Toward a Well-Calibrated Discrimination through Survival Outcome-Aware Contrastive Learning

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Outline

- o Introduction to Survival Analysis
- o Consideration
- o Objective
- o Challenges and Motivation
- o Proposed Method
- o Experiments

What is survival analysis?

- A very common outcome in medical studies is the time until an event occurs:
 - The time until a patient dies
 - The time until a patient suffers a heart attack
 - The time until a liver transplant patient needs a new liver
 - The time until the recurrence of cancer following treatment

 Data involving such an outcome is often called "time-to-event" data or "failure-time data" or "survival" data, and the branch of statistics that deals with analyzing these data is called survival analysis

Survival data

o Survival (a.k.a. time-to-event) data

 $\mathcal{D} = \{(\mathbf{x}_i, \tau_i, \delta_i)\}_{i=1}^N$

- x: Observed features (covariates)
- τ : Time-to-event or time-to-censoring elapsed since the baseline (e.g., the entry to a clinical trial)
- δ : Label indicating whether event the event or the censoring occurred

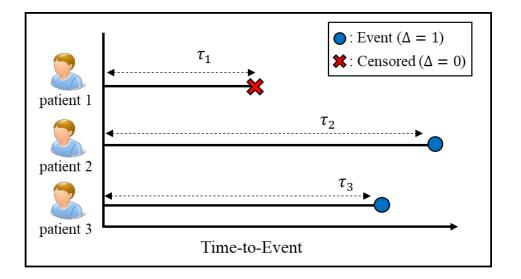


Figure. An illustration of survival data

Survival data

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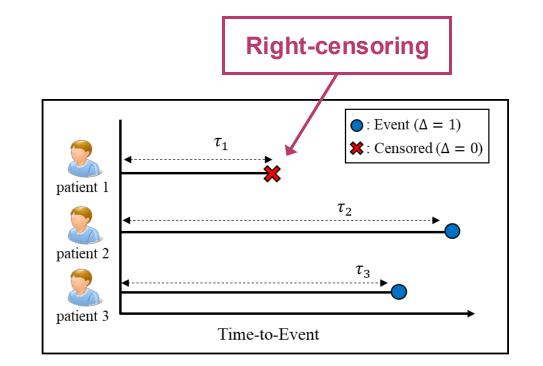


Figure. An illustration of survival data

• Distinct Characteristics: **Right-censoring**

Survival data

o Notations

- $T \in \mathbb{R}_+$ be the random variable for time-to-event
- $C \in \mathbb{R}_+$ be the random variables for time-to-censoring
- Right-censoring indicates when censoring occurs before the event of interest is observed. Denoting *t* and *c* be the realizations of r.v.s *T* and *C*, we have

$$\delta = \mathbb{I}(t \le c)$$
 $\tau = \min(t, c)$

- Often assume "independent censoring", i.e., $P(T, C|X = \mathbf{x}) = P(T|X = \mathbf{x})P(C|X = \mathbf{x})$

Solution: Survival Analysis

• Our goal

- Provides the probability an event occurring as a function of time and patient features
- Provides understanding of interactions between features and the time-to-event outcomes

- We want to use partial information from the right-censored samples:
 - Censoring implies that the event will occur after the censoring time

Important quantities : Survival / Risk function

 \circ Formally, we want to estimate the survival function given x

$$S(t|\mathbf{x}) = \mathbb{P}(T > t|\mathbf{x})$$

probability an event occurring after time t

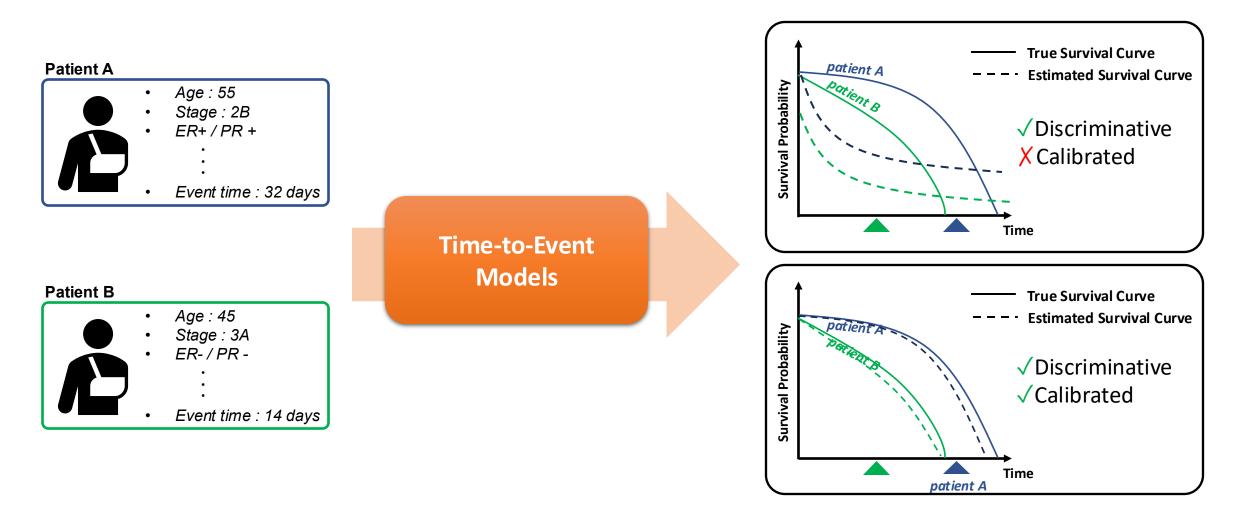
- $T \in \mathbb{R}_+$: Random variable for the time-to-event
- x: Patient input feature
- $\circ~$ Or equivalently, we want to estimate the risk function given x

$$R(t|\mathbf{x}) = 1 - S(t|\mathbf{x}) = \mathbb{P}(T \le t|\mathbf{x})$$

probability an event occurring before time t

Consideration

• Discriminate patients' risks of having an event of interest



Objective : Negative log-likelihood loss

- The log-likelihood of the time-to-events for survival dataset \rightarrow unbiased
 - Event is observed (i.e., $\delta_i = 1$), knowing that the event occurred at time τ_i
 - Event is not observed (i.e., $\delta_i = 0$), knowing that the event will occur after time τ_i

$$\mathcal{L}_{NLL} = -\log \prod_{i=1}^{N} \left[\hat{p}(\tau_i | \mathbf{x}_i)^{\delta_i} \cdot \hat{S}(\tau_i | \mathbf{x}_i)^{(1-\delta_i)} \right]$$
$$= -\sum_{i=1}^{N} \left[\frac{\delta_i \log \hat{p}(\tau_i | \mathbf{x}_i)}{\int for \text{ uncensored}} + \frac{(1-\delta_i) \log \hat{S}(\tau_i | \mathbf{x}_i)}{\int for \text{ censored}} \right]$$

