



***Precision Medicine in Diabetes Mellitus:  
Where Are We Now?***

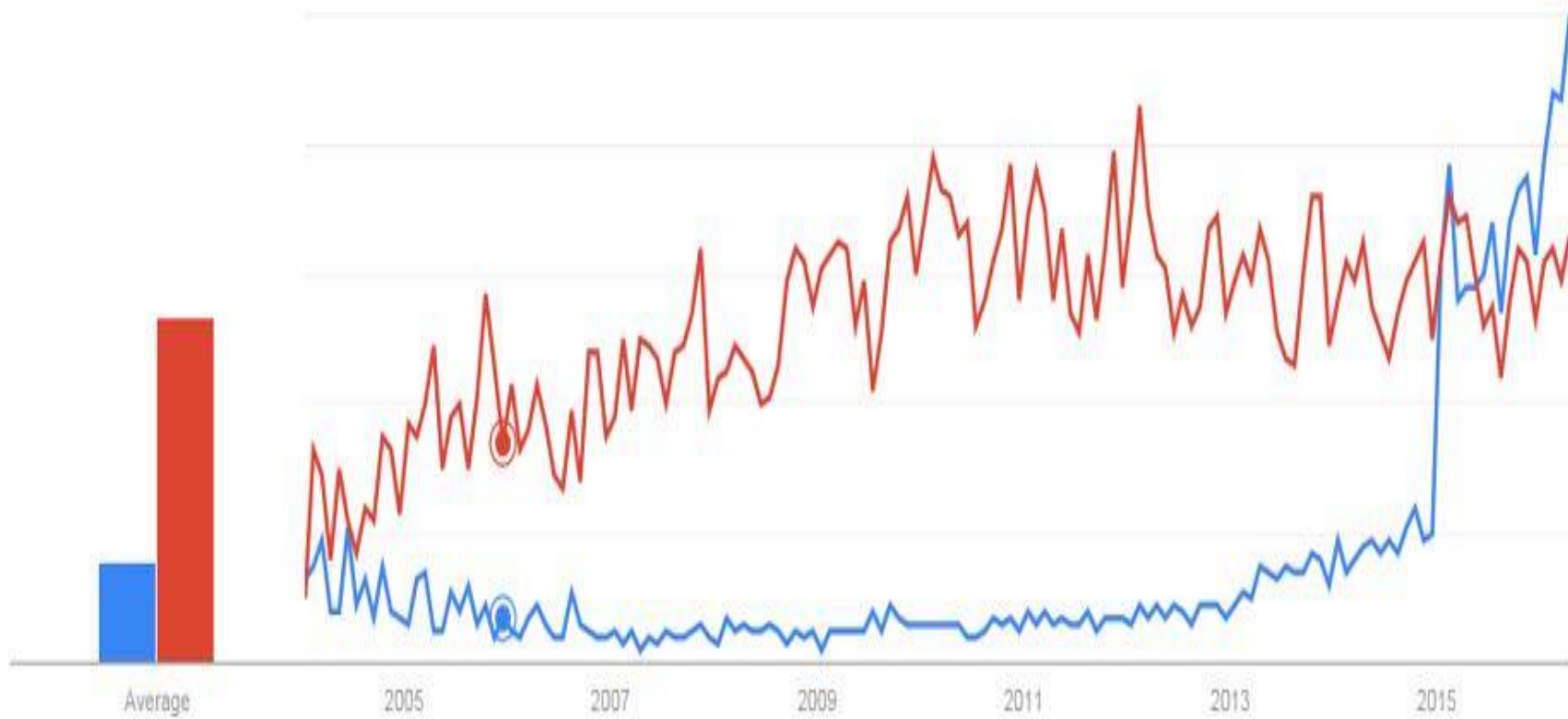
**IRAJ HEYDARI, MD**

**Iran University of Medical Science  
INSTITUTE OF ENDOCRINOLOGY AND METABOLISM**

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# AGENDA

- Precision medicine
- Precision medicine and diabetes mellitus
- Precision medicine and diabetic nephropathy



Relative Interest in Precision Medicine (blue) vs. Personalized Medicine (red), 2005-2016

(Reflected in Google Trend Search on April 10, 2016)

# Precision Medicine

- In PM, diagnostic testing is for selecting optimal therapies , based on the context of a patient's genetic content or other molecular or cellular analysis.
- Every person has a unique variation of the human genome.
- Individual's health stems from genetic variation and environment.
- The terms personalized medicine, precision medicine, stratified medicine and P4 medicine are used interchangeably to describe this concept.
- The right treatment, to the right patient, at the right time.

# P4 Medicine

A term coined by Dr. Leroy Hood

Personalized	Diagnosis and treatment are customized
Predictive	Determine the risk and outcome
Preventive	Take preventive action according to the risk
Participatory	Patients are involved



## Obama's Precision Medicine Initiative

*“ Most medical treatments have been designed for the “ average patient ”. As a result of ‘ this one- size- fits-all- approach ’, treatments can be very successful for some patients but not for others.”*

*“Tonight I’m launching a new Precision Medicine Initiative to bring us closer to curing diseases like cancer and diabetes.*

*And to give us all access to the personalized information we need to keep ourselves and our families healthier.”*

President Barack Obama

2015



In 2015, Barack Obama dedicated  
**\$215 million** to the Precision  
Medicine Initiative

## Development of concept

- Human Genome Project (HGP) , determining the DNA sequence of the entire euchromatic human genome , formally launched in 1990 and was complete in 2003.
- Using high throughput technologies, we are now able to perform an exhaustive measurements over a short period of time .
- New technologies and knowledge emerging from HGP , led to the omic revolution in the beginning of the 21th century .
  - Genomic, Transcriptomic , Epigenomic , Proteomic , Metabolomic...



