

19TH ANNUAL

**WORLD CONGRESS  
INSULIN RESISTANCE DIABETES  
& CARDIOVASCULAR DISEASE**

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*Exploring New Frontiers in Metabolism - Tomorrow's Clinical Science Today*



# Has the Beta Cell Been Forgotten? 2 December 2021

## Gene Mutations Play a Dominant Role in $\beta$ -cell Failure in Type 2 Diabetes

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# Dualities of Interest

## Advisory Board, Consulting and Lectures

Bayer

Merck

Boehringer Ingelheim

Novo Nordisk

Casma Therapeutics

Pfizer

Eli Lilly

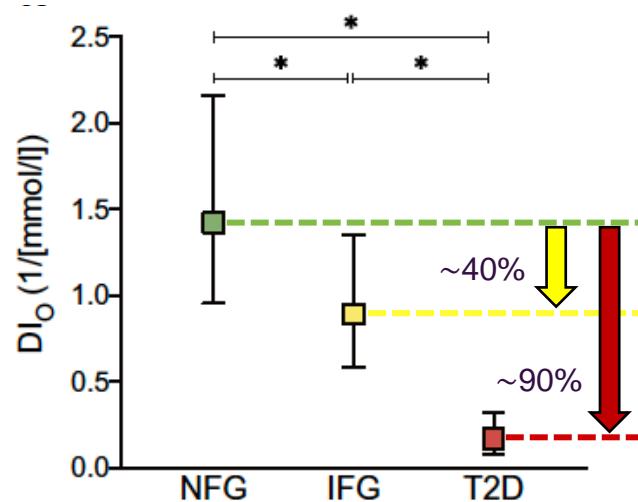
Third Rock Ventures

Intarcia

# Outline

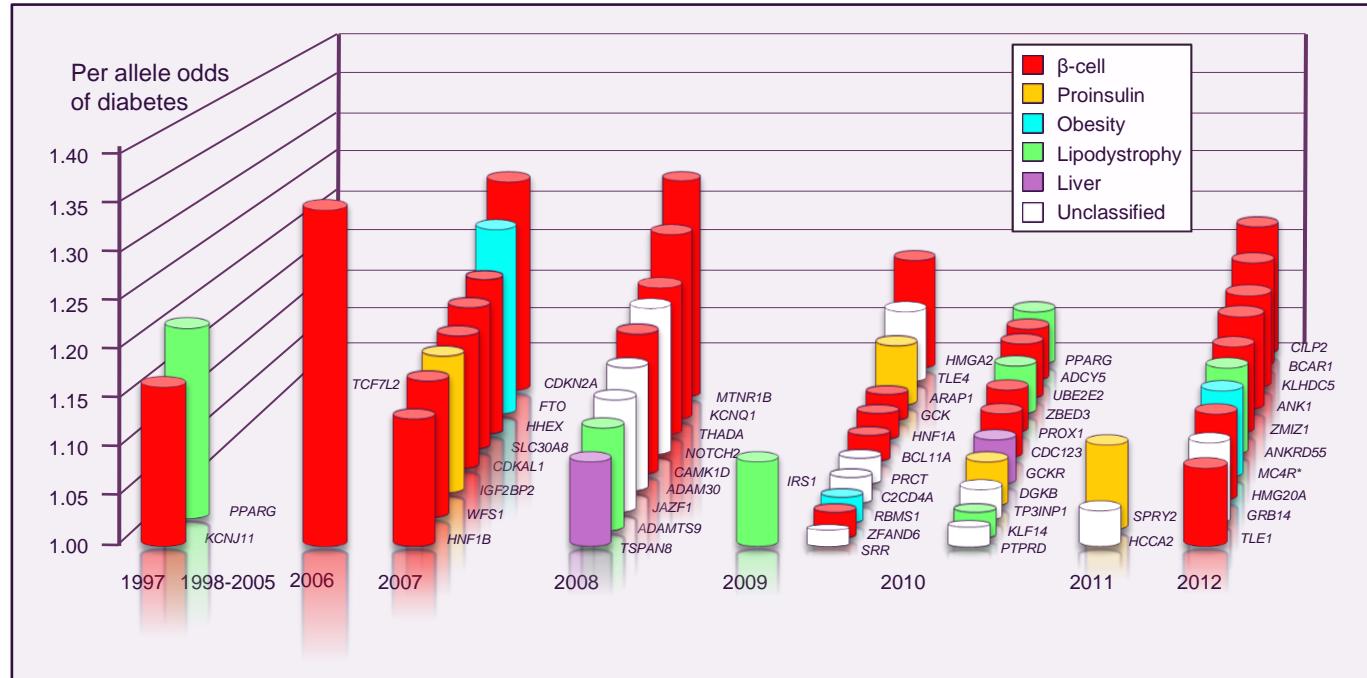
1. Phenotypic characteristics and genetics of type 2 diabetes in adults.
2. Genetics of type 2 diabetes in youth.