- Often augmented with the NLL loss to enhance the discriminative power
- Aim to maximize a relaxed proxy of the concordance index
 - Well-established metric for evaluating the quality of patient rankings based on the risk predictions of survival model

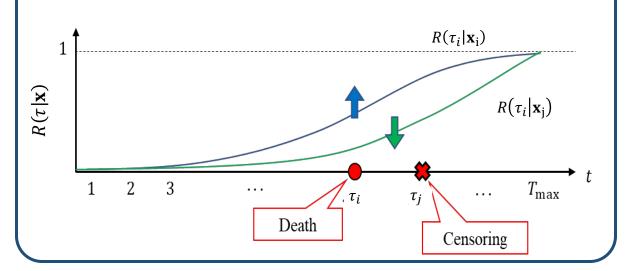
$$\mathcal{L}_{Rank} = \sum_{i \neq j} A_{i,j} \cdot \eta \left(\hat{R}(\tau_i | \mathbf{x}_i), \hat{R}(\tau_i | \mathbf{x}_j) \right)$$

Objective : Ranking loss

$$\mathcal{L}_{Rank} = \sum_{i \neq j} \underline{A_{i,j}} \cdot \eta \left(\widehat{R}(\tau_i | \mathbf{x}_i), \widehat{R}(\tau_i | \mathbf{x}_j) \right)$$

Case 1: Correctly ordered pairs

- $\hat{R}(\tau_i | \mathbf{x}_i) > \hat{R}(\tau_i | \mathbf{x}_j)$ (O)
- Rewards the estimated Risk Function $\hat{R}(\tau_i | \mathbf{x}_i)$ $\hat{R}(\tau_i | \mathbf{x}_i)$



re
$$A_{i,j} = \mathbb{I}(\delta_i = 1, \tau_i < \tau_j)$$

and $\eta(x, y) = \exp\left(\frac{-(x-y)}{\sigma}\right)$

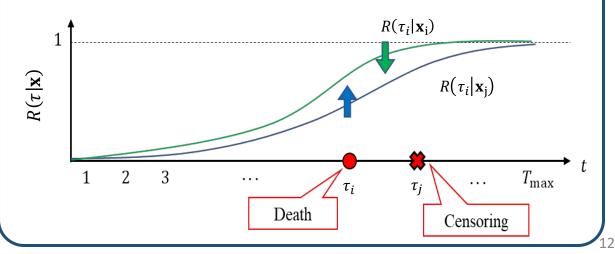
Case 2: Wrongly ordered pairs

• $\hat{R}(\tau_i | \mathbf{x}_i) < \hat{R}(\tau_i | \mathbf{x}_j)$ (X)

whe

Penalizes the estimated Risk Function





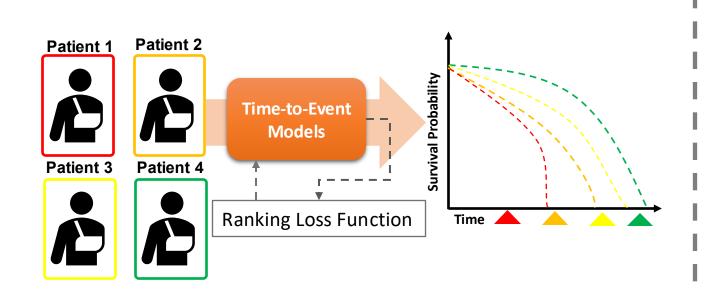
Challenges

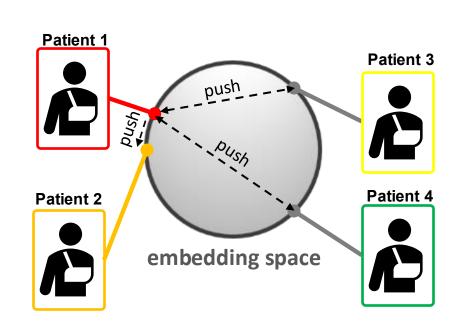
- Combining NLL with ranking loss enhances discrimination but compromises calibration, harming the clinical utility of predicted survival outcomes.
- Ranking loss directly modifies model outputs, potentially leading to misalignment with the actual risk distribution.
 - Typically based on exponential, log-sigmoid, or linear functions

Model	Ranking Loss
DeepHit	$\exp(-(\hat{R}(\tau_i \mathbf{x}_i) - \hat{R}(\tau_i \mathbf{x}_j)/\boldsymbol{\kappa}))$
DCS	$\exp(-(\hat{S}(\tau_i \mathbf{x_j}) - \hat{S}(\tau_i \mathbf{x_i}) / \boldsymbol{\kappa})$
LowerCI	$\log \sigma(\hat{R}(\tau_i \mathbf{x_i}) - \hat{R}(\tau_i \mathbf{x_j}))$
SSMTL	$\hat{h}(\tau_i \mathbf{x}_j) - \hat{h}(\tau_i \mathbf{x}_i)$

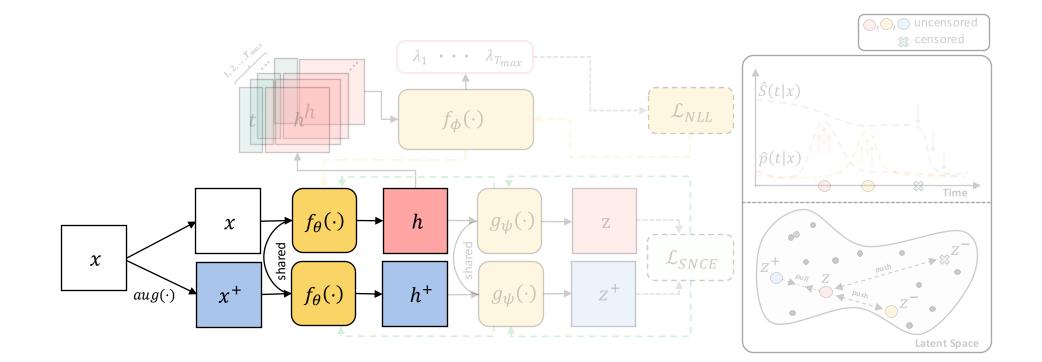
Motivation

- Propose a novel contrastive learning approach for deep survival model
 - Differentiate each sample by their survival outcome, leveraging contrastive learning framework
 - Overcomes ranking loss limitations from directly comparing model outcome in the form of risk/survival function.