# Some Facts

- **In human beings, 99.9% bases are same**
- **Remaining 0.1% makes a person unique**
  - Different attributes / characteristics / traits
    - how a person looks
    - diseases he or she develops
- **These variations can be:**
  - Harmless (change in phenotype)
  - Harmful (diabetes, cancer, heart disease, Huntington's disease, and hemophilia )
  - Latent (variations found in coding and regulatory regions, are not harmful on their own, and the change in each gene only becomes apparent under certain conditions e.g. susceptibility to heart attack)

# Genomic and GWAS Studies

- To know if a mutation is connected to a certain disease, researchers often do a study called a “genome-wide association study” (GWAS).
- A GWAS study will sequence the genome of many patients with particular disease to look for shared (SNPs) in the genomes.
- SNP : a difference in a single nucleotide and is the most common type of genetic variation among people, there is roughly 10 million SNPs in the human genome.
- The first GWAS, conducted in 2005, studied patients with age-related macular degeneration.
- Already identified SNPs related to diabetes, heart abnormalities, Parkinson disease, and Crohn disease.

# omics

- **Transcriptome** : study of all RNA molecules, produced in one or a population of cells.
  - Transcriptomic analyses have the potential to answer the question of functionality genes discovered in whole-genome studies.
  - Transcriptomes, performed on samples such as bone cells revealing the profiles of about 20,000 expressed genes which may differ in health and disease.

# Omics

## ➤ Proteomic studies :

- Represent the analysis of the gene products.
- Recognition of proteins differentially expressed between healthy and diseased individuals.

*Usually performed by difficult and quite expensive technologies like : 2-DE , MS , WB .*

## omics

- **Metabolomes** give us an insight into metabolites that differ between healthy and diseased individuals ,metabolites depend on both genetics and environment.
  - The small changes in gene and protein could be amplified in the metabolic level.
- **Pharmacogenomics** combines pharmacology and genomics , study of the role of genetic in drug response,..

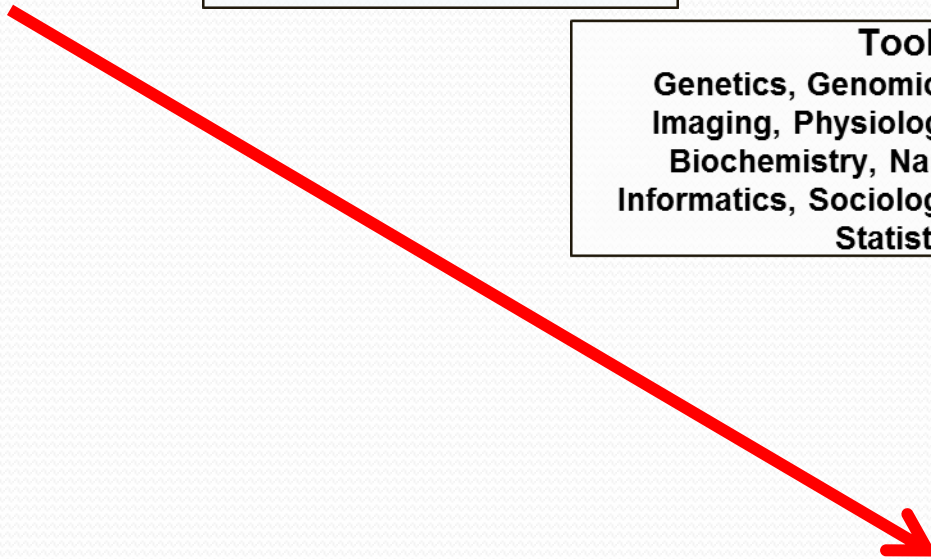
# The Personalized Medicine Research Cycle from Bench to Bedside

**Research  
Questions**

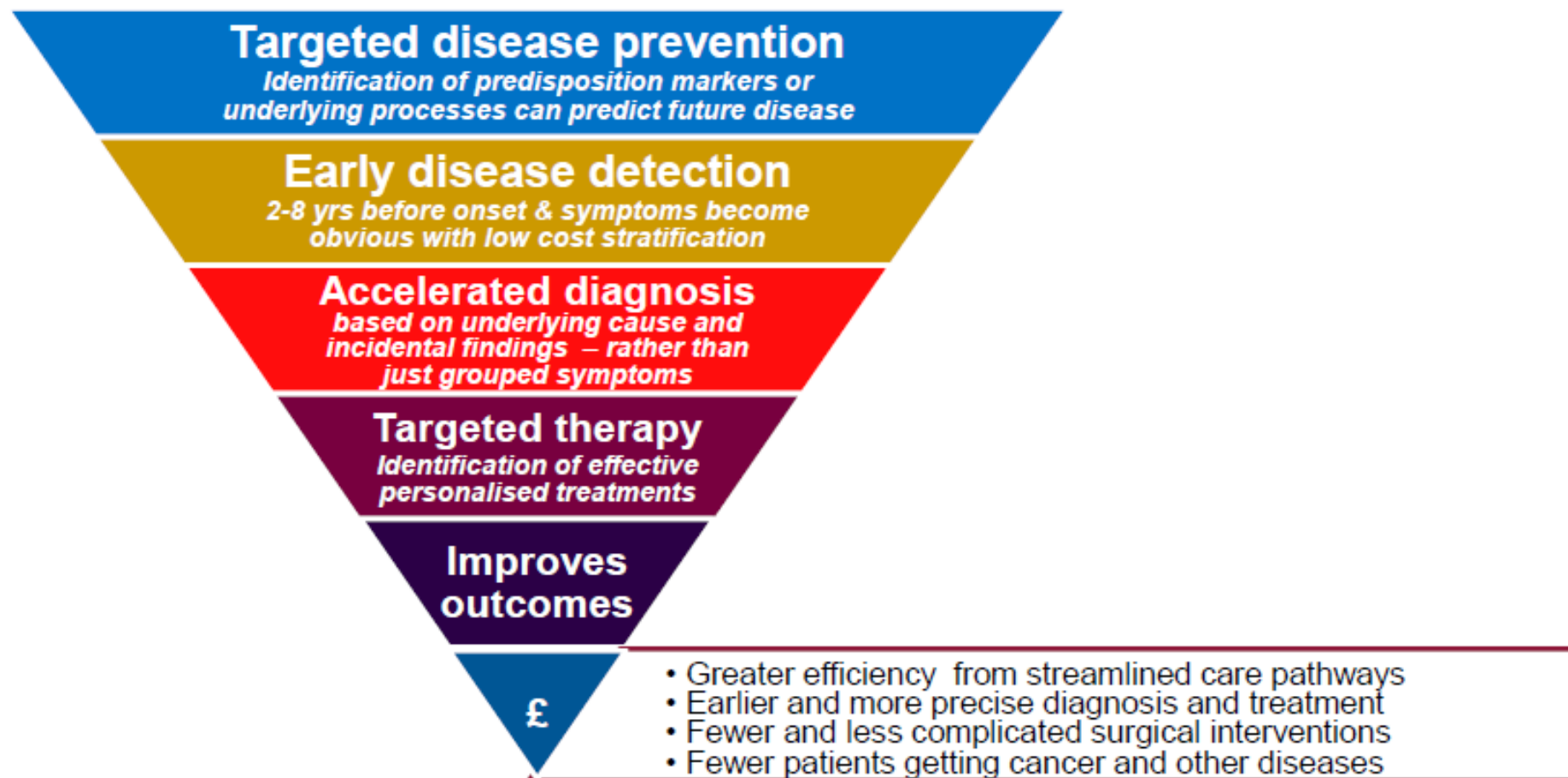
**Biobank**  
Tissues, Cells, Fluids, &  
Products and Dry Data

