



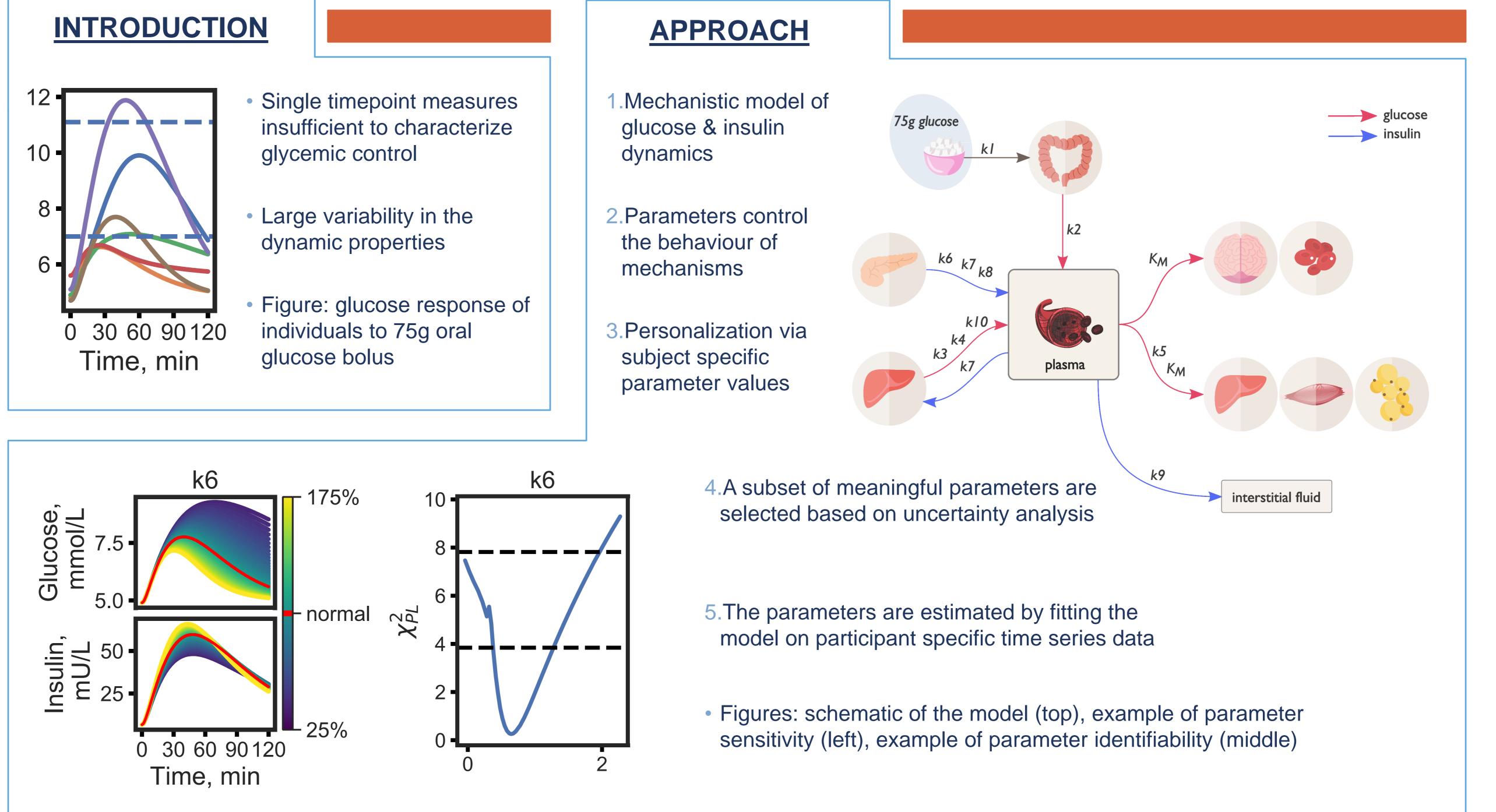
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INTRODUCTION

The evolution from healthy towards control glucose prediabetes and type 2 diabetes occurs continuously over years, characterized by deteriorations in the plasma glucose and insulin concentrations [1]. However, the heterogeneity large in the pathophysiology of type 2 diabetes and individual's glycemic control it difficult to categorize make participants into prevention target groups and necessitates the mechanistic characterization of the glucose and insulin dynamics on a personalized level. Ordinary differential equation (ODE) based mathematical models have been developed to describe the plasma glucose response in humans to a single dose of glucose These models [2,3]. are mathematical abstractions of the real biological system and they provide quantitative information on the interactions, dynamics and regulation of specific components of the system. Quantifying the using а modelling response approach facilitates the mechanistic understanding of the underlying physiology as well as the development of decision support systems for preventing diabetes. We aim to characterize an individual's glucose response to an oral glucose tolerance test (OGTT) using a personalized ODE model that describes glucose and insulin dynamics in the postprandial state [4].