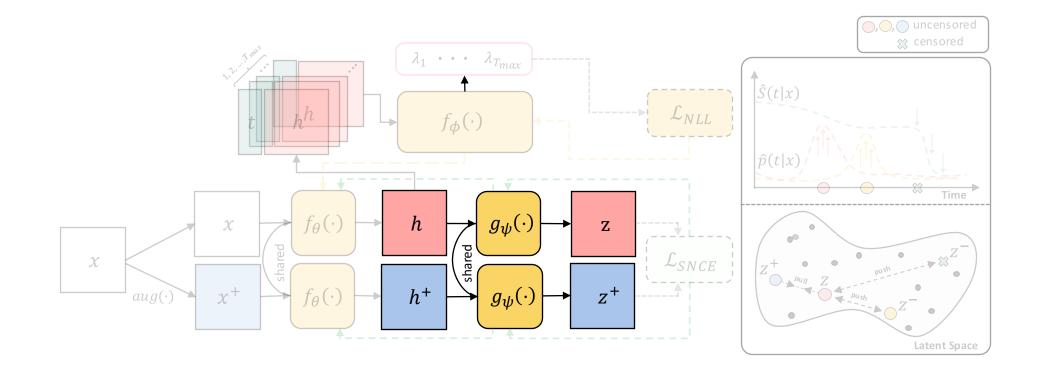




• **The encoder**, $f_{\theta} : \mathcal{X} \to \mathcal{H}$, takes features $\mathbf{x} \in \mathcal{X}$ as input and outputs latent representation, i.e., $\mathbf{h} = f_{\theta}(\mathbf{x})$.

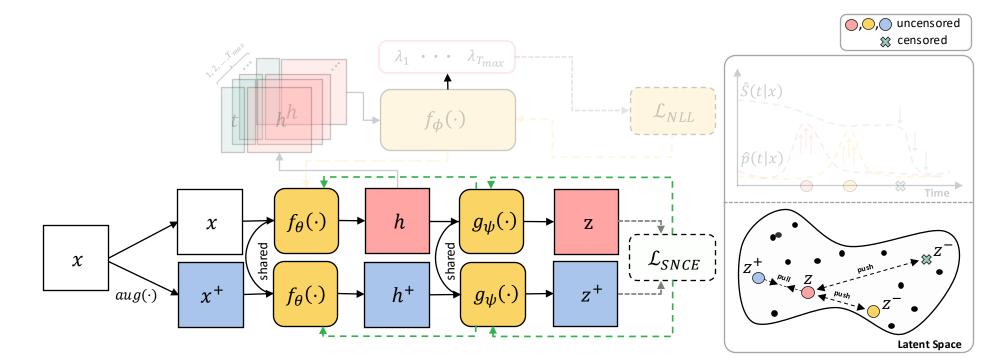


• The projection head., $g_{\psi} : \mathcal{H} \to \mathbb{R}^d$, maps latent representation **h** to the embedding space where contrastive learning is applied, i.e., $\mathbf{z} = f_{\theta}(\mathbf{h})$.



o Contrastive Learning Network

- By passing the original, positive, and negative samples through $f = g_{\psi} \circ f_{\theta}$, computing our survival outcome-based contrastive learning loss function \mathcal{L}_{SNCE}



- Goal : Aligns with our inductive bias that patients with similar survival outcomes should share similar clinical status, which manifests through similar representations.
- Noise Contrastive Estimation (NCE)
 - To learn mapping $f = g_{\psi} \circ f_{\theta}$ utilizing a positive sample $\mathbf{x}^+ \sim p_{X^+}$, and negative samples $\mathbf{x}^- \sim q$

$$\mathbb{E}_{\substack{\mathbf{x} \sim p_{X} \\ \mathbf{x}^{+} \sim p_{X^{+}}}} \left[-\log \frac{e^{s(\mathbf{x}, \mathbf{x}^{+})}}{M \cdot \mathbb{E}_{\mathbf{x}^{-} \sim q} \left[e^{s(\mathbf{x}, \mathbf{x}^{-})} \right]} \right]$$

- *M* : scaling term which is set to the batch size, $s(\mathbf{x}, \mathbf{x}') = \frac{f(\mathbf{x})^T f(\mathbf{x}')}{||f(\mathbf{x})|| \cdot ||f(\mathbf{x}')||}$

- omit the corresponding temperature v and write $e^{s(\mathbf{x}, \mathbf{x}^{-})}$ to denote $e^{s(\mathbf{x}, \mathbf{x}^{-})/v}$

- Key aspect of NCE : selecting negative samples to differentiate the anchor sample
- To reflect the difference in the time-to-events in the embedding space, we design a novel distribution q by utilizing the available information from survival outcomes.

