.5 _(0.4)

Model	Task 3: Length of Stay Prediction								
	MIMIC-III				MIMIC-IV				
	AUROC	Kappa	Accuracy	F1-score	AUROC	Kappa	Accuracy	F1-score	
GRU	78.3 _(0.1)	26.2 _(0.2)	40.3 _(0.3)	34.9 _(0.5)	78.7 _(0.1)	26.0 _(0.1)	35.2 _(0.1)	31.6 _(0.2)	
Transformer	78.3 _(0.2)	25.4 _(0.4)	40.1 _(0.3)	34.8 _(0.2)	78.3 _(0.3)	25.3 _(0.4)	34.4 _(0.2)	31.4 _(0.3)	
RETAIN	78.2 _(0.1)	26.1 _(0.4)	40.6 _(0.3)	34.9 _(0.4)	78.9 _(0.3)	26.3 _(0.2)	35.7 _(0.2)	32.0 _(0.2)	
GRAM	78.2 _(0.1)	26.3 _(0.3)	40.4 _(0.4)	34.5 _(0.2)	78.8 _(0.2)	26.1 _(0.4)	35.4 _(0.2)	31.9 _(0.3)	
DeepPr	77.9 _(0.1)	25.3 _(0.4)	40.1 _(0.6)	35.0 _(0.4)	79.5 _(0.3)	26.4 _(0.2)	35.8 _(0.3)	32.3 _(0.1)	
StageNet	78.3 _(0.2)	24.8 _(0.2)	39.9 _(0.2)	34.4 _(0.4)	79.2 _(0.3)	26.0 _(0.2)	35.0 _(0.2)	31.3 _(0.3)	
GRAPHCARE	w/ GAT	79.4 _(0.3)	28.2 _(0.2)	41.9 _(0.2)	36.1 _(0.4)	80.3 _(0.3)	28.4 _(0.4)	36.2 _(0.1)	33.3 _(0.3)
	w/ GINE	79.2 _(0.2)	28.3 _(0.3)	41.5 _(0.3)	36.0 _(0.4)	79.9 _(0.2)	27.5 _(0.3)	36.3 _(0.3)	32.8 _(0.2)
	w/ EGT	80.3 _(0.3)	28.8 _(0.2)	42.8 _(0.4)	36.3 _(0.5)	80.5 _(0.2)	28.7 _(0.3)	36.7 _(0.2)	33.5 _(0.1)
	w/ GPS	80.9 _(0.3)	28.8 _(0.4)	43.0 _(0.3)	36.8 _(0.4)	80.7 _(0.3)	28.8 _(0.4)	36.7 _(0.3)	33.9 _(0.3)
	w/ BAT	81.4 _(0.3)	29.5 _(0.4)	43.2 _(0.4)	37.5 _(0.2)	81.7 _(0.2)	29.8 _(0.3)	37.3 _(0.3)	34.2 _(0.3)