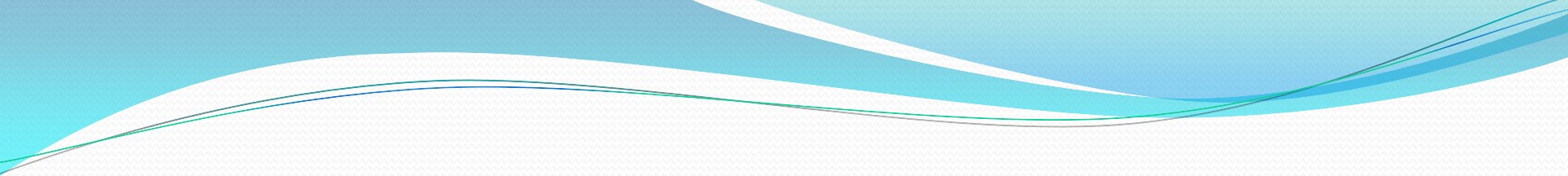
**Tools**  
Genetics, Genomics, Proteomics,  
Imaging, Physiology, Biophysics,  
Biochemistry, Nanotechnology,  
Informatics, Sociology, Epidemiology,  
Statistics

**Biomarker or Target  
Validation**  
Multi-population  
Assessment, High-  
throughput Screening  
Clinical Trials



# Improving outcomes through personalisation

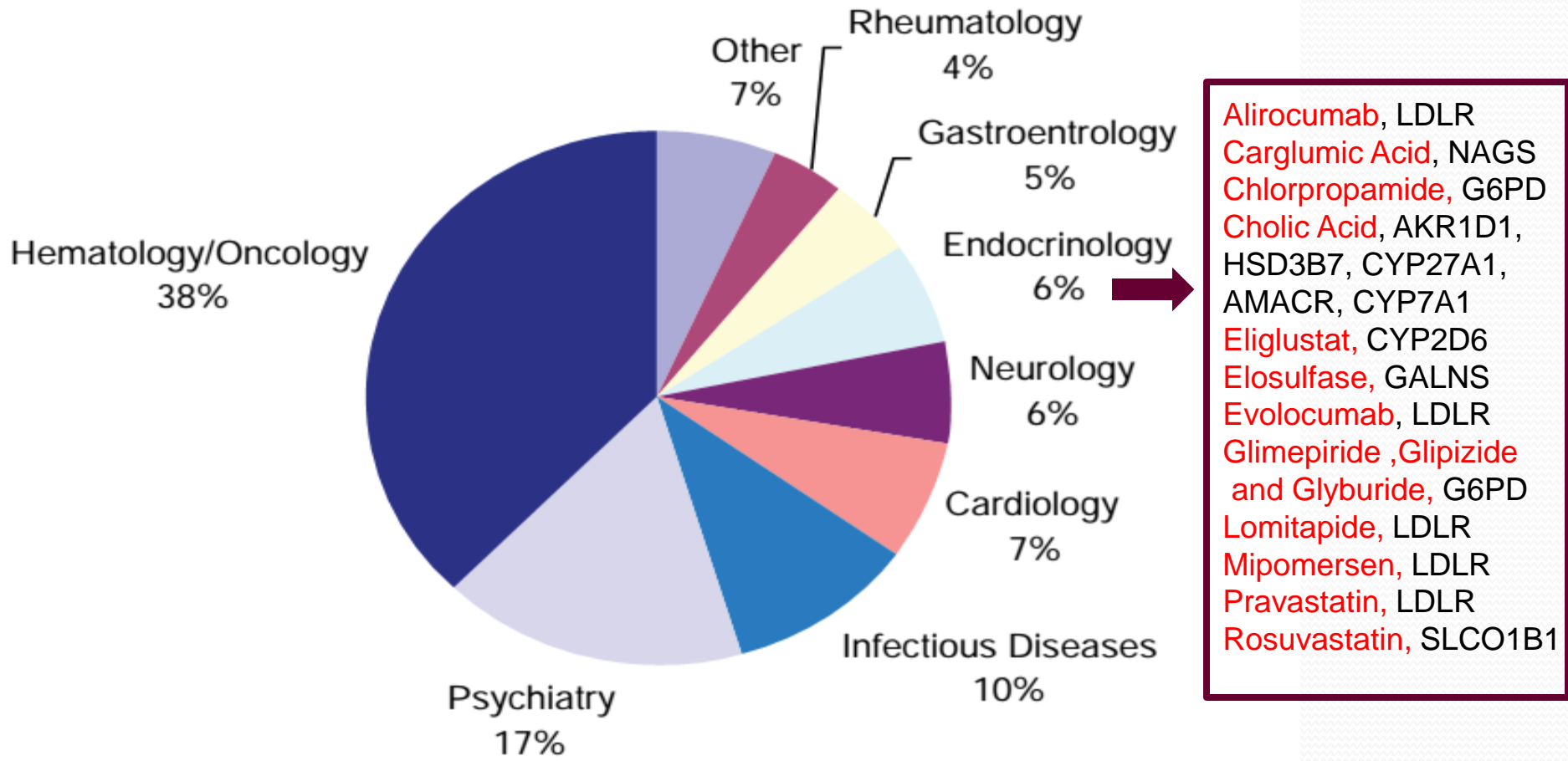




Diabetes care has remarkable growth  
whereas oncology pioneered the introduction of  
precision treatment based on genetics



# Pharmacogenomic Biomarker Information in Drug Labeling



افسردگی



آسم



آریتمی قلبی



دیابت



میگرن



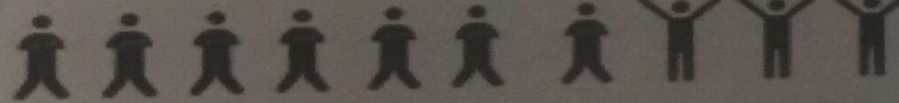
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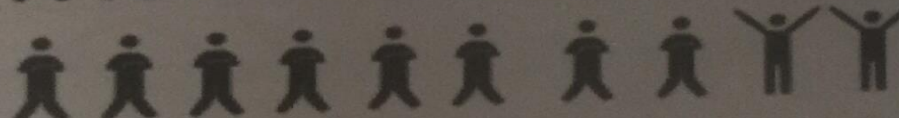
پوکی استخوان



