



# A new diabetes patient sub-classification: What's next?

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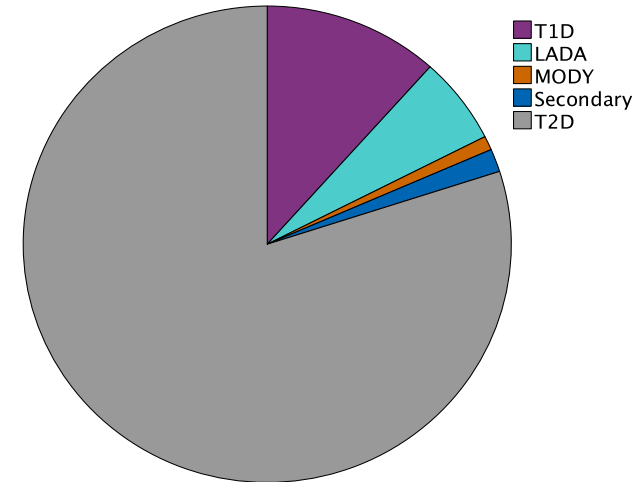
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# Current diabetes classification

- **Type 1 diabetes** – autoimmune diabetes in the young (<35 years)
- **LADA** – latent autoimmune diabetes of the adult (>35 years)
- **Monogenic diabetes** – rare mutations (MODY, neonatal)
- **Secondary**
- **Type 2 diabetes** – everyone else



# Is this not enough?

- Individuals with diabetes, especially type 2 diabetes, are **very different**
  - Risk of complications
  - Need for treatment
  - Response to diabetes medication
- Can we divide diabetes patients into **smaller more homogeneous groups** that are clinically useful for **predicting progression** and **need for medication**?



# A new diabetes sub-classification

March 2018, *Lancet Diabetes and Endocrinology*

## Novel subgroups of adult-onset diabetes and their association with outcomes: a data-driven cluster analysis of six variables



Emma Ahlqvist, Petter Storm, Annemari Käräjämäki\*, Mats Martinell\*, Mozghan Dorkhan, Annelie Carlsson, Petter Vikman, Rashmi B Prasad, Dina Mansour Aly, Peter Almgren, Ylva Wessman, Nael Shaat, Peter Spéjel, Hindrik Mulder, Eero Lindholm, Olle Melander, Ola Hansson, Ulf Malmqvist, Åke Lernmark, Kaj Lahti, Tom Forsén, Tiinamajja Tuomi, Anders H Rosengren, Leif Groop

### Summary

**Background** Diabetes is presently classified into two main forms, type 1 and type 2 diabetes, but type 2 diabetes in particular is highly heterogeneous. A refined classification could provide a powerful tool to individualise treatment regimens and identify individuals with increased risk of complications at diagnosis.

*Lancet Diabetes Endocrinol* 2018

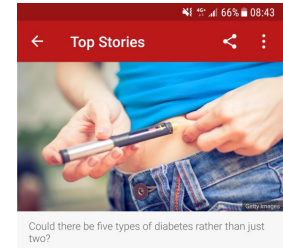
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<http://dx.doi.org/10.1016/>



