

Contrastive Learning of Temporal Distinctiveness for Survival Analysis in Electronic Health Records

Mohsen Nayebi Kerdabai, Arya Hadizadeh Moghaddam, Bin Liu, Mei Liu, Zijun Yao

CIKM 2023



Outline

- **Background and Motivation**
- **Proposed Method**
- **Evaluation**
- **Conclusion**

Background and Motivation

- **What is Survival Analysis (SA)?**

- Methods used to analyze **time-to-event data** => **Time Modeling**
- Applied in various fields such as **healthcare, economics, engineering**, etc.

- **Goals**

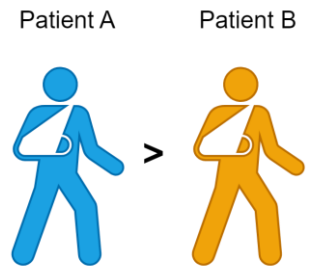
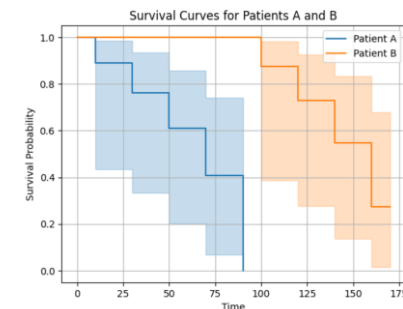
- **Rank** patients regarding their predicted survival time/chance.
- **Time prediction** ability

- **Semi-supervised Learning**

- Labeled Data : **Observed**
- Unlabeled Data : **Censored** (a key challenge)

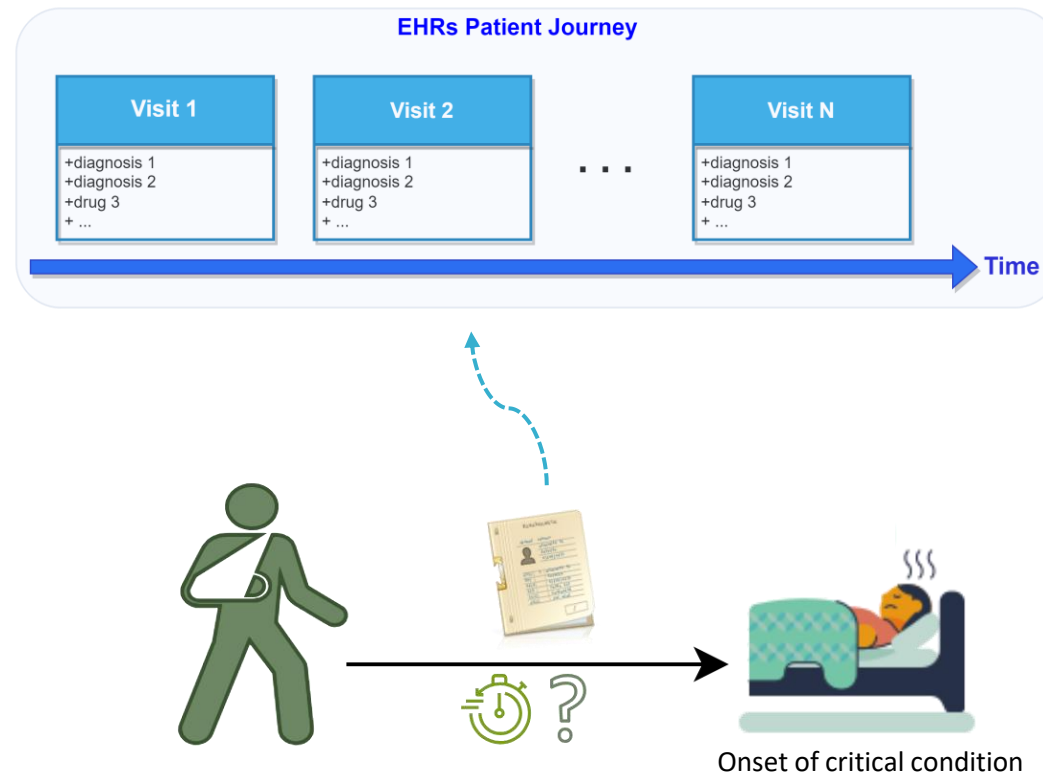
- **Challenges and Opportunities:**

- Traditionally, ranking and survival likelihood objectives are used.
- Learning **discriminative patient embedding based on survival time difference** was not explored in the literature