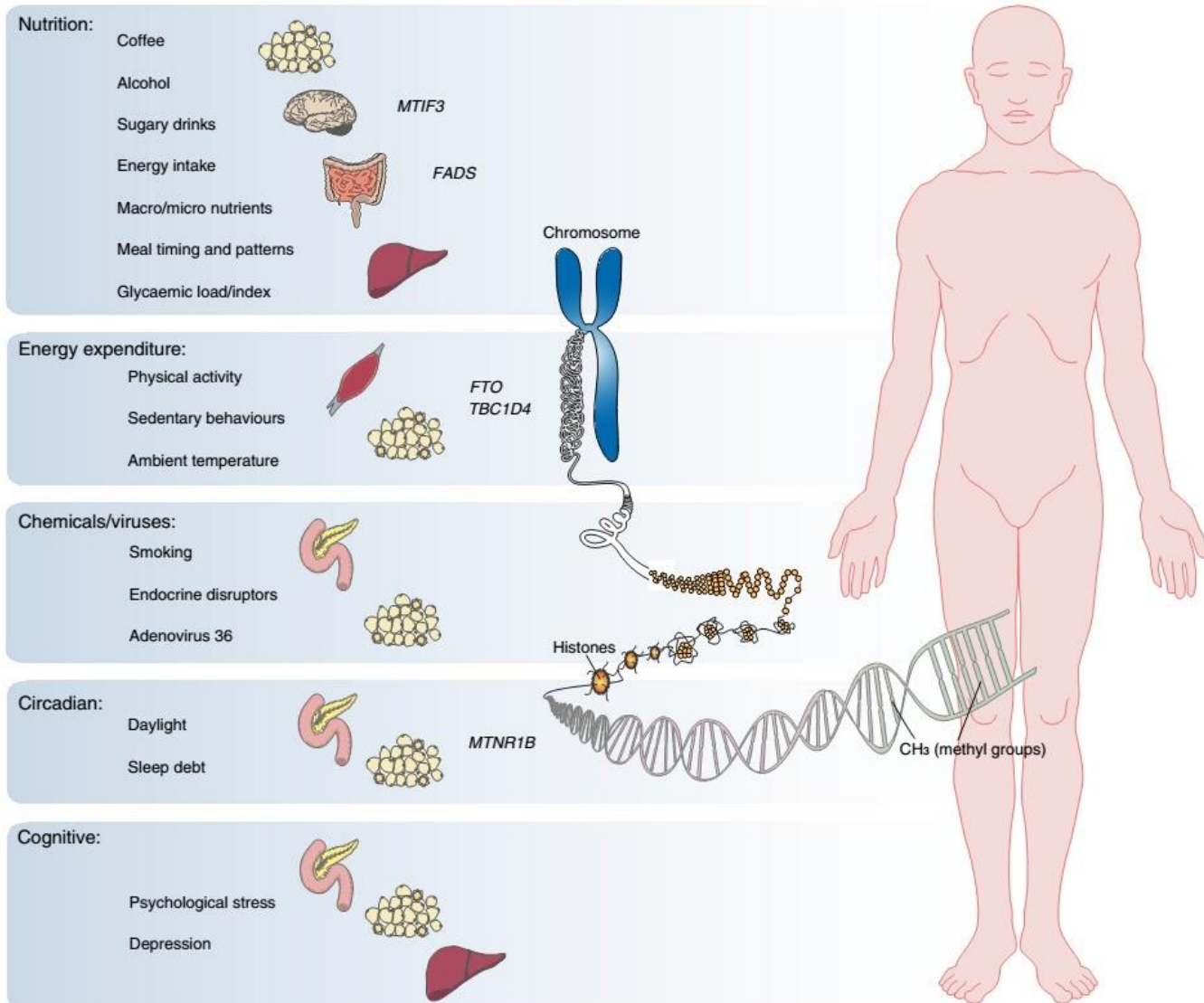
آلزایمر



سرطان










# Type 2 diabetes results from the complex interplay between environmental and genomic factors



# Type 2 Diabetes Genetics

Insulin secretion / beta cell or islet function				Unknown				Insulin resistance			
INRA4	CE	KCNQ1	EA	ADAMTS9	CE	AND1/ROHAF	AA	ANK1	CE	IRS1	CE
TCF7L2	CE	MAEA	EA	AP3S2	SA	SOC3	SA	BCAR1	CE	PCSK9	CE
GCK	CE	ZSXL	EA	CHCHD9P9	CE	SPRY2	EA	CCND2	CE	FTO	CE
INR10	CE	SLC30A8	CE	DNER	SA	SRR	EA	CEP2	CE	GRII4	SA
KCNJ11	CE	TNMD4	CE	FITM2/ROHOM1	EA	STGAL1	SA	KLFDC3	CE	HMG2A2	CE
WFS1	CE	UBE2E2	EA	GCC1	EA	TLE4	CE	TLE1	CE	KLF14	CE
ADAP1	CE	VPS26A	SA	GAK3	EA	TMEM167	SA	ZNF71	CE	PEPD	EA
BCL11A	CE	ZNF23	CE	NR2PM	SA	TSP3NP1	CE	COBLL1	CE	ROSV1	CE
CPCDM1/CPCDM2	EA	ZFAND3	EA	JAZF1	CE	TSPAN8/LGR5	CE	MACF1	CE	ARL15	TA
CDC123/CAMK1D	CE	GFSM1	EA	KCNK16	EA	ZFAND8	CE			LEP	EA
CDKAL1	CE	LPP	TA	LAMA1	CE	FAT1	TA			SLC16A11	M
CDKN2A/B	CE	SSR1/AREB1	TA	MOG2	CE	HLA-B	AA			GCCR	CE
DUSP9	CE	ADCY5	CE	NOTCH2	CE	KIF2	AA			ANKRD6A	CE
GLIS2	EA	DGR2	CE	PRCY	CE	MPYOSP99	TA			MC6R	CE
HNF1X/IDE	CE	M7M10B	CE	PSMD8	EA	POU5F1/TCF19	TA			TBC1D4	G
