

AMERICAN DIABETES ASSOCIATION

DIABETES AND PERSONALIZED HEALTHCARE: WHY GENES MATTER



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THE PRECISION MEDICINE INITIATIVE®



WHAT IS IT?

Precision medicine is an emerging approach for disease prevention and treatment that takes into account people's individual variations in genes, environment, and lifestyle.

The Precision Medicine Initiative® will generate the scientific evidence needed to **move the concept of precision medicine into clinical practice.**

WHY NOW?

The **time is right** because of:

Sequencing
of the human
genome

Improved
technologies for
biomedical analysis

New tools
for using large
datasets



PERSONALIZED HEALTHCARE

- Predictive genomic medicine, predictive Medicine, personalized medicine, individualized medicine
- Precisely applying prevention and treatment
- Highest risks of disease, complications, particular prognosis
- Maximize efficacy, minimize side effects

PARADIGM SHIFT

CURRENT MODEL

Patient becomes sick

History and exam

Differential diagnosis

Test selection/application

Refine differential or d

Treat

ALTERNATIVE MODEL

Test selection/application

History and exam

Differential diagnosis

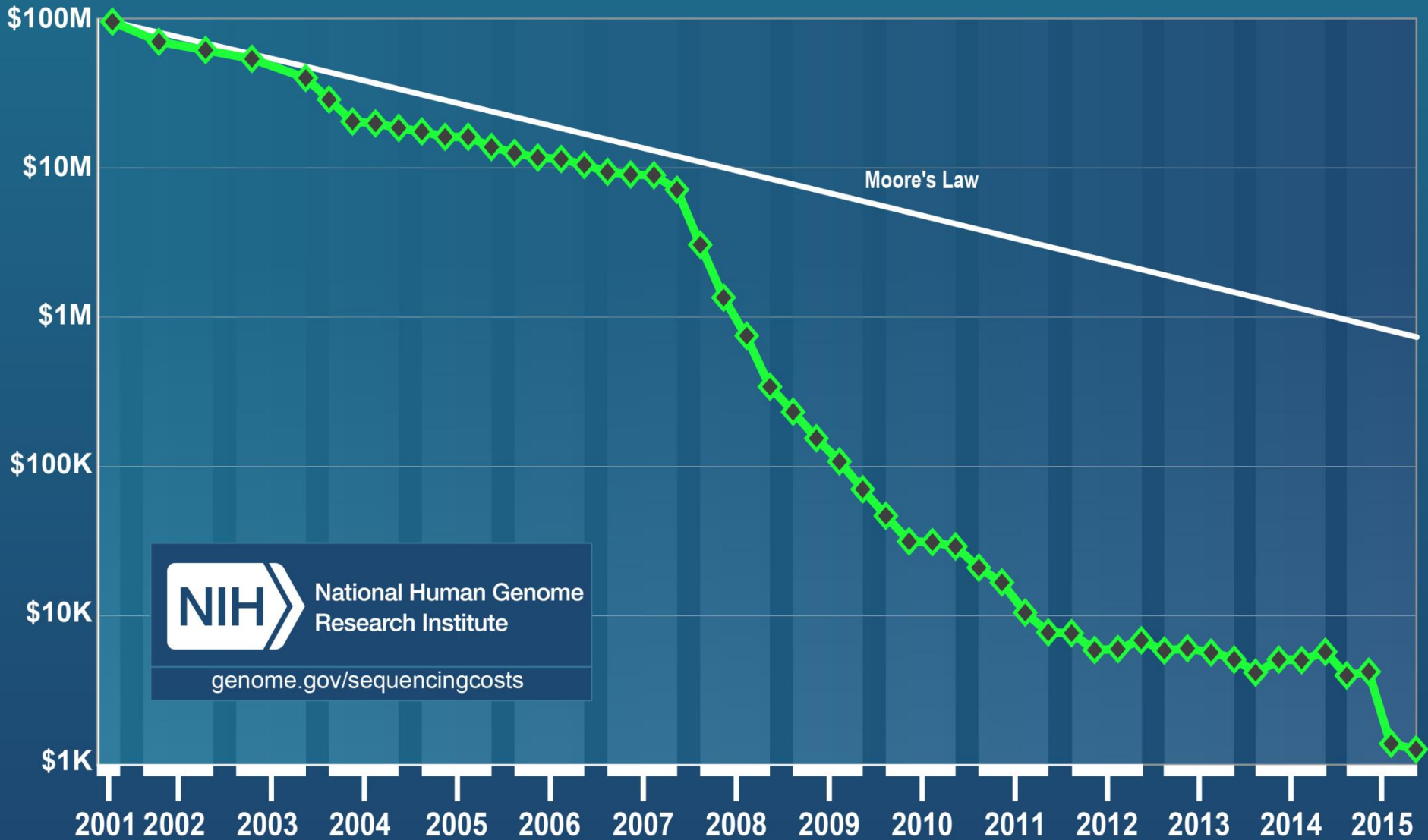
Refine differential
or diagnose

Treat or prevent

ALTERNATIVE MODEL

- Risk Reduction
- Early Detection
- Diagnosis/Prognosis
- Therapeutic Decision-Making
- Tailored Therapy

Cost per Genome





Getting Started with G2C2

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Find websites, books, articles and more - enhance your class content with peer-reviewed resources.

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Search By Topic

Discipline Genetic counselor, nurse, pharmacist, physician, physician assistant

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PERFORMANCE INDICATORS

Professional Practice

- Risk Assessment and Interpretation
- Genetic Education, Counseling, Testing, and Results Interpretation
- Clinical Management
- Ethical, Legal, and Social Implications (ELSI)

Professional Responsibilities

- Professional Role
- Leadership
- Research

<http://www.nursingworld.org/MainMenuCategories/EthicsStandards/Genetics-1/Essential-Genetic-and-Genomic-Competencies-for-Nurses-With-Graduate-Degrees.pdf>

http://www.aacn.nche.edu/education-resources/Genetics__Genomics_Nursing_Competencies_09-22-06.pdf



THE HUMAN GENOME PROJECT: THE BASICS AND BEYOND

Chromosomes



Change of chromosome number

Changes number in autosome chromosomes
Example: Down's syndrome

Changes number in sex chromosomes
Examples: Turner syndrome, Klinefelter syndrome, XYY condition.

Change of chromosome structure

Duplication

Genes sequences are **repeated** several to many hundreds or thousands of times.

Deletion

Loss of some segment of a chromosome

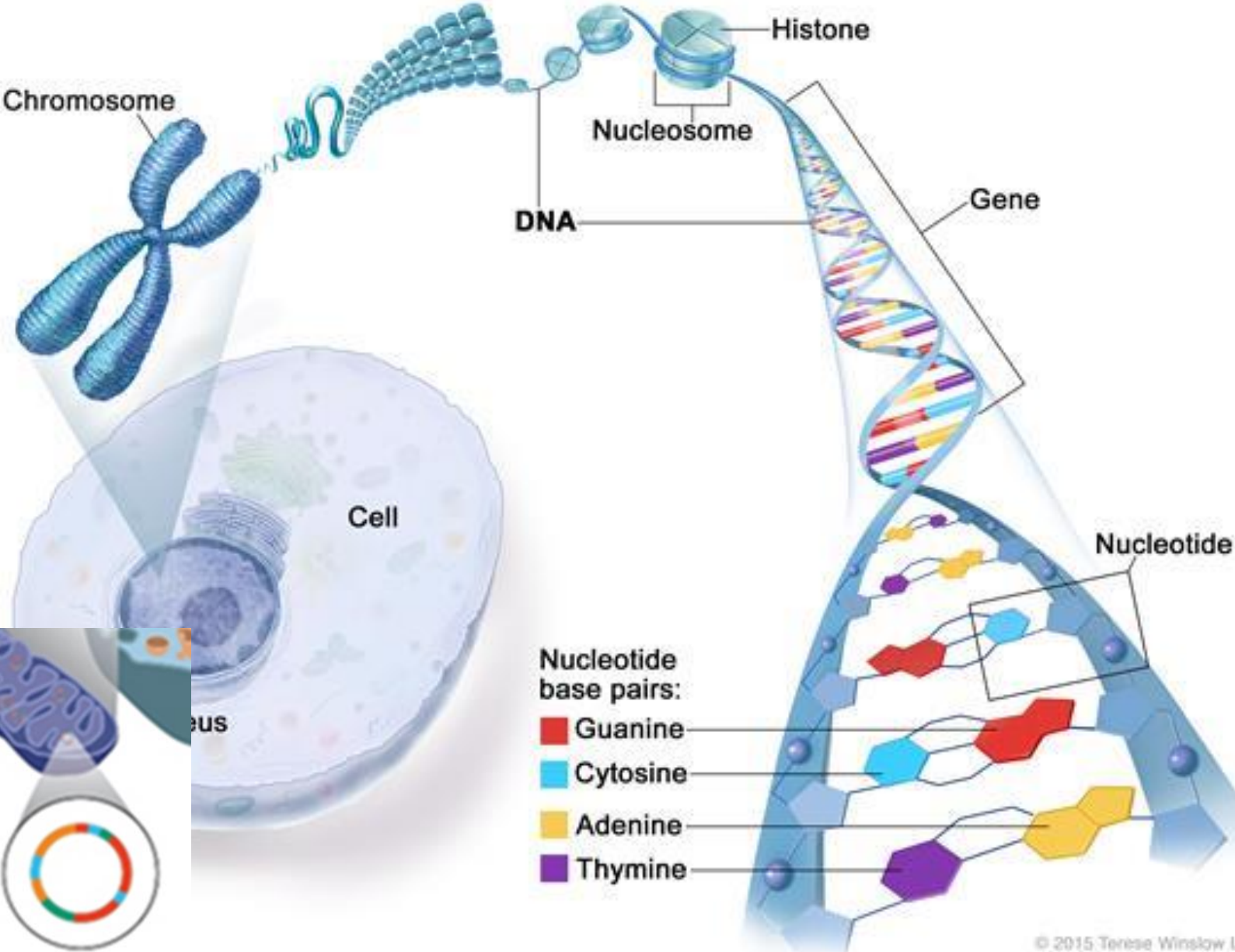
Inversion

A linear segment of DNA within the chromosome becomes oriented in the **reverse direction**, with no molecular loss.