Background and Motivation

- Learning the longitudinal and rich Electronic Health Records (EHRs) for comprehensive patient profiling

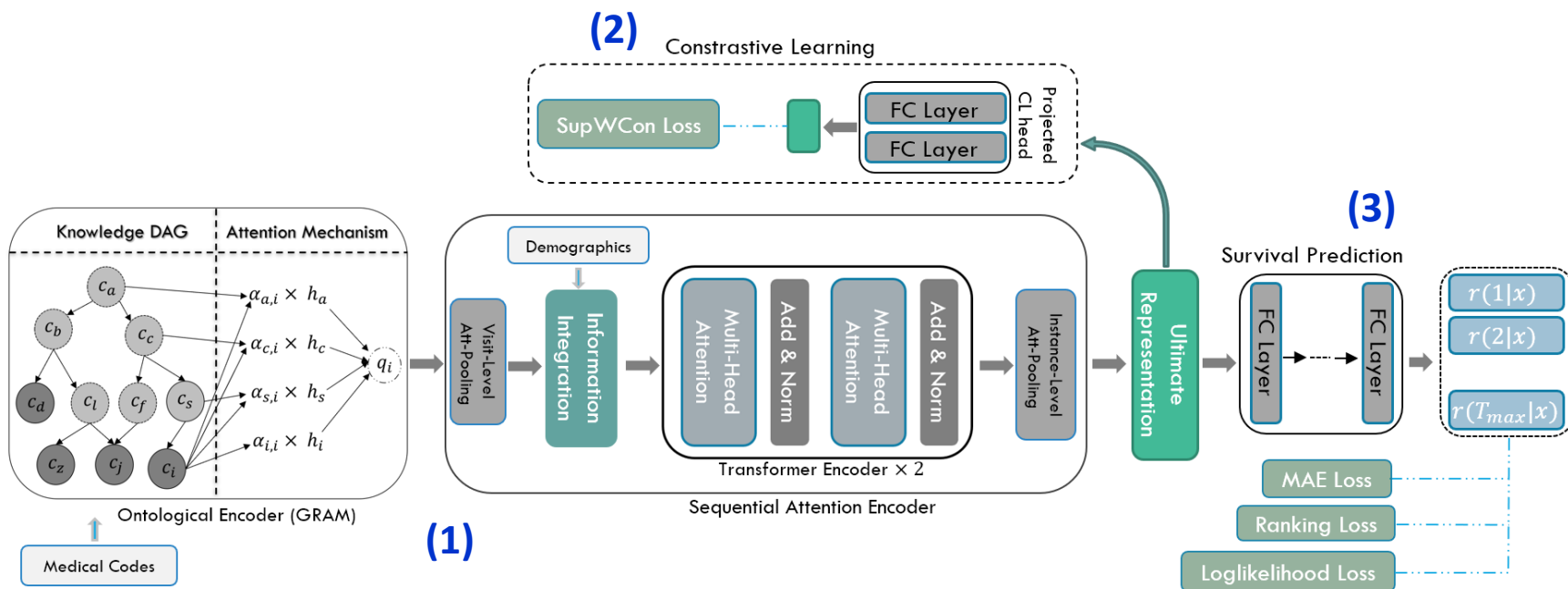


Outline

- Background and Motivation
- **Proposed Method**
- Evaluation
- Conclusion

Proposed Method

- We present **Ontology-aware Temporality-based Contrastive Survival (OTCSurv)** Analysis to predict time-to-event using patient history in EHRs.
- Three main components:
 - Sequential Attention-based Ontological Encoder**— encoding patient history with multiple attention-based components as well as domain knowledge from ontology
 - Contrastive Component** — utilizes survival durations to derive hardness-aware temporal distinctiveness
 - Survival Prediction** — outcomes necessary survival probabilities for each discrete time interval.



