

DIABETES MELLITUS – TYPES AND PATHOGENESIS

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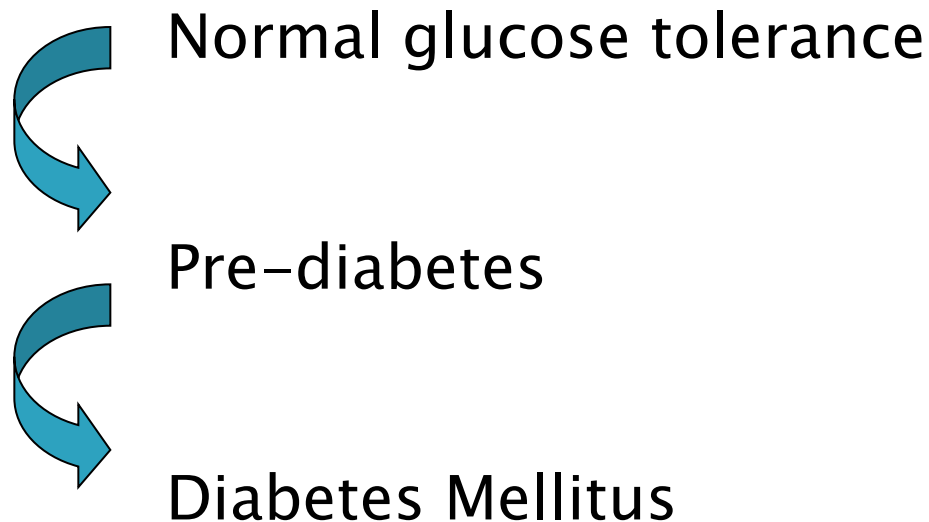
DIABETES MELLITUS

Group of metabolic disorders with hyperglycemia:

1. reduced insulin secretion
2. decreased glucose utilization
3. increased glucose production

CLASSIFICATION

- ▶ DM is classified on the basis of pathogenic process that causes hyperglycemia
- ▶ Common feature: transversion from



NORMOGLYCEMIA ↔ HYPERGLYCEMIA

	Normal glucose tolerance	Impaired fasting glucose or impaired glucose tolerance	Diabetes Mellitus
Fasting plasma glucose	< 5.6 mmol/L (100mg/dL)	5.6 – 6.9 mmol/L (100 – 125 mg/dL)	≥ 7.0 mmol/L (126 mg/dL)
2-h-plasma glucose	< 7.8 mmol/L (140 mg/dL)	7.8 – 11.1 mmol/L (140 – 199 mg/dL)	≥ 11.1 mmol/L (200 mg/dL)

These values do not apply to gestational DM!!!

Etiologic classification of DM

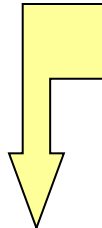
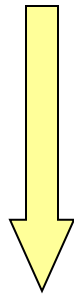
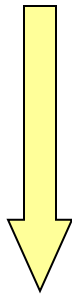
- I. Type 1 diabetes (destruction of beta cells)
- II. Type 2 diabetes
- III. Other specific types of diabetes
- IV. Gestational diabetes mellitus (GDM)

- ▶ Other types of DM include specific genetic defects in insulin secretion, action, metabolic disorders that impair insulin secretion or glucose tolerance, mitochondrial abnormalities
- ▶ **MODY** – *maturity onset diabetes of the young*, subtype of DM, autosomal dominant inheritance, early onset of hyperglycemia (<25 years), impaired of insulin secretion
- ▶ **GDM** – gestational DM, caused by insulin resistance during pregnancy, may revert to normal glucose tolerance; risk of developing DM later in life.

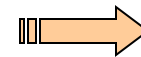
TYPE 1 DM – pathogenesis

GENETIC
PREDISPOSITION

ENVIRONMENTAL
FACTORS



IMMUNOLOGIC
ABNORMALITIES



IMMUNOLOGIC
MARKERS

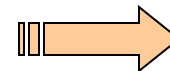
DESTRUCTION OF BETA CELLS



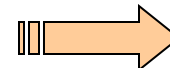
IMPAIRMENT OF INSULIN SECRETION



LACK OF INSULIN
(>80% BETA CELLS
DESTROYED)