Model	Task 4: Drug Recommendation								
	MIMIC-III				MIMIC-IV				
	AUPRC	AUROC	F1-score	Jaccard	AUPRC	AUROC	F1-score	Jaccard	
GRU	77.0 _(0.1)	94.4 _(0.0)	62.3 _(0.3)	47.8 _(0.3)	74.1 _(0.1)	94.2 _(0.1)	60.2 _(0.2)	44.0 _(0.4)	
Transformer	76.1 _(0.1)	94.2 _(0.0)	62.1 _(0.4)	47.1 _(0.4)	71.3 _(0.1)	93.4 _(0.1)	55.9 _(0.2)	40.4 _(0.1)	
RETAIN	77.1 _(0.1)	94.4 _(0.0)	63.7 _(0.2)	48.8 _(0.2)	74.2 _(0.1)	94.3 _(0.0)	60.3 _(0.1)	45.0 _(0.1)	
GRAM	76.7 _(0.1)	94.2 _(0.1)	62.9 _(0.3)	47.9 _(0.3)	74.3 _(0.2)	94.2 _(0.1)	60.1 _(0.2)	45.3 _(0.3)	
DeepPr	74.3 _(0.1)	93.7 _(0.0)	60.3 _(0.4)	44.7 _(0.3)	73.7 _(0.1)	94.2 _(0.1)	59.1 _(0.4)	43.8 _(0.4)	
StageNet	74.4 _(0.1)	93.0 _(0.1)	61.4 _(0.3)	45.8 _(0.4)	74.4 _(0.1)	94.2 _(0.0)	60.2 _(0.3)	45.4 _(0.4)	
SafeDrug	68.1 _(0.3)	91.0 _(0.1)	46.7 _(0.4)	31.7 _(0.3)	66.4 _(0.5)	91.8 _(0.2)	56.2 _(0.4)	44.3 _(0.3)	
MICRON	77.4 _(0.0)	94.6 _(0.1)	63.2 _(0.4)	48.3 _(0.4)	74.4 _(0.1)	94.3 _(0.1)	59.3 _(0.3)	44.1 _(0.3)	
GAMENet	76.4 _(0.0)	94.2 _(0.1)	62.1 _(0.4)	47.2 _(0.4)	74.2 _(0.1)	94.2 _(0.1)	60.4 _(0.4)	45.3 _(0.3)	
MoleRec	69.8 _(0.1)	92.0 _(0.1)	58.1 _(0.1)	43.1 _(0.3)	68.6 _(0.1)	92.1 _(0.1)	56.3 _(0.4)	40.6 _(0.3)	
GRAPHCARE	w/ GAT	78.5 _(0.2)	94.8 _(0.1)	64.4 _(0.3)	49.2 _(0.4)	74.7 _(0.5)	94.4 _(0.3)	60.4 _(0.3)	45.7 _(0.4)
	w/ GINE	78.2 _(0.1)	94.7 _(0.1)	63.6 _(0.4)	47.9 _(0.3)	74.8 _(0.3)	94.6 _(0.1)	60.6 _(0.4)	46.1 _(0.4)
	w/ EGT	79.6 _(0.2)	95.3 _(0.0)	66.4 _(0.2)	49.6 _(0.4)	75.4 _(0.4)	95.0 _(0.1)	61.6 _(0.3)	47.3 _(0.3)
	w/ GPS	79.9 _(0.3)	95.5 _(0.1)	66.2 _(0.3)	49.8 _(0.4)	75.9 _(0.5)	94.9 _(0.1)	62.1 _(0.3)	46.8 _(0.4)
	w/ BAT	80.2 _(0.2)	95.5 _(0.1)	66.8 _(0.2)	49.7 _(0.3)	77.1 _(0.1)	95.4 _(0.2)	63.9 _(0.3)	48.1 _(0.3)