Five categories for adult diabetes, not just type 1 and type 2, study shows



**Diabetes is actually five separate diseases, research suggests**

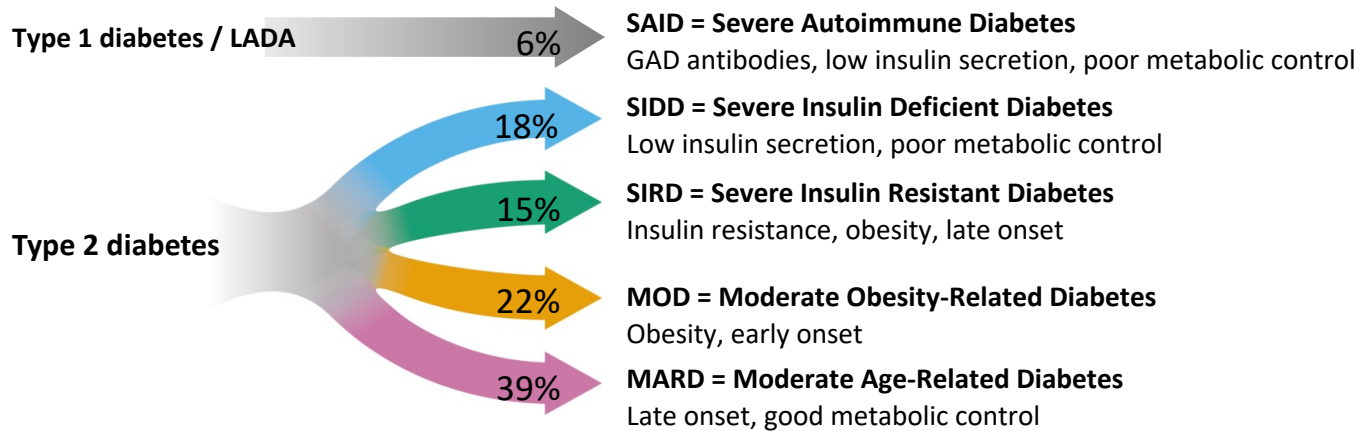
By James Gallagher

Health and science correspondent, BBC News  
6 hours ago | [Health](#)

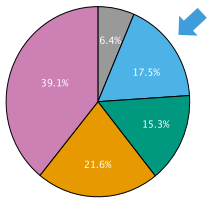
Scientists say diabetes is five separate diseases, and treatment could be tailored to each form.

# Novel subtypes

- We can reproducibly divide patients into **five subgroups** with **different characteristics and progression**

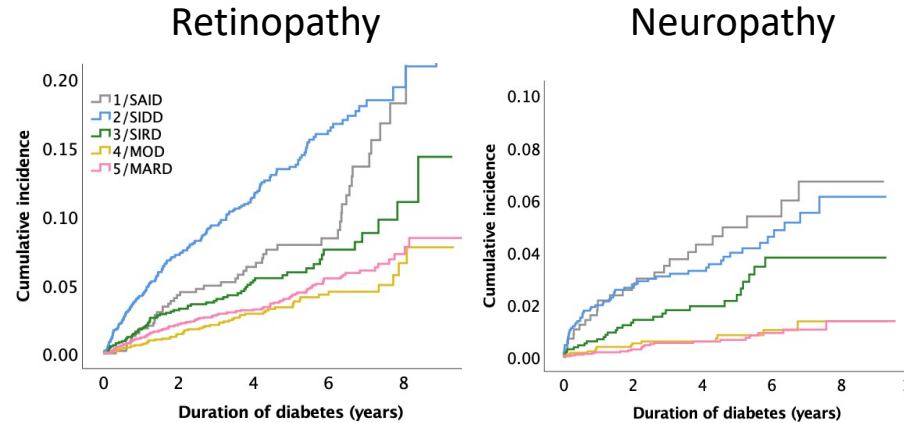


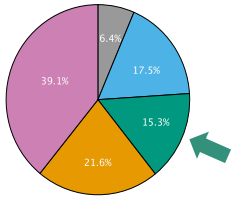
- This clustering approach has been **replicated in numerous cohorts**



# SIDD - Severe insulin-deficient diabetes

- 17.5% of patients
- Low insulin secretion
- High HbA1c at diagnosis
- Relatively early onset (mean 57 years)
- Receive the most treatment but still the last to reach treatment goals
- Increased risk of retinopathy and neuropathy