Characterizing postprandial glucose responses in individuals using a computational modelling approach

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RESULTS & DISCUSSION

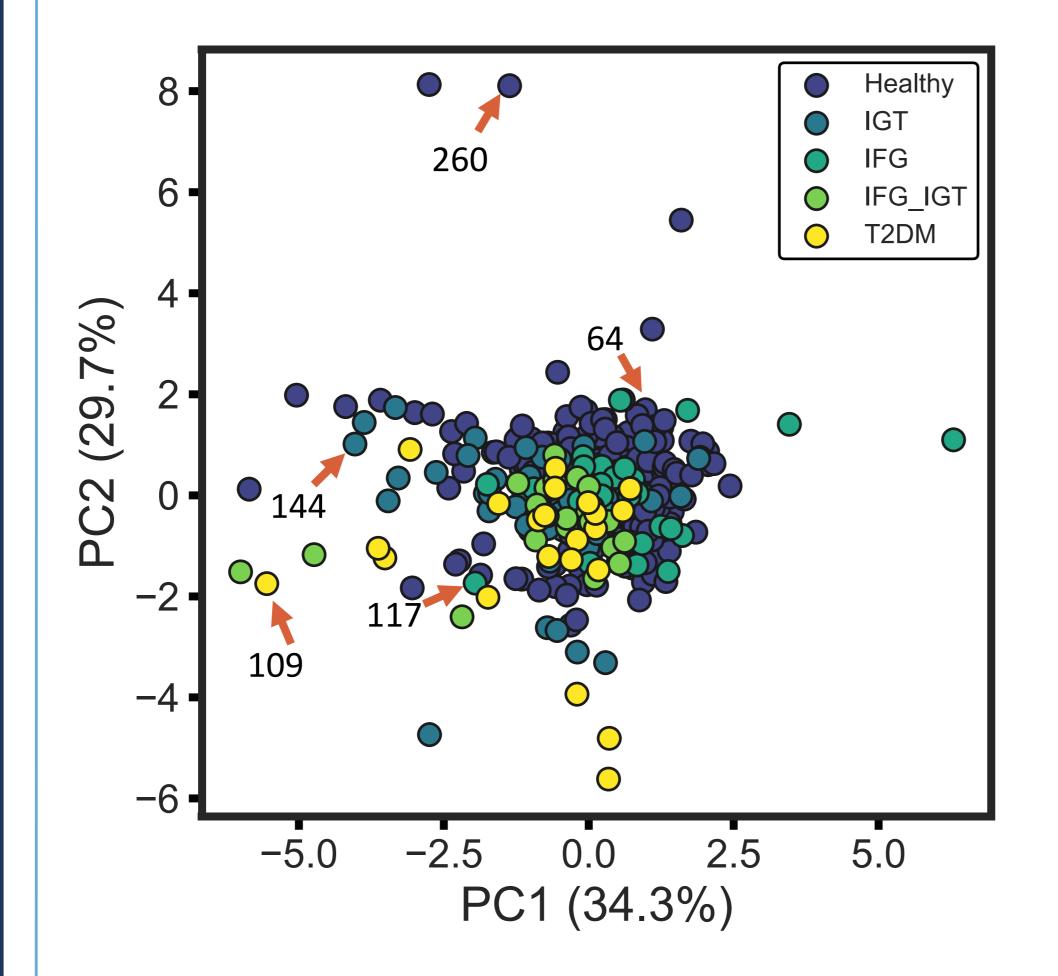
APPROACH

Data from the Diogenes [5] dietary intervention study, including a 5 time point OGTT was used in modelling the glucose responses.

In order to estimate participant's glucose and insulin dynamics we fit an ODE based mechanistic model describing glucose and insulin dynamics on the OGTT challenge test data. The model was adapted from [4] to allow personalization by selecting a subset of the parameters to be estimated. The candidate models then were carefully curated for best fitting while also maintaining model certainty (sensitivity and identifiability) in its parameter values. Finally, parameters of the model estimated were on individual's data. The model was implemented in MATLAB 2018b.

