



Predicting Adverse Events for Patients with Type-1 Diabetes Via Self-Supervised Learning

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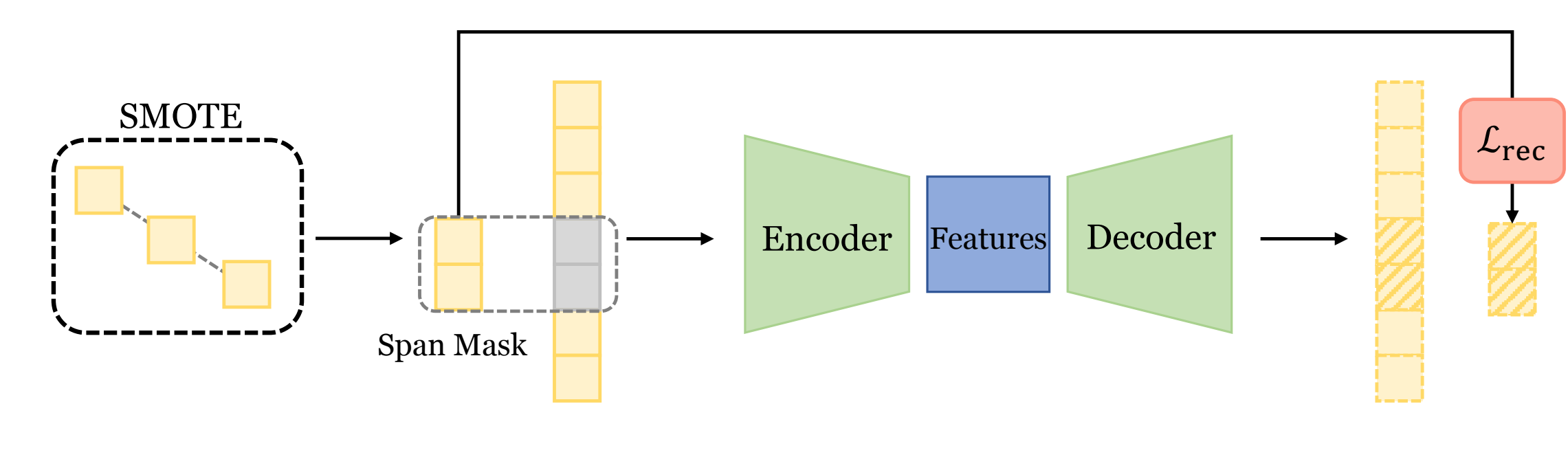
1 Introduction

Predicting blood glucose levels is fundamental for precise primary care of type-1 diabetes (T1D) patients. However, it is challenging to predict glucose levels accurately, not to mention the early alarm of adverse events (**hyperglycemia** and **hypoglycemia**), namely the *minority class*. In this paper, we propose **BG-BERT**, a novel self-supervised learning framework for blood glucose level prediction.

2 Contributions

- Contextual Modeling:** **BG-BERT** effectively models contextual information in blood glucose monitoring data, capturing trends and the influence of observed data on future readings.
- Addressing Limited Data:** **BG-BERT** addresses the limited availability of blood glucose data within adverse events by employing data augmentation techniques and a bias-free training process.
- Improved Performance:** **BG-BERT** outperforms existing models in predicting blood glucose levels within adverse events, achieving higher accuracy.
- Open-source Framework:** **BG-BERT** is an open-source framework available on GitHub: <https://github.com/aiot-lab/BG-BERT>.

