



Towards a Digital Twin for Personalized Diabetes Prevention: The PRAESIIDIUM Project

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PRAESIIDIUM - PHYSICS INFORMED MACHINE LEARNING-BASED PREDICTION AND REVERSION OF IMPAIRED FASTING GLUCOSE MANAGEMENT

Project Idea

To develop a tool aimed at providing a **real-time prediction of the prediabetic risk** of an individual.

The prediction algorithm will be based on a "*physics-informed machine learning*" approach: a rich dataset of real-life data, obtained from already existing previous and a new clinical trial with continuous data ingestion through wearable sensors, will be **combined with mathematical models and eXplainable AI (XAI) techniques**, to overcome the limits of "*black-box*" ML approaches, while improving the prediction performances and reducing the computational time of the risk calculation based on the simulations of the mathematical model.

The final algorithm will be implemented in **a web-based platform**, where medical doctors and patients can inject data from several sources (acquisition form connected sensors and manual insertion) and obtain a **real time analysis of the risk** to develop the prediabetic condition over time.

PRAESIIDIUM will develop easy-to-use, reliable, efficient and innovative tools (e.g., web/ mobile app, mathematical models, wearable sensors) for the **personalized management and real-time prediction of the prediabetic risk**. This goal will result in benefits for affected citizens, their families, hospitals, different industries, scientific community, governments/ policy-makers, and environment.

https://praesiidium.spindoxlabs.com/









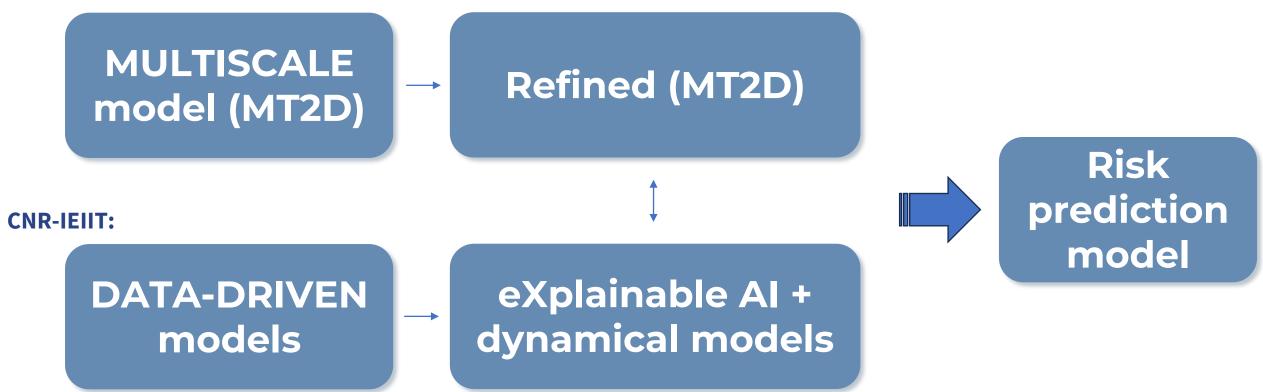






PERSONALIZED RISK PREDICTION AND PREVENTION

CNR-IAC:





Whole-body, multi-scale model of metabolic homeostasis during exercise An agent-based simulator of the human immune system

F. Castiglione, F. Celada. Immune system modelling and simulations. Boca Raton: CRC Press; 2015

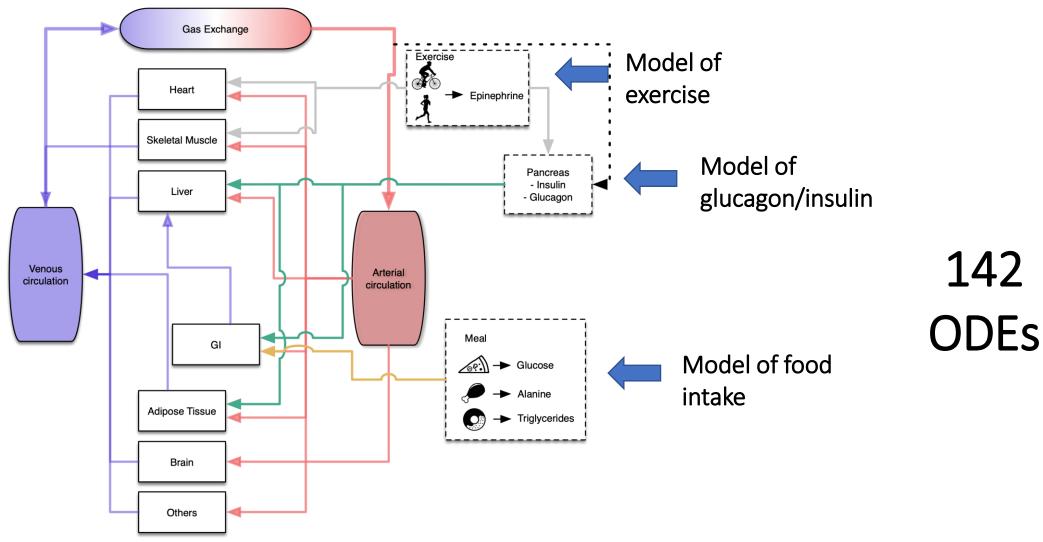
J. Kim, G. M. Saidel, and M. E. Cabrera. Multi-scale computational model of fuel homeostasis during exercise: effect of hormonal control. Annals of Biomedical Engineering, 2007, 35: 69-90.

