



# Blood Glucose Forecasting based on CGM data

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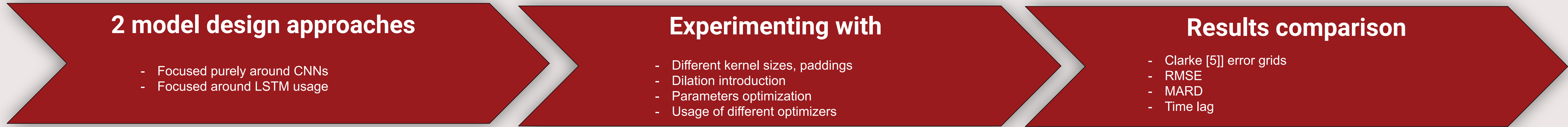
## Introduction

This project is about forecasting blood glucose (BG) based on continuous (taken every 5 minutes) glucose monitoring (CGM) [1], exogenous insulin and carbohydrate (CHO) intake from meals. This 30 minute forecast is shown to the user who can then more easily regulate their BG levels and prevent hypo- (below 70 mg/dL) or hyper- (above 180 mg/dL) glycemic states.

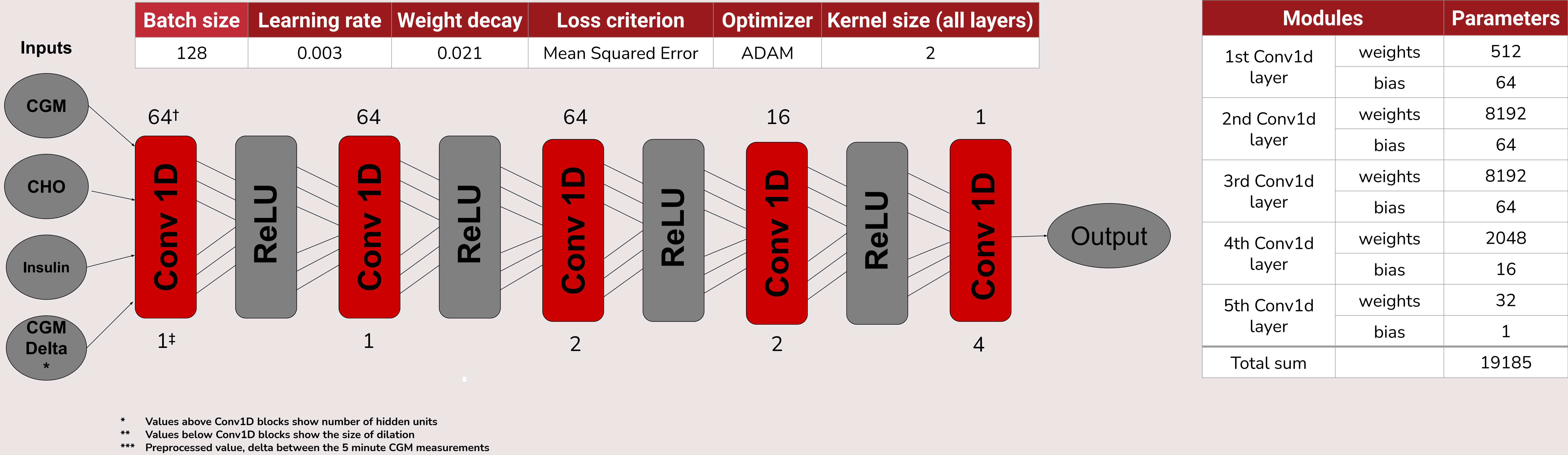
## Data & Tools

The team was given the *in vivo* Ohio T1D dataset [2] and a deep learning framework from Hedia which preprocesses the data, trains the model and evaluates the results. Part of the framework was a simple model with five dilated convolutional layers. The attempted models were inspired by this simple model (which is in turn based on [3]) and by an architecture presented in [4].

## Workflow



## Model



## Results

Model	RMSE	MARD	Time lag (min)	A+B	C+D+E
Hedia - Best	20.027	9.97%	20.45	99.57%	0.43%
Ours - Best Personalized	20.401	10.57%	20.41	97.85%	2.15%
Ours - Best Generalized	20.492	10.66%	20.49	97.83%	2.17%
GluNet* [3]	19.28	8.73%	8.03	-	-
CRNNfGP* [4]	21.07	-	-	-	-

\* These results are not directly comparable to other results in the table as they are taken from the papers where it is not described in detail how were these values obtained.

Fig. 1: The results comparison between Hedia's top model, our model and research papers

The models that are proposed are two, a generalized and a personalized. The generalized one is a model that is used for all case samples. On the other hand, the personalized one it's tuned for each individual case. Taking a look at the results, as expected the model that is tuned for each patient performs better than the generalized one.



