

Explicitly modeling risk dynamics improves health status representations



August 12, 2026
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RESEARCH
TRACK

HazardFlow: Enhancing Health Status Representations via Score-based Energy Modeling

by:

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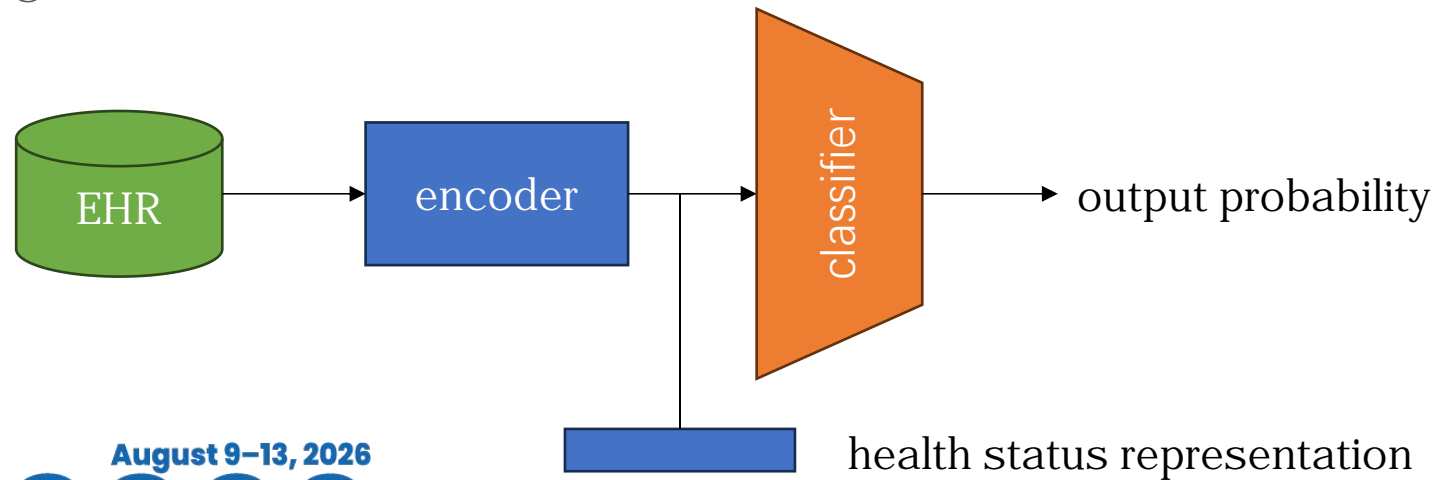
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Health Risk Prediction with Electronic Health Records

- EHRs digitize diverse clinical data, including diagnoses, laboratory results, and treatments, providing a comprehensive view of patient health.
- Health risk prediction plays a critical role in early intervention, resource allocation, and clinical decision support by anticipating adverse clinical events.

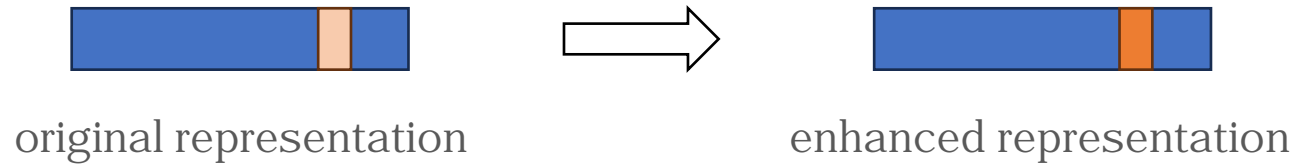


Subtle risk signals are overwhelmed in EHR representations

Real-world EHR (Electronic Health Record) data are sparse, irregularly sampled, and highly imbalanced. As a result, learned representations tend to be dominated by majority low-risk patterns, while subtle but critical risk signals remain underrepresented.

Task	MIMIC-III		MIMIC-IV	
	#label=1	#label=0	#label=1	#label=0
mortality	662	9,055	2,083	123,631

Intuition



If we can identify a risk-related direction in the latent space, we can move health representations along this direction to make risk signals more explicit.