HNF1A	CE	PROX1	CE	PFP9D	EA	SLC16A13	EA				
KIF20P2	CE	GPR	CE	RASGRP1	EA	TMEM154	TA				

T2D susceptibility genes primarily found through:

-  Linkage studies
-  Candidate genes studies
-  GWAS and GWAS meta-analyses
-  GWAS and GWAS meta-analyses of both imputed and genotyped SNPs
-  GWAS and GWAS meta-analyses for T2D-related quantitative traits
-  GWAS based on MetaboChip custom beadchips
-  WES

# Why Personalized medicine for T2DM

- In tight glycemic control, some individuals develop complications,
- Others with poor control seem to not develop complications.
- Patients with impaired glucose tolerance are at greater risk for macrovascular complications than those with impaired fasting glucose

# Why Personalized medicine for T2DM

## ➤ Prediabetes identification (ADA)

1. Fasting plasma glucose (FPG) of 100– 125 mg/dL (**impaired fasting glucose**)
2. Plasma glucose 2 hours after a 75-g oral glucose challenge of 140–199 mg/dL (**impaired glucose tolerance**)
3. Hemoglobin A1c (HbA1c) test of 5.7%–6.4%.

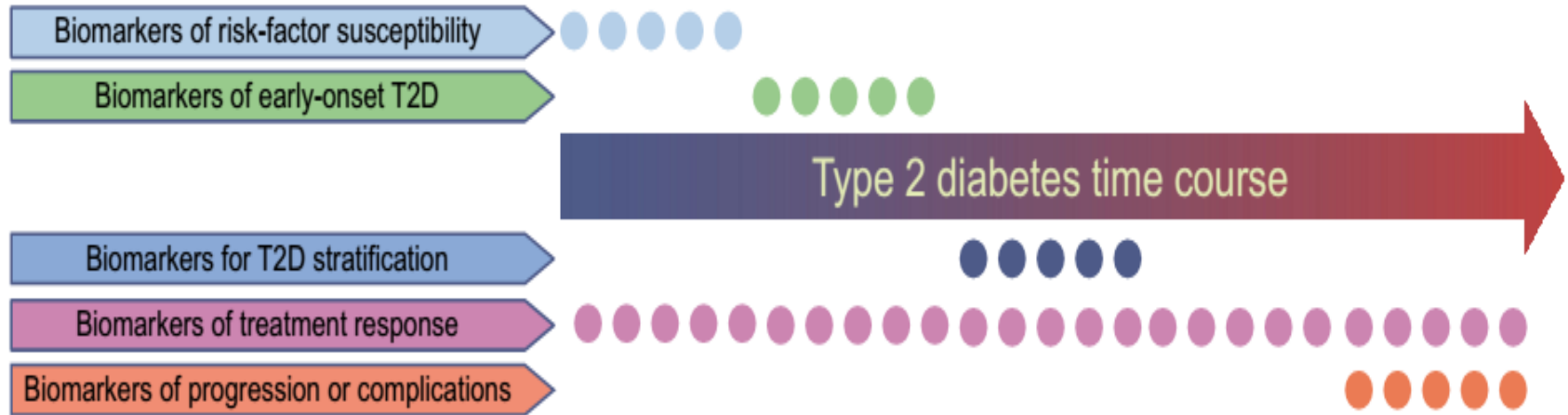
# Why Personalized medicine for T2DM

Problem ....!??!

