TEXAS A&M UNIVERSITY Statistics

Background

According to the International Diabetes Federation, 463 million adults worldwide have diabetes [1]. Glucose management is a critical component of diabetes care. **Continuous glucose monitoring (CGM)** is a technology that enables patients to track their glucose levels at **short regular intervals**. Coupled with a **predictive model**, CGM can help people with diabetes manage their blood glucose levels either by providing **feedback** or as a part of **artificial pancreas** (AP) that can regulate blood glucose automatically.

Main limitation of the models: poor uncertainty quantification or poor distributional fit.

Data generation: consider a **toy example**, where the **data is generated** following:

$$\mathbf{y} \sim \mathcal{N}(\boldsymbol{\mu}, \Sigma)$$
$$\boldsymbol{\mu}_{1:n} \sim \mathcal{N}(\mathbf{0}, I_n) \qquad \boldsymbol{\mu}_{n:2n} \sim \frac{1}{2} \mathcal{N}(\mathbf{0}, I_n) + \frac{1}{2} \mathcal{N}(\mathbf{1}, I_n)$$

Why is it relevant to the CGM data?

- . Patients typically adhere to daily schedules (periodicity).
- 2. Fixed part corresponds to fixed portions of the schedule.
- 3. Mixture part corresponds to the variable portion of the schedule.

Setup: given past observations, $y_{1:t}$, we want to form a predictive distribution $p(\mathbf{y}_{t:t_l}|\mathbf{y}_{1:t})$. Our training data consists of randomly subsampled chunks: $\{ y_{1:t}, y_{t:t+l} \}.$

We consider 2 models:

- 1. Transformer (DL) trained with $MSE \Leftrightarrow$ additive Gaussian assumption on the noise,
- 2. Transformer (DL) trained with our infinite mixture model **(IMM)** objective.



Figure 1. Top: model trained with MSE, bottom: our model trained with IMM.

Gluformer: Transformer-Based Personalized Glucose Forecasting with Uncertainty Quantification

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We change model loss by conditioning on latent variables z: $oldsymbol{y}_{t:t+l} | oldsymbol{y}_{1:t}, oldsymbol{z} \sim \mathcal{N}(oldsymbol{\hat{\mu}}, \hat{\sigma}^2 I_l) \qquad (oldsymbol{\hat{\mu}}, \hat{\sigma}^2) = f_{ heta}(y_{1:t} | oldsymbol{z}) \qquad oldsymbol{z} \sim p_{oldsymbol{z}}.$ We optimize model parameters θ to minimize negative log-likelihood:

 $heta^{\star} = rg\min\left(-\log\int p(oldsymbol{y}_{t:t+l}|oldsymbol{y}_{1:t},oldsymbol{z})p(oldsymbol{z})doldsymbol{z}
ight).$

During training, we approximate the integral using Monte Carlo, which amount to several stochastic passes through the network.

During inference, we form the predictive distribution as a finite mixture using the stochastic passes:

$$p(\boldsymbol{y}_{t:t+l}|\boldsymbol{y}_{1:t}) \approx \frac{1}{n} \sum_{1}^{n} \mathcal{N}(\boldsymbol{\hat{\mu}}_{i}, \hat{\sigma}_{i}^{2} I_{l}) \qquad (\boldsymbol{\hat{\mu}}_{i}, \hat{\sigma}_{i}^{2}) = f_{\theta}(\boldsymbol{y}_{1:t}|z_{i}) \qquad z_{i} \sim p_{\boldsymbol{z}}$$

Overall, the **computational cost grows linearly** with the number of estimated components.



Figure 2. Predictive distribution for a sample of CGM curves based on **IMM**.

Key takeaway: The effect of unobserved variables on future glucose trajectories presents a challenge to traditional approaches for uncertainty quantification. We propose a novel approach that explicitly incorporates the impact of latent covariates on glucose dynamics, resulting in accurate uncertainty quantification.

We use a publicly available CGM data set [2], which contains information on glucose levels of **38 subjects** tracked continuously throughout multiple disjoint intervals with the measurement frequency of 5 minutes. Similar to the previous studies, we apply the following pre-processing steps:

- 1. Interpolation of small gaps;
- 2. Drop readings where measurements fluctuate by more than 40 mg/dl in 5 minutes;
- 3. Split into train / validation / test in chronological order in 20:1:1 proportion.

Total: 399,302 measurements with 30 uninterrupted sequences per subject.

Model	Full	Event	Нуро	Hyper	Likelihood
ARIMA	9.85 / 17.65	8.91 / 19.86	19.94 / 14.53	8.51 / 22.17	-14.93
RF:Rec	9.04 / 17.15	8.97 / 20.36	18.84 / 12.43	8.68 / 23.41	-14.58
RF:MO	10.22 / 18.27	8.61 / 19.90	21.64 / 17.36	7.99 / 21.58	-15.34
PolySeqMO	8.55 / 15.68	8.27 / 18.81	22.86 / 21.87	6.77 / 18.30	-15.61
RNN	8.17 / 15.67	8.29 / 19.37	18.72 / 16.26	6.99 / 19.22	-13.50
TFT	7.80 / 15.78	8.03 / 18.23	16.23 / 14.62	6.87 / 18.98	_
Our: IMM	7.78 / 15.40	7.89 / 17.85	15.75 / 14.03	7.08 / 19.58	-2.67
Our: Gaussian	7.82 / 14.73	7.93 / 18.62	18.28 / 13.55	6.81 / 18.94	-12.82

Table 1. APE/RMSE for 60-minute prediction window (Full), hypoglycemia, hyperglycemia, and event (hypo-or hyperglycemia), and model log-likelihood on test data.

Further considerations that we have addressed to make the model and fitting more stable:

- and add it to the model input.
- gradient descent are taken from different patients or different CGM curves for a patient.

Computational cost of training the model on the data set:

- 1. GPU: NVIDIA RTX 2080 with 12GB
- 2. Each epoch of SGD: \approx 10 minutes
- 3. Total time for 100 epochs: 16 hours

Future work:

- 2. Learn number of *active* mixture components;
- 3. Incorporate different data sources;
- 4. Conditional prediction based on future activity.
- Federation, 2021.
- strategies for T1DM patients. 33(6):e2833.
- pages 11106-11115, 2021.



Results

Personalization. The model is trained simultaneously on the data from all patients. We learn an embedding of patient-specific demographic information

2. Long-term temporal dependence. To be able to accommodate larger context window size, i.e., accommodate large t, we use an adaptation of the Transformer that down-samples the input in-between attention layers [4]. Within-batch dependence. During training, samples for each mini-batch of

1. Learn latent distribution p_z to optimally explore different regimes;

[1] International Diabetes Federation. International Diabetes Federation Diabetes Atlas. International Diabetes

[2] Ian Fox, Lynn Ang, Mamta Jaiswal, Rodica Pop-Busui, and Jenna Wiens. Deep Multi-Output Forecasting: Learning to Accurately Predict Blood Glucose Trajectories. ACM SIGKDD, pages 1387–1395, July 2018. [3] Silvia Oviedo, Josep Vehí, Remei Calm, and Joaquim Armengol. A review of personalized blood glucose prediction

[4] Haoyi Zhou, Shanghang Zhang, Jieqi Peng, Shuai Zhang, Jianxin Li, Hui Xiong, and Wancai Zhang. Informer: Beyond efficient transformer for long sequence time-series forecasting. In Proceedings of the AAAI, volume 35,