M.C. Palumbo, M. Morettini, P. Tieri, F. Diele, M. Sacchetti, and F. Castiglione. Personalizing physical exercise in a computational model of fuel homeostasis. *PLoS Computational Biology*, 2018, 14: e1006073.

M.C. Palumbo, A.A. de Graaf, M. Morettini, P. Tieri, S. Krishnan, and F. Castiglione (2023). A computational model of the effects of macronutrients absorption and physical exercise on hormonal regulation and metabolic homeostasis. *Computers in Biology and Medicine*, 107158.



MULTISCALE MODEL MT2D – THE METABOLIC MODEL





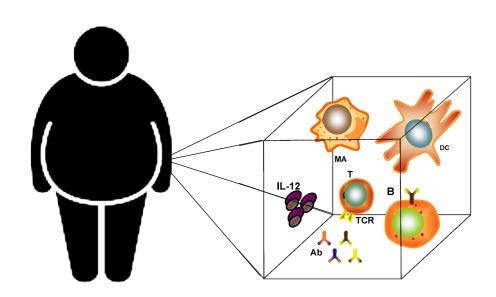
Personalization of model predictions

Input Parameter	Meaning
S	Sex (male/female)
А	Age in years
BM	Body mass in kg
Н	Height in m
VO _{2max}	Maximal oxygen uptake in ml·kg ⁻¹ ·min ⁻¹
Τ _v	Target value of exercise intensity in percentage of VO _{2max}
Fitness status	Cardio-respiratory fitness classification from "poor" to "superior"
t _{ex,start} , t _{ex,end}	Start/end of the exercise session in minutes
C _{fast,glu}	Fasting glucose level in mmol/l



MULTISCALE MODEL MT2D – THE IMMUNE SYSTEM

C-IMMSIM: an agent-based model of the immune system



Immune cells are individually represented. They follow behavioral rules coding for known immunological mechanisms

B (B-1, B-2), PLB, TH (Th1, Th2, Th17, Treg), CTL, Treg, NK, MA, DC

EP, ADIP

lgM, lgG (lgG1, lgG2), lC

IL-2, Danger, IL-12, IFN-g, IL-4, TNF-a, TGF-b, IL-10, IL-6, IL-10, IL-18, IL-23, IFN-b, IL-1b

LPS, Leptine

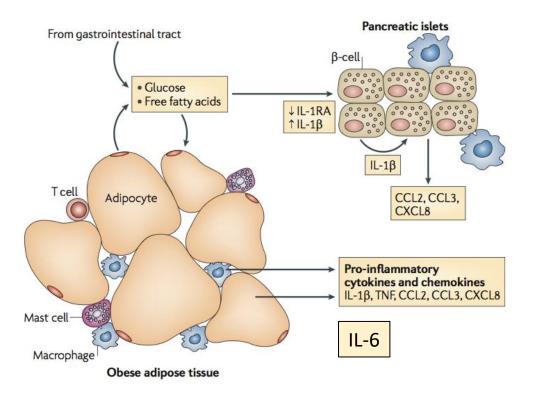
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MULTISCALE MODEL MT2D – THE IMMUNE SYSTEM