Survival Analysis

Survival Function



Survival Function: $S(t) = 1 - F(t)$. The probability that a person or machine or a business lasts longer than t time units. ($F(t)$: the cumulative distribution function)

Hazard Function: $\lambda(t) = f(t)/S(t)$. The probability that the person or machine or business dies in the next instant, given that it survived to time t . ($f(t)$: the probability density function)

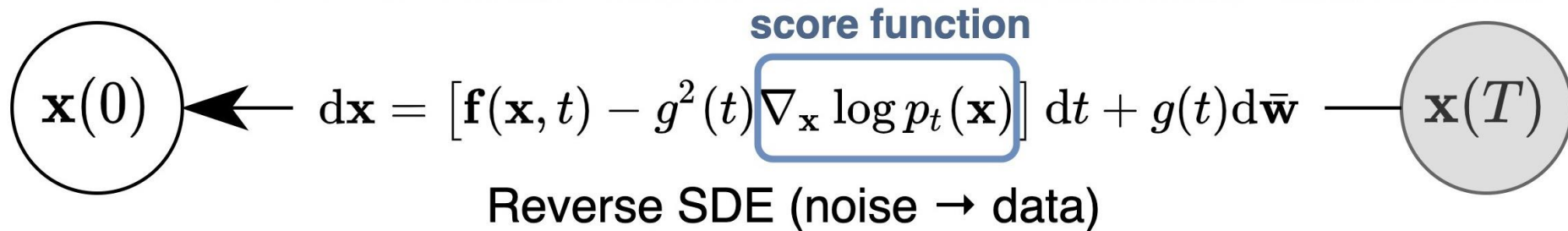
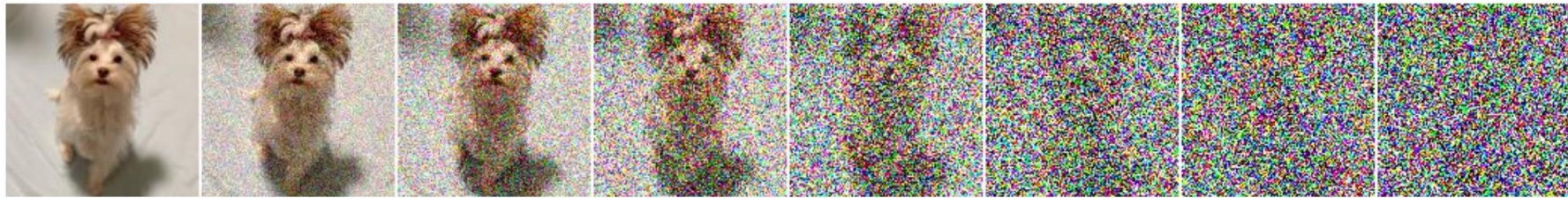
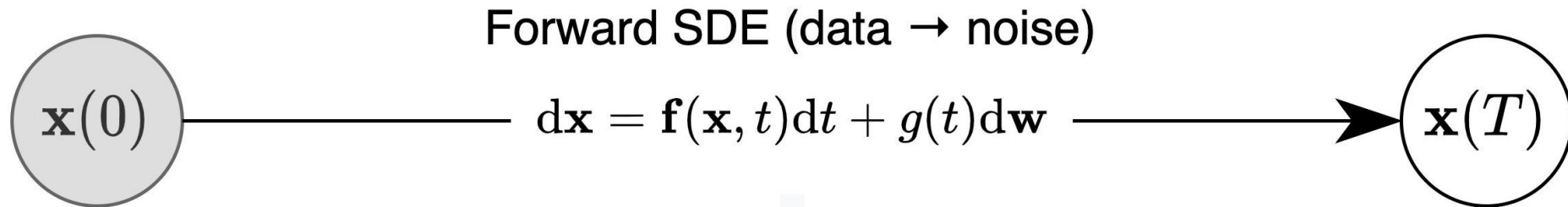
The score function provides a hazard-related direction

Proposition 1 (Score-Hazard Correspondence). Let $\mathbf{x}_t \in \mathbb{R}^{1 \times d}$ denote a health status representation at time t , with marginal distribution $p_t(\mathbf{x})$. Suppose the survival function $S(t)$ is learned by $1 - \psi(\mathbf{x}_t)$, and there exists a learned mapping from $\mathbf{x}_t$ to a dynamic hazard vector $\mathbf{h}_t \in \mathbb{R}^{1 \times d}$, which denotes the corresponding hazard function $h(t)$. Then, the time-dependent score function $\nabla_{\mathbf{x}} \log p_t(\mathbf{x})$ is approximately proportional to the hazard,

$$\mathbf{h}_t \approx \alpha(t) \nabla_{\mathbf{x}} \log p_t(\mathbf{x}),$$

where $\alpha(t)$ is a scalar function of time.