Proposed Method - Ontological Encoder

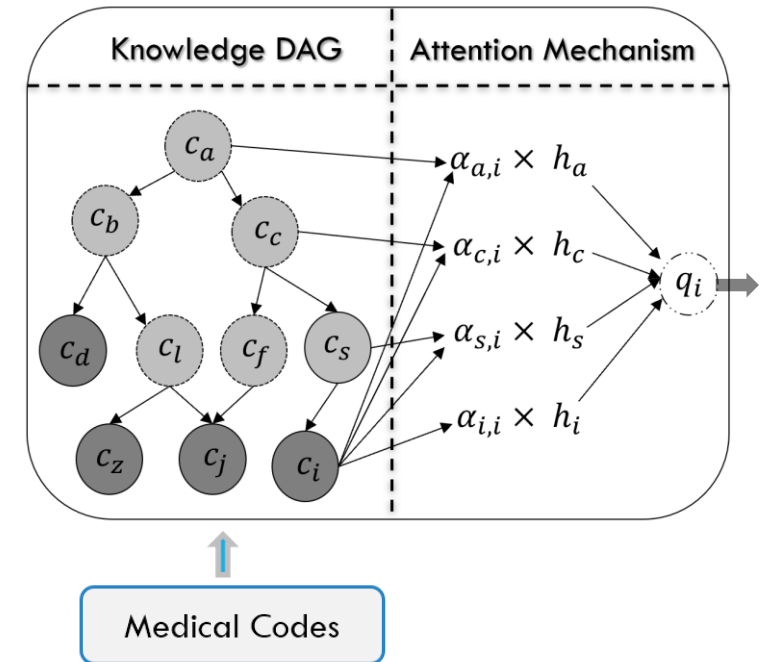
Ontology based EHR code encoding using GRAM*

- For each code (EHRs feature) find its ancestors in the ontology
- Assign each of them a vector $h_j \in \mathbb{R}^{d_c}$
- Use **attention mechanism** to generate attention weights for each of them $\alpha_{ij} \in \mathbb{R}^+$
- Use the attention weights to produce a convex combination of the initial embeddings as a final vector $q_i \in \mathbb{R}^{d_c}$

attention mechanism

$$f(h_i, h_k) = \omega_\alpha^T \tanh(W_\alpha \text{Concat}(h_i; h_k) + b_\alpha) \quad f(.) \Rightarrow \text{MLP operator}$$
$$\alpha_{ij} = \frac{\exp(f(h_i, h_j))}{\sum_{k \in A(i)} \exp(f(h_i, h_k))} \quad \text{Softmax function}$$

$$q_i = \sum_{j \in A(i)} \alpha_{ij} h_j, \quad \sum_{j \in A(i)} \alpha_{ij} = 1, \quad \alpha_{ij} \geq 0 \text{ for } j \in A(i) \quad \text{Convex combination}$$



Proposed Method - Sequential Attention Encoder

- **Attention-pooling**

- **Visit-level** : compress and combine the information of different codes of a visit using the attention mechanism and produce the visit representation
- **Instance-level** - compress and combine the information of different visits of an instance using the attention mechanism and produce the instance representation

Information Integration: simple concatenation

- **Transformer encoder** – Using self-attention mechanism extract the interactions of medical visits to produce a rich representation of a patient