Experiments

Effect of Knowledge Graph Size

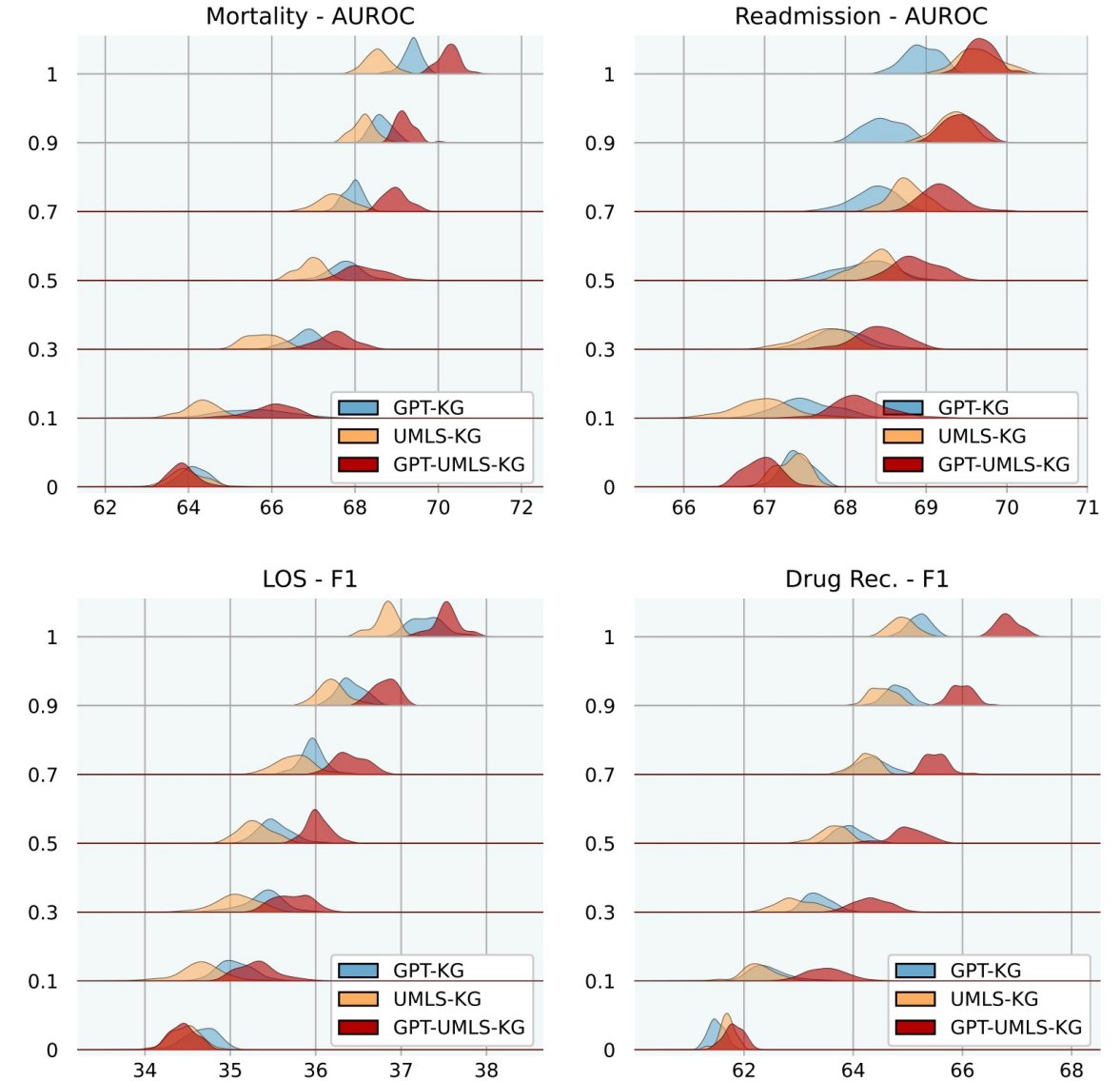
Statistics of Extracted KGs

KG	Hyperparameter	# Nodes	# Edges	# Triples
<u>GPT-KG</u>	$\chi=3$	4599	752	31325
<u>UMLS-KG</u>	$\kappa=1$	3053	40	12421
UMLS-KG	$\kappa=2, \epsilon=5$	10805	54	81073
<u>GPT-UMLS-KG</u>	$\chi=3, \kappa=1$	6355	774	40496
GPT-UMLS-KG	$\chi=3, \kappa=2, \epsilon=5$	12284	785	104460

Findings:

1. Better performance with larger KGs.
2. Combined GPT-UMLS-KG outperforms both GPT-KG and UMLS-KG consistently.
3. GPT-KG contributes more to mortality and LOS predictions, while UMLS-KG edges out in readmission prediction.
4. Lower KG ratios are associated with larger standard deviations, due to the reduced likelihood of vital knowledge being contained.