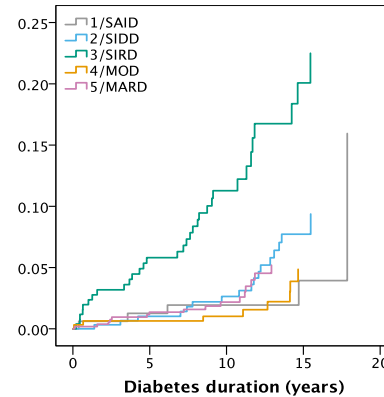




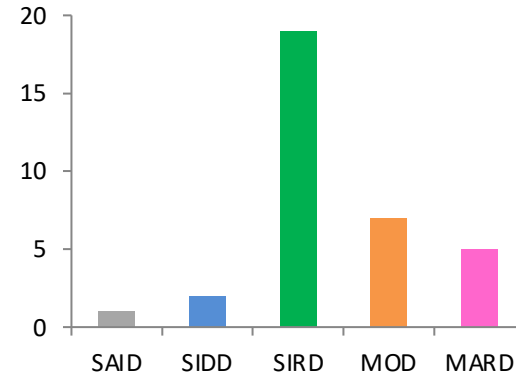
# SIRD - Severe insulin-resistant diabetes

- 15.3% of patients
- Very insulin resistant
- Obese (mean BMI 34)
- Late onset (mean 65 years)
- Low HbA1c
- Higher risk of kidney complications
- Higher prevalence of non-alcoholic fatty liver disease

End-stage renal disease



Hepatocellular lipid content



# What is next?

Are the subtypes different diseases?

When can they be implemented in the clinic?



- Clinical characterisation
- Replication in other ethnicities
- Genetics
- Epigenetics
- Response to medication
- Biomarkers



# Replication in other ethnicities

- The same subtypes have been identified in cohorts from all over the world
- Most find the same five subtypes
- Some have identified additional subtypes e.g. in India
- Comparisons are complicated by use of slightly different methods and cluster variables