Score-based Generative Modeling



Survival Energy Hypothesis (SEH) [Shimizu et al., 2021]. SEH models mortality as the depletion of a latent quantity termed *survival energy*, which can be modeled as a drifted Brownian motion:

$$E_t = E_c + f_c t + g_c w_t.$$



Score-based Energy Modeling. To characterize the Brownian motion in SEH, we use $\mathbf{x}_t$ to encode the survival energy, and model $d\mathbf{x}_t$ using a diffusion process, governed by the following SDE:

$$d\mathbf{x}_t = f(\mathbf{x}_t, t)dt + g(t)dw_t.$$



Probability Flow Ordinary Differential Equation. We express the reverse process using a probability-flow ODE:

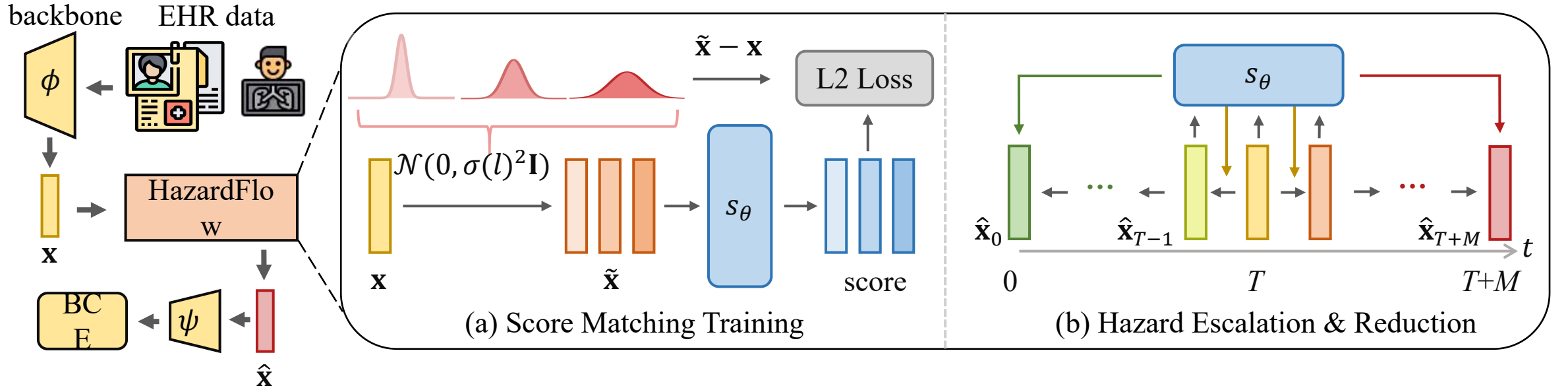
$$d\mathbf{x}_t = [f(\mathbf{x}_t, t)dt - g(t)^2 \nabla_{\mathbf{x}} \log p_t(\mathbf{x})]dt.$$

We omit the drift term and discretize the continuous denoising process using annealed Langevin dynamics:

$$x_{i+1} = x_i + \frac{\epsilon}{2} g(i)^2 \nabla_{\mathbf{x}} \log p_i(\mathbf{x}).$$

This iterative step can be regarded as enhancing the representation toward a higher-risk state.

HazardFlow



HazardFlow refines backbone representations through two steps:

Step 1: Train a score network using denoising score matching.

$$\mathcal{L}_{SM} = \mathbb{E}_l \left\{ \lambda(l) \mathbb{E}_{p(\mathbf{x})} \mathbb{E}_{q_l(\tilde{\mathbf{x}}|\mathbf{x})} \left[\left\| s_\theta(\tilde{\mathbf{x}}, l) + \frac{\tilde{\mathbf{x}} - \mathbf{x}}{\delta(l)^2} \right\|_2^2 \right] \right\}$$

Step 2: Iteratively add or subtract the learned score to obtain higher- or lower-risk representations.