Translocation

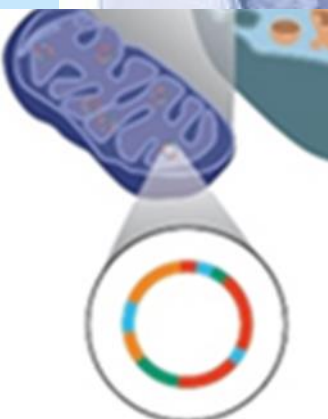
Exchange of segments DNA between **non-homologous chromosomes**

DNA Structure



Nucleotide base pairs:

- Guanine
- Cytosine
- Adenine
- Thymine



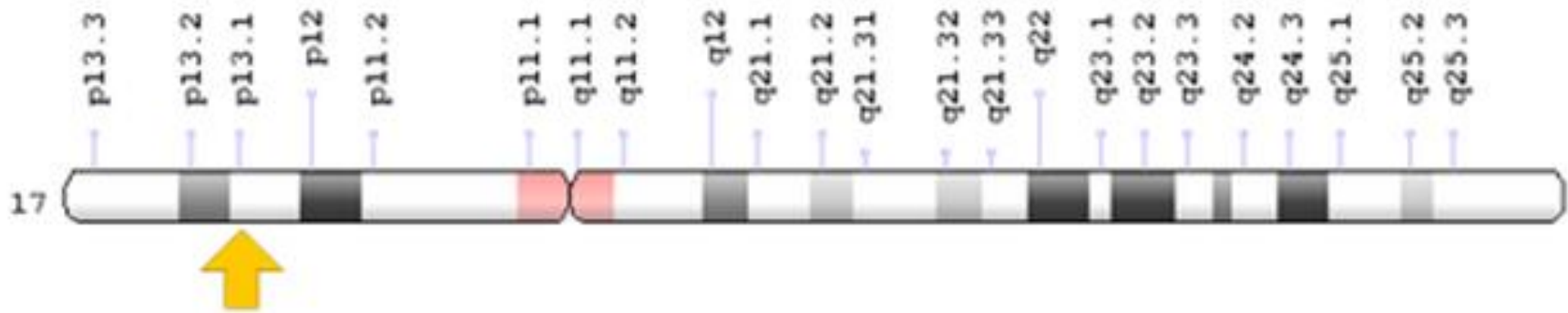
Mitochondrial DNA

GENE: SLC2A4 (GLUT4)

Chromosomal Location

Cytogenetic Location: 17p13, which is the short (p) arm of chromosome 17 at position 13

Molecular Location: base pairs 7,281,735 to 7,288,048 on chromosome 17 (Homo sapiens Annotation Release 108, GRCh38.p7) ([NCBI](#))



Credit: [Genome Decoration Page/NCBI](#)

<https://ghr.nlm.nih.gov/gene/SLC2A4#location>

17

- Canavan disease
- Ovarian cancer
- Miller-Dieker syndrome
- Retinitis pigmentosa
- Tumor protein p53
- Colorectal cancer
- Li-Fraumeni syndrome
- Cystinosis, nephropathic
- Diabetes mellitus, noninsulin-dependent
- Cone dystrophy
- Myasthenic syndrome
- Deafness, autosomal recessive
- Smith-Magenis syndrome
- VLCAD deficiency
- Maturity Onset Diabetes of the Young, type V
- Hypertension, essential, susceptibility to
- T-cell immunodeficiency, alopecia, and nail dystrophy
- Chondrosarcoma, extraskeletal myxoid
- Neurotransmitter transporter, serotonin (anxiety-related)
- Neurofibromatosis, type 1
- Watson syndrome
- Leukemia, juvenile myelomonocytic
- HIV-1 disease, delayed progression of
- Meesmann corneal dystrophy
- Muscular dystrophy, limb-girdle
- Epidermolysis bullosa simplex, recessive
- Pachyonychia congenita, Jackson-Lawler type
- Steatocystoma multiplex
- Wilms tumor, type 4
- Glycogen storage disease (von Gierke disease)
- Parkinsonism-dementia
- Epidermolytic hyperkeratosis
- Patella aplasia or hypoplasia
- Osteogenesis imperfecta
- Ehlers-Danlos syndrome, types I and VIIA
- Osteoporosis, idiopathic
- Ovarian carcinoma antigen
- Neuroblastoma
- Glanzmann thrombasthenia, type A
- Thrombocytopenia, neonatal alloimmune
- CLL/lymphoma, B-cell
- Retinitis pigmentosa
- Pituitary tumor, invasive
- Myocardial infarction, susceptibility to
- Alzheimer disease, susceptibility to
- Myotonia congenita, atypical
- Cramps, familial
- Fetal Alzheimer antigen
- Lung cancer, small-cell
- Campomelic dysplasia with autosomal sex reversal
- Apoptosis inhibitor
- Diabetes mellitus, type II
- Radical fringe

81 million base pairs



- Bernard-Soulier syndrome
- Breast cancer-related regulator of TP53
- Hypermethylated in cancer
- Lissencephaly
- Subcortical laminar heterotopia
- Leber congenital amaurosis, type I
- Medulloblastoma
- Cataract, anterior polar
- Myasthenia gravis, familial infantile
- Bruck syndrome
- Sjogren-Larsson syndrome
- Charcot-Marie-Tooth neuropathy
- Dejerine-Sottas disease
- Van der Woude syndrome modifier
- Choroidal dystrophy, central areolar
- Huntingtin-associated protein
- Psoriasis susceptibility
- Epidermolysis bullosa
- Alzheimer disease, susceptibility to
- Van Buchem disease
- Malignant hyperthermia susceptibility
- Leukemia, acute promyelocytic
- Epidermolytic palmoplantar keratoderma
- Pachyonychia congenita, Jadassohn-Lewandowsky type
- Keratoderma, nonepidermolytic palmoplantar
- Sclerosteosis
- Muscular dystrophy, Duchenne-like, type 2
- Adhalinopathy, primary
- Breast cancer, early onset
- Ovarian cancer
- Leukemia, myeloid/lymphoid or mixed-lineage
- Breast cancer, sporadic
- Gliosis, familial progressive subcortical
- Pseudohypoadosteronism type II
- Spherocytosis, hereditary
- Hemolytic anemia
- Renal tubular acidosis, distal
- T-cell leukemia virus (I and II) receptor
- Dementia, frontotemporal, with Parkinsonism
- Trichodontoosseous syndrome
- Glanzmann thrombasthenia, type B
- Symphalangism, proximal
- Synostoses syndrome, multiple
- Mulibrey nanism
- Growth hormone deficiency
- Myeloperoxidase deficiency
- Cataracts
- Tylosis with esophageal cancer
- Adrenoleukodystrophy, pseudoneonatal
- Deafness, autosomal dominant
- Leukemia, acute myeloid, therapy-related
- Myasthenic syndrome, slow-channel congenital
- Sanfilippo syndrome, types A and B

TRIPLET REPEATS

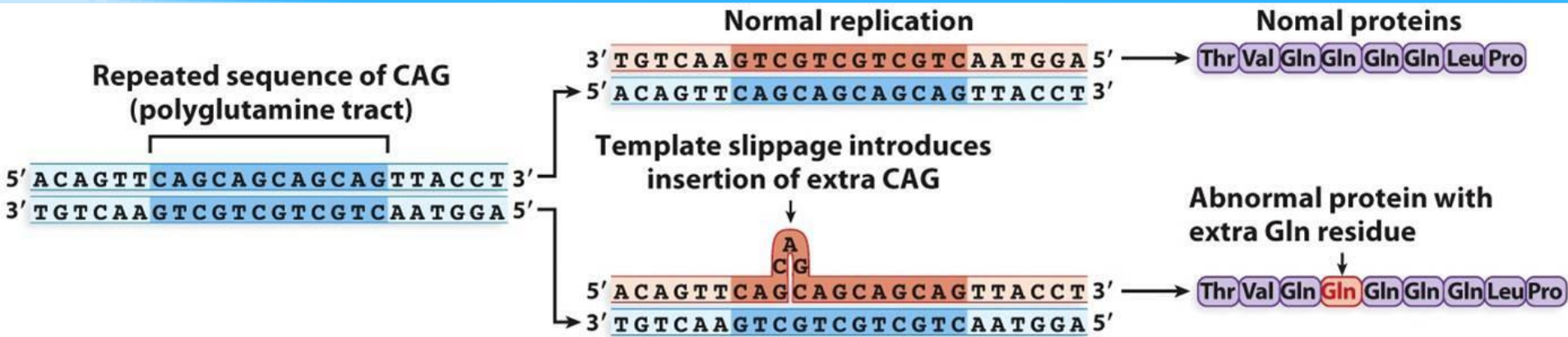


Figure 12-4
Molecular Biology: Principles and Practice
 © 2012 W. H. Freeman and Company

	Normal	Disease	Gene
➤ Huntington disease	CAG 9-35	37-100	Huntingtin
➤ Kennedy disease	CAG 17-24	40-55	androgen receptor
➤ Spino-cerebellar Ataxia	CAG 19-36	43-81	Ataxin 1
➤ Machado Joseph D	CAG 12-36	67-75	SCA
➤ Myotonic dystrophy	CTG 5-35	50-400	DM
➤ Fragile X	CGG CCG GCC 6-50	200-1000	FMR1