# Genetic risk score (GRS) analysis

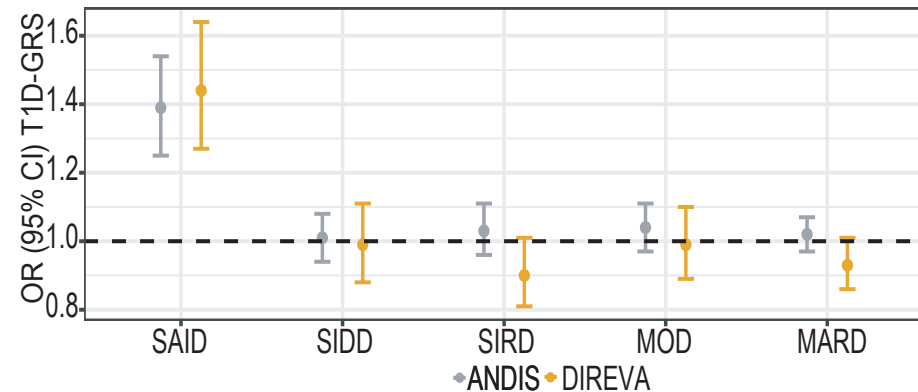


Dina  
Mansour Aly

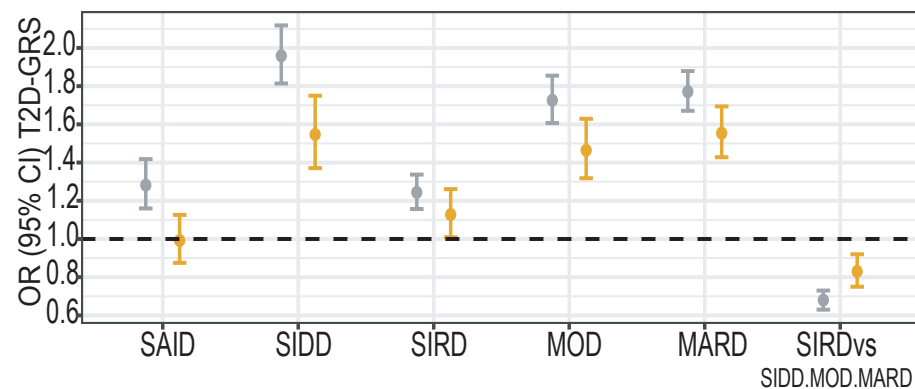


Om Prakash  
Dwivedi

## T1D GRS

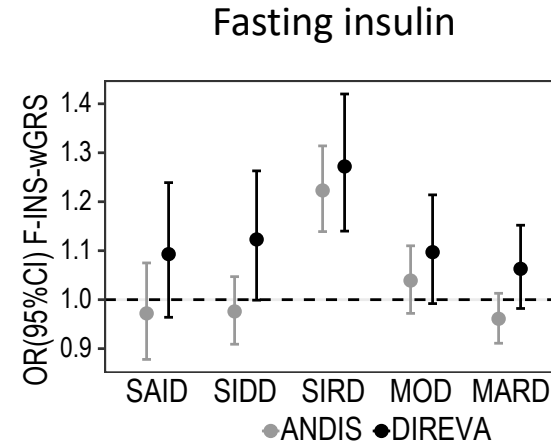
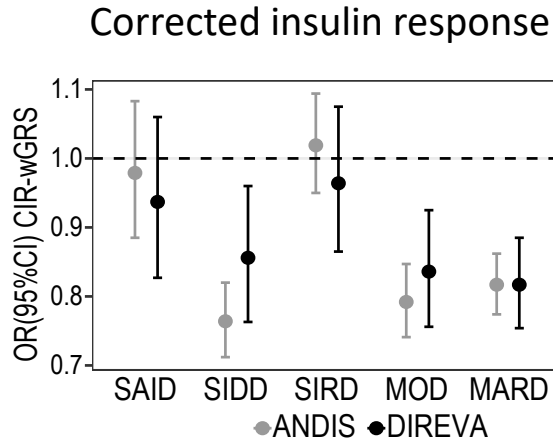


## T2D GRS



Mansour Aly et al, MedRxiv, 2020

# GRS for insulin secretion and fasting insulin



- SIRD is associated with fasting insulin but not insulin secretion

*Mansour Aly et al, MedRxiv, 2020*

# Subtype specific genetic risk variants

	ANDIS		DIREVA		META-ANALYSIS	
	OR	p	OR	P	OR	P
SAID	1.03	0.65	0.94	0.45	0.99	0.85
SIDD	1.08	0.18	1.10	0.95	1.05	0.28
SIRD	1.13	0.01	1.06	0.47	1.11	0.02
<b>MOD</b>	<b>1.35</b>	<b>1.3x10<sup>-9</sup></b>	<b>1.14</b>	<b>0.04</b>	<b>1.27</b>	<b>4.3x10<sup>-9</sup></b>
MARD	1.10	0.01	1.03	0.55	1.07	0.03

- Genetic variants in the LRMDA gene increase the risk of developing MOD but not other subtypes

*Mansour Aly et al, MedRxiv, 2020*



# Response to treatment

- SIDD would likely benefit from early treatment with insulin or insulin secretagogues
- SIRD would likely benefit from insulin sensitizers
- Needs to be proven before clinical implementation
- **Randomized trial** – SIDD and SIRD ongoing in ANDIS
- **Analysis of existing trials** in collaboration with AstraZeneca and Sanofi



# Clinical implementation

Single-patient analysis | [Multiple-patient analysis](#) | [For the clinician](#)

## Single-patient analysis

Please fill in all fields in the form below to analyze your patient.

ID:

Sex:  Male  Female

GADA-positive?  No  Yes

Age at diagnosis:

BMI at diagnosis:

HbA1c at diagnosis(mmol/mol):

Calculate HOMA2IR/HOMA2B

HOMA2IR at diagnosis:

HOMA2B at diagnosis:

## Report for ID Patient4

class: **2/SIDD**, score = 1  
second best match = 3/SIRD, score = 0

density

value

group  
2/SIDD  
Overall

- Information about subtype will soon be returned to the treating doctor for newly registered ANDIS participants
- Treatment cannot be recommended yet but SIDD and SIRD could benefit from screening for complications

# Conclusions

- Diabetes patients can be reproducibly subdivided into **five subtypes**
- The insulin deficient and insulin resistant subtypes have the **highest risk of complications**
- Better science today – better treatment in the near future!
- Learn more on the **RHAPSODY outcomes webpages**:  
<https://imi-rhapsody.eu/outcomes/>



# Thank you for your attention!

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