$$e^n = l(v_n) = W_2 \sigma(v_n W_1 + \beta_1), \quad 1 \leq n \leq N$$

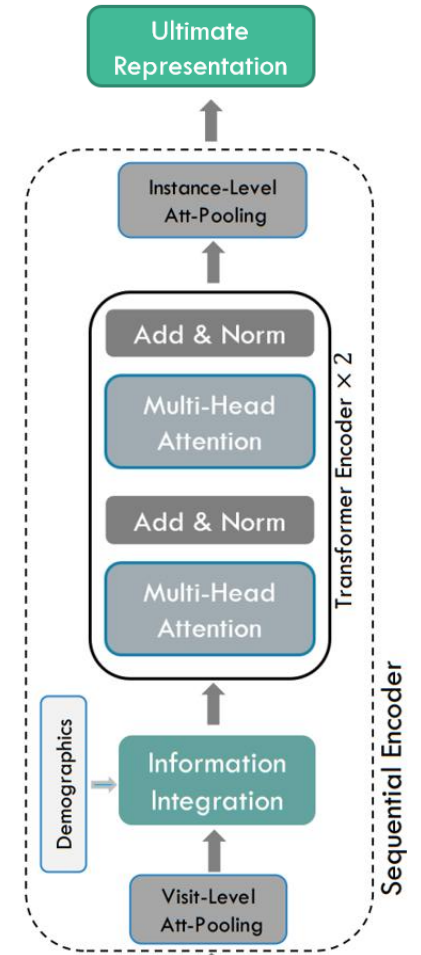
$$\alpha^n = \text{softmax}(e^n)$$

$$p^n = \sum_{i=1}^M \alpha_m^n q_m^n$$

$$\text{head}_i = \text{Attention}(QW_i^Q, KW_i^K, VW_i^V)$$

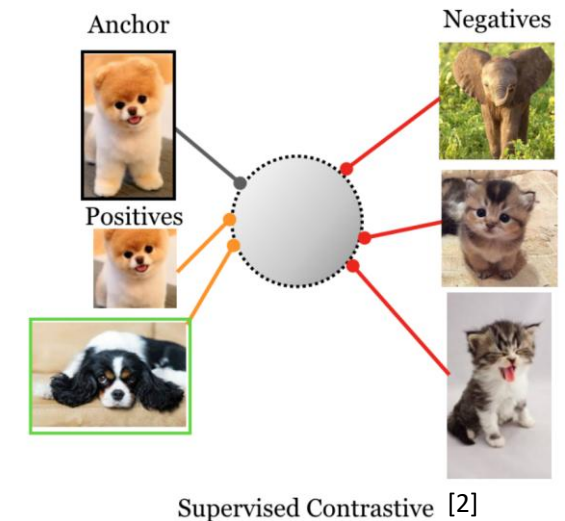
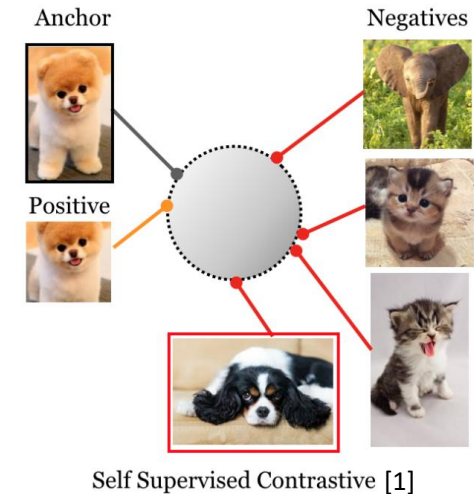
$$\text{SelfAttention}(Q, K, V) = \text{softmax}\left(QK^T / \sqrt{d_k}\right) V$$

$$U = \text{Concat}(\text{head}_1, \text{head}_2, \dots, \text{head}_h) W^O$$



Proposed Method – Contrastive Component

- What is contrastive learning?
 - Representation Learning
 - Contrasting positive samples with negative samples.
- Originally, self-supervised, but later, supervised and semi-supervised
- Objective: bring positive samples closer together in the embedding space while pushing negative samples farther apart.



[1] A Simple Framework for Contrastive Learning of Visual Representations, Chen *et al.*, PMLR 2020

[2] Supervised Contrastive, Khosla *et al.*, Learning, NeurIPS 2020

Proposed Method – Contrastive Component

- **Nonlinear Transformation**

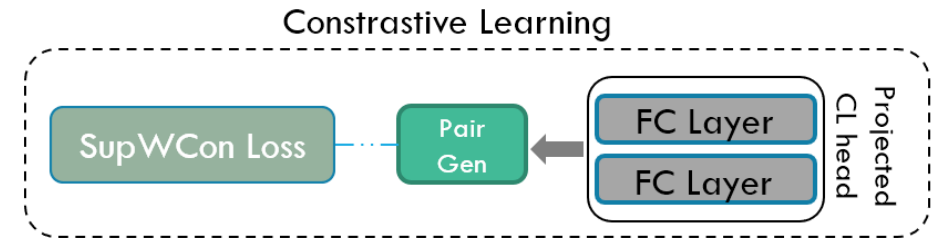
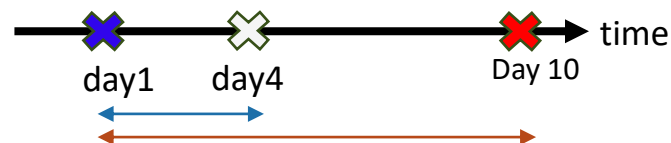
FNN with a nonlinear AF--> **transfers** the representation to a different latent space