Performance



(Based on 30 runs; Y-axis: ratio of KG used)

Experiments

Effect of BAT GNN

Statistics of MIMIC-III/IV

	#patients	#visits	#visits/patient	#conditions/patient	#procedures/patient	#drugs/patient
MIMIC-III	35,707	44,399	1.24	12.89	4.54	33.71
MIMIC-IV	123,488	232,263	1.88	21.74	4.70	43.89

Ablation Study of BAT GNN's Variants

Case	Variants	MIMIC-III				MIMIC-IV			
		MT.	RA.	LOS	Drug.	MT.	RA.	LOS	Drug.
#0	w/ <i>all</i>	70.3	69.7	37.5	66.8	73.1	68.5	34.2	63.9
#1	w/o α	68.7 \downarrow _{0.6}	68.5 \downarrow _{1.2}	36.7 \downarrow _{0.8}	64.6 \downarrow _{2.2}	72.2 \downarrow _{0.9}	67.8 \downarrow _{0.7}	33.1 \downarrow _{1.1}	61.6 \downarrow _{2.3}
#2	w/o β	69.9 \downarrow _{0.4}	68.7 \downarrow _{1.0}	37.2 \downarrow _{0.3}	66.5 \downarrow _{0.3}	72.1 \downarrow _{1.0}	67.0 \downarrow _{1.5}	33.5 \downarrow _{0.7}	63.2 \downarrow _{0.7}
#3	w/o $w_{\mathcal{R}}$	69.8 \downarrow _{0.5}	68.4 \downarrow _{1.3}	36.8 \downarrow _{0.7}	66.3 \downarrow _{0.5}	72.9 \downarrow _{0.2}	67.9 \downarrow _{0.6}	33.7 \downarrow _{0.5}	63.1 \downarrow _{0.8}
#4	w/o <i>AttnInit</i>	69.5 \downarrow _{0.8}	69.2 \downarrow _{0.5}	37.2 \downarrow _{0.3}	65.5 \downarrow _{1.3}	72.5 \downarrow _{0.6}	68.1 \downarrow _{0.4}	34.1 \downarrow _{0.1}	62.4 \downarrow _{1.5}
#5	w/o #(1,2,3,4)	67.4 \downarrow _{2.9}	68.1 \downarrow _{1.6}	36.0 \downarrow _{1.5}	64.0 \downarrow _{2.8}	71.7 \downarrow _{1.4}	67.5 \downarrow _{1.0}	32.9 \downarrow _{1.3}	60.5 \downarrow _{3.4}

Observations:

1. Excluding node-level attention (α) results in a general drop across all tasks/datasets.
2. Excluding visit-level attention (β) affects more on MIMIC-IV, as it has a higher #visits/patient.
3. Readmission prediction is more sensitive to the visit-level attention.
4. Drug recommendation is more sensitive to the attention initialization.

Experiments

Appendix: Patient Representation Learning

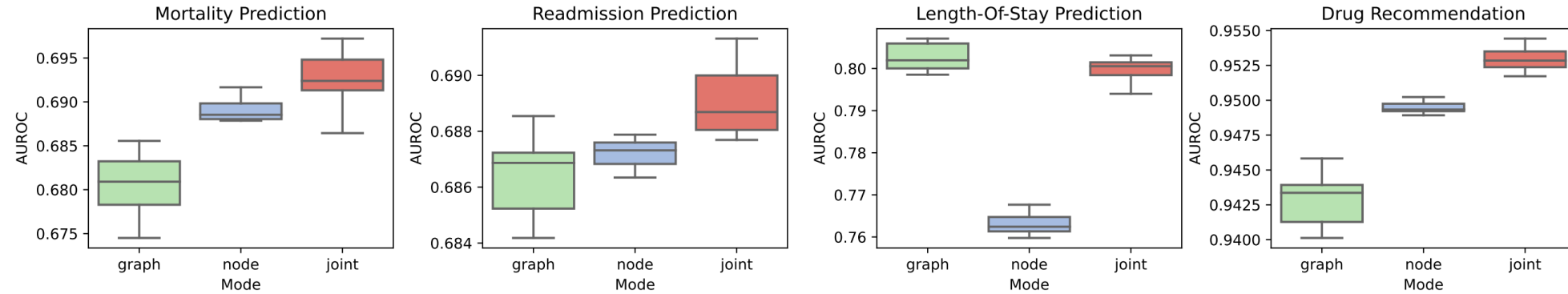


Figure 8: **Performance of healthcare predictions with three types of patient representations (§3.3):** (1) **graph** - patient graph embedding obtained through mean pooling of node embedding; (2) **node** - patient node embedding connected to the direct EHR node; (3) **joint** - embedding concatenated by (1) and (2). We use GPT-KG to perform this analysis.