- To accurately distinguish patients based on their time-to-event outcomes, we fully utilize the time-to-event information
- Hence, given an anchor (\mathbf{x}, τ) and a negative $(\mathbf{x}^-; \tau^-)$, we define the weight function, $\sigma > 0$ is a temperature coefficient.
 - This function is a variant of the Laplacian Kernel, which assigns larger weights to samples with large differences in time-to-event outcomes, and smaller weights to samples with small differences

$$w(\tau^{-};\tau) = 1 - e^{|\tau - \tau^{-}|/\sigma}$$

- Designing *q* based on the following inductive bias : similar patients are more likely to experience the event at similar time points than the ones who are not.
- We will slightly abuse the notation $w(\mathbf{x}^-; \mathbf{x})$ to denote $w(\tau^-; \tau)$

$$q(\mathbf{x}^-; \mathbf{x}) = \frac{1}{Z} w(\mathbf{x}^-; \mathbf{x}) p(\mathbf{x}^-)$$

normalizing constant $Z = \frac{1}{M} \sum_{j=1}^{M} w(x_j^{-}; x)$

Importance sampling using survival outcomes

$$E_{x^{-} \sim q} \left[e^{s(x,x^{-})} \right] = E_{x^{-} \sim p} \left[\left(\frac{q(x^{-};x)}{p(x^{-})} \right) \cdot e^{s(x,x^{-})} \right]$$

$$= E_{x^{-} \sim p} \left[\left(\frac{w(x^{-};x)}{Z} \right) \cdot e^{s(x,x^{-})} \right]$$

$$\approx \frac{1}{Z \cdot M} \sum_{j=1}^{M} w(x_j^-; x) \cdot e^{s(x,x_j^-)}$$

normalizing constant $Z = \frac{1}{M} \sum_{j=1}^{M} w(x_j^{-}; x)$

• Survival outcome-aware NCE (SNCE) loss

$$\mathcal{L}_{SNCE} = \sum_{i=1}^{N} \left[-\log\left(\frac{e^{s(x_i, x_i^+)}}{\frac{1}{Z}\sum_{j=1}^{M} w(x_j^-; x_i) \cdot e^{s(x_i, x_j^-)}}\right) \right]$$

case 1 : Both samples are uncensored(i.e., have observed events)

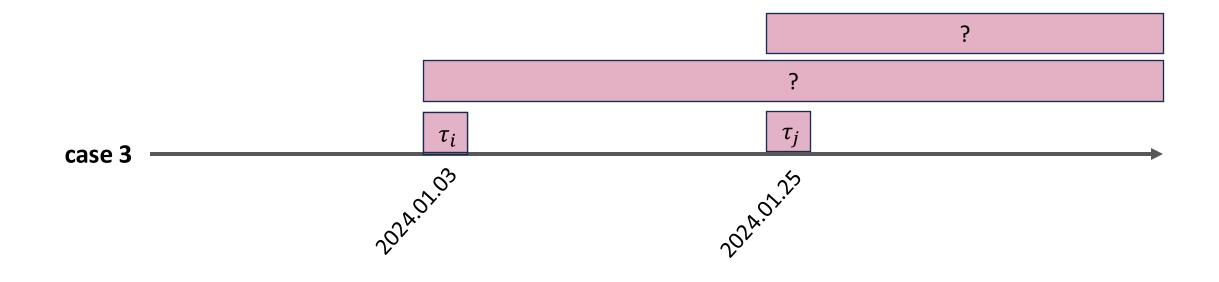
- o case 2 : Both samples are censored
- o case 3 : One is uncensored and the other is censored.



case 1 : Both samples are uncensored(i.e., have observed events)

o case 2 : Both samples are censored

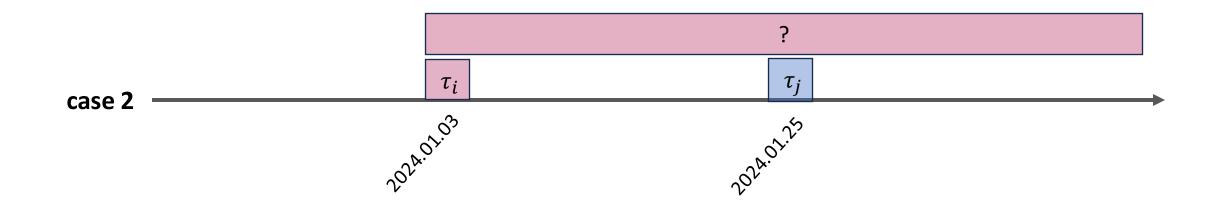
o case 3 : One is uncensored and the other is censored.



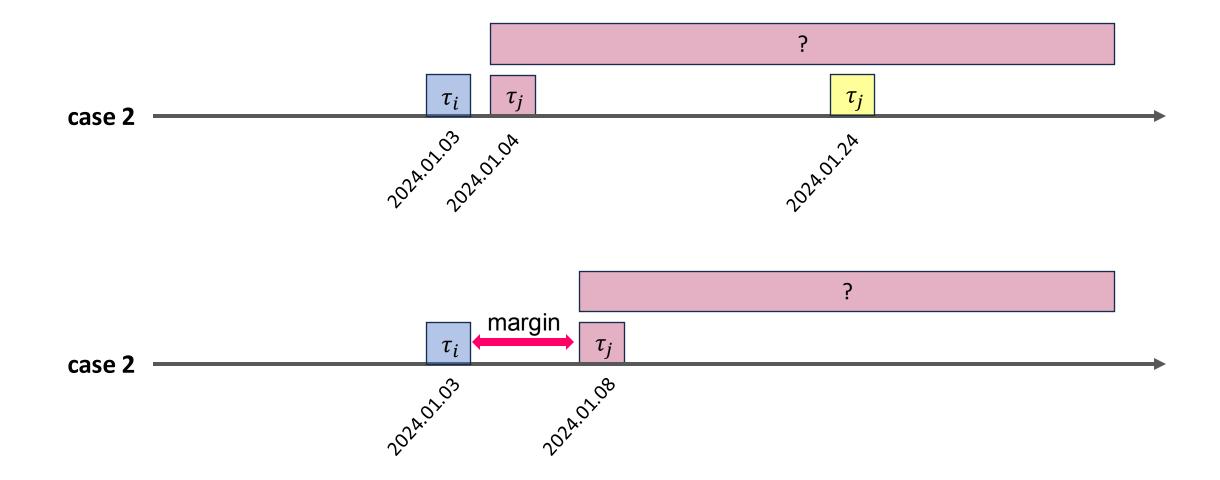
- o case 1 : Both samples are uncensored(i.e., have observed events)
- o case 2 : Both samples are censored
- o case 3 : One is uncensored and the other is censored.