Immunology of adipose tissue inflammation

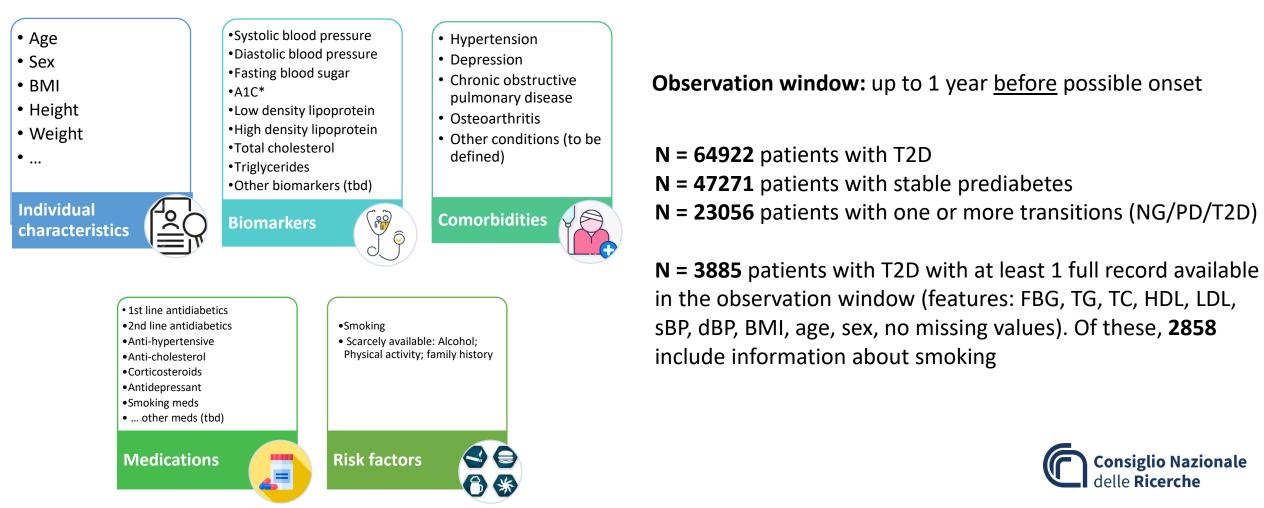
- Adipose Tissue (AT) largely composed of adipocytes + pre-adipocytes, endothelial cells and immune cells
- In obesity AT undergoes expansion and remodeling (↑pro-, ↓anti-inflammatory cytokines and infiltration of immune cells)
- This obesity-associated pattern of events in AT → chronic state of low-grade inflammation





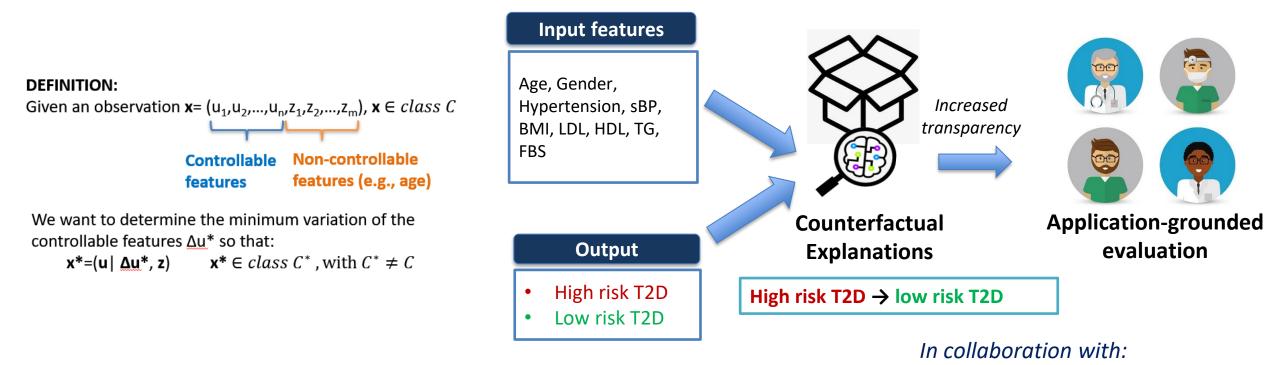
DATA-DRIVEN MODELS: RETROSPECTIVE DATASET

Canadian Primary Care Sentinel Surveillance Network (CPCSSN) EMR database (<u>http://cpcssn.ca/</u>)



DATA-DRIVEN MODELS: COUNTERFACTUAL EXPLANATIONS

GOAL: find minimum yet meaningful changes in biomarkers that may reduce the risk of developing T2D on an individual basis