1. Masked Self-supervised Learning Phase



2. Glucose Level Predicting Phase

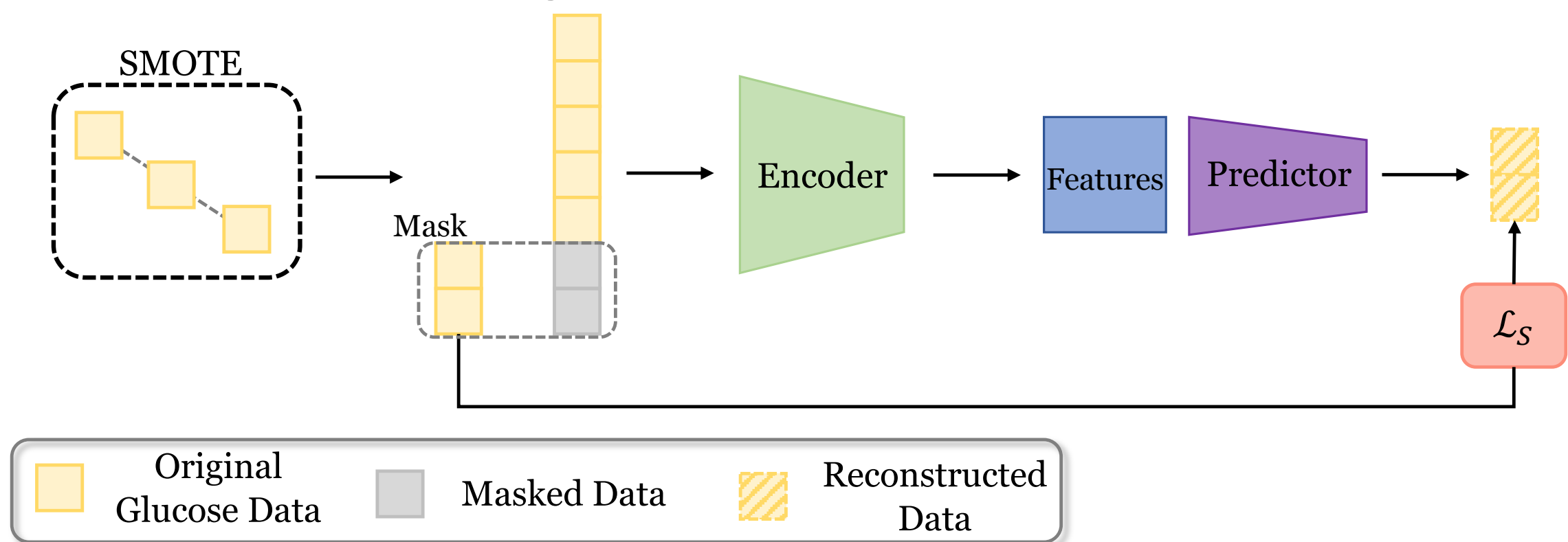


Figure 1: System Workflow

3 Methodology

- Masked Self-supervised Learning Phase:** Automatically learn rich contextual information from glucose readings.
- Glucose Level Prediction Phase:** Forecast glucose levels based on learned representations.

In addition, to address the imbalanced data issue within the range of hyperglycemia and hypoglycemia. We incorporate SMOTE data augmentation and a customized shrinkage loss function.

3.1 SMOTE Data Augmentation

the synthetic instances are generated by combining a specific sample from adverse events with one of its k nearest neighbors. The calculation for generating the synthetic object is as follows:

$$\mathbf{X}_s = \mathbf{X}_i + r \cdot (\mathbf{X}_b - \mathbf{X}_i), \quad (1)$$

where \mathbf{X}_s is the synthetic instance, \mathbf{X}_i is one sample from adverse events, and \mathbf{X}_b denotes the neighbor sample of \mathbf{X}_i .

3.2 Shrinkage Loss

We amplify large loss values and diminish small loss values during back-propagation, implementing the concept of focal loss for regression tasks. The formula of the shrinkage loss is given as:

$$\mathcal{L}_s = \frac{\|\hat{\mathbf{g}} - \mathbf{g}\|^2}{1 + \exp(a \cdot (c - \|\hat{\mathbf{g}} - \mathbf{g}\|^2))}, \quad (2)$$

where $\hat{\mathbf{g}}$ is the golden standard, \mathbf{g} is the predictions. a and c are hyper-parameters of shrinkage loss.

4 Results

We evaluate **BG-BERT** on two benchmark datasets: OhioT1DM and Diatrend, with two SOTA baseline models: DRTF and MT-NB-L (supervised-learning).