# Relationship of Glucose Tolerance and Fasting Glucose with $\beta$ -cell Function



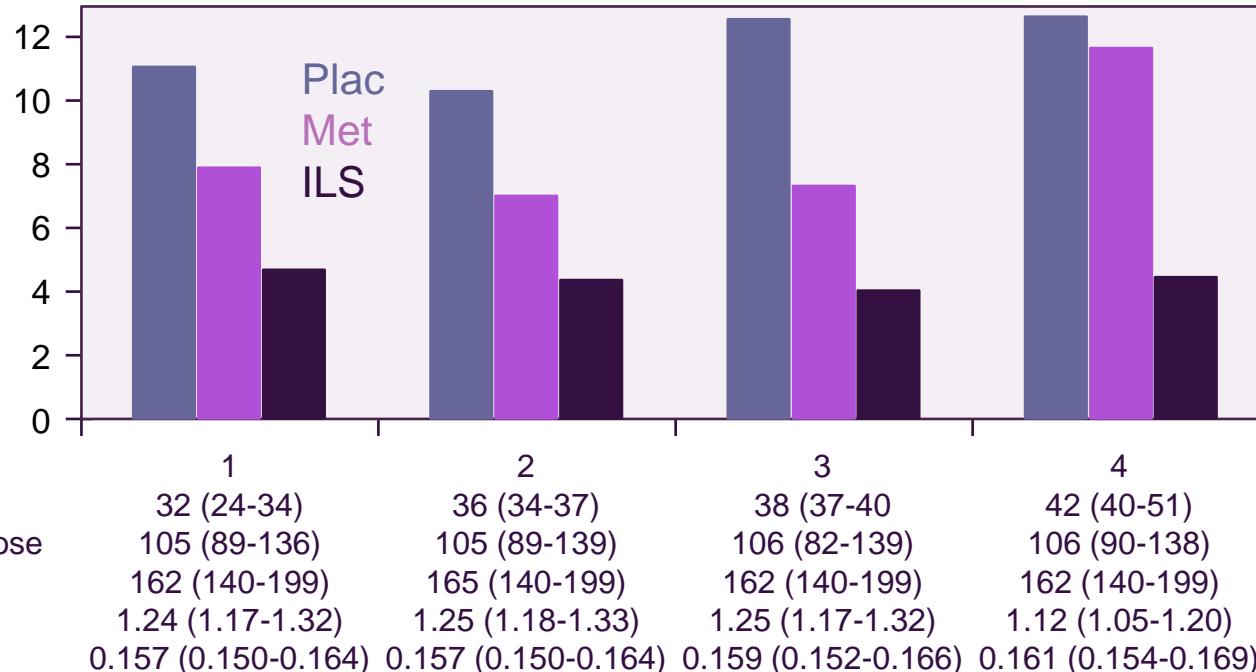
Number of individuals: 374 168 71

# Type 2 Diabetes Genes Associate Most Commonly with $\beta$ -cell Dysfunction



Adapted from Franks PW et al: Diabetes Care 36:1413-1421; 2013, Simons-Bik 59:293-301; 2010, Dimas AS et al: Diabetes 63:2158-2171; 2014, Thomsen SK et al: Diabetes 65:3805-3811; 2016 and Udler MS et al: PLoS Med 2018 Sep 21;15(9):e1002654