- **Pair Generator**

Negative and positive pair generation criteria

- **Supervised Weighted Contrastive (SupWCon) loss**

- Constant temperature parameter for + pairs $\tau \in \mathbb{R}^+$
- **Adaptive** temperature parameter for - pairs $\tau_{ia} \in \mathbb{R}^+$
 - The inverse of their difference in survival duration
 - Encourage our model to regulate the amount of dissimilarity.



$$L^{\text{SupWCon}} = \sum_{i \in I} \frac{-1}{|P(i)|} \sum_{p \in P(i)} \log \frac{\exp(z_i \cdot z_p / \tau)}{\sum_{a \in A(i)} \exp(z_i \cdot z_a / \tau_{ia})}$$

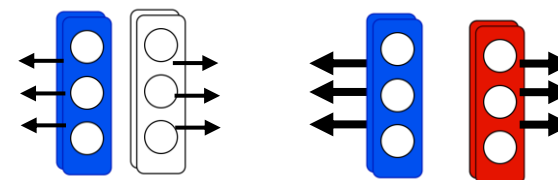
+ pair similarity (points to $\exp(z_i \cdot z_p / \tau)$)

pair similarity (points to $\exp(z_i \cdot z_a / \tau_{ia})$)

the set of instances that make a positive pair with anchor (points to $A(i)$)

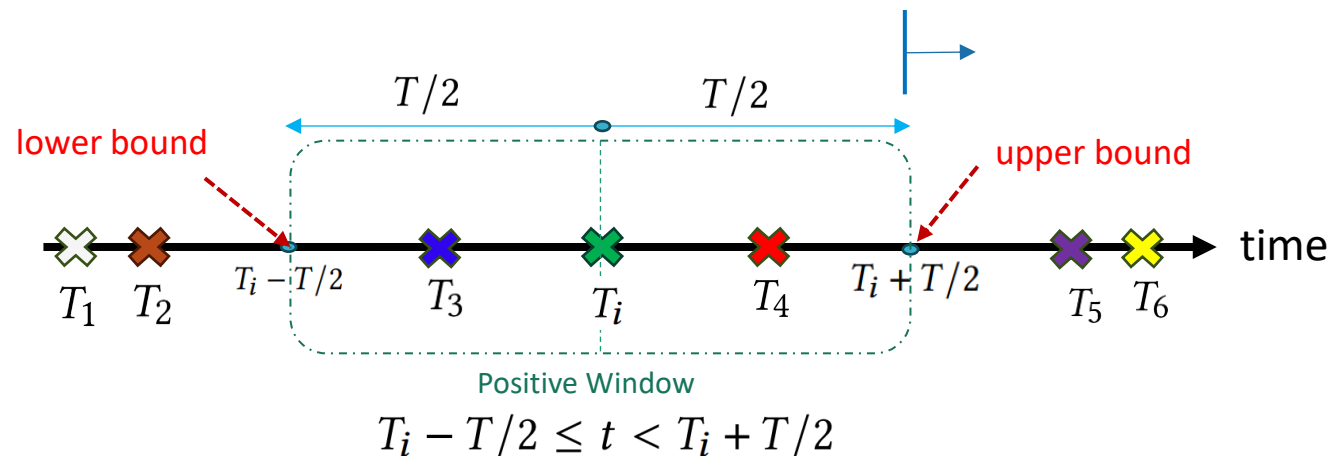
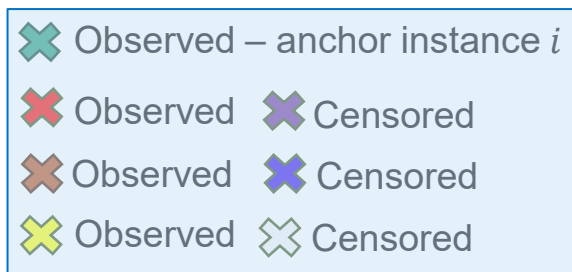
$$\tau_{i,a} = |T_i - T_a|^{-1}$$

hardness – aware (points to $\tau_{i,a}$)