Horizon	30 mins							
Dataset	OhioT1DM				Diatrend			
Metric	RMSE (mg/dL)	TG (mins)	Sen Hype (%)	Sen Hypo (%)	RMSE (mg/dL)	TG (mins)	Sen Hype (%)	Sen Hypo (%)
DRTF	18.21	15.54	80.67	53.58	15.23	15.07	80.28	39.12
MT-NB-L	21.50	14.74	61.36	34.05	19.80	13.88	75.16	39.89
w/o aug	14.38	16.19	81.49	64.41	15.01	16.27	79.53	56.69
w/o Ls	13.92	15.72	81.16	70.75	15.13	16.16	80.75	57.42
BG-BERT	14.02	16.56	82.54	73.24	14.85	16.47	81.34	62.27

Table 1: Evaluation Results (30mins). TG: temporal gain, which indicates the amount of average time gained for early detection of a potential adverse event; Sen: sensitivity.

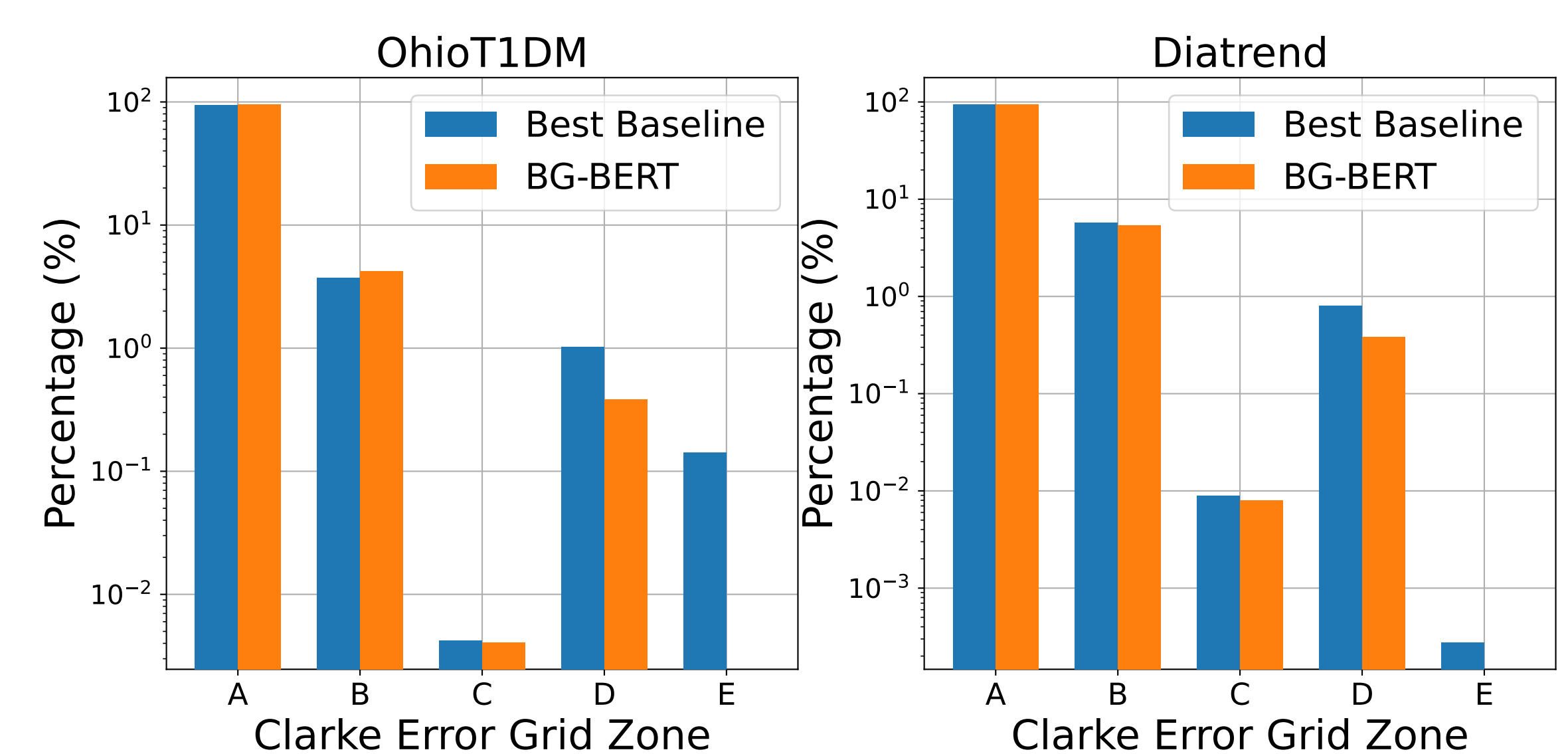


Figure 2: Clarke error grid analysis. (A: medically accurate result, B: medically acceptable, C: unnecessary treatment, D: failure to detect a dangerous condition, E: mistaking adverse events.)