# DPP: Diabetes Incidence Rate by Weighted Genetic Risk Score of 34 T2D Gene Loci



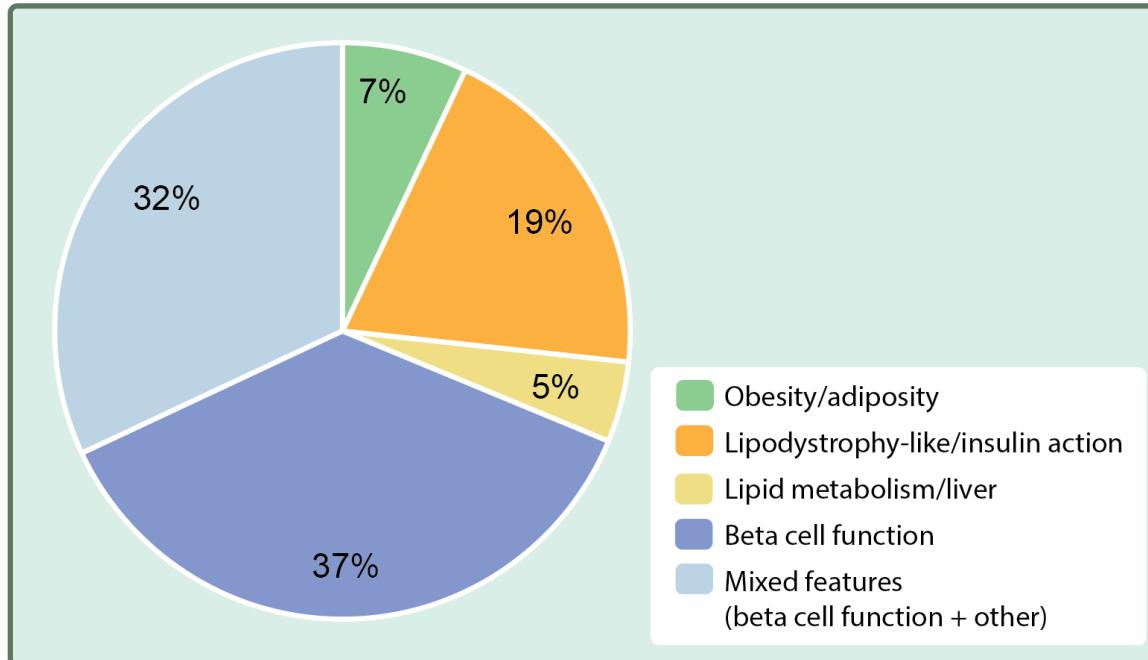
# Partitioned Polygenic Score Clusters Capturing Etiological Heterogeneity in T2D

|  |                                 | Cluster Name                                      |                                    |                                      |  |
|--|---------------------------------|---|------------------------------------|--------------------------------------|--|
| Physiological Impact                           |                                 | Phenotypic Features                               | Udler <i>et al.</i> ,<br>2018 (38) | Mahajan <i>et al.</i> ,<br>2018 (20) | Examples of T2D Loci                                 |
| Adverse impact<br>on $\beta$ -cell<br>function | High proinsulin                 | Low fasting insulin (+ high proinsulin)           | $\beta$ -Cell                      | Insulin secretion 1                  | ABO, ADCY5, HNF1A, HNF1B, MTNR1B, SLC30A8, TCF7L2    |
|  | Low proinsulin                  | Low fasting insulin (+ low proinsulin)            | Proinsulin                         | Insulin secretion 2                  | IGF2BP2, CENTD2/ARAP1, CCND2                         |
| Reduced insulin<br>sensitivity                 | Mediation with fat distribution | High fasting insulin + low BMI + low WC + high TG | Lipodystrophy                      | Insulin action                       | MACF1, GRB14, IRS1, PPARG, ANKRD55, KLF14, LPL, CMIP |
|  | Mediation via obesity           | High fasting insulin + high BMI + high WC         | Obesity                            | Adiposity                            | NRXN3, FTO, MC4R                                     |
|  | Mediation via lipid metabolism  | Low TG  | Liver/lipid                        | Dyslipidemia                         | GCKR, TM6SF2   |
| Undetermined                                   |                                 | No striking phenotype association                 | No assignment                      | Mixed features                       | BCL11A, TLE1, PLEKHA1, HMGA2, MTMR3                  |

Comparison of pPS clusters identified by Mahajan *et al.* (20) and Udler *et al.* (38).

