

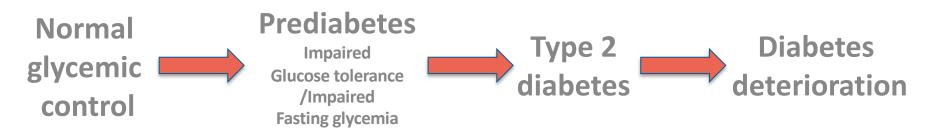
The RHAPSODY project: Biomarkers in type 2 diabetes

Bernard Thorens

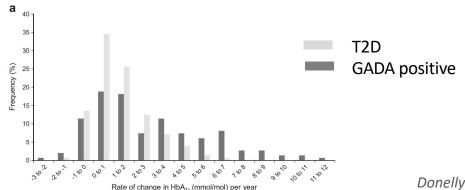
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Development of type 2 diabetes: A stepwise process



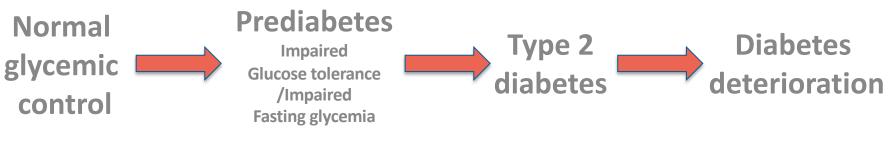
Rate of glycemic deterioration in T2D patients from the GoDarts cohort (HbA_{1C}/year)

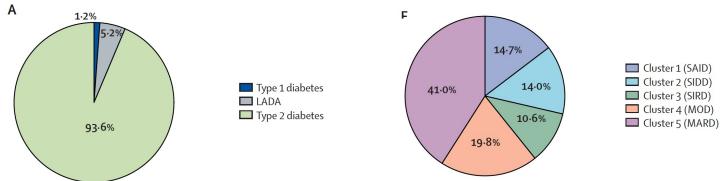


Donelly LA, et al., Diabetologia, 2018



Development of type 2 diabetes: A stepwise process





Ahlqvist E et al., The Lancet Diabetes Endocrinology, 2018

Identification of biomarkers



Specific Questions:

- Can we identify biomarkers that are prognostic of T2D susceptibility and T2D deterioration?
- Can such biomarkers predict dysfunctions in ß-cells or in insulin target tissues?
- Can we identify the tissues and metabolic pathways controlling the production of these biomarkers?

Biomarkers

 Molecules that can be objectively measured and evaluated as an indicator of normal biological or pathogenic processes, as well as pharmacological response to a therapeutic intervention

http://www.fda.gov/drugs/scienceresearch/researchareas/pharmacogenetics/ucm083378.htm

Plasma Biomarkers

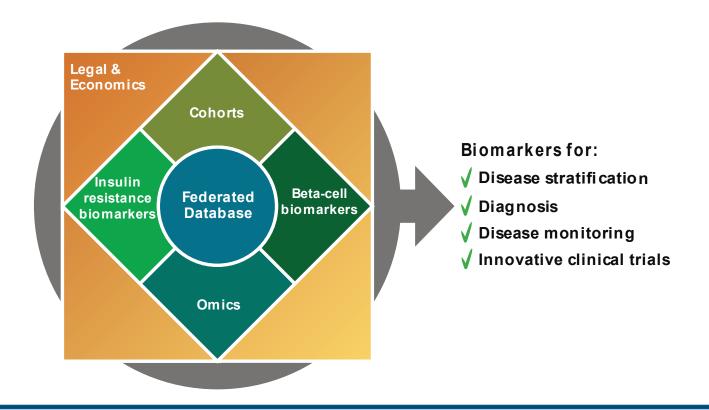
- Lipids
- Polar metabolites
- Proteins
- RNAs

Plasma Biomarkers

Requirements for biomarker discovery:

- Technological platforms for quantitative measurements of molecular species
- Patient cohorts with detailed phenotyping and biosamples (plasma)
- Preclinical models of prediabetes/diabetes
- Federated database for all data storage and analysis

RHAPSODY Plasma Biomarkers Strategy

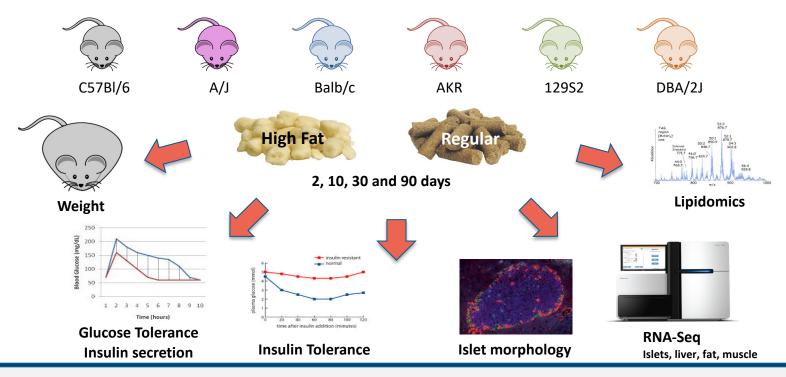


Use of preclinical models to identify biomarkers of diabetes susceptibility and of ß-cell dysfunction

