Limits of a Glucose-Insulin Model to Investigate Intestinal Absorption in Type 2 Diabetes IARIA BIOTECHNO 2021

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Danilo Dursoniah

PhD student, 2nd year, BioComputing research team, CRIStAL Lab. **Education:**

- BSc Biotechnology, Grenoble Alpes Univ. (2014 2017);
- MSc Bioinformatics, Univ. of Paris-Saclay (2017 2019);
- PhD thesis, Univ. of Lille (2019 ongoing).

Thesis topic:

Computational Models of Intestinal Glucose Absorption for Diabetes Prediction.

Directors: Cedric Lhoussaine (Prof.), François Pattou (Prof.) **Supervisor:** Maxime Folchette (Assoc. Prof.) **Interests:**Formal modeling - Clinical study - Medical biotechnology **Contact:** danilo.dursoniah@univ-lille.fr

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Type 2 Diabetes (T2D)

Definition

Chronic metabolic disease characterized by:

- lack of insulin secretion → hypoinsulinemia;
- unability of the body to use insulin → hyperglycemia;
- perturbated glucose and insulin homeostasis;
- high prevalence: 400 millions affected \rightarrow 90% of diabetics;
- limited therapeutic solutions.

Challenges

- Type 2 diabetes is a major public health issue.
- Identifying new therapeutic targets.

Type 2 Diabetes (T2D)

Hypothetical Causes

- High-carbonhydrate diet.
- Sedentary lifestyle.
- \blacksquare Excess of intestinal glucose absorption \rightarrow High rate of glucose appearance.

Problematic

How much impact does glucose intestinal absorption have on T2D?

 \rightarrow Hypothesis based on clinical observations.

- How can a standard glucose-insulin model take into account such hypothesis?
 - \rightarrow Limits of the standard model.

Roux-en-Y Gastric Bypass (RYGB)

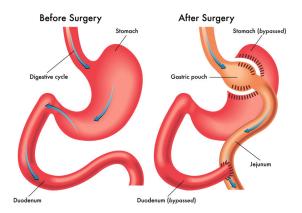


Figure: Roux-en-Y Gastric Bypass (RYGB). UCLA Health: http://surgery.ucla.edu/bariatrics-gastric-bypass

RYGB leads to weight loss and glucose-insulin homeostasis restoration.

Clinical Observations After a Meal

Dataset

Clinical datasets collected and provided by Inserm collaborators, from obese patients, before and after surgery.

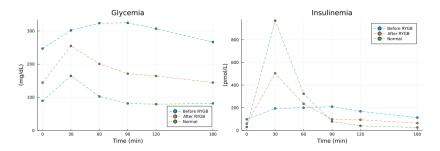


Figure: Average glycemia and insulinemia on 180 minutes, after a meal.

 \implies After surgery: good glucose and insulin homeostasis restoration.

Reference Model

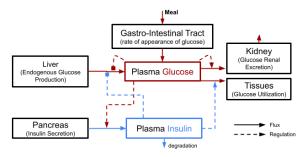
Standard, highly cited model from:

Meal Simulation Model of the Glucose-Insulin System, Dalla Man et al., 2007.

System of ordinary differential equations (ODEs):

- 12 dependent ODEs, functions of time;
- 36 pairs of parameters values:
 - non-diabetics;
 - diabetics.
- $\blacksquare \mbox{ No original dataset provided by the modellers} \rightarrow \mbox{ low reproductibility } \rightarrow \mbox{ model validation from our dataset.}$
- Equations distributed in **7 biological modules** corresponding to biological functions: gastro-intestinal tract, liver, pancreas, etc.

Graph representation of the reference model



 $\mathsf{RYGB} \text{ simulation } \iff \mathsf{Modifying \ gastro-intestinal \ parameters \ values}$

- 1. How much does each biological module contribute to glucose-insulin homeostasis restoration?
- 2. Can the model predict our own dataset of diabetic patients?
- 3. Can the model predict glucose-insulin homeostasis restoration after RYGB?

Parameters estimation (optimization problem):

- Estimated parameter values: all or parts;
- Fitted variables: Glycemia and Insulinemia.
- Objective function: maximum likelihood or MSE etc.

initial parameters $\stackrel{\text{estimation}}{\longrightarrow}$ inferred parameters $\stackrel{\text{model}}{\longrightarrow}$ inferred variables.

1. How much does each biological module contribute to glucose-insulin homeostasis restoration?

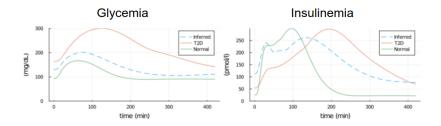
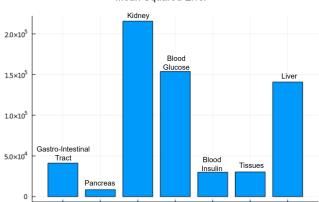


Figure: Inferred glycemia and insulinemia for gastro-intestinal tract

Initial parameters: diabetic parameters of [DallMan2007]; Fitted variables: non-diabetic glycemia and insulinemia of [DallMan2007]; Inferred parameters: only gastro-intestinal tract Errors computed between healthy objective model and best estimated fit. 1. How much does each biological module contribute to glucose-insulin homeostasis restoration?

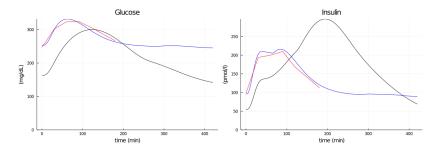


Mean Squared Error

Contribution of each functional module to glucose homeostasis (low error = good contributor)

Parameters Estimation Results

2. Can the model predict our own dataset of diabetic patients?

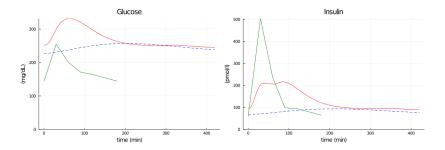


Initial parameters: diabetic parameters from [DallaMan2007]; **Fitted variables**: own dataset of pre-surgery glycemia and insulinemia; **Inferred parameters**: all parameters (36).

 \Longrightarrow good fitting despite very different diabetic population than [DallaMan2007].

Parameters Estimation Results

3. Can the model predict glucose-insulin homeostasis restoration after RYGB?



Initial parameters: previous pre-surgery estimated parameters; **Fitted variables**: own dataset of post-surgery glycemia and insulinemia; **Inferred parameters**: only gastro-intestinal tract (\implies simulation of RYGB).

$$\implies$$
 bad fitting.

- 1. From *Inserm* collaboration: animal datasets (Mini Pigs). Exploring populations dataset, **before** and **after** different experiments:
 - * Pancreatectomy
 - * Small bowel resection
 - * Metabolic modulation for obesity

More homogenous values. Better reproductibility for model validation.

- 2. New variables to fit
 - * D-xylose \rightarrow rate of glucose appearance
 - * C-peptide \rightarrow insulin secretion

Would increase the parameters identifiability.

- Struggle to fit our datasets.
- Shifting model validation criteria: from accurate fitting to qualitative validation.
- Building up a new model based on experimental data: more qualitative and explicative for the glucose intestinal absorption on type 2 diabetes prediction.
- Integrating more equations in a new model to explain intestinal glucose absorption.

Questions?