Abbreviations: TG, triglyceride; WHR, waist/hip ratio.

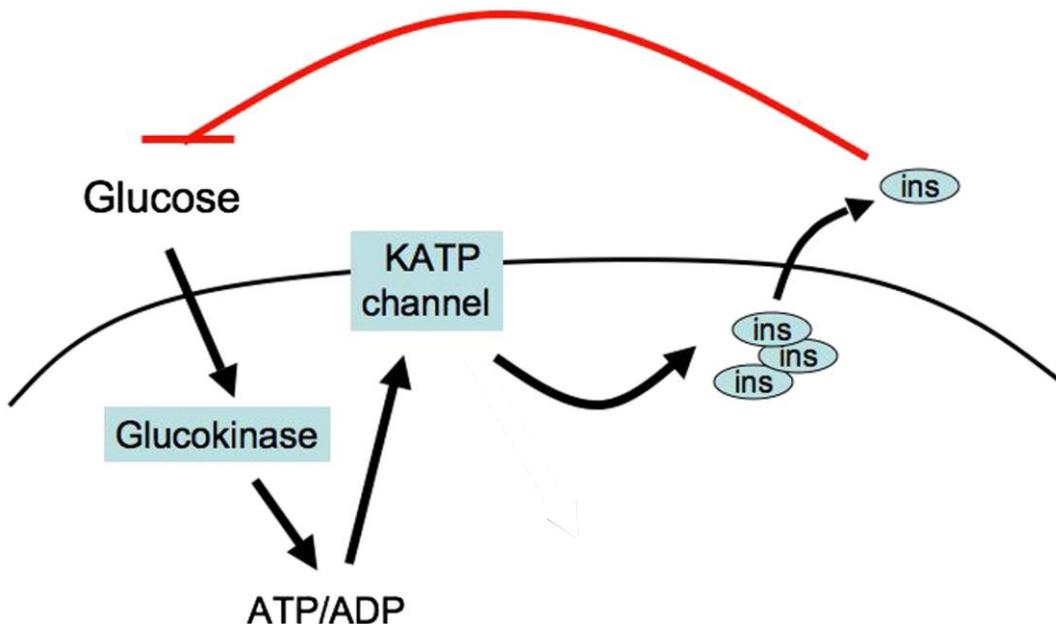
# Phenotypes Associated with 128 Gene Variants Linked to Type 2 Diabetes



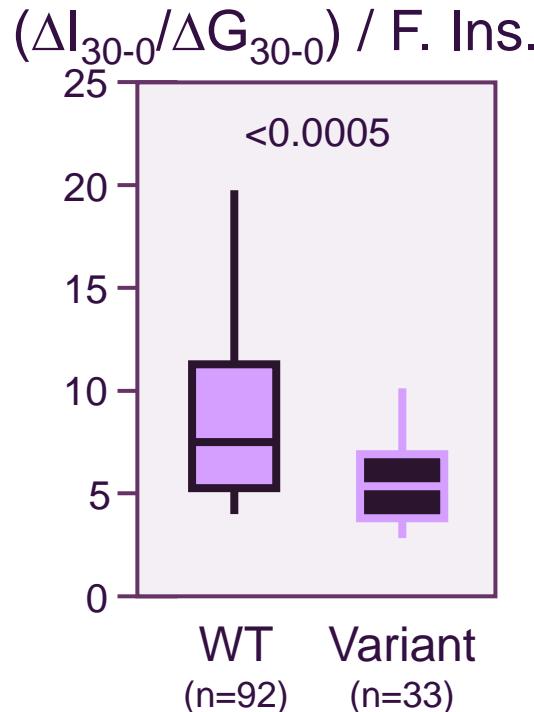
# Function of Select Gene Products Linked to the β-cell in Type 2 Diabetes and MODY