SINGLE NUCLEOTIDE POLYMORPHISMS

DNA molecule

Version 1

C T A **A** G T A

Version 2

C T A **C** G T A

Version 3

C T A **G** G T A

Version 4

C T A **T** G T A

▲
SNP

Linked SNPs
outside of gene

no effect on
protein production
or function

Causative SNPs
in gene

Non-coding SNP:
● changes amount of
protein produced

Coding SNP:
● changes amino
acid sequence

Regulatory sequences

Gene

Coding region

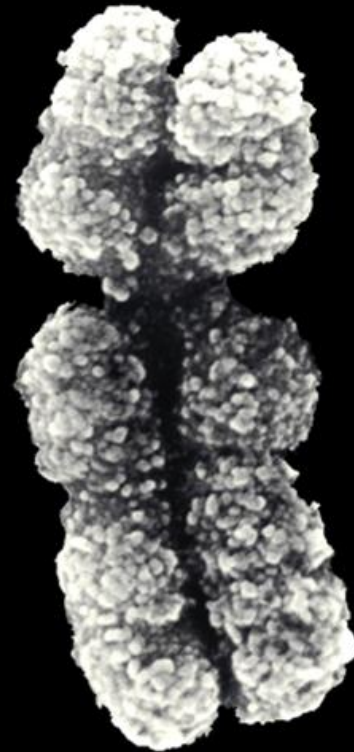
Protein

EPIGENOME

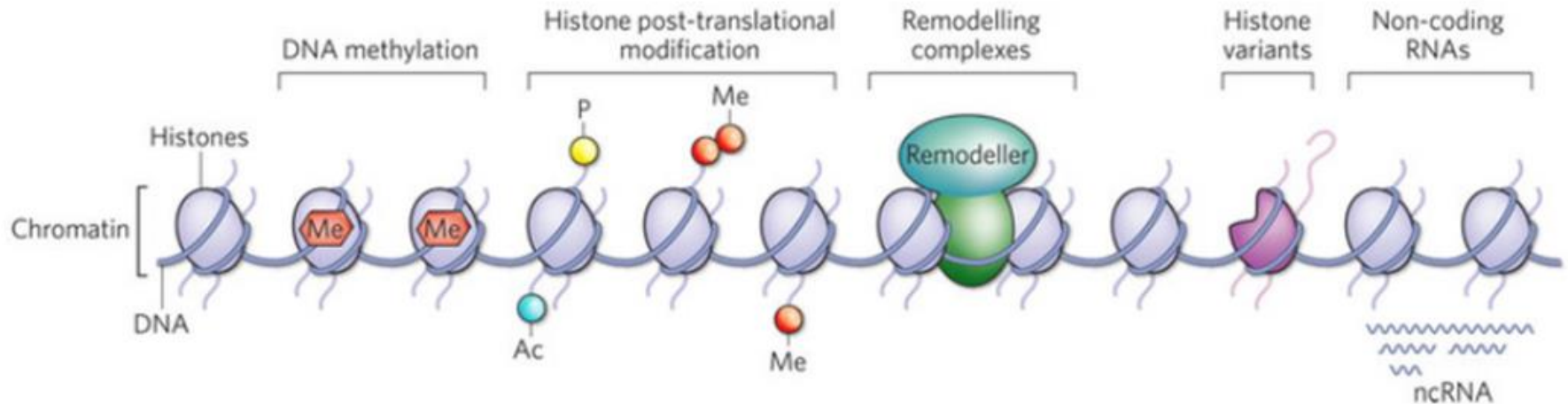
Nature (genome)
Inherited



Nurture (epigenome)
Acquired



EPIGENETIC MODIFICATIONS



C. Dulac, *Nature* 465:7299 (2010)

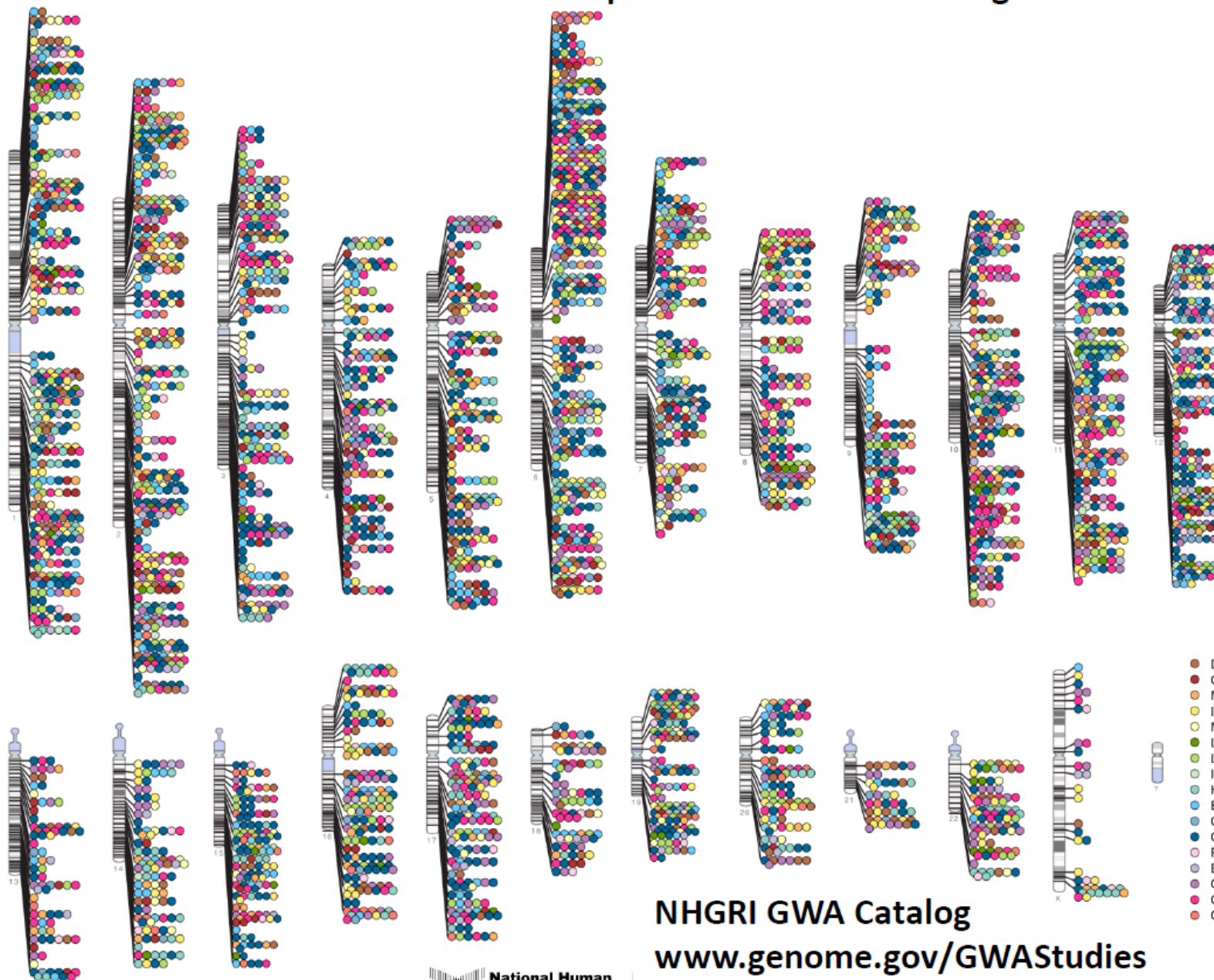
Factors Influencing Epigenetic Modifications

- **Developmental program (includes aging)**
- **Genetic variation**
- **Nutrition**
- **Environment**
- **Drugs**
- **Others**

- Digestive system disease
- Cardiovascular disease
- Metabolic disease
- Immune system disease
- Nervous system disease
- Liver enzyme measurement
- Lipid or lipoprotein measurement
- Inflammatory marker measurement
- Hematological measurement
- Body measurement
- Cardiovascular measurement
- Other measurement
- Response to drug
- Biological process
- Cancer
- Other disease
- Other trait

Published Genome-Wide Associations through 12/2013

Published GWA at $p \leq 5 \times 10^{-8}$ for 17 trait categories

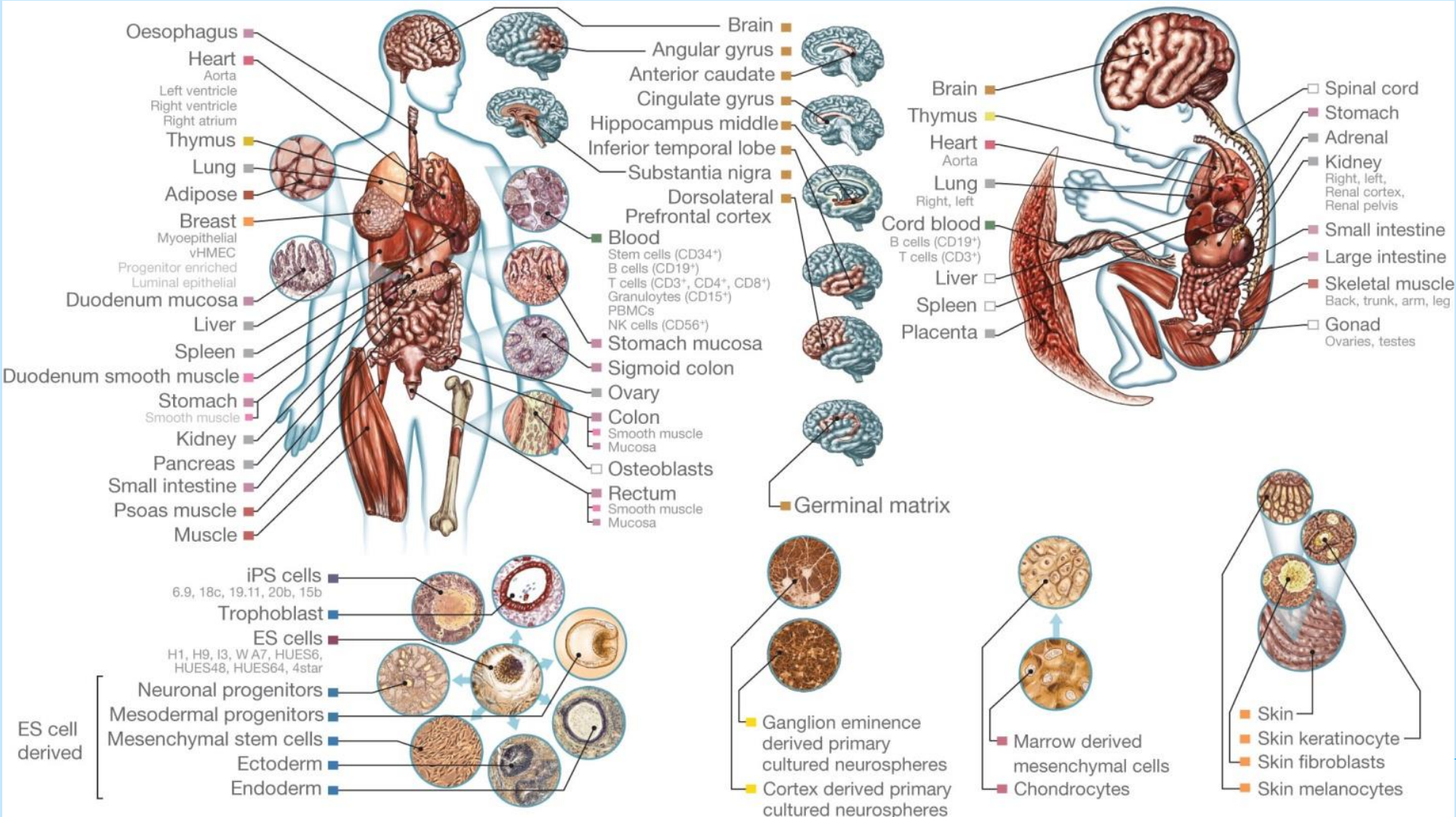


- Digestive system disease
- Cardiovascular disease
- Metabolic disease
- Immune system disease
- Nervous system disease
- Liver enzyme measurement
- Lipid or lipoprotein measurement
- Inflammatory marker measurement
- Hematological measurement
- Body measurement
- Cardiovascular measurement
- Other measurement
- Response to drug
- Biological process
- Cancer
- Other disease
- Other trait

NHGRI GWA Catalog
www.genome.gov/GWAStudies
www.ebi.ac.uk/fgpt/gwas/



Tissues and cell types profiled in the Roadmap Epigenomics Consortium.