Observations:

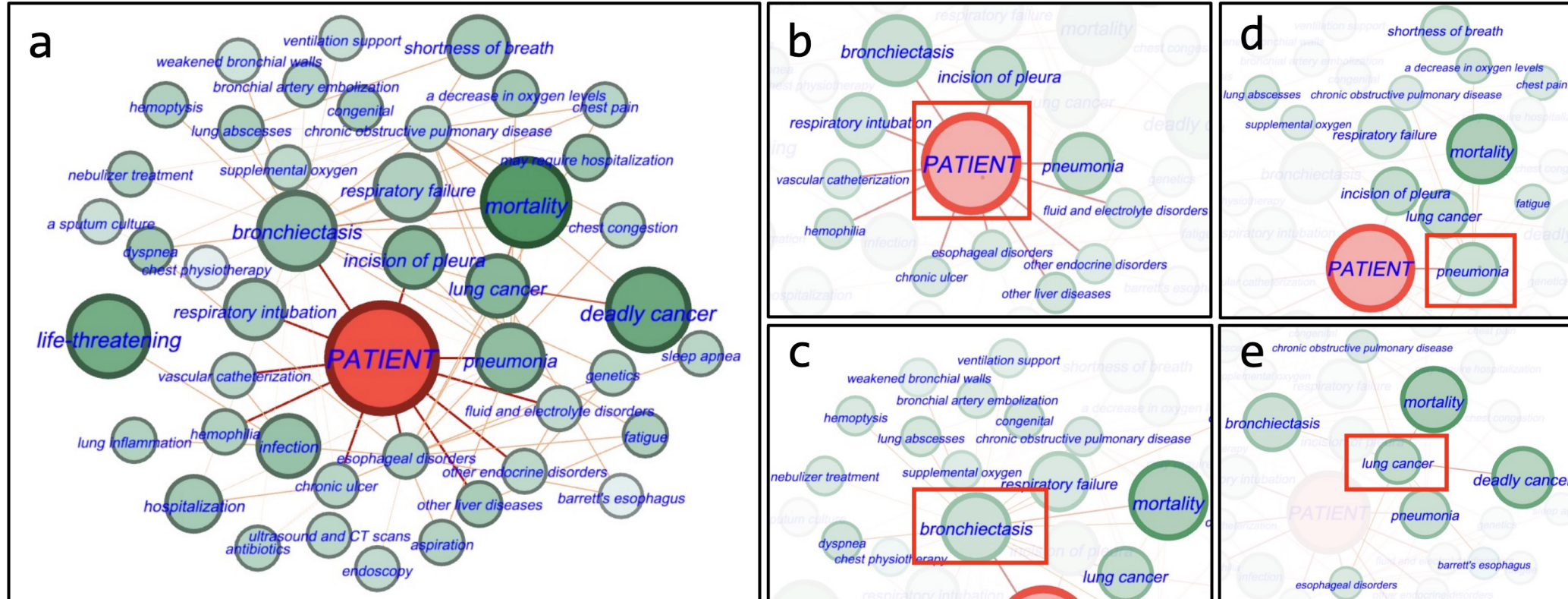
The patient graph contains more information as well as more noise.

The patient node contains more accurate information, as it directly links the EHR nodes.

A joint representation of them is a balance.

Experiments

Interpretability of GraphCare



Observations:

Removing the indirect node “lung cancer” connecting crucial nodes “mortality” and “deadly cancer” would result in a failure of mortality prediction.

➔ Highlight the importance of the personalized KG.

Thank you!

Preprint: <https://arxiv.org/pdf/2305.12788.pdf>

Code: <https://github.com/pat-jj/GraphCare>