$$\hat{\mathbf{x}}_{T+i+1} = \hat{\mathbf{x}}_{T+i} + \frac{\sigma^{2i}\epsilon}{2} s_\theta(\hat{\mathbf{x}}_{T+i}, i)$$

$$\hat{\mathbf{x}}_{i-1} = \hat{\mathbf{x}}_i - \frac{\sigma^{2i}\epsilon}{2} s_\theta(\hat{\mathbf{x}}_i, i)$$

Method	mortality				readmission			
	MIMIC-III		MIMIC-IV		MIMIC-III		MIMIC-IV	
	PRAUC	AUROC	PRAUC	AUROC	PRAUC	AUROC	PRAUC	AUROC
ConCare	11.76 $\pm$ 1.13	61.08 $\pm$ 2.47	4.18 $\pm$ 0.34	71.00 $\pm$ 0.80	67.75 $\pm$ 3.47	64.44 $\pm$ 2.05	68.12 $\pm$ 0.39	67.54 $\pm$ 0.36
ConCare w/ HF	25.22$\pm$7.47	75.65$\pm$7.47	4.59$\pm$0.68	72.60$\pm$1.37	71.82$\pm$3.96	68.37$\pm$4.90	69.18$\pm$2.25	68.26$\pm$1.54
StageNet	34.21 $\pm$ 34.83	69.08 $\pm$ 18.59	3.26 $\pm$ 0.32	66.19 $\pm$ 1.97	62.56 $\pm$ 2.40	61.06 $\pm$ 1.88	68.70 $\pm$ 2.99	67.72 $\pm$ 2.39
StageNet w/ HF	39.29$\pm$18.92	77.80$\pm$13.48	3.54$\pm$0.21	67.28$\pm$1.99	66.46$\pm$3.81	64.88$\pm$3.88	70.76$\pm$2.91	69.48$\pm$2.22
TCN	13.45 $\pm$ 5.77	61.32 $\pm$ 7.78	3.09 $\pm$ 0.33	65.66 $\pm$ 2.22	63.53 $\pm$ 2.23	60.70 $\pm$ 1.59	67.12 $\pm$ 0.48	66.70 $\pm$ 0.38
TCN w/ HF	17.46$\pm$9.21	65.08$\pm$8.03	3.15$\pm$0.26	66.77$\pm$2.18	65.36$\pm$3.32	62.34$\pm$3.40	67.29$\pm$0.40	66.81$\pm$0.42
GRASP	12.34 $\pm$ 5.09	60.24 $\pm$ 6.50	3.48 $\pm$ 0.31	67.84 $\pm$ 2.31	64.77 $\pm$ 2.81	61.76 $\pm$ 2.40	67.57 $\pm$ 0.44	67.07 $\pm$ 0.35
GRASP w/ HF	14.68$\pm$5.35	63.67$\pm$7.33	5.03$\pm$1.78	70.09$\pm$3.45	66.47$\pm$1.96	63.68$\pm$1.80	68.87$\pm$2.29	68.12$\pm$1.56
AdaCare	51.42 $\pm$ 36.06	77.52 $\pm$ 18.31	3.10 $\pm$ 1.22	64.79 $\pm$ 3.63	64.60 $\pm$ 2.28	63.45 $\pm$ 2.77	66.78 $\pm$ 0.78	66.62 $\pm$ 0.59
AdaCare w/ HF	58.88$\pm$31.77	79.99$\pm$16.83	4.60$\pm$2.72	66.06$\pm$5.41	68.75$\pm$5.72	68.05$\pm$5.14	68.07$\pm$1.37	67.71$\pm$1.77