| Function                     | Gene Product  | Function                          | Gene Product   |
|------------------------------|---|-----------------------------------|--|
| Growth and development       | CDKAL1, GLIS3, HHEX,<br><i>HNF1A</i> , <i>HNF1B</i> ,<br><i>HNF4A</i> , IGF2BP2 | Transcription factor              | <i>PAX4</i> , PROX1, RREB1,<br><b>TCF7L2</b> , ZBED3 |
| Insulin production           | INS   | Peptide processing                | SLC30A8  |
| Insulin secretion            | <i>ABCC8</i> , <i>KCNJ11</i> ,<br>KCNQ1   | Potentiation of insulin secretion | GIPR   |
| Glucose metabolism           | G6PC2, GCK  | Peptide degradation               | IDE  |
| Golgi vesicles and structure | AP3S2, ARAP1  | Cell cycle regulation             | CCND2, CDC123,<br>CDKN2A                             |
| Cell adhesion and migration  | BCAR1   | Steroid hormone metabolism        | HSD17B12   |

# Glucokinase is Critical In Glucose-stimulated Insulin Secretion by the $\beta$ -cell



# -30 (G>A) β-cell GCK Promoter Variant is Associated with Reduced β-cell Function



| NGT / IGT               | Wild Type<br>55 / 37 | Variant<br>10 / 23 | p     |
|-------------------------|----------------------|--------------------|-------|
| Fasting Insulin<br>(pM) | 60<br>(45-78)        | 60<br>(47-104)     | NS    |
| Fasting Glucose<br>(mM) | 5.5<br>(5.3-5.9)     | 5.7<br>(5.4-6.1)   | 0.02  |
| 2-h Glucose<br>(mM)     | 7.4<br>(6.4-8.5)     | 8.6<br>(7.4-9.5)   | 0.001 |

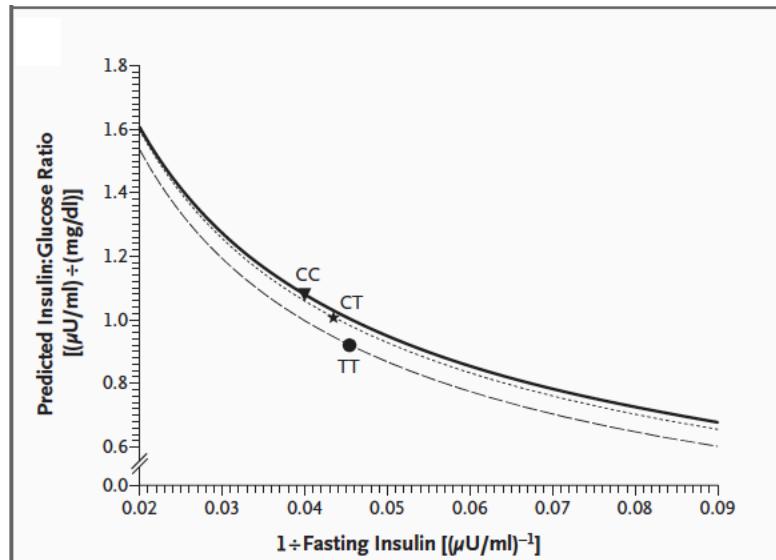
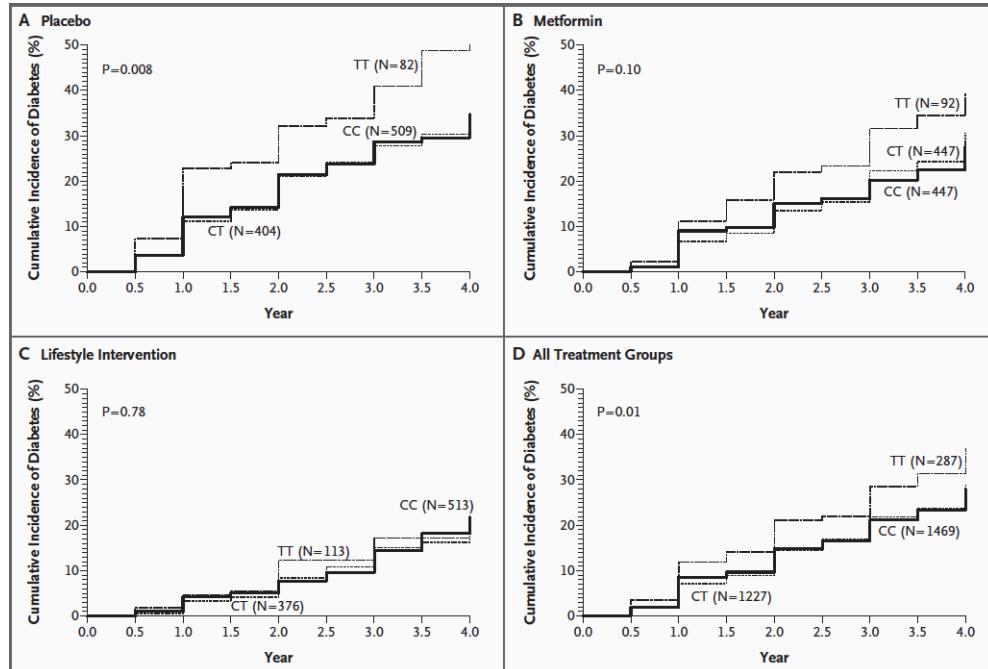
Data are medians (interquartile range)