These criteria do not identify identical groups of people at increased risk for DM2, and their pathophysiology and susceptibility to complications

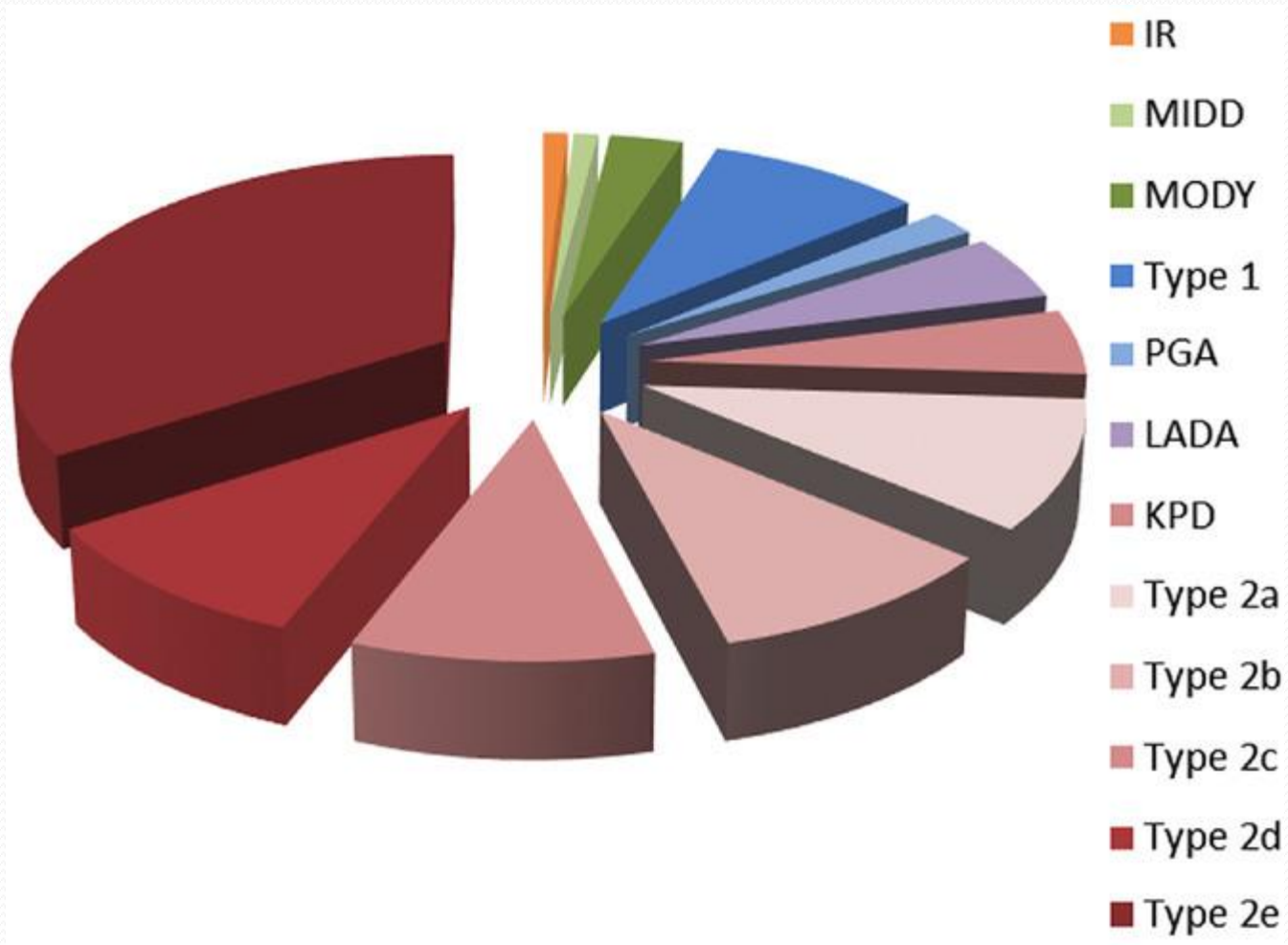
So We need PERSONALIZED MEDICINE FOR T2DM

# Precision Medicine T2DM Time Course



A schematic showing key time points for intervention in the course of type 2 diabetes pathophysiology where precision lifestyle medicine might play a role





## Many important resources for precision diabetes medical research have been launched.

- NIH and pharmaceutical companies with the aim of precision preventive and treatment , collected the results from 28 large (GWAS)

([www.type2diabetesgenetics.org](http://www.type2diabetesgenetics.org))

- (IRAS) by supporting of (NHLBI) have found a possible biomarker for the incidence of T2DM (16).

# search for the genetic basis of T2DM

- GWAS studies: >150 risk alleles
  - Responsible for only 10% of the variation in T2DM tendency (53,54).
  - Most of them encode intracellular proteins that mediate the secretion of insulin from beta cells.
- Linkage study : A susceptibility gene, which has been found was TCF7L2
- Candidate gene approach : The association of PPARG and KCNJ11- ABCC8 with T2DM have been found (47).