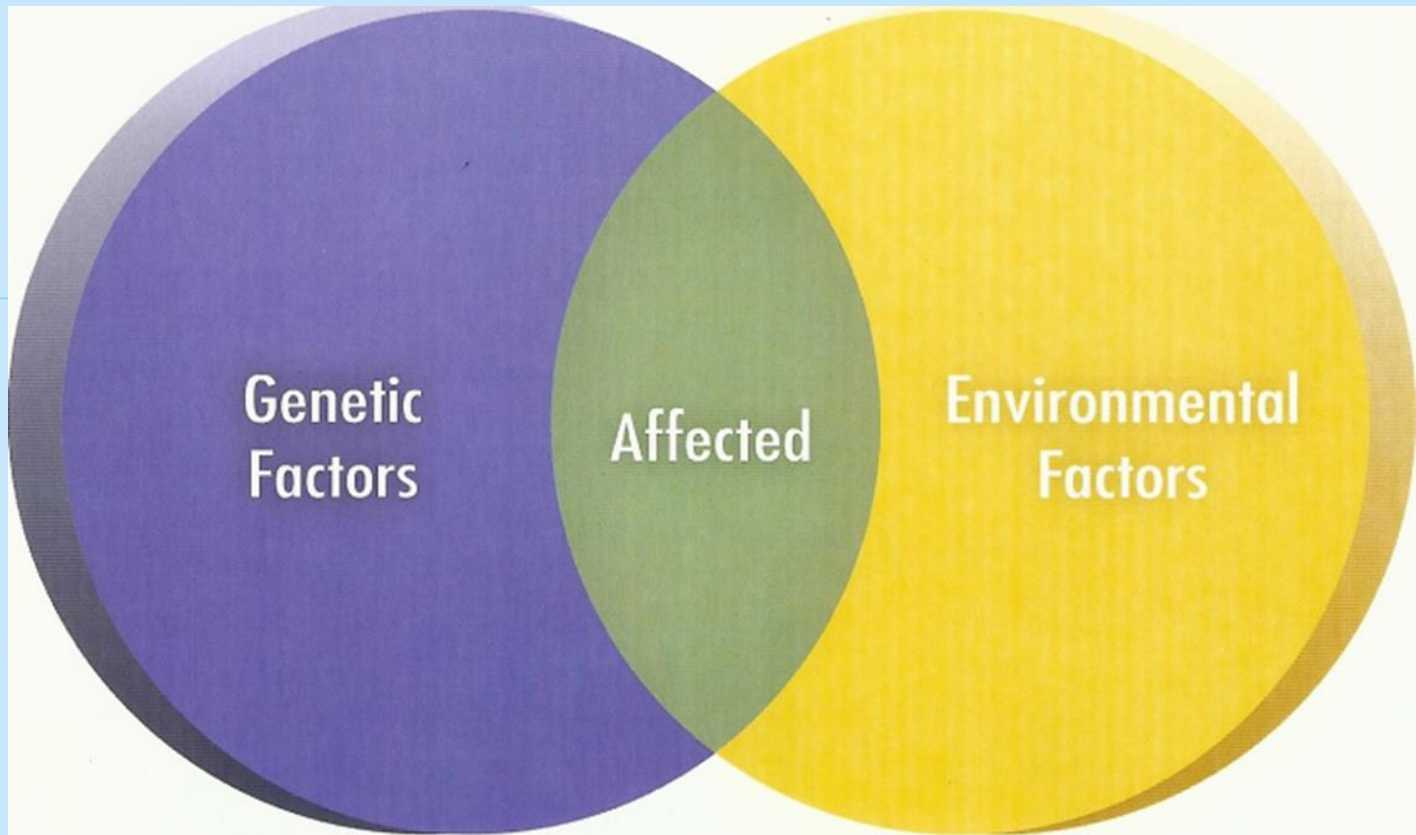


Roadmap Epigenomics Consortium *et al. Nature* **518**, 317-330 (2015) doi:10.1038/nature14248

THE REST OF THE STORY

Term	Definition	Examples of Techniques Used
Transcriptomics	“The quantitative study of all genes expressed in a given biological state” ²⁵	Gene expression microarrays; RNA sequencing ²⁵
Proteomics	Large-scale analysis of all the proteins in an organism, tissue type, or cell (called the proteome). Proteomics can be used to reveal specific, abnormal proteins that lead to diseases	Matrix-assisted laser desorption/ionization ²⁸ ; mass spectroscopy; electrospray ionization ²⁹
Metabolomics (metabolic profiling)	“Measurements of the metabolome, which represents the entire collection of all small-molecule metabolites present in any biological organism” ³⁶	Nuclear magnetic resonance; mass spectrometry ³⁶
Pharmacogenomics	“Pharmacogenomics is the study of an individual’s interaction with a specific drug based upon the genetic make-up of the individual” ³⁹	“Pharmacogenomics studies the influence of genetic variations on the patient’s response to specific drugs, such as the correlation between the efficacy or toxicity of a certain drug and a specific gene expression or a single-nucleotide polymorphism” ³⁹
Bioinformatics	“Information technology as applied to the life sciences, especially the technology used for the collection and analysis of genomic data” ¹¹⁸	

DIABETES AND PERSONALIZED HEALTHCARE PERSPECTIVES

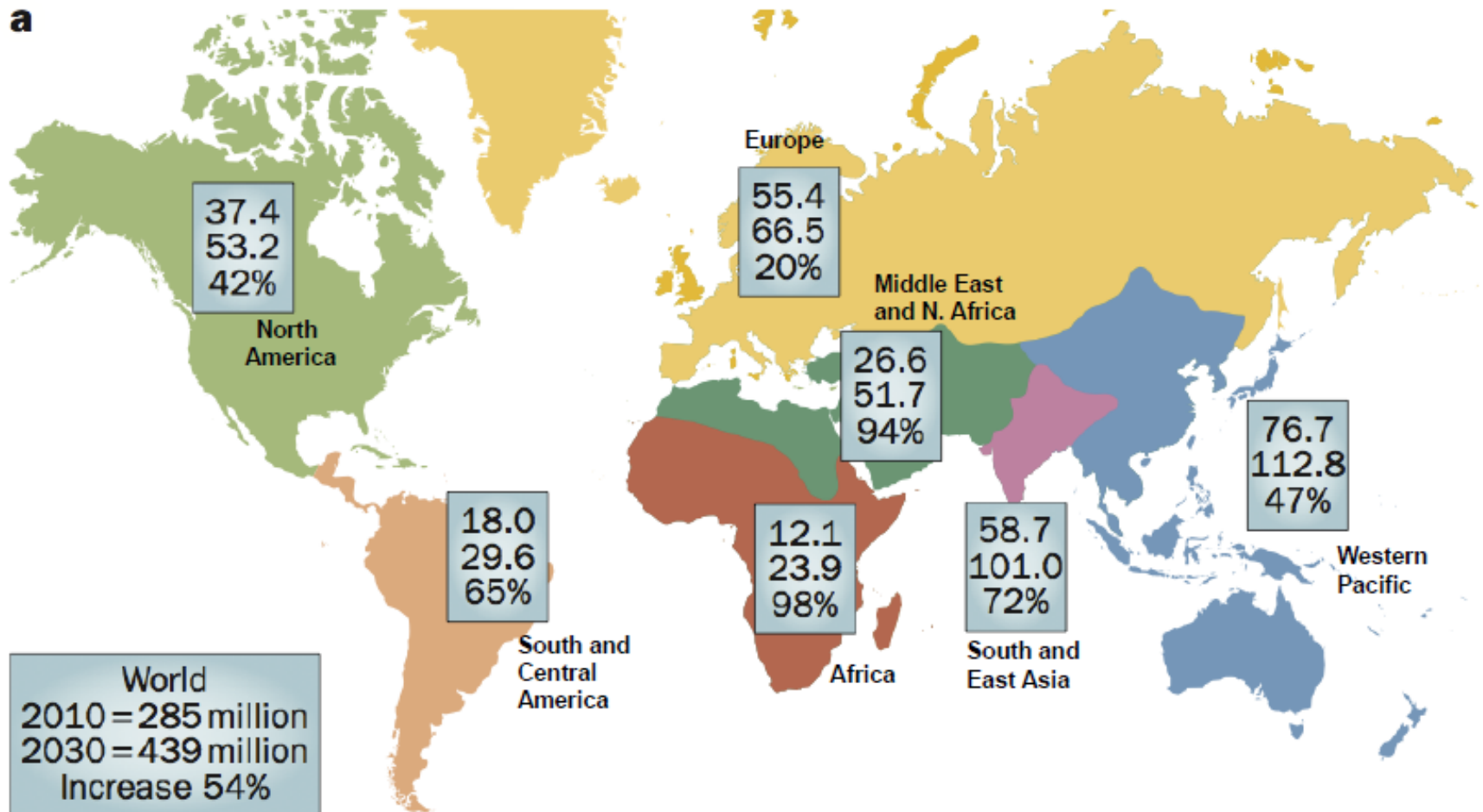


10 LEADING CAUSES OF DEATH, U.S.

- ❖ Heart disease: 599,413
- ❖ Cancer: 567,628
- ❖ Chronic lower respiratory diseases: 137,353
- ❖ Stroke: 128,842
- ❖ Accidents: 118,021
- ❖ Alzheimer's disease: 79,003
- ❖ Diabetes: 68,705
- ❖ Influenza and Pneumonia: 53,692
- ❖ Kidney diseases: 48,935
- ❖ Intentional self-harm: 36,909

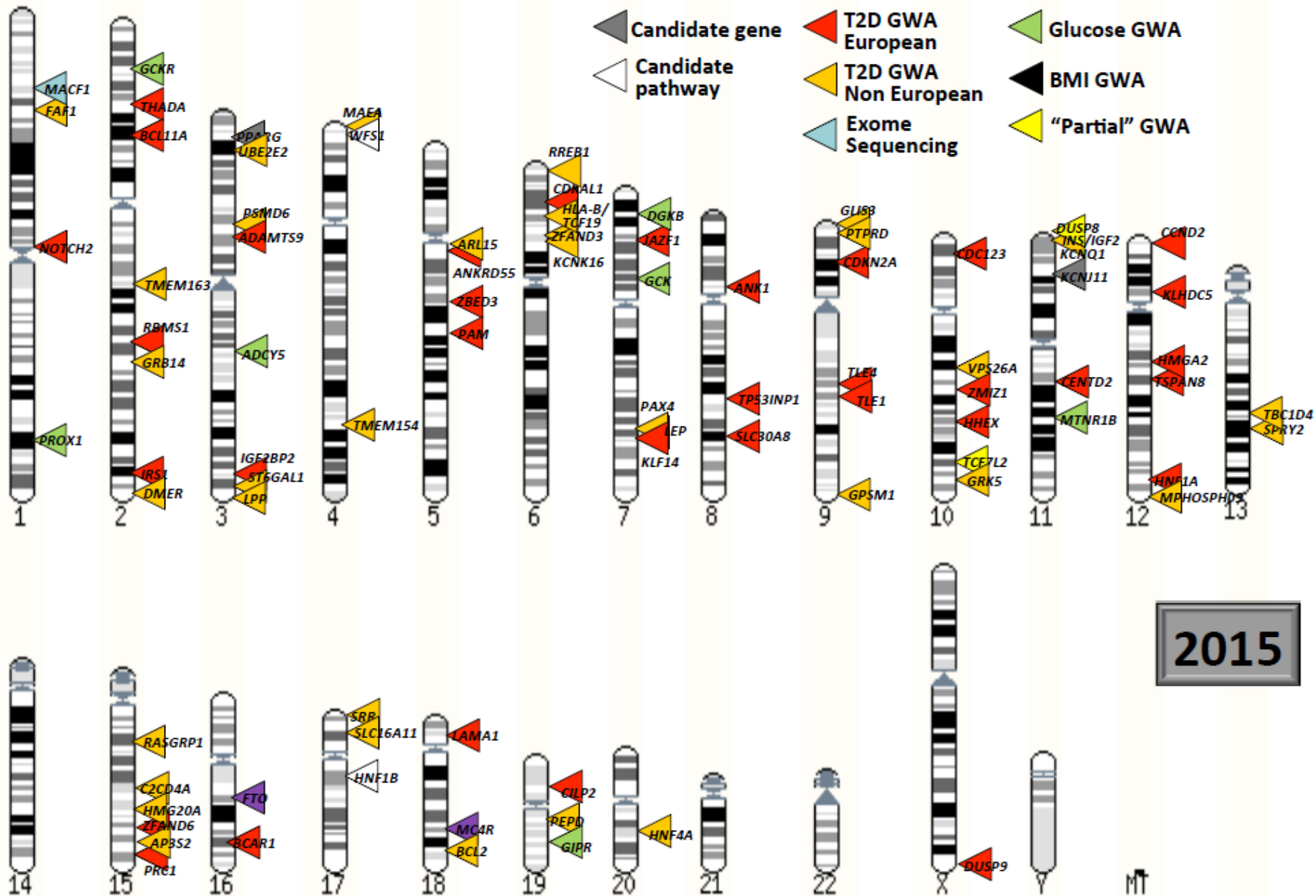
Global projections for the diabetes epidemic: 2010–2030

a



Asia: “diabetes epicenter”