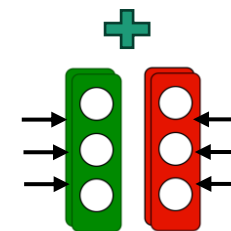
Proposed Method – Contrastive Component

Pair generation for contrastive learning



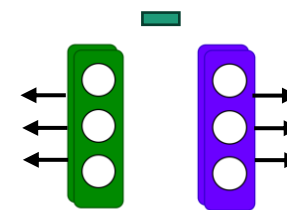
Positive pairs criteria:

- Any observed subject with survival time inside the positive window.
 - Positive pairs example : (T_i, T_4)



Negative pairs criteria :

- Any observed subject with survival time outside the positive window:
 - Pair example: (T_i, T_2) , (T_i, T_6)
- Any censored subject with survival time larger than the *upper bound* of the positive window:
 - Pair example: (T_i, T_5)



Proposed Method – Other Loss Components

1. Loglikelihood Loss

Maximize the summation the survival probabilities before T

Minimize the summation the survival probabilities after T

$$\begin{aligned} L_{\text{ob}}^{\text{Loglikelihood}} &= - \sum_{t=1}^{T-1} \log \hat{S}(t|x) - \sum_{t=T}^{T_{\text{max}}} \log(1 - \hat{S}(t|x)) \\ L_{\text{cen}}^{\text{Loglikelihood}} &= - \sum_{t=1}^T \hat{S}(t|x) \\ L^{\text{Loglikelihood}} &= L_{\text{cen}}^{\text{Loglikelihood}} + L_{\text{ob}}^{\text{Loglikelihood}} \end{aligned}$$

2. Pairwise Ranking Loss

- Based on concordance
- penalize the discordant pairs

$$L^{\text{Ranking}} = \max(0, (T_j - T_i) - (\hat{T}_j - \hat{T}_i))$$

penalize the discordant pairs

3. Mean Squared Error (MSE) Loss

penalizes wrong predicted survival duration times for only observed data points

$$\text{MSE} = \frac{1}{N_{\text{ob}}} \sum_{i=1}^{N_{\text{ob}}} (T_i - \hat{T}_i)^2$$

Outline

- Background and Motivation
- Proposed Method
- **Evaluation**
- Conclusion

Evaluation

- **EHRs Dataset**

- A real-world **EHR dataset** from the University of Kansas Medical Center (KUMC) gathered for studying **Acute Kidney Injury (AKI)**

- **Medical Ontologies:**

- Drug codes: Anatomical Therapeutic Chemical (**ATC**) classification
Diagnosis codes : The International Classification of Diseases (**ICD-9**)

- **Evaluation Metrics**

- The time-dependent discrimination index C^{td}
 - Extension of Harrell's **concordance index (c-index)**
 - **Evaluating ranking ability**
- The Mean Absolute Error (MAE)
 - **Evaluating time prediction ability**

Table 1: Dataset Statistics

Number of patinets	56779
Number of censored patients	47773 (84.1%)
Number of observed patients	9006 (15.8%)
The average age of patients	59.47
Sex of patients distribution	(52% M, 48% F)
Average number of medical codes in a visit	13.29
Average number of medical codes in a patient	66.46