PRE-DIABETES



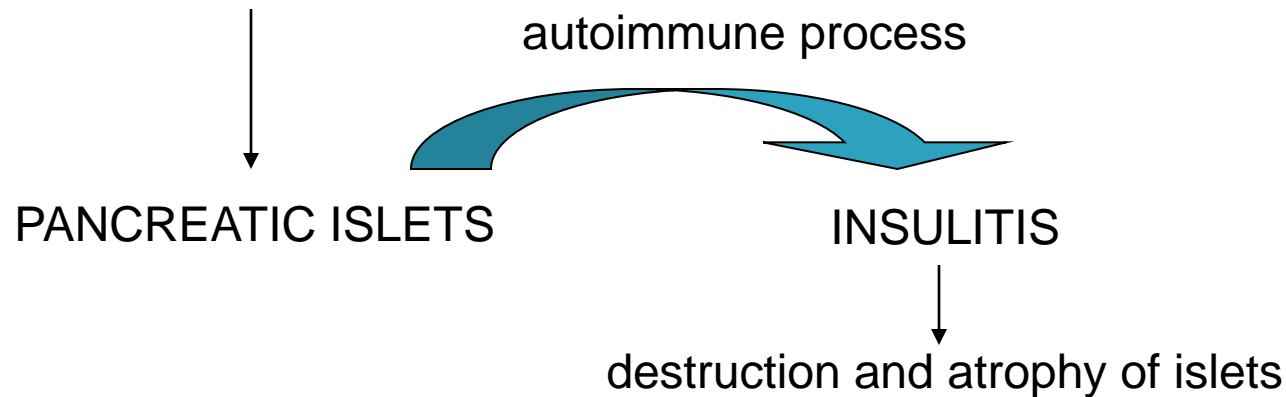
DIABETES

GENETIC PREDISPOSITION

- ▶ Involves multiple genes
- ▶ Major susceptibility gene located in the HLA region on chromosome 6
- ▶ Polymorphisms in the HLA complex account for about 50% of the genetic risk factors of developing DM1
- ▶ Haplotype DQA1*0301, DQB1*0302, DQB1*0201 are strongly associated with type 1 DM.
- ▶ Concordance of type 1 DM in identical twins is variable (30 – 70%) – role of environmental factors

Pathophysiology of type 1 DM

Infiltration with lymphocytes T



Immunologic abnormalities – islet cell autoantibodies (ICAs):

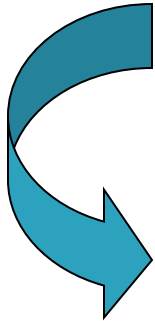
1. Insulin
2. Glutamic acid decarboxylase (GAD)
3. Tyrosine phosphatases (ICA-512/IA-2)
4. Phogrin (insulin secretory granule protein)

} immunologic markers
present in >75% new-onset
type 1 DM
disappear after all beta cells
are destroyed

Environmental factors

- ▶ Trigger an autoimmune process in genetically susceptible individuals
- ▶ Viral infections (coxsackie, rubella)
- ▶ Bovine milk proteins
- ▶ Nitrosourea compounds
- ▶ Event may precede the onset of DM by several years

Pathogenesis of type 2 DM



INSULIN RESISTANCE WITH HYPERINSULINEMIA

Obesity (visceral) → insulin resistance → compensatory increased insulin output

INSULIN SECRETORY DEFECT

Decline in insulin secretion – beta cells failure
(islet amyloid deposits and decreased beta cell mass
in long-standing DM type2)

DIABETES DEVELOPES WHEN INSULIN SECRETION IS INADEQUATE

1. Strong genetic component (concordance in identical twins 70 – 90%)
2. Both parents with type 2 DM = 40% risk of DM !
3. Responsible genes are not completely specified
4. DM type 2 is polygenic and multifactorial (environmental factors: obesity, nutrition, physical activity)

INSULIN RESISTANCE

characteristic feature of type 2 DM

Insulin sensitive tissues
(muscle, fat)

LIVER
Not suppressed
gluconeogenesis

Impaired glucose utilization
Increased FFA flux from
adipocytes

Increased hepatic
glucose output

Postprandial
hyperglycemia

Increased lipid
synthesis in hepatocytes

Increased FPG levels

↑ TGL
↓ HDL

Steatosis of liver
(Nonalcoholic fatty liver disease)



Genetically defined, monogenic forms of DM

- ▶ 6 variants of **MODY** – mutation in genes encoding islet-enriched transcription factors (e.g. HNF – *hepatic nuclear transcription factors*) or glucokinase
- ▶ These factors affect islet development or the expression of genes responsible for glucose-stimulated insulin secretion
- ▶ Clinical features:
progressive decline in insulin secretion and control of glycemia, in some types they respond to **sulfonylureas**
- ▶ Mutation in MODY-associated genes are rare (<5% of type 2 DM)