“An integrated approach, taking into account genetic and epigenetic determinants is required for the effective prevention of T2DM beginning from the start of life”



2015

Search <https://ghr.nlm.nih.gov/search?query=diabetes&start=0&tab=gene>

diabetes

X

Filters

- All (162)
- Health Conditions (65)
- Genes (89)
- Help Me Understand Genetics (8)

[MTFE gene](#)

diabetes and deafness (MIDD). People with this condition have *diabetes* and sometimes hearing loss

[IRS1 gene](#)

in this gene are associated with type II *diabetes* and susceptibility to insulin resistance. [provided

[GCGR gene](#)

glucose levels. Defects in this gene are a cause of non-insulin-dependent *diabetes* mellitus (NIDDM



[PDX1 gene](#)

lead to early-onset insulin-dependent *diabetes* mellitus (NIDDM), as well as maturity onset *diabetes* of

[HNF1B gene](#)

the embryonic pancreas. Mutations in this gene result in renal cysts and *diabetes* syndrome and

ESTABLISHED TYPE 2 DIABETES SUSCEPTIBILITY LOCI

Index SNP	Chromosome	Position	Region/gene	Identification	λ_s^*
rs10010131	4	6343816	<i>WFS1</i>	Candidate gene	1.004
rs1801282	3	12368125	<i>PPARG</i>	Candidate gene	1.005
rs757210	17	33170628	<i>HNF1B (TCF2)</i>	Candidate gene	1.002
rs5219	11	17366148	<i>KCNJ11</i>	Candidate gene	1.005
rs7901695	10	114744078	<i>TCF7L2</i>	Linkage peak fine-mapping	1.022
rs10811661	9	22124094	<i>CDKN2A/B</i>	GWA	1.003
rs10946398	6	20769013	<i>CDKAL1</i>	GWA	1.002
rs13266634	8	118253964	<i>SLC30A8</i>	GWA	1.003
rs4402960	3	186994381	<i>IGF2BP2</i>	GWA	1.002
rs5015480	10	94455539	<i>HHEX/IDE</i>	GWA	1.002
rs8050136	16	52373776	<i>FTO[†]</i> 	GWA	1.009
rs2237892	11	2796327	<i>KCNQ1</i>	GWA	1.031
rs10830963	11	92348358	<i>MTNR1B[‡]</i> 	GWA	1.001
rs10923931	1	120319482	<i>NOTCH2</i>	GWA meta-analysis	1.001
rs12779790	10	12368016	<i>CDC123/CAMK1D</i>	GWA meta-analysis	1.002
rs4607103	3	64686944	<i>ADAMTS9</i>	GWA meta-analysis	1.002
rs7578597	2	43586327	<i>THADA</i>	GWA meta-analysis	1.002
rs7961581	12	69949369	<i>TSPAN8/LGR5</i>	GWA meta-analysis	1.001
rs864745	7	28147081	<i>JAZF1</i>	GWA meta-analysis	1.001

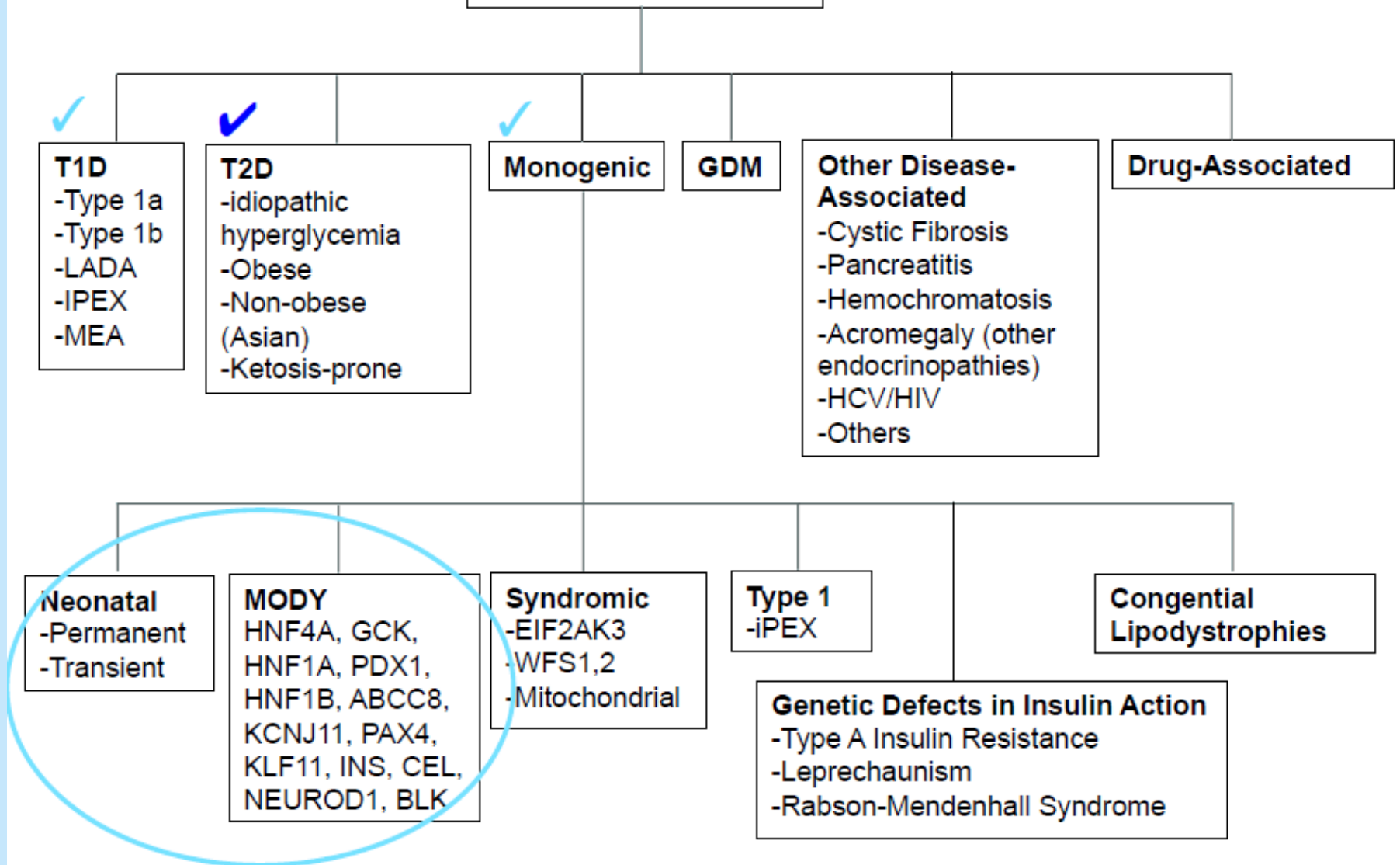
*The sibling recurrence risk ratio calculated in European populations, with the exception of the *KCNQ1* locus, which was based on East Asian populations.

[†]The primary association for this locus is with body mass index.

[‡]The primary association for this locus is with fasting glucose levels.

GWA—genome-wide association; SNP—single nucleotide polymorphism.

Diabetes Mellitus (2015)

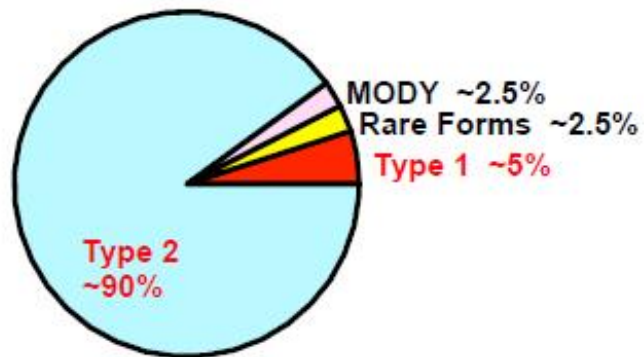


Optimal Treatments for Monogenic Diabetes by Subtype

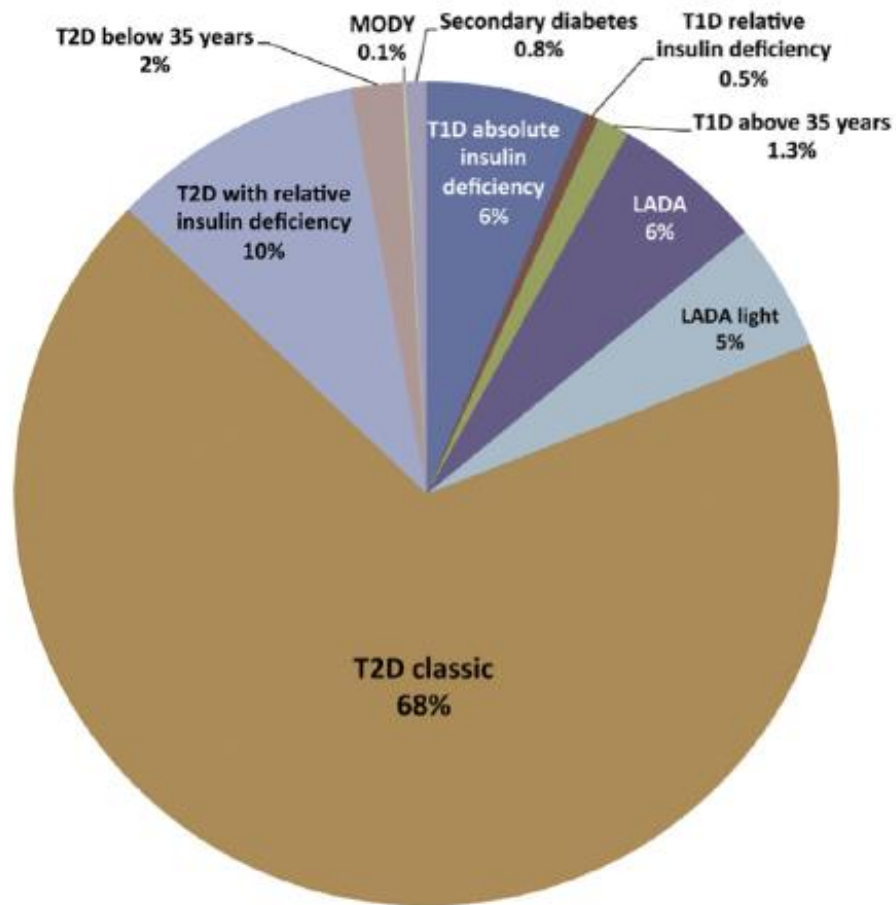
Table 1. Optimal treatments for monogenic diabetes by subtype

Monogenic diabetes subtype	Distinguishing clinical features	Examples of causal genes	Optimal treatment
Examples of more common subtypes			
GCK-MODY	Mild fasting hyperglycaemia	<i>GCK</i>	Diet alone
HNF1A-MODY	Young onset diabetes	<i>HNF1A</i>	Low dose sulphonylurea
Neonatal diabetes	Diabetes diagnosed before 6 months	<i>KCNJ11, ABCC8</i>	High dose sulphonylurea Insulin
HNF4A-MODY	Young onset diabetes, increased birth weight and macrosomia	<i>INS</i> <i>HNF4A</i>	Low dose sulphonylurea
Examples of rarer subtypes with extrapancreatic features			
HNF1B-MODY	Renal cysts, genitourinary abnormalities, exocrine pancreatic insufficiency	<i>HNF1B</i>	Early insulin
Mitochondrial diabetes	Deafness, short stature, pigmentary retinopathy	<i>MTTL1</i>	Sulphonylurea initially but rapid progression to insulin requirement
Wolfram syndrome	Optic atrophy, diabetes insipidus, deafness, renal tract abnormalities, neurological abnormalities	<i>WFS1</i>	Insulin
TRMA syndrome	Megaloblastic anaemia, deafness	<i>SLC19A2</i>	Thiamine ± sulphonylurea ± early insulin

GCK, glucokinase; HNF4A, hepatocyte nuclear factor 4A; MODY, maturity-onset diabetes of the young; TRMA, thiamine responsive megaloblastic anaemia.