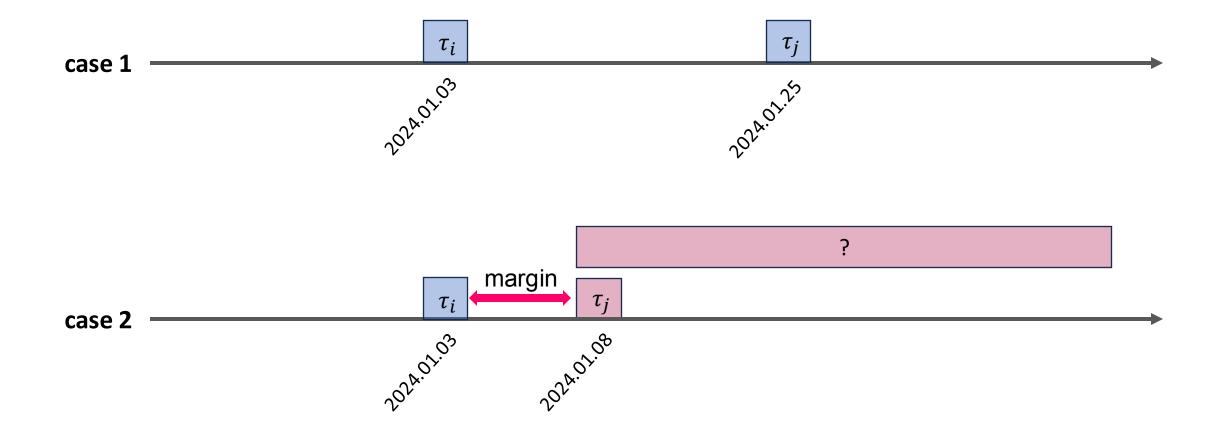
- o case 1 : Both samples are uncensored(i.e., have observed events)
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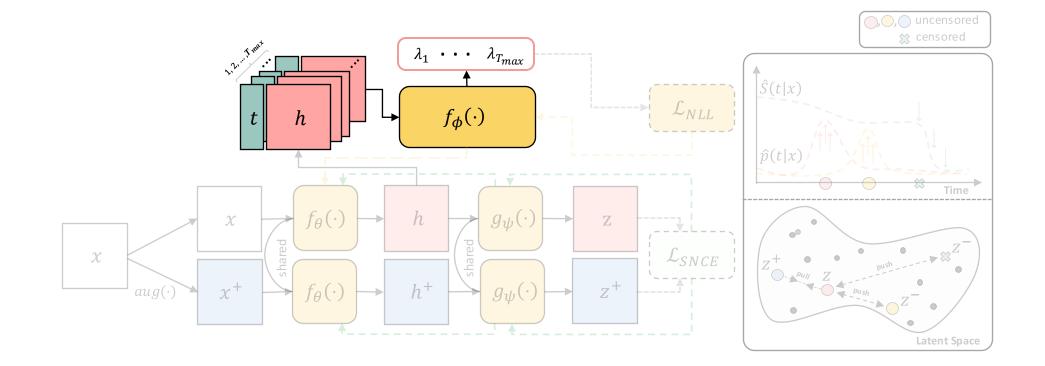
o case 3 : One is uncensored and the other is censored.



• Redefine the weight function considering the right-censoring as



• The hazard network., $f_{\phi} : \mathcal{H} \times \mathcal{T} \to [0,1]$, predicts the hazard rate at each time point $t \in \mathcal{T}$ given input latent representation **h**, i.e., $\hat{\lambda}(t|\mathbf{x}) = f_{\phi}(\mathbf{h}, t) = f_{\phi}(f_{\theta}(\mathbf{x}), t)$



Important quantities : Hazard function

• The hazard function, $\lambda(t)$, is the instantaneous rate of failure at time t, given that an individual has survived until at least time t:

$$\lambda(t|\mathbf{x}) = P(T = t|T \ge t, \mathbf{x}) \quad t \in \{1, 2, \dots\}$$

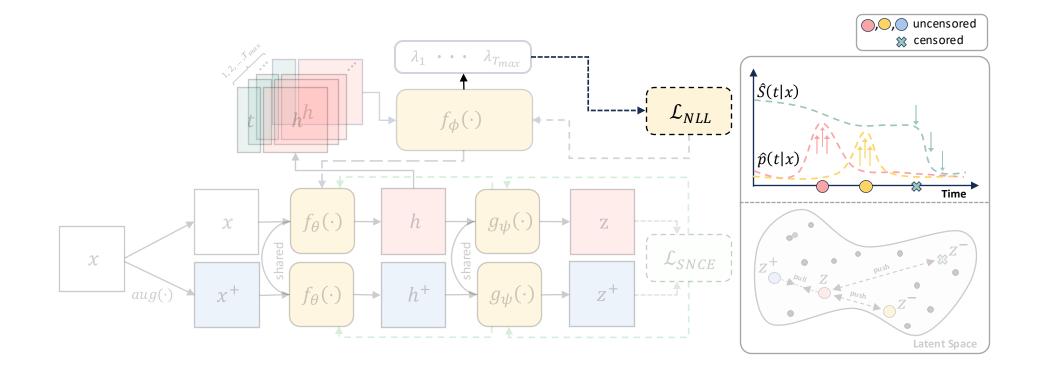
• There is an important relationship between the survival and hazard functions:

$$S(t|\mathbf{x}) = P(T > t | \mathbf{x})$$

= $P(T \neq 1 | \mathbf{x}) \cdot P(T \neq 2 | T > 1, \mathbf{x}) \cdots P(T = t | T > t - 1, \mathbf{x})$
= $P(1 - \lambda(1 | \mathbf{x})) \cdot P(1 - \lambda(2 | \mathbf{x})) \cdot \cdots \cdot P(1 - \lambda(t | \mathbf{x}))$
= $\prod_{t' \leq t} (1 - \lambda(t'|\mathbf{x}))$

• Negative Log-likelihood

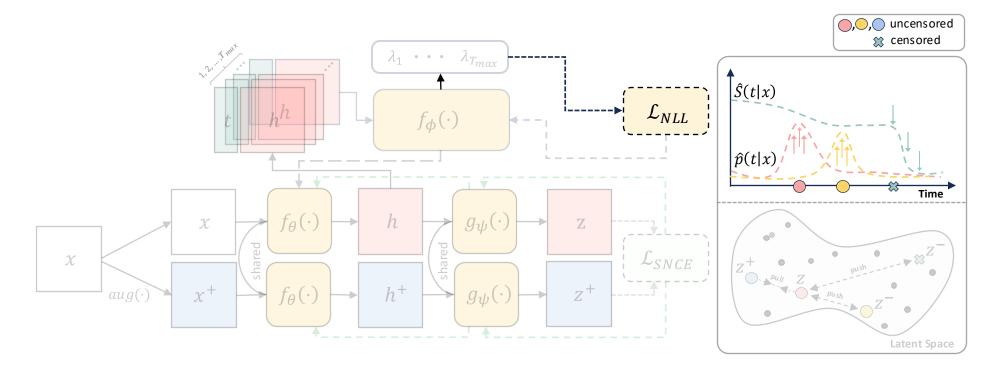
 Hazard estimate is defined as a function of time given an input feature, we can naturally model the time-varying effect of input features on risk/survival functions.