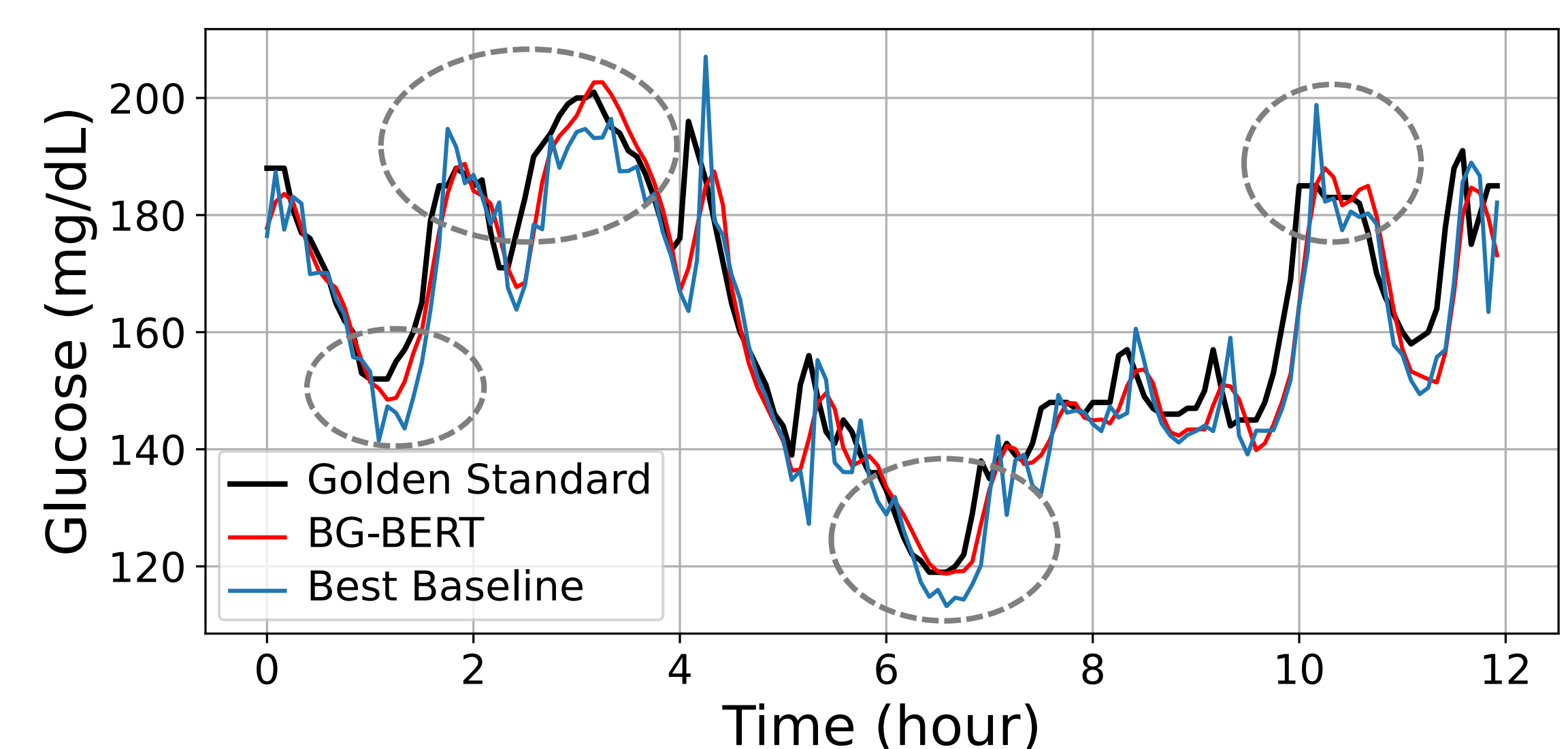


Figure 3: Visualization of half-day glucose prediction. The gray circles highlight the better performance on turning points.