The Revised Diabetes Pie (2013)



ANDIS project (Sweden):
Reclassification based
on genetic markers and
biomarkers

PRECISION MEDICINE DIABETES CLINIC

- Prevention
- Diagnosis
- Treatment
- Monitoring



Klonoff, D. C. (2015). Precision medicine for managing diabetes. *Journal of Diabetes Science and Technology*, 9(1), 3-7. doi:10.1177/1932296814563643

GREATEST RISK CONSIDERATIONS

- Fasting plasma glucose 100-125 mg/dl (impaired fasting)
- Plasma Glucose 2 hours after 75g. oral glucose challenge of 140-199 mg/dl (impaired glucose tolerance)
- Hemoglobin A1c

Not All persons at the Same RISK for microvascular and macrovascular complications



THE GENETIC EYE

FAMILY HISTORY:

IN GENETICS, THE FAMILY IS THE PATIENT

- A comprehensive family history is an important first step in the analysis of any disorder, whether or not the disorder is known to be genetic
- It can be critical in diagnosis
- May show that a disorder is hereditary
- Can provide information about the natural history of a disease and variation in its expression
- Clarify the pattern of inheritance
- The diagnosis of a hereditary condition allows the risk in other family members to be estimated, so that proper management, prevention and counseling can be offered to the patient AND the family

Look for these Red Flags

- 🚩 Family history of multiple affected members
- 🚩 Onset of disease at age earlier than expected
- 🚩 Condition in the less-often affected sex
- 🚩 Disease in the absence of known risk factors
- 🚩 Ethnic predisposition of genetic disorders
- 🚩 Close biological relationship between parents
- 🚩 Developmental delays
- 🚩 Unexplained mental retardation
- 🚩 One or more major malformations
- 🚩 Recurrent pregnancy losses (>2)
- 🚩 Unexpected drug reactions/responses

QUESTIONS: MONOGENIC DIABETES

1. Diabetes diagnosed before one year of age; or
2. Type 1 diabetes and a parent or child with type 1 diabetes; or
3. Diabetes other than Type 1 diagnosed at 30 years of age or less; or
4. Type 2 diabetes diagnosed by 45 years of age, not extremely overweight at diagnosis, and 2 or more relatives diagnosed with diabetes by 50 years of age; or
5. Diabetes along with other features, such as birth defects, intellectual disability, deafness or blindness; or
6. Lean and have or have had gestational diabetes; or
7. Diabetes suspected by your physician to be monogenic or unusual in some way

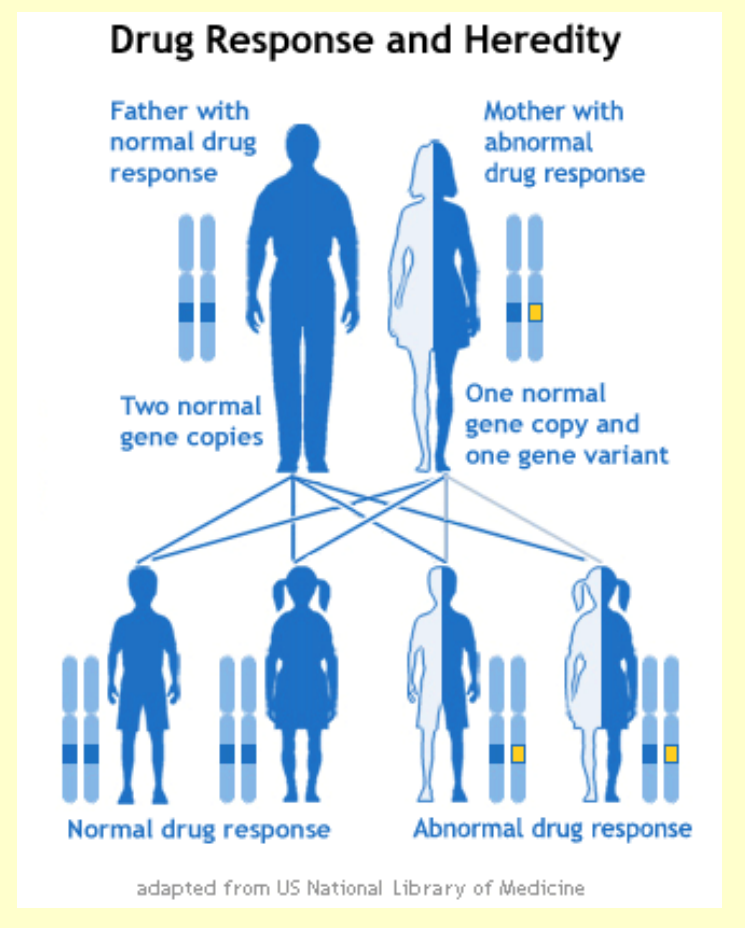


Think Genomics and Drugs!

Pharmacogenomics is how a person's genomic makeup influences their response to drugs

ANY OF THESE MEDICATION RESPONSES at RECOMMENDED DOSES

- ★ Unexpected reactions
- ★ Toxicity
- ★ No Response
- ★ Limited response
- ★ Some decreased efficacy
- ★ Decreased efficacy



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

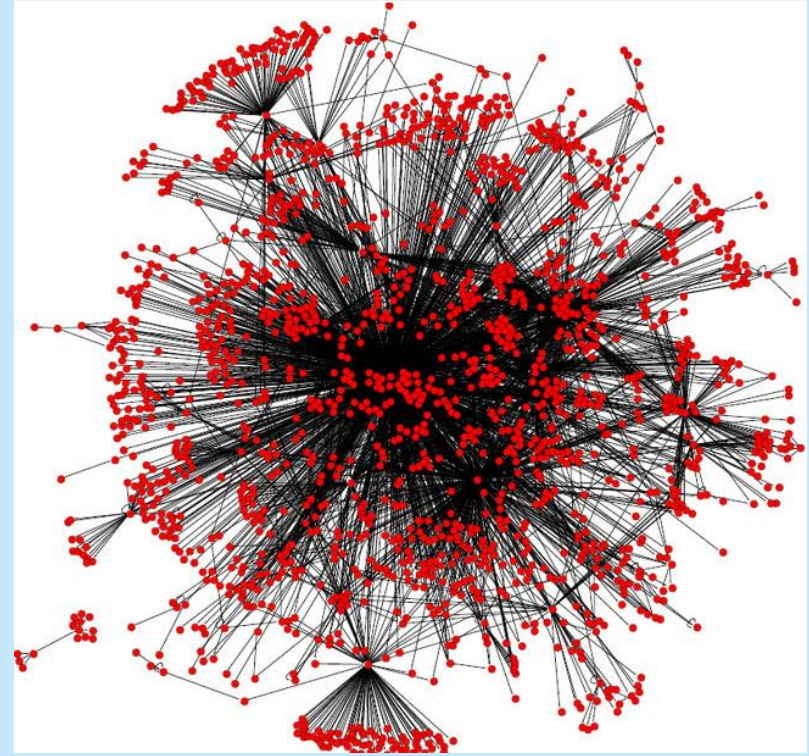
<https://www.pharmgkb.org/>

DRUG RESPONSES



Spear B, Heath-Chiozzi M, Huff J (2001). Clinical application of pharmacogenetics. *Trends in Molecular Medicine*. 2001;7(5): 201-204.

<http://healthyblackmen.org/2012/03/11/physicians-patients-pills-part-i/>



Genetics and Complex Disease

CASE STUDY

GENETICS AND COMPLEX DISEASE

- Although almost every disease has a genetic component, most diseases are not inherited in predictable, single-gene patterns (dominant, recessive, x-linked). That is why most of the conditions that burden us from a public health perspective – **caries, diabetes, heart disease** – are called "complex."
- Complex diseases arise from a combination of factors, including the interaction of multiple genes, lifestyle choices, and environmental exposure.
- Although these conditions are more frequent, their patterns of transmission are more elusive because the disease traits don't segregate neatly from generation to generation, as do single-gene disorders.
- Instead, complex traits aggregate or cluster within families, and it is difficult to predict who will be affected and how the disease will express itself.

A DENTAL PRACTICE SERVING AN ADULT POPULATION OF 2,000 CAN EXPECT TO ENCOUNTER 40-80 PERSONS WITH DIABETES, ABOUT HALF OF WHOM WILL BE UNAWARE OF THEIR CONDITION