• Negative Log-likelihood

- Then, compute $\mathcal{L}_{NLL}^{\theta,\phi}$ by plugging in $f_{\phi}(f_{\theta}(\mathbf{x}), t)$ into \hat{p} and \hat{S}

 $- \hat{p}(\tau | \mathbf{x}) = f_{\phi}(f_{\theta}(\mathbf{x}), \tau) \prod_{t' \leq \tau - 1} \left(1 - f_{\phi}(f_{\theta}(\mathbf{x}), t') \right), \quad \hat{S}(\tau | \mathbf{x}) = \prod_{t' \leq \tau} \left(1 - f_{\phi}(f_{\theta}(\mathbf{x}), t') \right)$



• Overall, we can estimate the hazard function by training ConSurv with a loss function that combines the NLL loss and the SNCE loss, where β is a balancing coefficient

$$\mathcal{L}_{Total}^{\theta,\phi,\psi} = \mathcal{L}_{NLL}^{\theta,\phi} + \beta \mathcal{L}_{SNCE}^{\theta,\psi}$$

Experiments Setup : Datasets & Benchmarks & Metrics

• Datasets

Dataset	No. Uncensored	No. Censored	No. Features (Real, Binary, Category)
METABRIC	888 (55.2%)	1093 (44.8%)	21 (6, 0, 15)
NWTCO	571 (14.2%)	3457 (85.5%)	6 (1, 4, 1)
GBSG	1267 (56.8%)	965 (43.2%)	7 (4, 2, 1)
FLCHAIN	4562 (69.9%)	1962 (30.3%)	8 (4, 2, 2)
SUPPORT	6036 (68.1%)	2837 (31.9%)	14 (8, 3, 3)
SEER	604 (1.11%)	53940 (98.9%)	12 (4, 5, 3)

o Benchmarks

Loss Function	Туре	Model
Partial Log-likelihood	ML	CoxPH
Fartial Log-likelihood	DL	DeepSurv
Ranking Loss	DL	DeepHit
	DL	DRSA
Collibration Loop	DL	DCS
Calibration Loss	DL	X-CAL

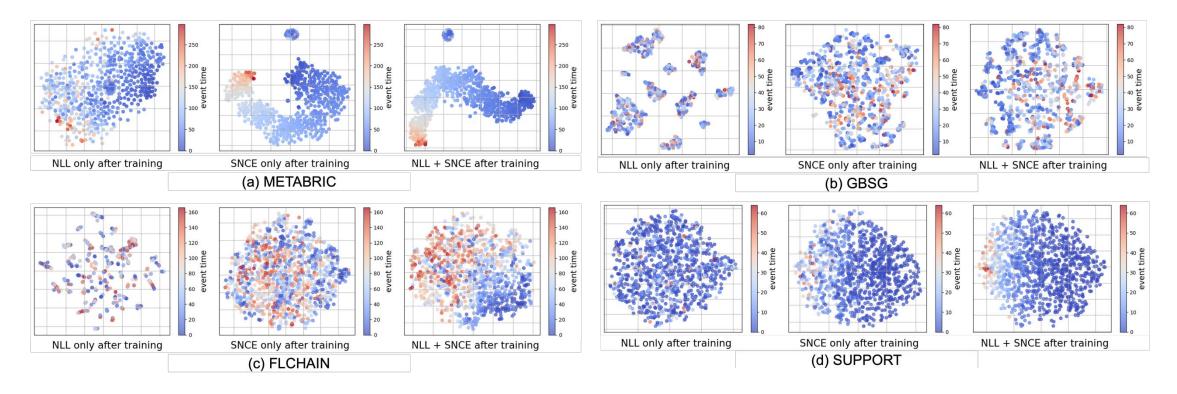
• Metrics

Evaluation Metric	Туре	Range	
Concordance Index (CI)	Discrimination	0.000 ~ 1.000	ſ
Integrated Brier Score (IBS)	Calibration	0.000 ~ 1.000	↓
Distribution Divergence for Calibration (DDC)	Calibration	0.000 ~ 1.000	↓
D-calibration (D-CAL)	Calibration	P-value>0.05	×