Evaluation - Results

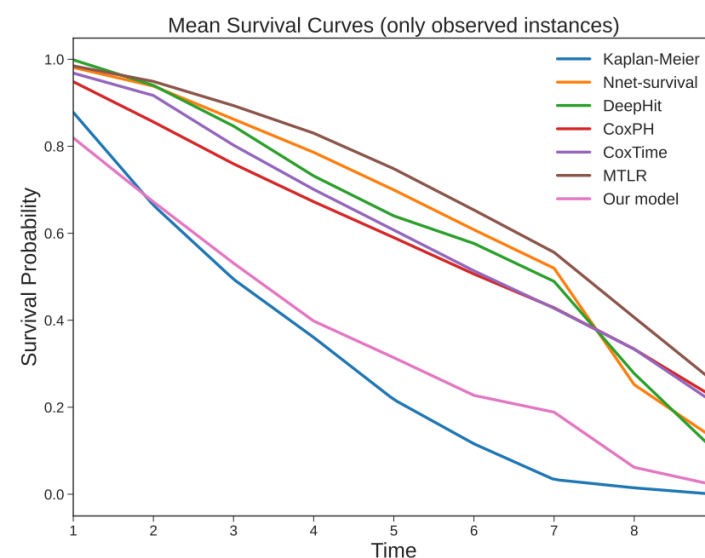
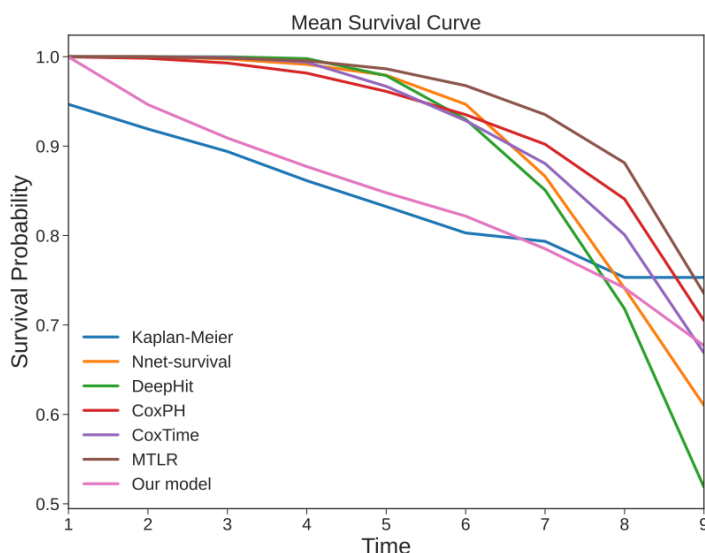
- **Baseline Comparison**

- OTCSurv **outperforms** other baselines regarding **both** evaluation metrics

Model Name	C^{td}	MAE
Nnet-survival	0.6332	3.165
MTLR (N-MTLR)	0.6712	4.081
DeepHit	0.6929	3.012
CoxTime	0.6912	2.980
DeepSurv (CoxPH)	0.6898	3.007
Our model	0.6990	1.890

- **Survival Curves Comparison**

- OTCSurv's mean survival curve is the closest to the Kaplan-Meier curve => accurately captures the survival behavior and provides survival predictions that are **more consistent with the actual outcomes**.



Results - Ablation

- We experiment with different combinations of loss components.
- **SupWCon** is effective on improving **both** evaluation metrics.
- With all four loss components, we achieve the best **trade-off** in the performance.
- We utilized **a weighted summation** of these four losses as follows:

$$L^{\text{Total}} = \lambda_1 L^{\text{Loglikelihood}} + \lambda_2 L^{\text{Ranking}} + \lambda_3 L^{\text{SupWCon}} + \lambda_4 L^{\text{MSE}}$$
- Ontological Encoder and attention-poolings also play a significant role in improving the final results

Loss functions	C^{td}	MAE
$L^{\text{Loglikelihood}}$	0.6647	2.30
L^{Ranking}	0.6907	2.80
$L^{\text{Loglikelihood}}, L^{\text{SupWCon}}$	0.6808	1.90
$L^{\text{Loglikelihood}}, L^{\text{Ranking}}$	0.6951	2.43
$L^{\text{Loglikelihood}}, L^{\text{Ranking}}, L^{\text{SupWCon}}$	0.7030	1.91
$L^{\text{Loglikelihood}}, L^{\text{Ranking}}, L^{\text{SupWCon}}, L^{\text{MSE}}$	0.6990	1.89

Window size	C^{td}	MAE
w/o ontology	0.6888	2.11
w/o attention-pooling	0.6795	2.21
Full model	0.6990	1.89