# TCF7L2: Transcription Factor 7 Like 2

UniProtKB/Swiss-Prot:

Participates in the Wnt signaling pathway and modulates MYC expression by binding to its promoter in a sequence-specific manner. Acts as repressor in the absence of CTNNB1, and as activator in its presence.

# TT Genotype at rs7903146 of TCF7L2 Increases the Risk of Developing Diabetes

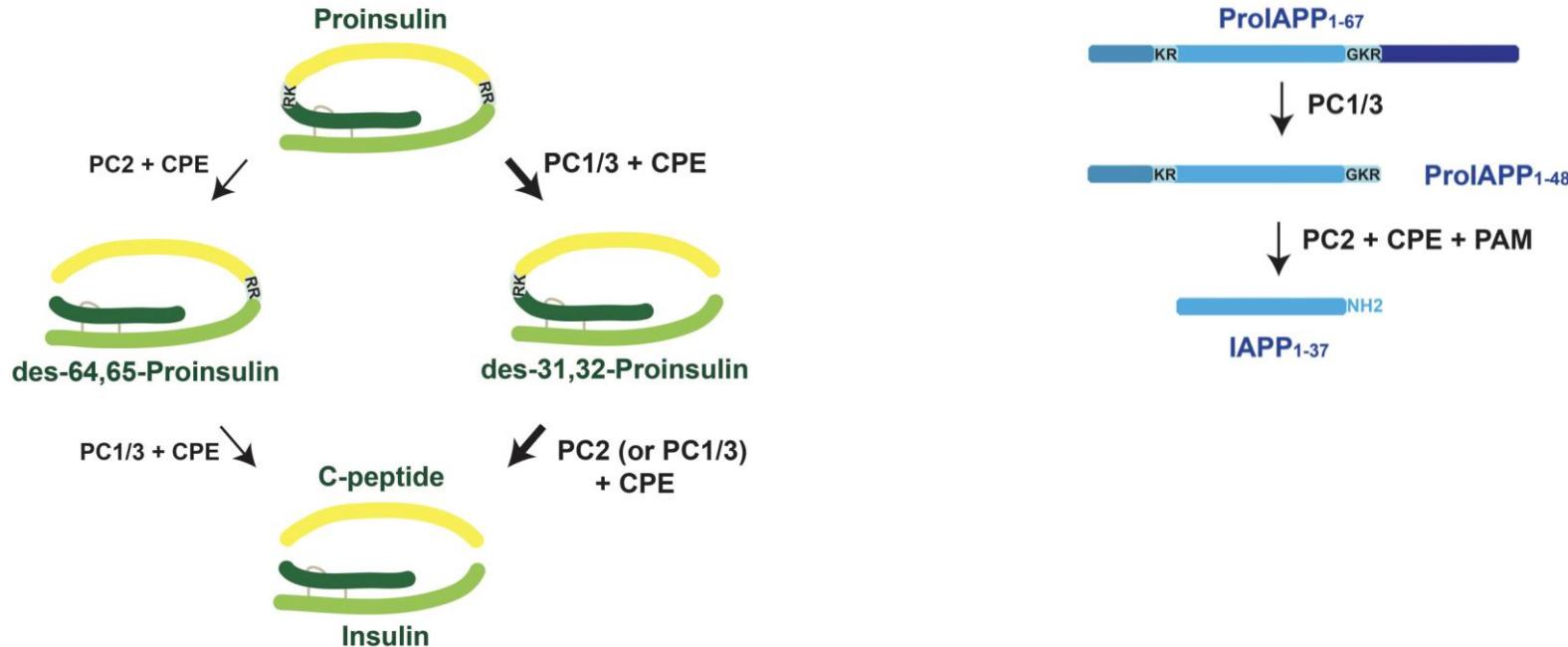