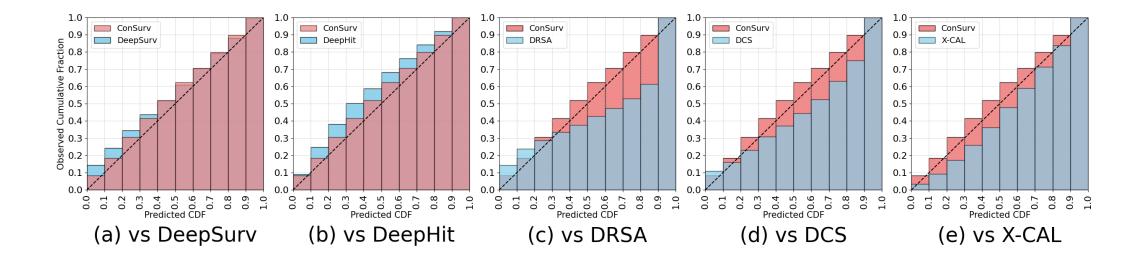
		МЕТАВ	BRIC		NWTCO			
Метнор	CI↑	IBS \downarrow	DDC \downarrow	D-CAL	CI ↑	IBS \downarrow	$DDC\downarrow$	D-CAL
CoxPH DeepSurv DeepHit DRSA DCS X-CAL	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	$\begin{array}{c} \textbf{0.175}_{\pm \textbf{0.028}} \\ 0.183_{\pm 0.029} \\ 0.204_{\pm 0.018} \\ 0.249_{\pm 0.038} \\ 0.206_{\pm 0.043} \\ 0.182_{\pm 0.023} \end{array}$	$\begin{array}{c} 0.111 {\scriptstyle \pm 0.024} \\ 0.103 {\scriptstyle \pm 0.026} \\ 0.292 {\scriptstyle \pm 0.017} \\ 0.178 {\scriptstyle \pm 0.060} \\ \textbf{0.054} {\scriptstyle \pm 0.039} \\ 0.065 {\scriptstyle \pm 0.037} \end{array}$	25 25 0 0 2 2	$\begin{array}{c} 0.716 _{\pm 0.025} \\ 0.640 _{\pm 0.080} \\ 0.717 _{\pm 0.028} \\ 0.709 _{\pm 0.019} \\ 0.642 _{\pm 0.036} \\ 0.622 _{\pm 0.037} \end{array}$	$\begin{array}{c} 0.108 _{\pm 0.008} \\ 0.117 _{\pm 0.011} \\ 0.143 _{\pm 0.024} \\ 0.281 _{\pm 0.041} \\ 0.119 _{\pm 0.018} \\ 0.128 _{\pm 0.025} \end{array}$	$\begin{array}{c} 0.515_{\pm 0.022} \\ 0.792_{\pm 0.011} \\ 0.657_{\pm 0.024} \\ 0.218_{\pm 0.065} \\ 0.209_{\pm 0.043} \\ \textbf{0.191}_{\pm 0.079} \end{array}$	25 24 12 0 19 12
L _{NLL} L _{NLL} & L _{NCE} L _{NLL} & L _{Rank}	$ \begin{vmatrix} 0.642_{\pm 0.022} \\ 0.659_{\pm 0.020} \\ 0.652_{\pm 0.022} \end{vmatrix} $	$\begin{array}{c} 0.197 _{\pm 0.030} \\ 0.193 _{\pm 0.029} \\ 0.247 _{\pm 0.030} \end{array}$	$\begin{array}{c} 0.077_{\pm 0.020} \\ 0.080_{\pm 0.022} \\ 0.177_{\pm 0.020} \end{array}$	13 21 0	$\begin{array}{c} 0.707_{\pm 0.024} \\ 0.715_{\pm 0.024} \\ 0.717_{\pm 0.027} \end{array}$	$\begin{array}{c} 0.109_{\pm 0.008} \\ 0.108_{\pm 0.009} \\ 0.137_{\pm 0.008} \end{array}$	$\begin{array}{c} 0.556 _{\pm 0.041} \\ 0.563 _{\pm 0.054} \\ 0.653 _{\pm 0.050} \end{array}$	23 22 0
CONSURV	$0.665_{\pm 0.023}$	$0.186_{\pm 0.021}$	$0.110_{\pm 0.024}$	23	$\textbf{0.718}_{\pm 0.025}$	$\textbf{0.107}_{\pm 0.008}$	$0.554_{\pm 0.045}$	24
	SUPPORT				SEER			
		SUPPO	ORT			SEE	R	
Метнор	CI↑	SUPPC IBS↓	DRT DDC↓	D-CAL	CI↑	SEE IBS↓	R DDC↓	D-CAL
METHOD CoxPH DeepSurv DeepHit DRSA DCS X-CAL	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$			D-CAL 0 0 0 0 0 0 0	$\begin{array}{c} CI\uparrow\\ 0.858{\scriptstyle\pm0.018}\\ 0.814{\scriptstyle\pm0.020}\\ 0.840{\scriptstyle\pm0.033}\\ 0.834{\scriptstyle\pm0.078}\\ 0.860{\scriptstyle\pm0.020}\\ 0.837{\scriptstyle\pm0.040}\end{array}$			D-CAL 25 25 0 0 21 18
CoxPH DeepSurv DeepHit DRSA DCS	$ \begin{vmatrix} 0.604_{\pm 0.006} \\ 0.603_{\pm 0.090} \\ 0.503_{\pm 0.009} \\ 0.570_{\pm 0.009} \\ 0.598_{\pm 0.008} \end{vmatrix} $	$\begin{array}{c} \text{IBS} \downarrow \\ 0.191_{\pm 0.005} \\ 0.192_{\pm 0.007} \\ 0.272_{\pm 0.003} \\ 0.259_{\pm 0.015} \\ 0.207_{\pm 0.012} \end{array}$	$\begin{array}{c} \text{DDC}\downarrow\\ 0.262_{\pm0.013}\\ 0.245_{\pm0.036}\\ 0.337_{\pm0.006}\\ 0.486_{\pm0.084}\\ 0.175_{\pm0.032}\end{array}$	0 0 0 0 0	$\begin{array}{c} 0.858 _{\pm 0.018} \\ 0.814 _{\pm 0.020} \\ 0.840 _{\pm 0.033} \\ 0.834 _{\pm 0.078} \\ 0.860 _{\pm 0.020} \end{array}$	$IBS \downarrow \\ 0.009_{\pm 0.005} \\ 0.010_{\pm 0.000} \\ 0.020_{\pm 0.001} \\ 0.021_{\pm 0.015} \\ 0.010_{\pm 0.001}$	$\begin{array}{c} \text{DDC} \downarrow \\ 0.966_{\pm 0.003} \\ 1.000_{\pm 0.000} \\ 0.836_{\pm 0.003} \\ \textbf{0.671}_{\pm 0.135} \\ 0.911_{\pm 0.044} \end{array}$	25 25 0 0 21