Results - Interpretability

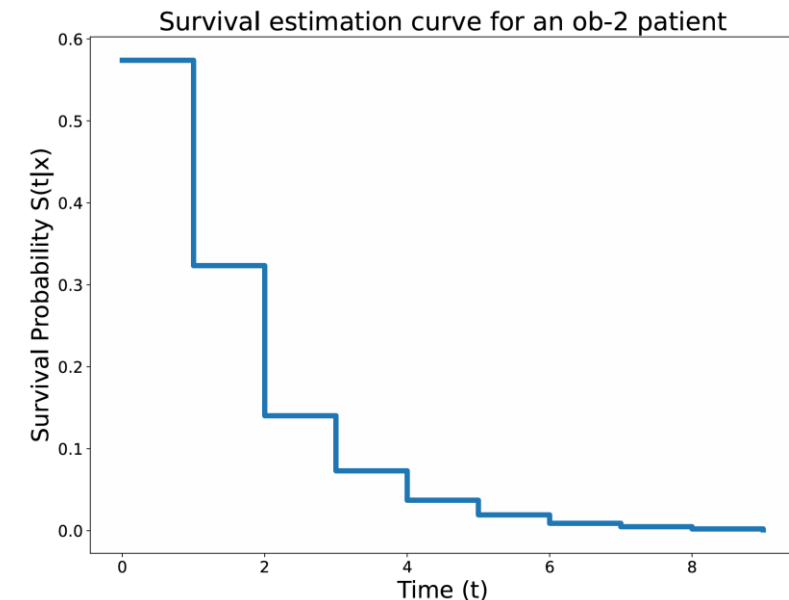
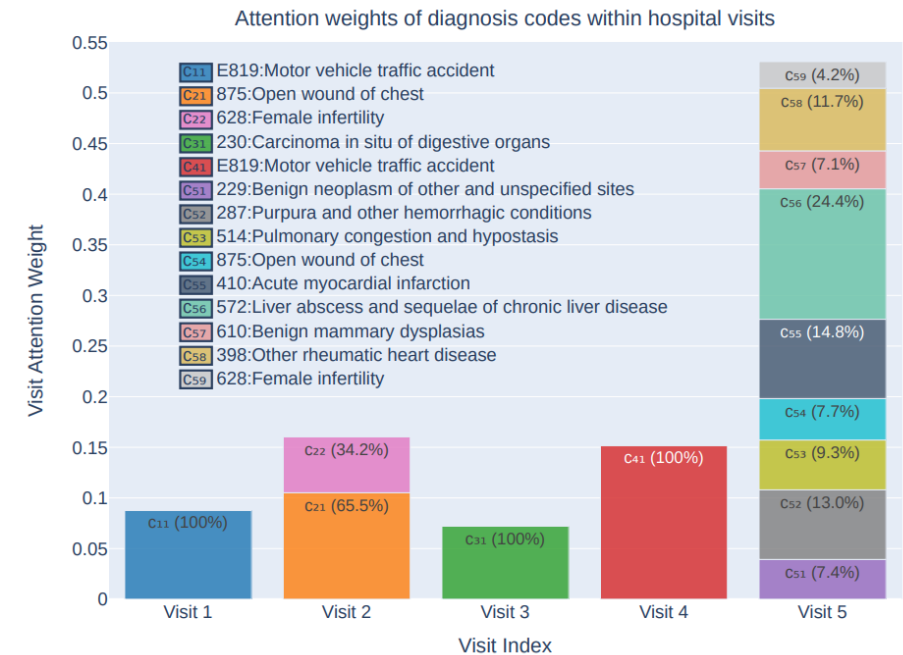
Interpretability by Attention Analysis:

Attention weights in the **visit-level** attention-pooling

- the relative significance of each feature (code)
 - The height of each stacked bar
- Example
 - c_{56} : "572" : liver abscess and sequelae of chronic liver disease
 - c_{55} : "410" : acute myocardial infarction (AMI)

Attention weights in the **instance-level** attention-pooling

- The relative importance of each of the visit
 - Each bar inside the stacked bars presents a medical code
Inside the visit and its attention weight.



Outline

- Background and Motivation
- Proposed Method
- Evaluation
- **Conclusion**

Conclusion

Ontology-aware Temporality-based Contrastive Survival (OTCSurv) analysis framework

- **We design a Supervised Weighted Contrastive (SupWCon) Learning loss function**
 - Based on survival duration as its contrastive pairing criteria
 - Handles and takes into account censored data
 - Features an adaptive hardness-aware temperature parameter
- **We used a sequential attention-based ontological encoder**
 - Incorporate knowledge domain from medical ontology
 - Learn embedding with rich context using multiple attention-based components
- **We optimize survival prediction with a configuration of SupWCon, along with three more loss functions, focusing on two key goals**
 - Precisely ranking the survival chances of patients
 - Accurately predicting survival duration time

Thank you

Questions ?