# SLC30A8: Solute Carrier Family 30 Member 8

UniProtKB/Swiss-Prot:

Facilitates the accumulation of zinc from the cytoplasm into intracellular vesicles, being a zinc-eﬄux transporter. May be a major component for providing zinc to insulin maturation and/or storage processes in insulin-secreting pancreatic  $\beta$ -cells.

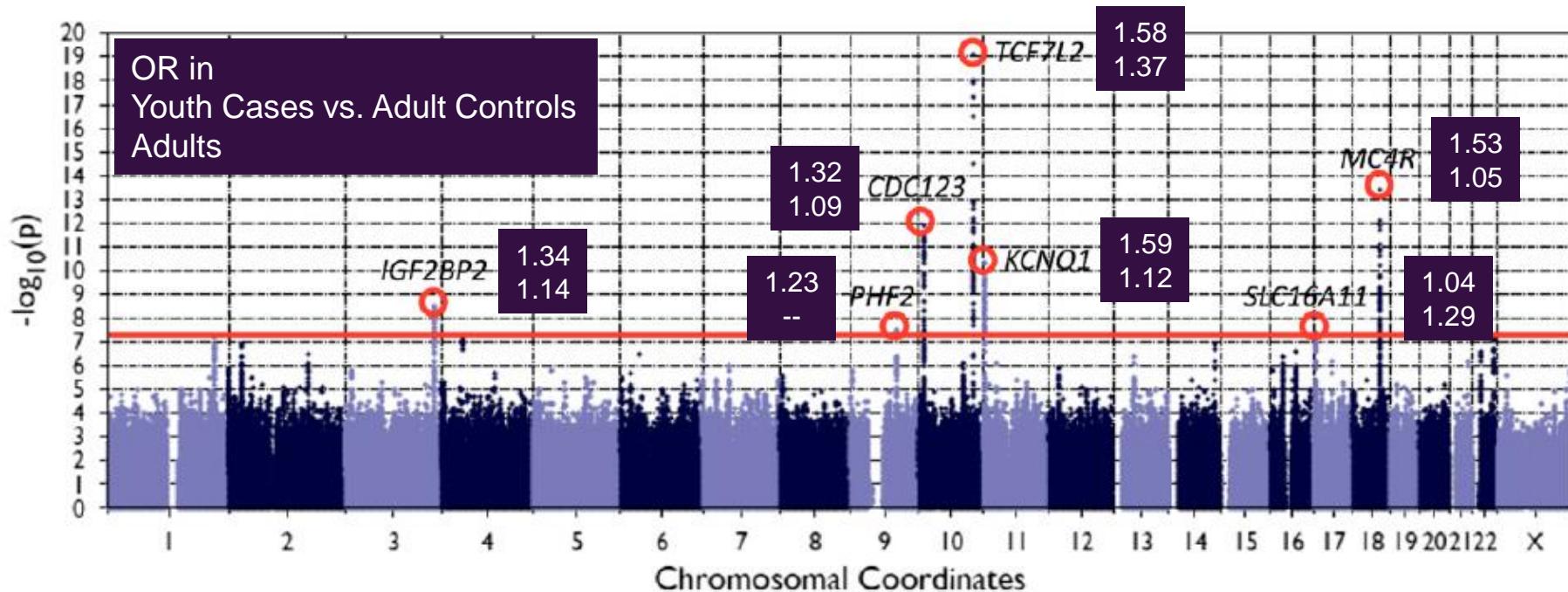
# Pathways for Processing Proinsulin and ProAPP in the $\beta$ -cell