		МЕТАВ	RIC		NWTCO			
Метнор	CI ↑	IBS \downarrow	$DDC\downarrow$	D-CAL	CI ↑	IBS \downarrow	$DDC\downarrow$	D-CAL
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X-CAL	$0.632_{\pm 0.027}$	$0.182_{\pm 0.023}$	$0.065_{\pm 0.037}$	2	0.642 ± 0.036 0.622 ± 0.037	$0.119 \pm 0.018 \\ 0.128 \pm 0.025$	$0.191_{\pm 0.079}$	12
$egin{array}{c} \mathcal{L}_{NLL} & \mathcal{L}_{NCE} \ \mathcal{L}_{NLL} & \mathcal{L}_{Rank} \end{array}$	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	$\begin{array}{c} 0.197_{\pm 0.030} \\ 0.193_{\pm 0.029} \\ 0.247_{\pm 0.030} \end{array}$	$\begin{array}{c} 0.077_{\pm 0.020} \\ 0.080_{\pm 0.022} \\ 0.177_{\pm 0.020} \end{array}$	13 21 0	$\begin{array}{c} 0.707_{\pm 0.024} \\ 0.715_{\pm 0.024} \\ 0.717_{\pm 0.027} \end{array}$	$\begin{array}{c} 0.109_{\pm 0.008} \\ 0.108_{\pm 0.009} \\ 0.137_{\pm 0.008} \end{array}$	$\begin{array}{c} 0.556_{\pm 0.041} \\ 0.563_{\pm 0.054} \\ 0.653_{\pm 0.050} \end{array}$	23 22 0
CONSURV	$0.665_{\pm 0.023}$	$0.186_{\pm 0.021}$	$0.110_{\pm 0.024}$	23	$0.718 \scriptstyle \pm 0.025$	$0.107 \scriptstyle \pm 0.008$	$0.554_{\pm 0.045}$	24
		SUPPO	ORT			SEE	R	
Метнор	CI↑	SUPPC IBS↓	DRT DDC↓	D-CAL	CI↑	SEE IBS↓	R DDC↓	D-CAL
METHOD CoxPH DEEPSURV DEEPHIT DRSA DCS X-CAL	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$			D-CAL 0 0 0 0 0 0 0	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$			D-CAL 25 25 0 0 21 18
CoxPH DeepSurv DeepHit DRSA DCS	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	$\begin{array}{c} \text{IBS} \downarrow \\ 0.191_{\pm 0.005} \\ 0.192_{\pm 0.007} \\ 0.272_{\pm 0.003} \\ 0.259_{\pm 0.015} \\ 0.207_{\pm 0.012} \end{array}$	$\begin{array}{c} \text{DDC}\downarrow\\ 0.262_{\pm0.013}\\ 0.245_{\pm0.036}\\ 0.337_{\pm0.006}\\ 0.486_{\pm0.084}\\ 0.175_{\pm0.032}\end{array}$	0 0 0 0 0	$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	$IBS \downarrow \\ 0.009_{\pm 0.005} \\ 0.010_{\pm 0.000} \\ 0.020_{\pm 0.001} \\ 0.021_{\pm 0.015} \\ 0.010_{\pm 0.001} \\ 0.010_{\pm 0.001}$	$\begin{array}{c} \text{DDC} \downarrow \\ 0.966_{\pm 0.003} \\ 1.000_{\pm 0.000} \\ 0.836_{\pm 0.003} \\ \textbf{0.671}_{\pm 0.135} \\ 0.911_{\pm 0.044} \end{array}$	25 25 0 0 21

- o Effect of Contrastive Learning
 - \mathcal{L}_{NLL} only, \mathcal{L}_{SNCE} only, and ConSurv (i.e., $\mathcal{L}_{NLL} \& \mathcal{L}_{SNCE}$)
 - significantly improves the alignment of representations with event time information



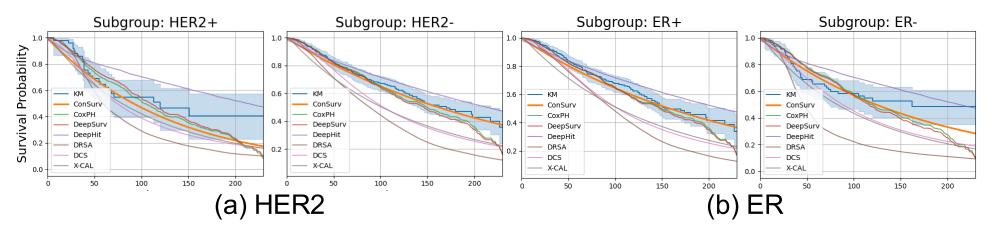
- Comparing calibration plot of ConSurv with the DL-based survival models
 - The x=y line represents the ideal state where predicted probabilities perfectly match the observed outcome

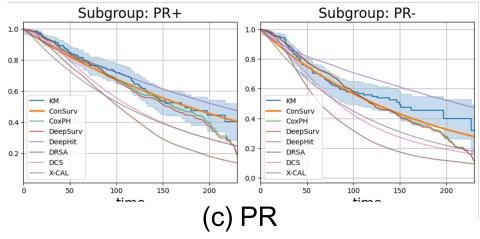


Subgroup Analysis

- To confirm the calibration performance of survival models, compare their survival plots with the Kaplan-Meier (KM) curve
- KM curve provides a non-parametric estimate of survival function at population level
- Examine three binary hormone receptor status in the METABRIC dataset: estrogen receptor (ER) ,human epidermal growth factor receptor 2 (HER2), and progesterone receptor (PR)

• Subgroup Analysis





• Subgroup Analysis

 To quantitatively assess calibration performance, compare the survival predictions of each model with the KM curves for each subgroup using the Wasserstein distance

Subgroup	ER		HER2		Size		PR	
	+	-	+	-	+	-	+	-
CoxPH	0.030	0.108	0.063	0.089	0.049	0.067	0.076	0.044
DeepSurv	0.033	0.115	0.066	0.101	0.054	0.074	0.105	0.089
DeepHit	0.063	0.082	0.146	0.156	0.043	0.102	0.233	0.033
DRSA	0.181	0.293	0.233	0.328	0.205	0.276	0.118	0.234
DCS	0.130	0.146	0.087	0.178	0.154	0.126	0.091	0.124
X-CAL	0.136	0.165	0.105	0.180	0.159	0.148	0.060	0.176
ConSurv	0.024	0.077	0.044	0.089	0.043	0.042	0.060	0.025

- Survival data often lacks clear event times complicating learning due to censored data.
 - Potential avenues include modifying models to account for uncertainties or developing alternative learning approaches
 - Need for new evaluation metrics that consider the characteristics
- Limited augmentation techniques in tabular data reduce model robustness
 - Explore augmentation methods suitable for survival datasets to enhance contrastive learning performance

Thank you