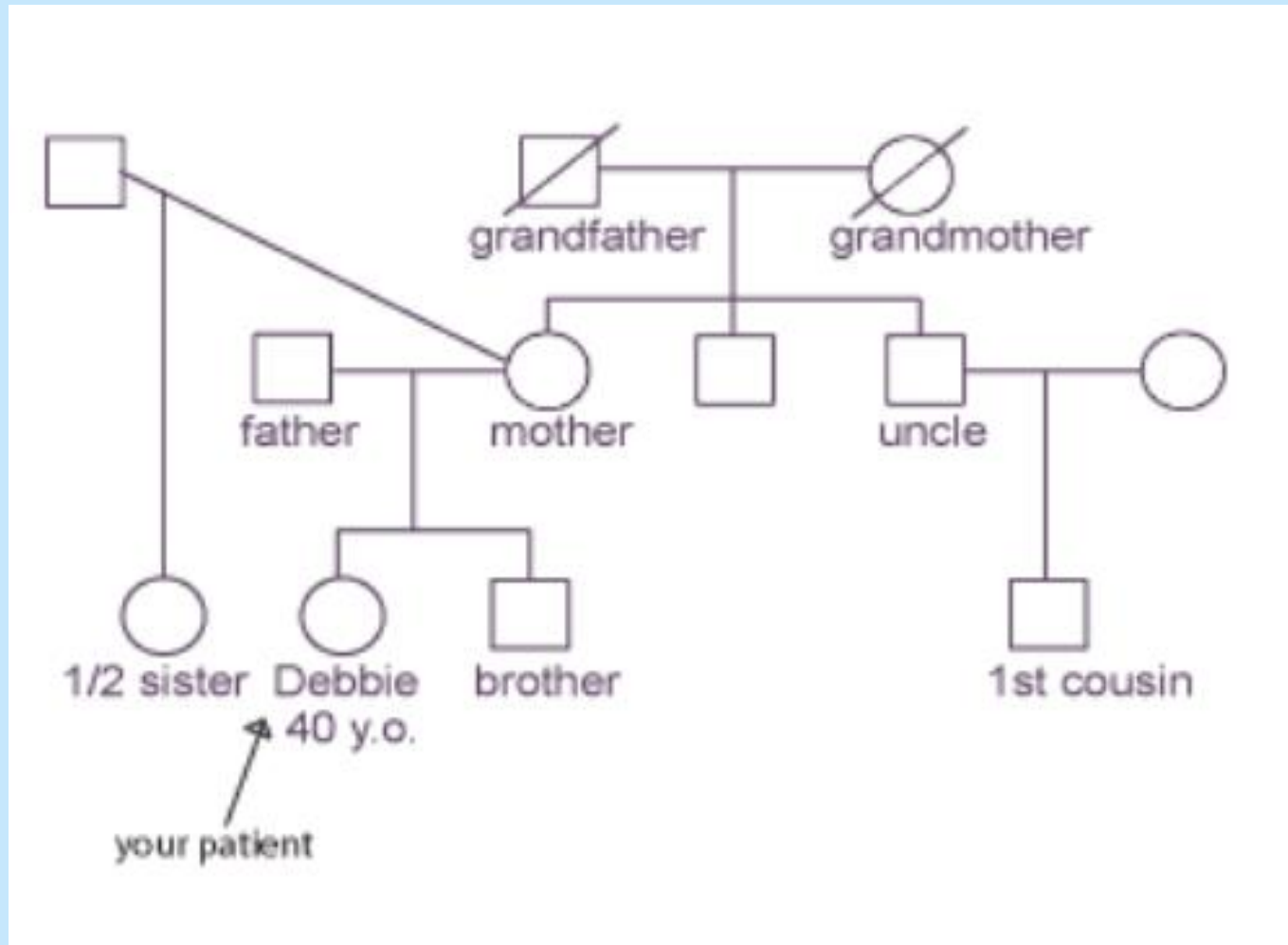
The most marked oral complications seen in uncontrolled diabetic patients includes:

- periodontal disease (which is more severe and has a higher prevalence than that seen in the non-diabetic),
- xerostomia,
- burning mouth syndrome,
- candidiasis,
- delayed and abnormal wound healing,
- increased propensity to infection,
- diminished salivary flow, and
- salivary gland enlargement.

DEBBIE

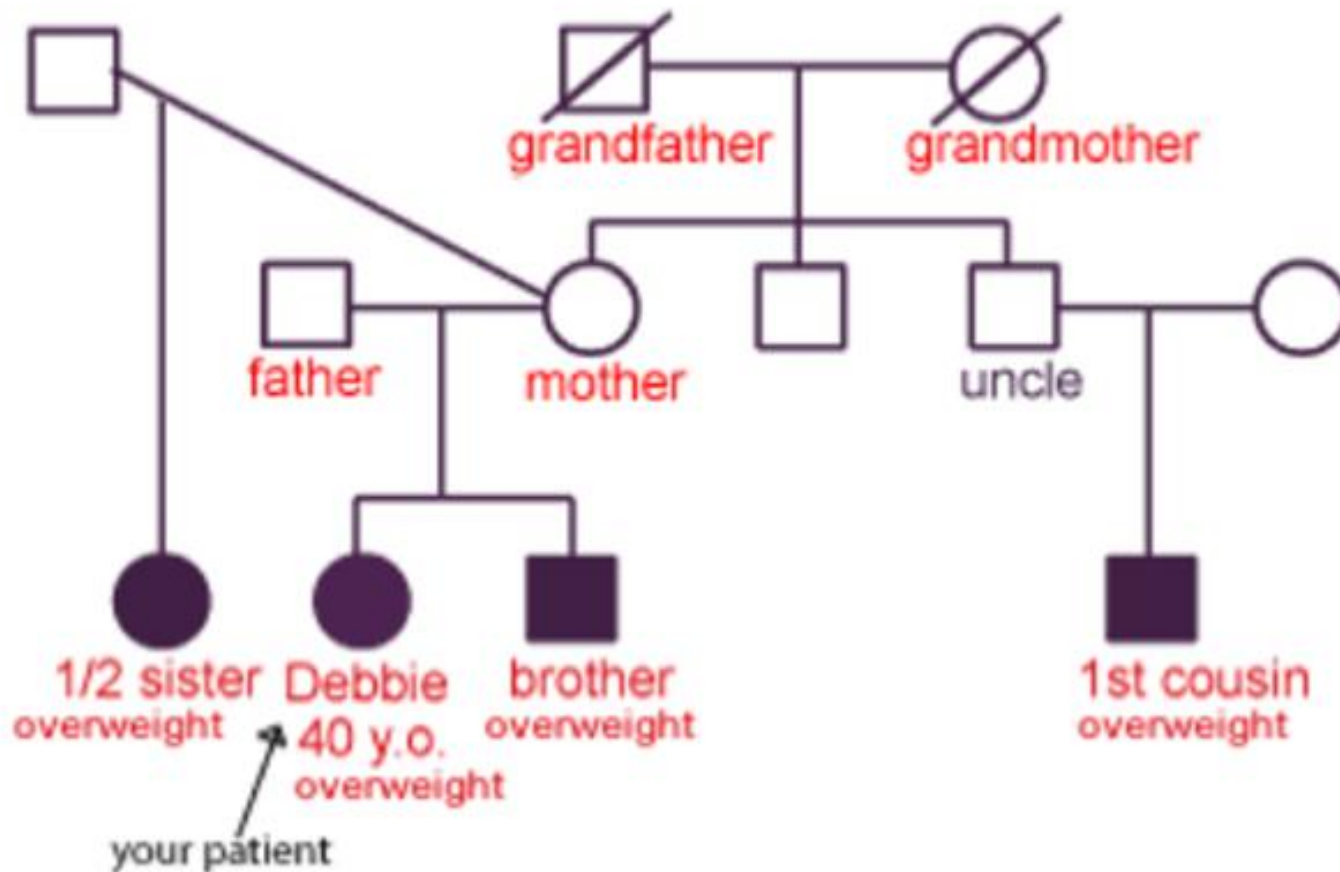
A 40-year-old Hispanic woman who comes to you to replace a crown. She has never been to your office before, and during oral examination, you find periodontal disease, several canker sores, and a carious lesion, even though she claims to have excellent home care. She makes an appointment for a restoration the following week. At her next appointment, she also complains of xerostomia, but attributes it to allergies.

TAKING THE FAMILY HISTORY WILL HELP PUT DEBBIE'S INTRAORAL FINDINGS INTO A BROADER PERSPECTIVE. GO THROUGH THE QUESTIONS BELOW TO CONSTRUCT DEBBIE'S GENETIC FAMILY HISTORY.



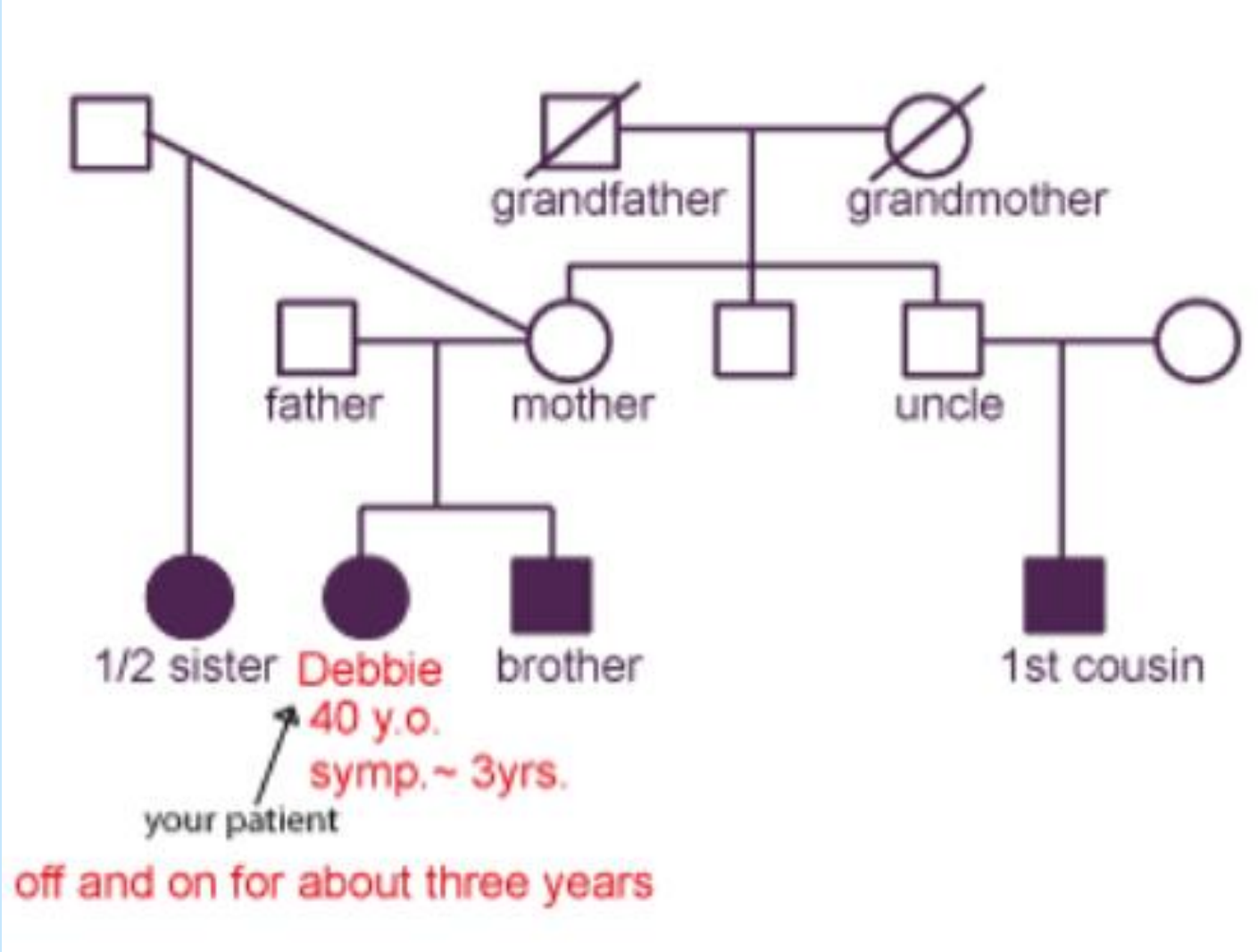
IS DEBBIE OVERWEIGHT?

IS THIS COMMON IN THE FAMILY?