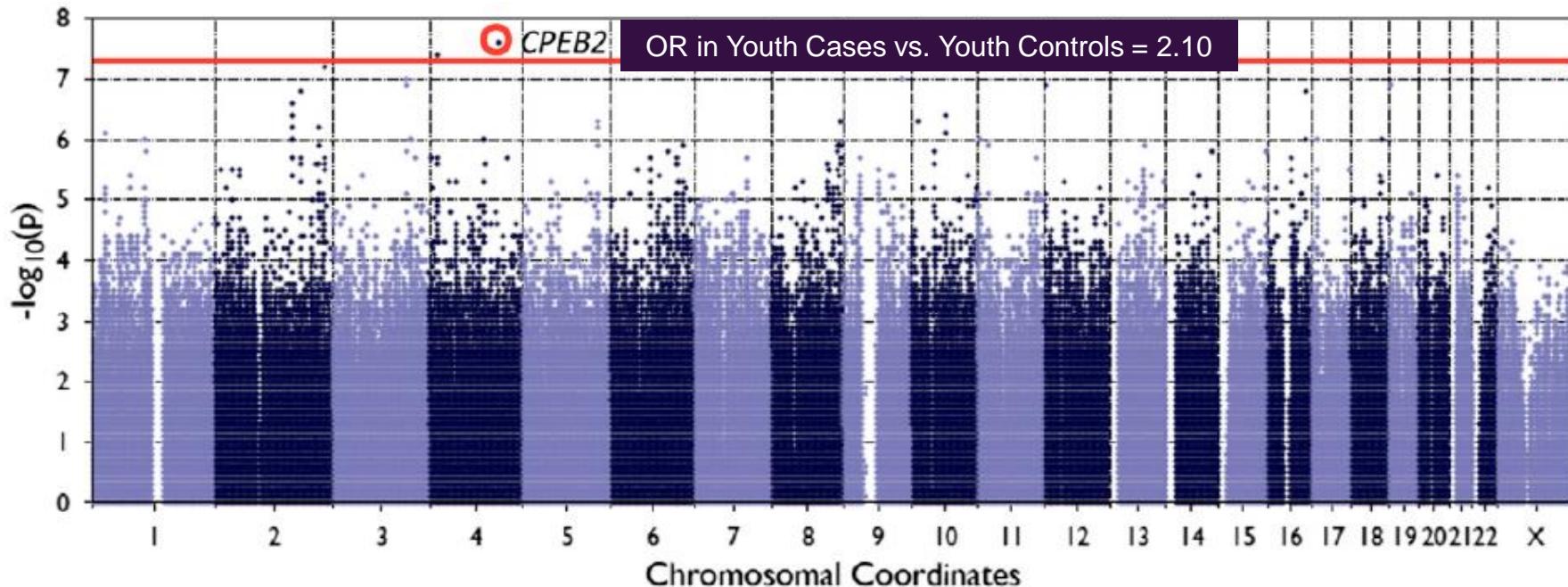
# Outline

1. Phenotypic characteristics and genetics of type 2 diabetes in adults.
2. Genetics of type 2 diabetes in youth.

# Manhattan Plot for 3,006 Youth Cases With T2D vs. 6,061 Adult Controls Without Diabetes



# Manhattan Plot for 3,006 Youth Cases With T2D vs. 856 Youth Controls Without Diabetes



# Summary: Gene Mutations and the $\beta$ -cell

$\beta$ -cell dysfunction is a key component of the pathogenesis of type 2 diabetes and can be demonstrated in individuals who still have normal fasting glucose or normal glucose tolerance.

$\beta$ -cell dysfunction is the predominant phenotype associated with gene variants linked to type 2 diabetes. These genotypes affect different components of the  $\beta$ -cell's function.

Genes linked to type 2 diabetes in youth are similar to those in adults.

# Epigenetic Modifications Occurring in the $\beta$ -cell in Diabetes