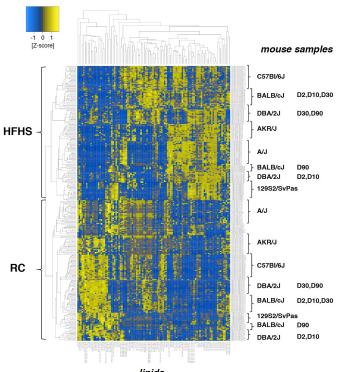
- Use of preclinical models to identify candidate biomarkers for the progression to type 2 diabetes and validation in human cohorts
- Use of preclinical models to identify plasma biomarkers predictive of ß-cell function and to identify the metabolic pathways involved in biomarker production

Use of preclinical models to identify biomarkers of diabetes susceptibility and of ß-cell dysfunction

6 mouse strains



Integrative analysis of plasma lipidomics with mouse physiological traits



Lipid profiles across all conditions (6 strains, 4 time points, 2 diets)



Correlation, hierarchical clustering

Modules of lipids with similar profiles

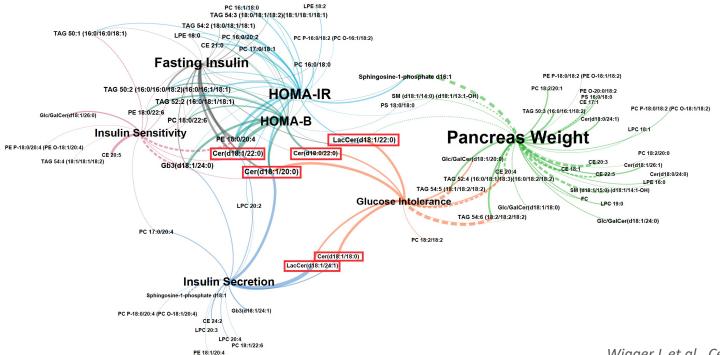


Correlation of module profiles with physiological traits



Investigation of trait-associated lipid modules

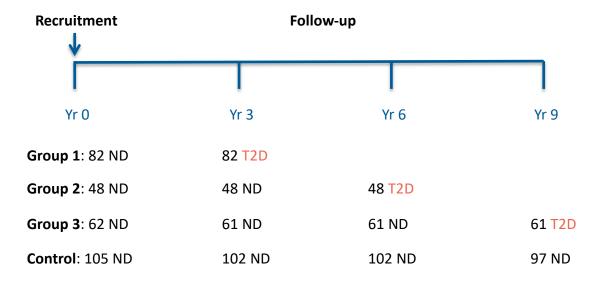
Ceramides are correlated to glucose intolerance and insulin sensitivity in metabolically challenged mouse strains



Wigger L et al., Cell Reports, 2017

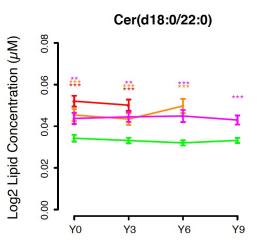
Analysis of ceramides in the plasma of individuals from the DESIR cohort

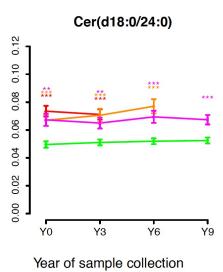
- Prospective cohort of > 5000 people followed for > 9 years
- Summary of plasma analysis

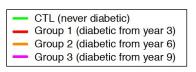


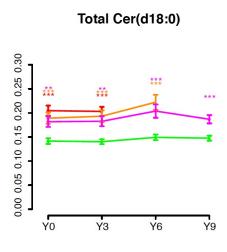
Dihydroceramides are enriched in the plasma of T2D patients up to 9 years before diagnostic

The DESIR cohort









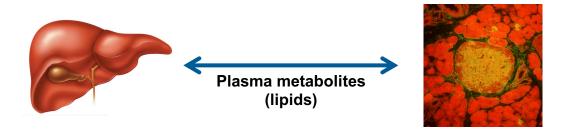
Wigger L et al., Cell Reports, 2017

Use of preclinical models to identify biomarkers of diabetes susceptibility and of ß-cell dysfunction

- Use of preclinical models to identify candidate biomarkers for the progression to type 2 diabetes and validation in human cohorts
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A systems biology analysis of the crosstalk between liver and pancreatic ß-cell function through plasma lipids