# Diabetes Complications

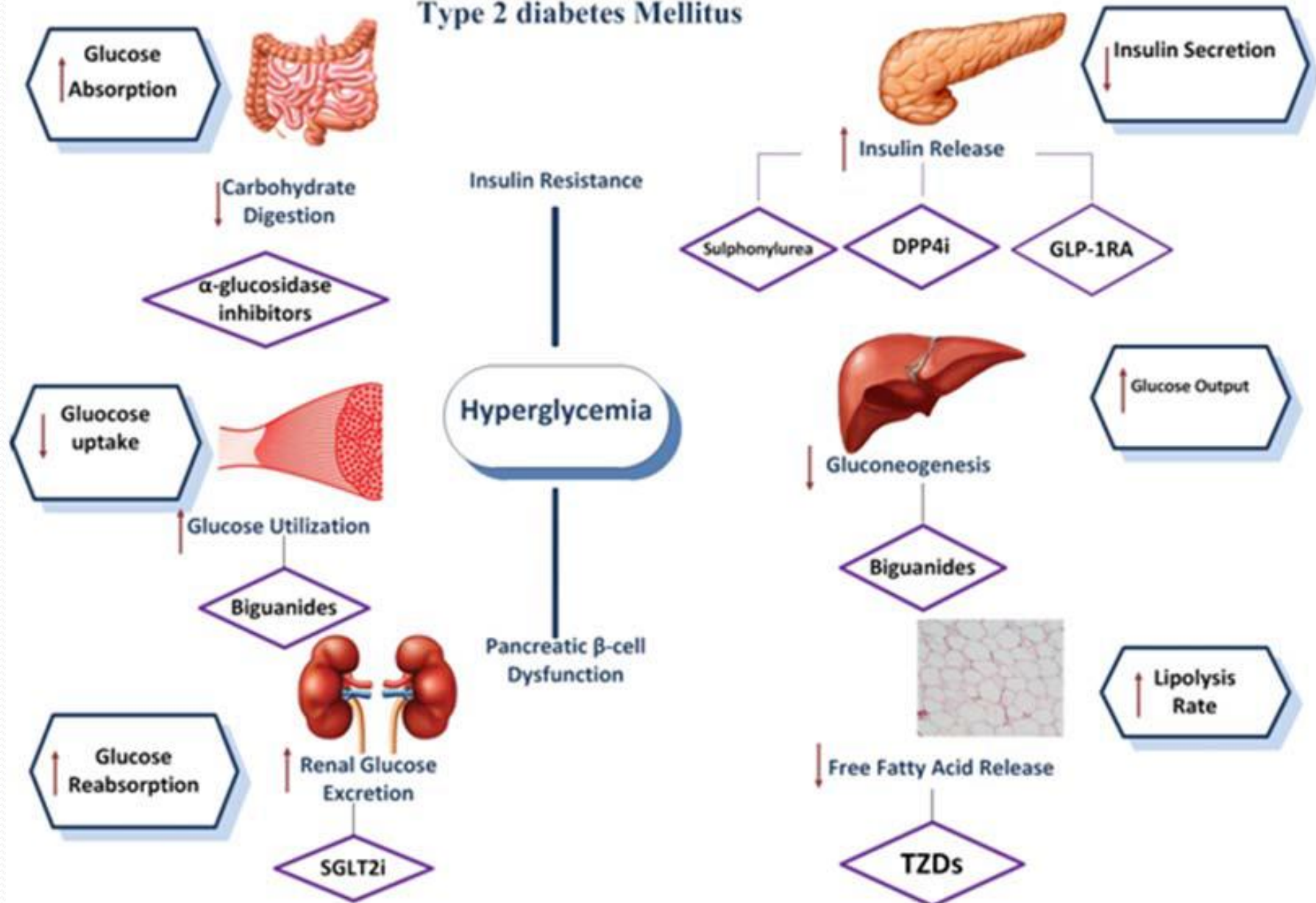
- Occurrence and progression of diabetes complications are influenced by:
  - Modifiable risk factors , such as HTN and HLP.
  - Non modifiable risk factors : genetic predisposition variants. (58).
- GWAS in (ACCORD) trial in patients with T2DM , under intensive glycemic therapy, two genetic variants (*rs9299870, rs57922 on chromosome 10 and 5* ) predict the cardiovascular effects (56).

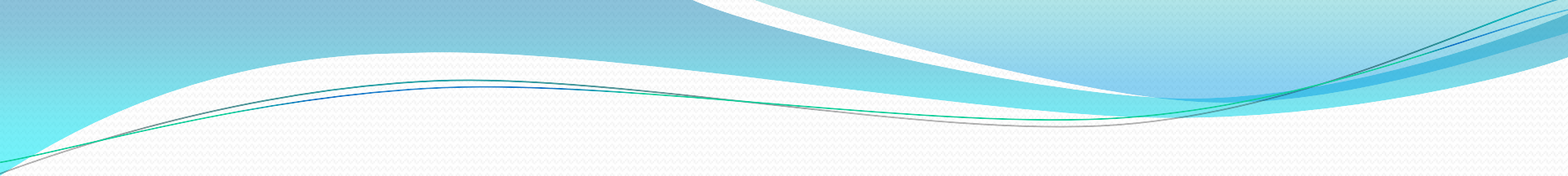
## After 15 to 20 years diagnosis of DM:

- 50%- 80% of patients show retinopathy(two candidate genes VEGFA and AKR1B1)
- 30% have shown an early stage of nephropathy(A well-known GWAS for DKD is related to variation in the gene encoding ACE.