Fig. 2: The plot presenting the actual (light blue) and predicted (dark blue) BGL measurements

For this application it is very important that the model can predict accurately the valleys and the hills because those are the danger zones for the patient. As shown from the results the personalized model outperforms the generalized one across the board. That would indicate that each patient has his own unique patterns and characteristics. A real world approach to this problem would be to use the generalized model on a patient for a period of time and after optimize it for that individual. A flexible strategy like this will ensure that any changes to behaviour/pattern in the future can be compensated and the model can be re-optimized for that individual.

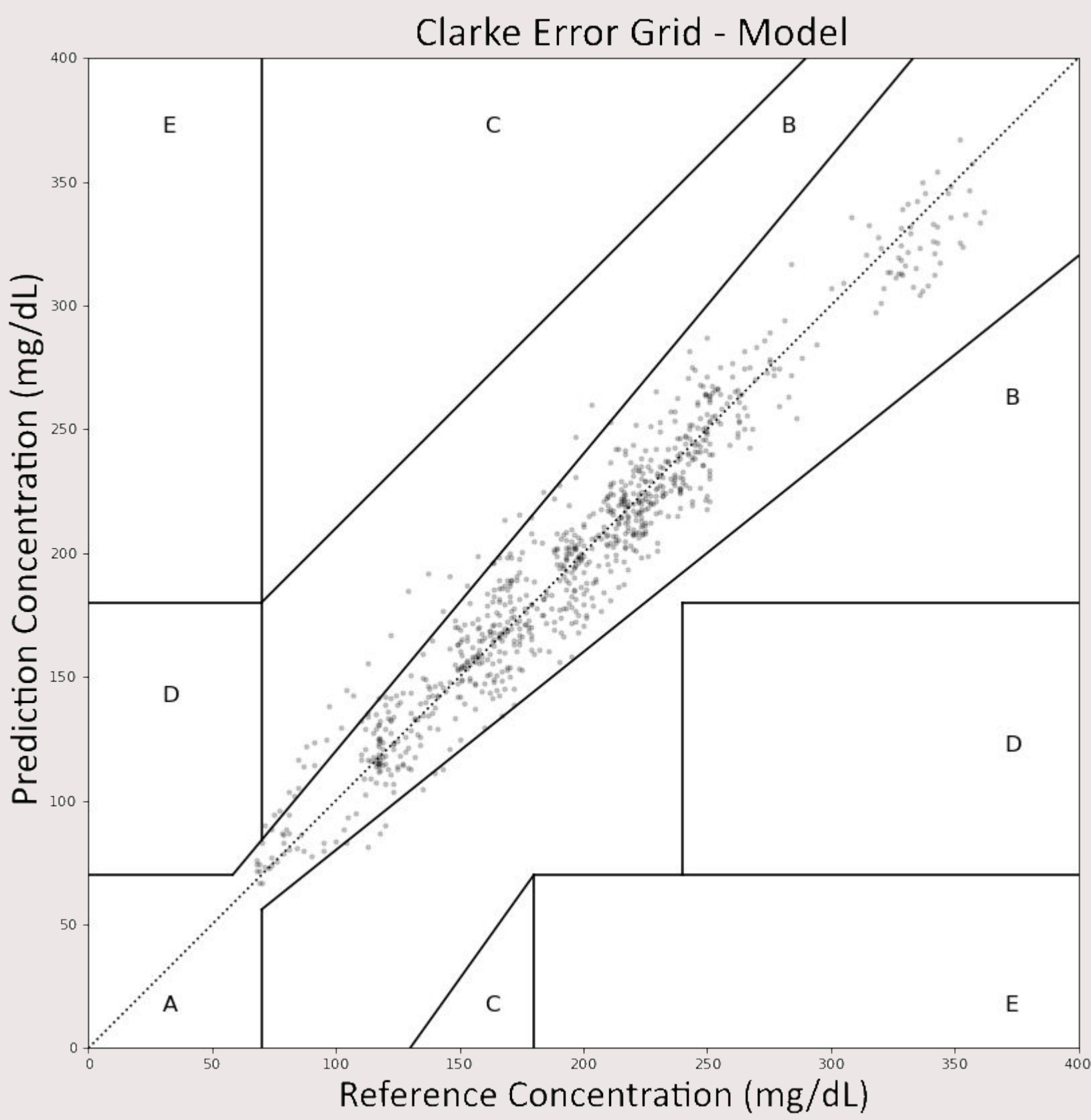


Fig. 3: Clarke Error Grid for our results

Both models performed similarly with an average A+B value close to 98%. A+B ranges in the Park graphs are the ones that are clinically acceptable, with A being very good and B being good. Both models have 86% in the A region which is acceptable. For comparison the model from Hedia has 89%.

## References

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