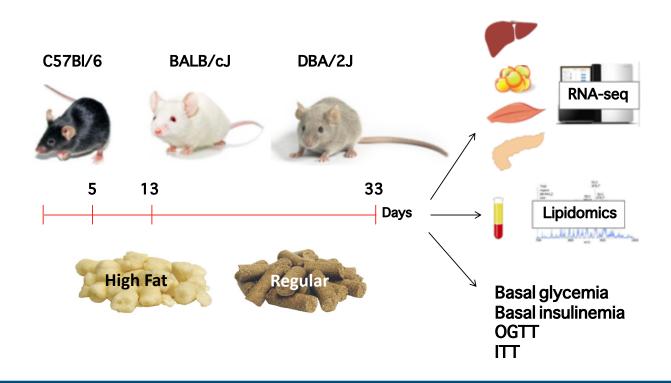
Progressive loss of ß-cell secretion capacity over time



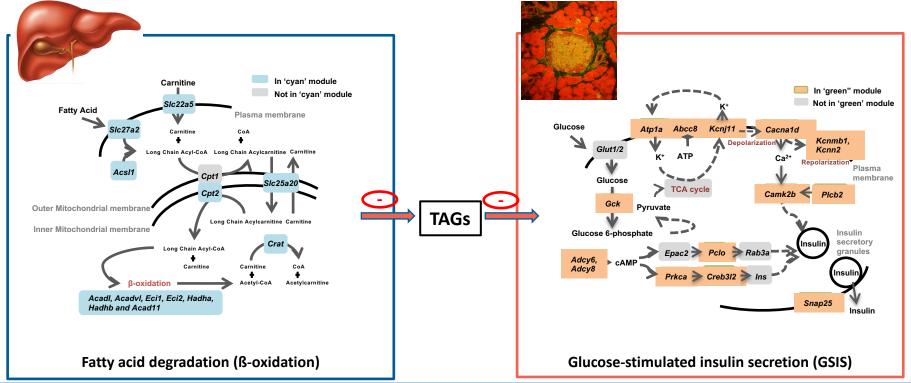
Questions:

- Can we find plasma biomarkers predictive of ß-cell function?
- Can we identify the tissue and metabolic pathway that produce the biomarker?
- Can we establish a link between a tissue metabolic pathway plasma biomarkers – and ß-cell function?

Search for plasma lipids as potential biomarkers: experimental design



Plasma Triglycerides (TAGs) correlate with ß-cell insulin secretion genes and liver fatty acid degradation pathway

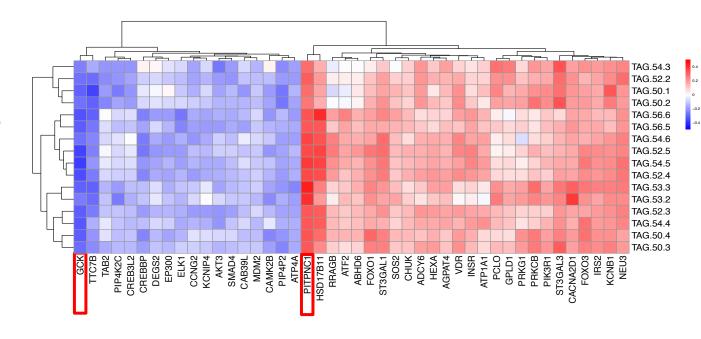




IN HUMANS: Same correlation between plasma TAGs and insulin secretion genes as in mice

- Islets from partially pancreatectomized patients
- Plasma lipids from the same patients

Wigger L. (...) Solimena, M., Nat. Metabolism, 2021



→ Identification of PITPNC1 as a novel regulator of insulin secretion



Conclusions

- The use of preclinical models allowed to identify plasma biomarkers for type 2 diabetes susceptibility
- Such biomarkers were found to be prognostic biomarkers also in humans
- These biomarkers could be demonstrated to correlate with the function of pancreatic ß-cells
- Comparative analysis in mice and humans of the correlation between plasma TAGs and islet gene expression allowed to characterise a so far unknown regulator of insulin secretion

RHAPSODY biomarker identification in humans (See presentation by R. Slieker)

- Biomarker discovery was assayed for:
 - Proteomics
 - Lipids
 - Polar metabolites
- Phase 1: Discovery of biomarkers of T2D progression using samples from the three RHAPSODY discovery cohorts (NT = 9900)
- Phase 2: Replication of biomarkers of T2D progression (NT= 4000)
- Phase 3: Establishment of a biomarker shortlist

Biomarker shortlist includes the same lipids identified in preclinical models

Lipidomics:

- TAG class (namely 50.1.0, 46.1.0, 46.2.0, 48.1.0, 51.1.0, 48.2.0, 48.3.0 and 49.1.0)
- Sphingomyelin (SM 42.2.2)



Next steps

Evaluate the utility of the biomarkers in a **prospective clinical trial** to assess their utility for:

- Stratification of diabetic patients to more precisely identify high-risk subjects at baseline more likely to respond to a specific intervention
- Monitoring of diabetes progression to improve understanding of the course of the disease, or specific symptoms of the disease





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