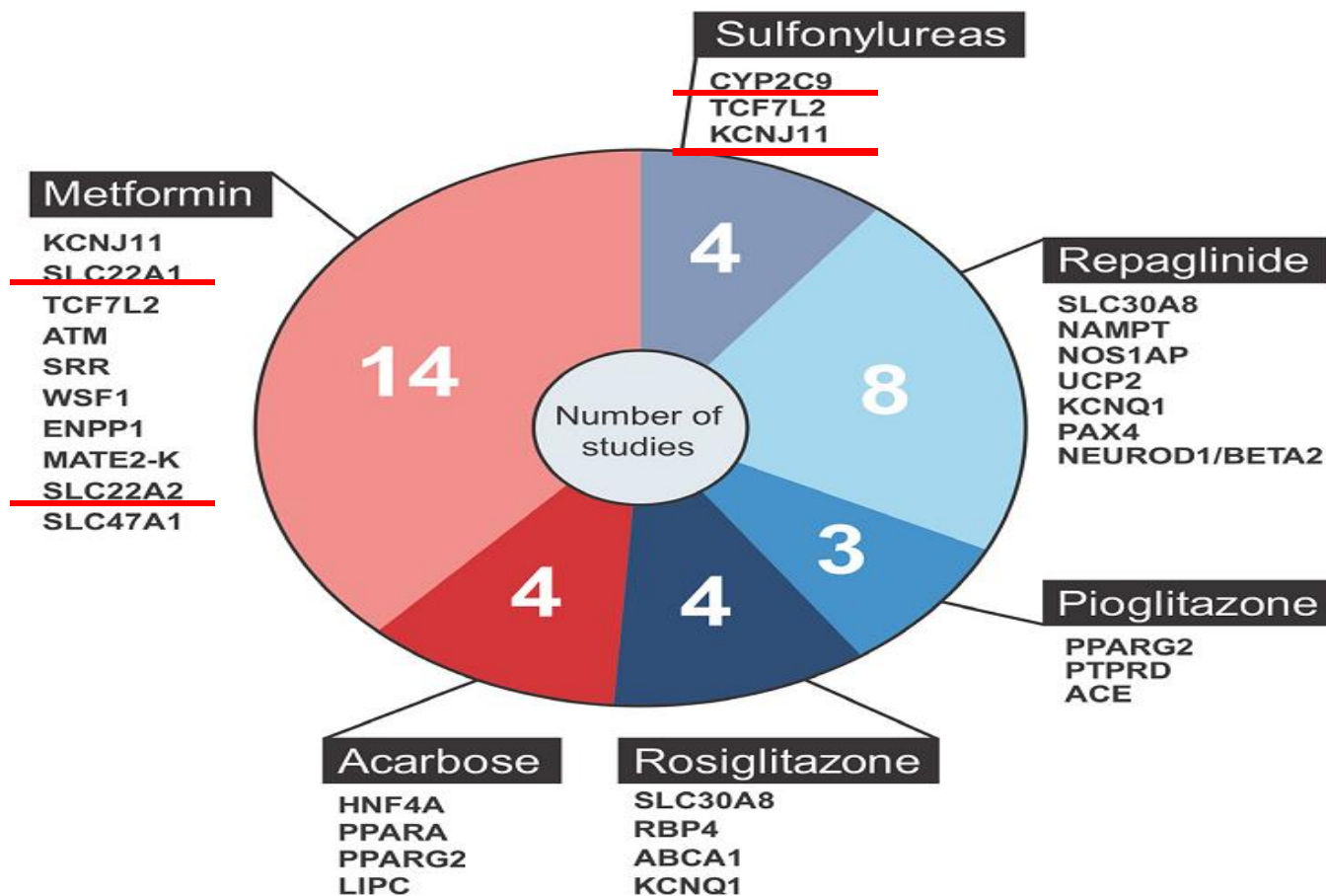
# Management of T2DM from a pathophysiological point of view

## Type 2 diabetes Mellitus



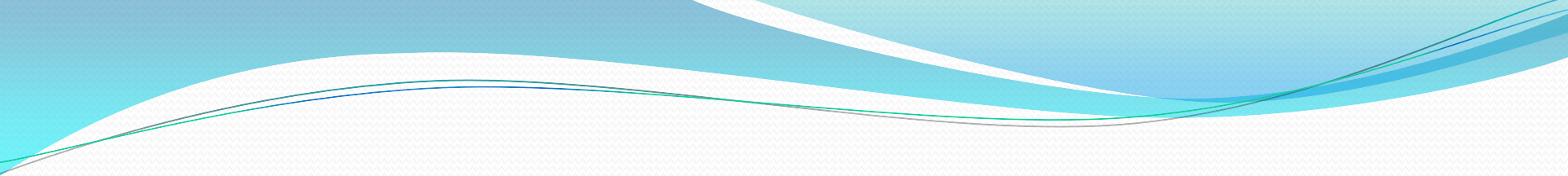
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- An essential concern on pharmacogenomic studies is the genetic makeup of drug response
  - The pharmacokinetic approach is focused on some candidate genes related to:
    - Drug targets
    - Drug metabolism
    - Drug distribution
  - Few strong pharmacogenetics studies have been done in DM treatment

# Regulatory genes in significant gene–drug interaction for anti-diabetic medication



Kaixin Zhou, Pharmacogenomics in diabetes mellitus: insights into drug action and drug Discovery. Nature Reviews Endocrinology 12, 337–346 (2016)





In spite of newly recognized diabetes molecular pathways, there is a large gap between molecular knowledge of diabetes and using it in the clinic or at the bedside.

## DKD

- Mechanistic heterogeneity of diabetic kidney disease (DKD) causes reduced capability to identify risk variants. Some risk variants from APOL1, UMOD genes have been reported, and APOL1 is independent of diabetes status. A well-known GWAS for DKD is related to insertion/deletion (I/D) variant in the gene encoding ACE.

# Conclusion & future perspective

The ideal of personalized care is that each patient receives the management plan best suited to him or her.

This means implementing a treatment strategy that is concordant with the patient's preferences, specific risks and unique underlying disease pathophysiology and drug metabolism profile. Although the benefit remains to be proven, such an approach holds the potential to substantially improve the care of patients with Type 2 diabetes.



**Thank  
you  
for  
Attention**