Toronto

Metropolitan

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University

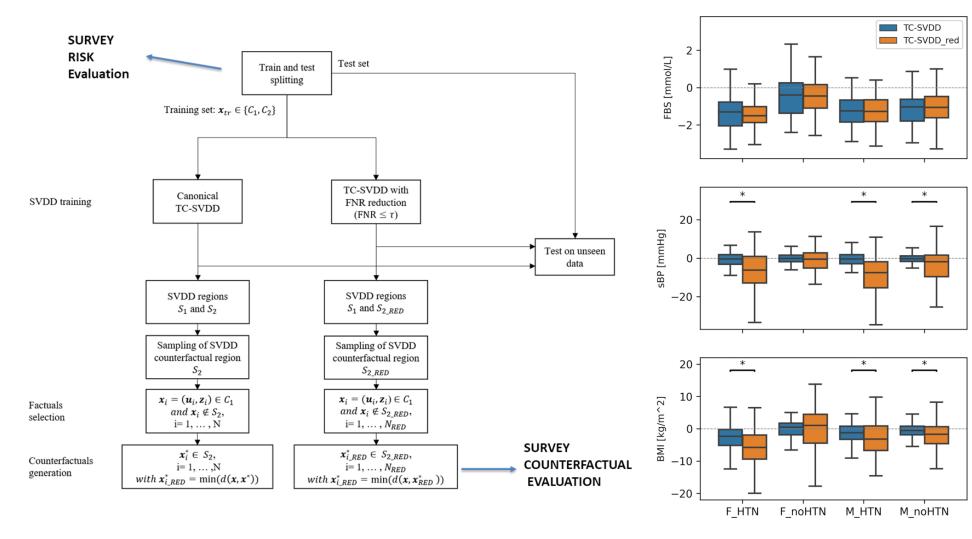
UNIVERSITY OF

TORONTO

- Lenatti M., Carlevaro A., Keshavjee K., Guergachi A., Mongelli M., Paglialonga A. "*A novel method to derive personalized minimum viable recommendations for type 2 diabetes prevention based on counterfactual explanations*," PLOS ONE, vol. 17(11): e0272825, 2022. https://doi.org/10.1371/journal.pone.0272825

- Carlevaro A., Lenatti M., Paglialonga A., Mongelli M., "*Multi-class counterfactual explanations using support vector data description," IEEE Transactions on Artificial Intellig*ence, 2° revision submitted on Oct 03, 2023. Preprint available at: https://doi.org/10.36227/techrxiv.22221007.v1

DATA-DRIVEN MODELS: COUNTERFACTUAL EXPLANATIONS





- Personalized definition of controllable features (e.g., if drug.resistant)
- Investigation of causal models
- Characterization of changes as a function of risk



DATA-DRIVEN MODELS: PATIENT TRAJECTORIES

GOAL: monitor the risk of developing T2D based on routinely collected biomarkers up to 8 years before the onset

19 Inputs (t1):	7 Outputs (t2):
 Medications (categorical) Comorbidities (categorical) Biomarkers BMI Age 	• Biomarkers

Algorithm: Multi-input multi-output gaussian process regression model (adapted from: Chen, Z., B. Wang, and A. N. Gorban. "Multivariate gaussian and student-t process regression for multi-output prediction, 2017")

• 5 fold cross validation on training data (80%) : isotropic kernel choice • Hyperparameter optimization $\begin{cases}
28 \text{ output covariance matrix hyperparameters} \\
+ \\
2 \text{ isotropic input kernel hyperparameters} \\
+ \\
1 \text{ noise variance hyperparameter}
\end{cases}$

• Computation of predictive distribution on testing data (20%)

- Simeone D., Lenatti M., Lagoa C., Keshavjee K, Guergachi A., Dabbene F., Paglialonga A., "*Multi-Input Multi-Output Dynamic Modelling of Type 2 Diabetes Progression*," Studies in Health Technology and Informatics, Proceedings of the European Federation of Medical Informatics Special Topic Conference (EFMI STC 2023), Oct 25-27, 2023, Torino, Italy. In press.

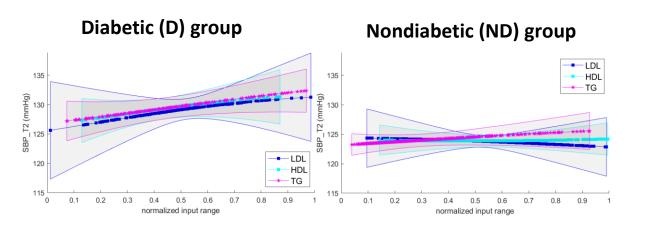
In collaboration with:



Toronto Metropolitan University

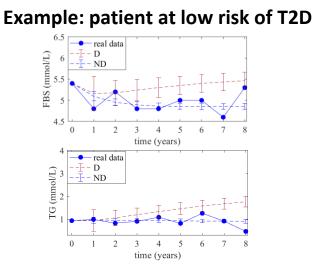


DATA-DRIVEN MODELS: PATIENT TRAJECTORIES

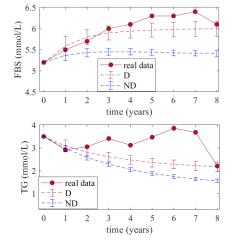


Example of findings from D and ND groups and from sub-groups:

- Stronger positive association between lipids and sBP for D patients compared to ND
- Stronger positive association between age and systolic blood pressure for females compared to males
- Stronger positive association between LDL and sBP for D smokers compared to D nonsmokers
- Stronger negative association between LDL and HDL for D smokers compared to D nonsmokers, for both males and females



Example: patient at high risk of T2D



NEXT STEPS

- Analysis and validation of models from subgroups (vs. sex, risk factors, medications)
- Analysis of different time windows w.r.t. possible onset
- Development of several MIMO models using sub-sets of input/outputs
- Modeling intervention (e.g. physical activity)







Thank you for your attention

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