RESULTS & DISCUSSION

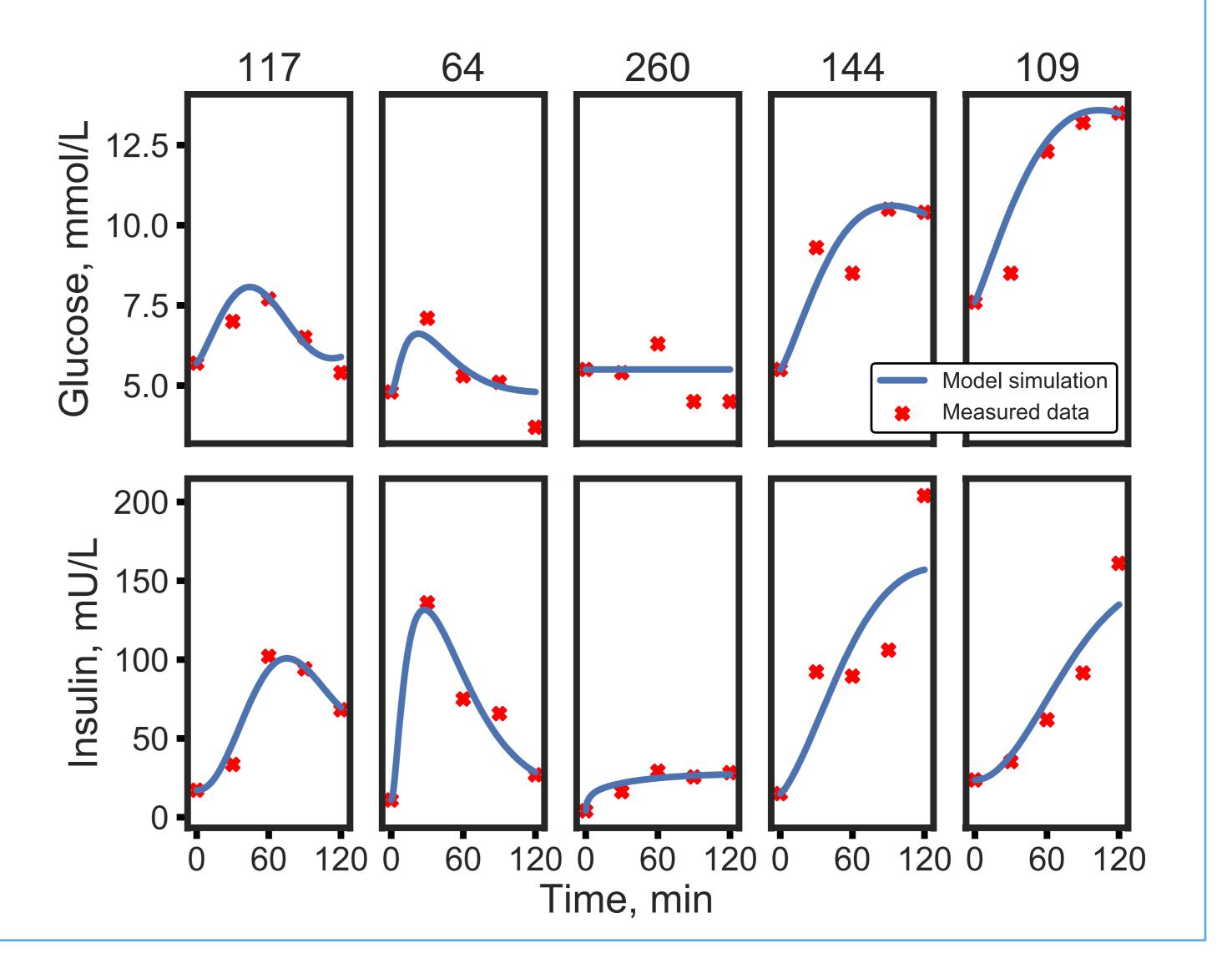
The screening of candidate models resulted in a model containing four sensitive and identifiable parameters. The model was successfully individualized by fitting it to subject specific time resulting data. The series personalized models were capable of capturing a wide range of glucose and insulin dynamics including normal, prediabetic and type 2 diabetic responses. The participant's responses can be characterized by their place in parameter space. This the approach also allows observation of the continuous trajectory between healthy diabetic states, to contributing to the mechanistic understanding of changing between states.



• The personalized models may assist in the understanding of the differences between metabolic states and the trajectories between them

 A model with 4 parameters was selected for parameter estimation after screening of candidate models

• The personalized models were capable of describing a wide range of responses including healthy, prediabetic and type 2 diabetic but also responses of intermediate states as well



Figures: Personalized models in the reduced parameter space, colored by classification of diagnosis by the American Diabetes Association criteria (left), examples of personalized model simulations (right)

REFERENCES

[1] Bergman, M. (2013). Pathophysiology of prediabetes and treatment implications for the prevention of type 2 diabetes mellitus. Endocrine, 43(3), 504–513. [2] Bergman, R. N. et al. (1979). Quantitative estimation of insulin sensitivity. American Journal of Physiology-Endocrinology and Metabolism, 236(6), E667. [3] Dalla Man, C. et al. (2007). Meal Simulation Model of the Glucose-Insulin System. IEEE Transactions on Biomedical Engineering, 54(10), 1740–1749. [4] Maas, A. et al. (2015). A Physiology-Based Model Describing Heterogeneity in Glucose Metabolism: The Core of the Eindhoven Diabetes.... Journal of Diabetes Science and Technology, 9(2), 282–292. [5] Larsen, T. M. et al. (2010). The Diet, Obesity and Genes (Diogenes) Dietary Study in eight European countries – a comprehensive design for long-term intervention. Obesity Reviews, 11(1), 76–91.

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