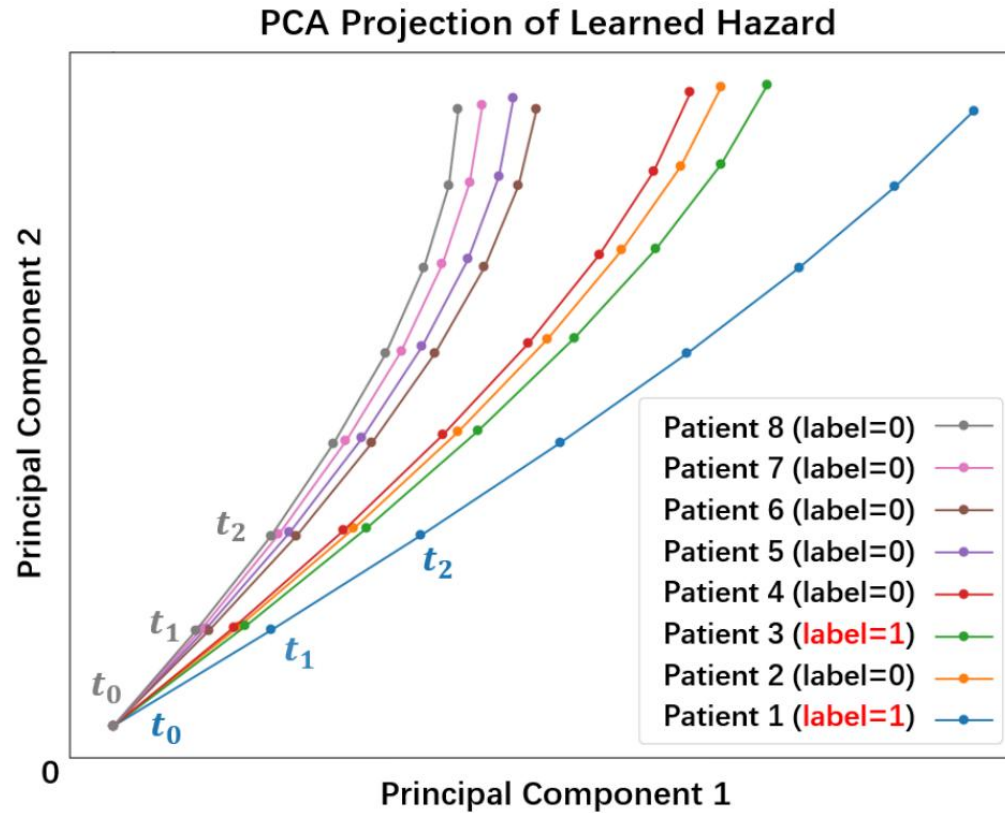
HazardFlow consistently improves five uni-modal backbones across mortality and readmission tasks.

Method	mortality		phenotyping	
	PRAUC	AUROC	PRAUC	AUROC
MedFuse	49.39 $\pm$ 1.41	85.16 $\pm$ 0.21	41.78 $\pm$ 0.26	76.31 $\pm$ 0.18
MedFuse w/ HF	51.58$\pm$2.22	85.51$\pm$0.38	41.91$\pm$0.13	76.44$\pm$0.09
DrFuse	41.70 $\pm$ 1.96	79.18 $\pm$ 1.54	39.02 $\pm$ 0.18	72.38 $\pm$ 0.22
DrFuse w/ HF	42.77$\pm$2.49	80.42$\pm$1.37	39.00 $\pm$ 0.21	72.45$\pm$0.09
DDL-CXR	44.92 $\pm$ 0.68	80.01 $\pm$ 0.74	43.58 $\pm$ 1.20	71.71 $\pm$ 0.70
DDL-CXR w/ HF	46.26$\pm$1.36	80.59$\pm$0.41	44.53$\pm$0.20	72.49$\pm$0.31

11 of 12 multi-modal metrics improved.

	PRAUC	AUROC
MedFuse	49.39 $\pm$ 1.41	85.16 $\pm$ 0.21
MedFuse w/ HF	50.20 $\pm$ 0.72	85.87$\pm$0.26
MeduFuse w/ HF+	50.70$\pm$0.86	85.73 $\pm$ 0.24
DrFuse	41.70 $\pm$ 1.96	79.18 $\pm$ 1.54
DrFuse w/ HF	48.24 $\pm$ 0.74	83.39$\pm$0.27
DrFuse w/ HF+	48.36$\pm$0.73	83.37 $\pm$ 0.28
DDL-CXR	44.92 $\pm$ 0.68	80.01 $\pm$ 0.74
DDL-CXR w/ HF	44.99 $\pm$ 0.49	80.44 $\pm$ 0.39
DDL-CXR w/ HF+	45.42$\pm$1.01	80.66$\pm$0.30

HazardFlow also improves frozen backbones without retraining their encoders.



current	t2	t3	t4	t5	t6	t7	t8	t9
PC	-0.21	-0.09	-0.07	-0.17	-0.12	-0.41	-0.35	-0.39

Higher learned hazard consistently corresponds to poorer lactate clearance.

Although the base model misranks Patients 2 and 3, the learned hazard assigns a larger magnitude to the positive Patient 3.

THANK YOU!