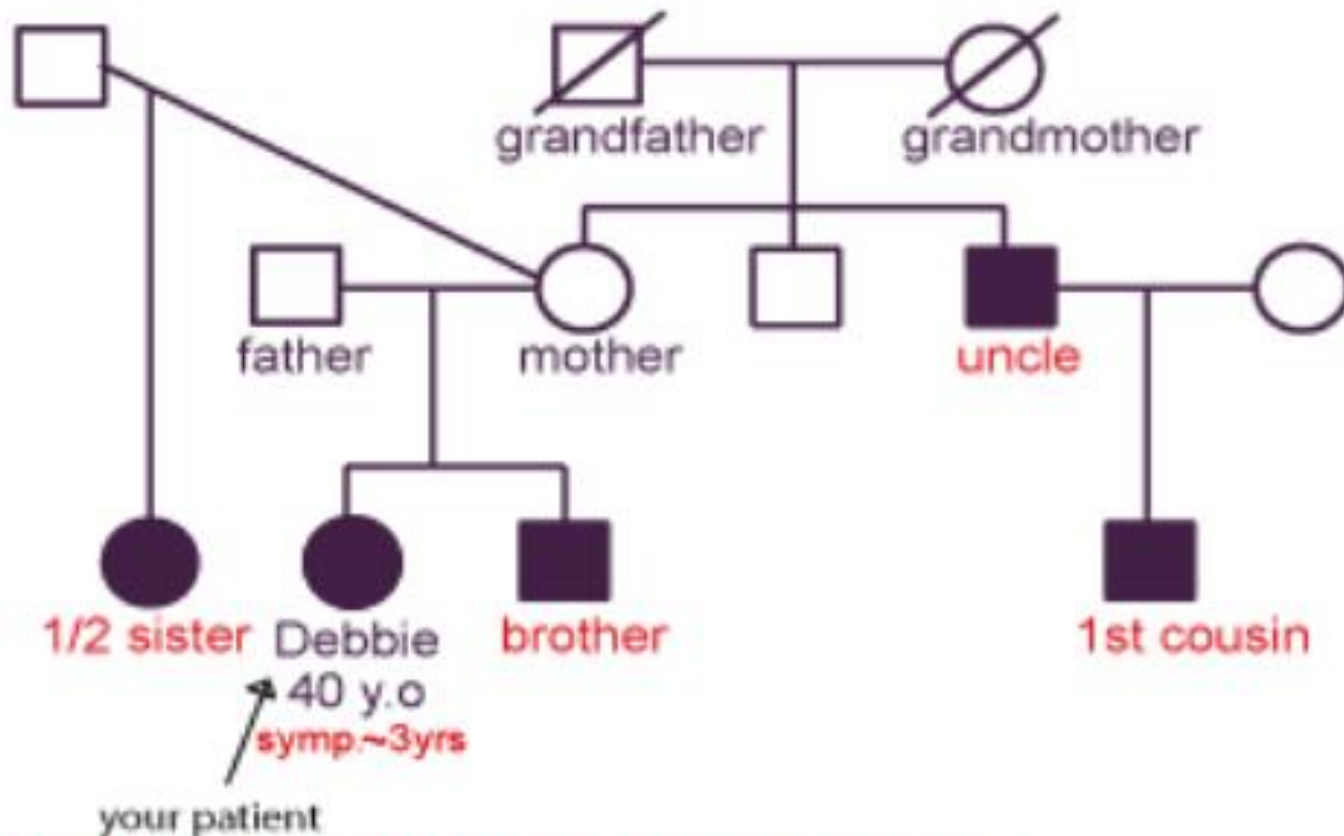


virtually all, except the uncle, who is a health nut.

HOW LONG HAS DEBBIE HAD THESE SYMPTOMS?

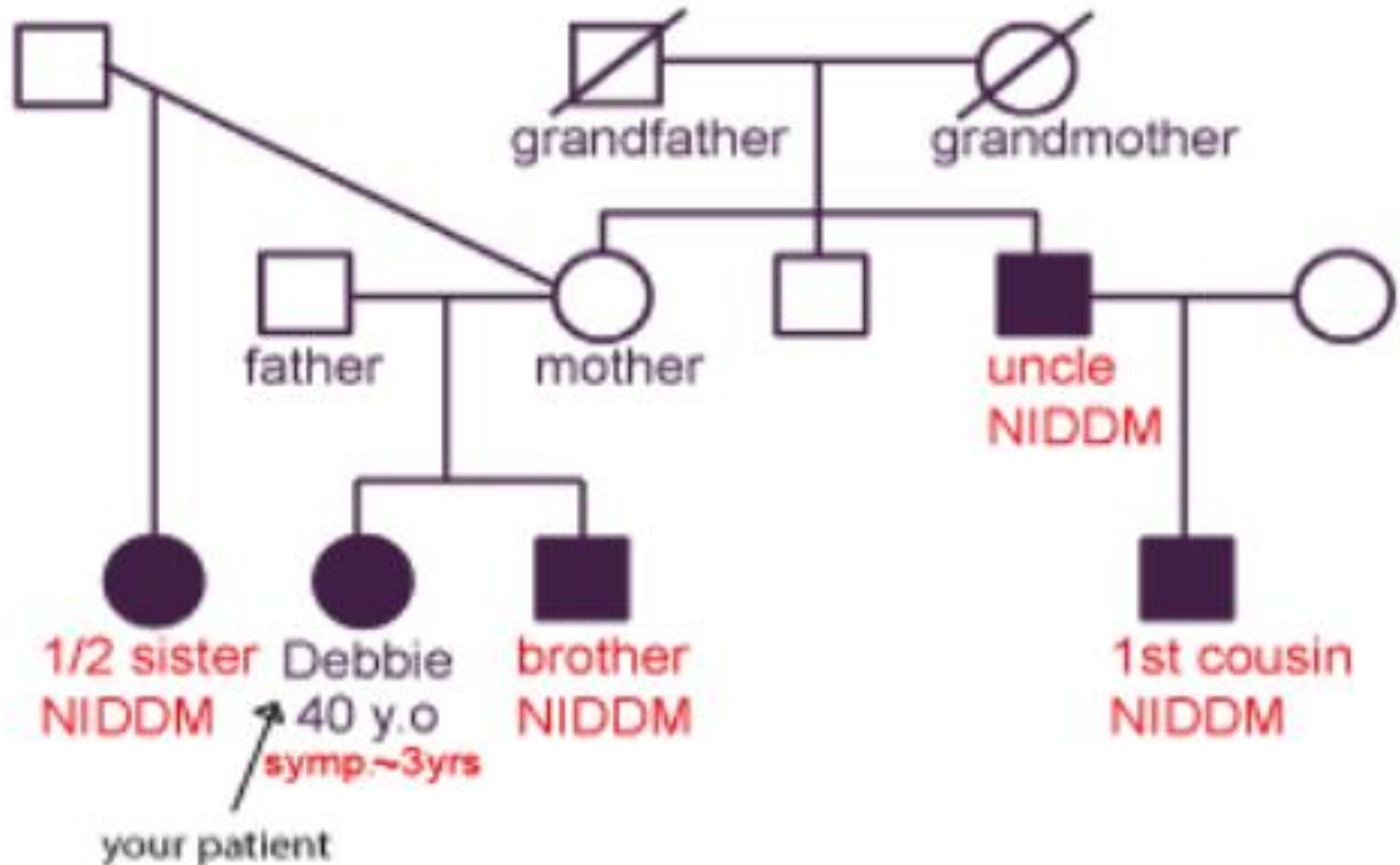


HAS DEBBIE OR ANYONE ELSE IN HER FAMILY BEEN DIAGNOSED WITH DIABETES?

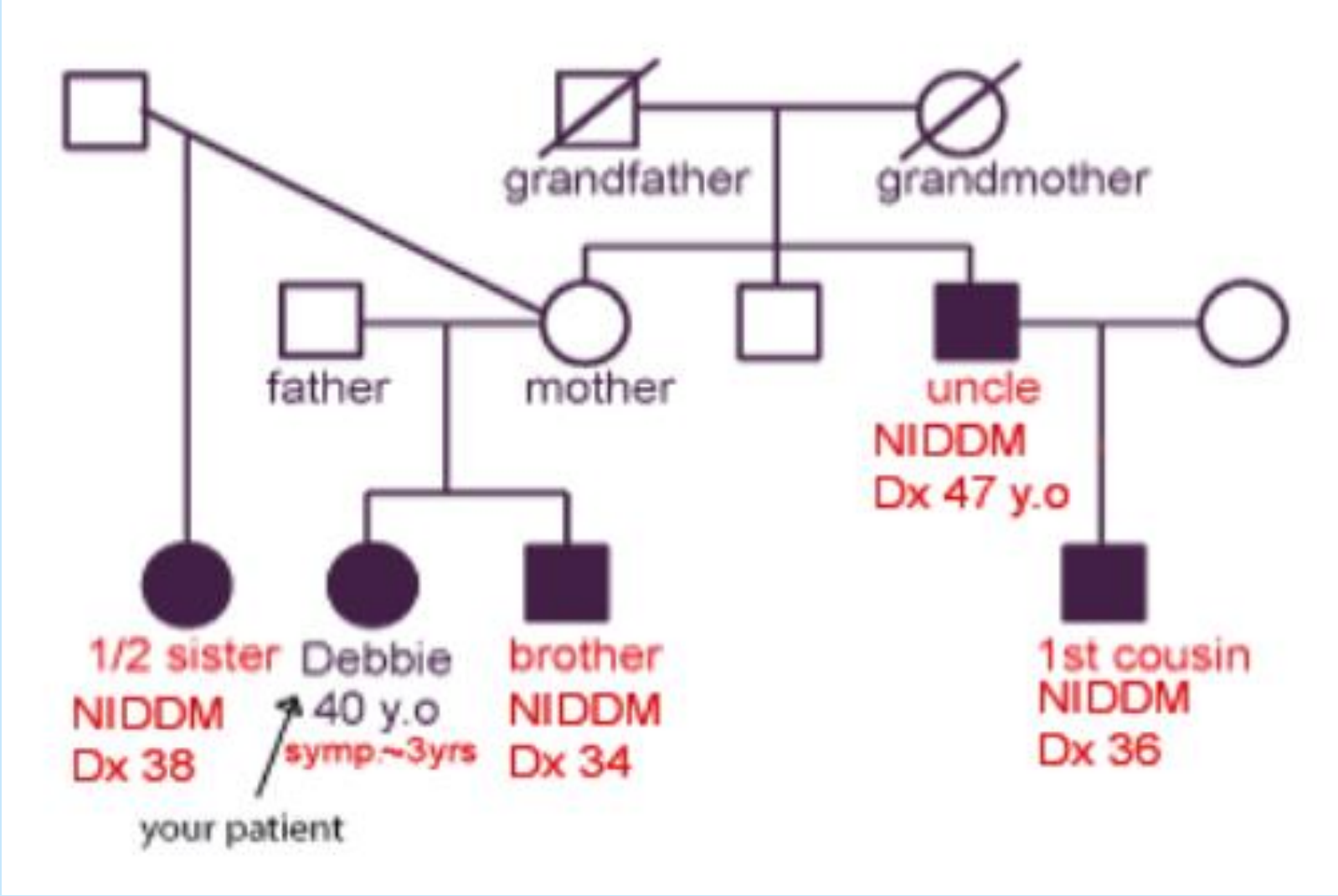


brother, uncle, and 1st cousin on mother's side,
1/2 sister from mother's second marriage

WHAT TYPE - INSULIN DEPENDENT OR NON-INSULIN DEPENDENT?



HOW OLD WERE THEY WHEN THEY WERE DIAGNOSED?

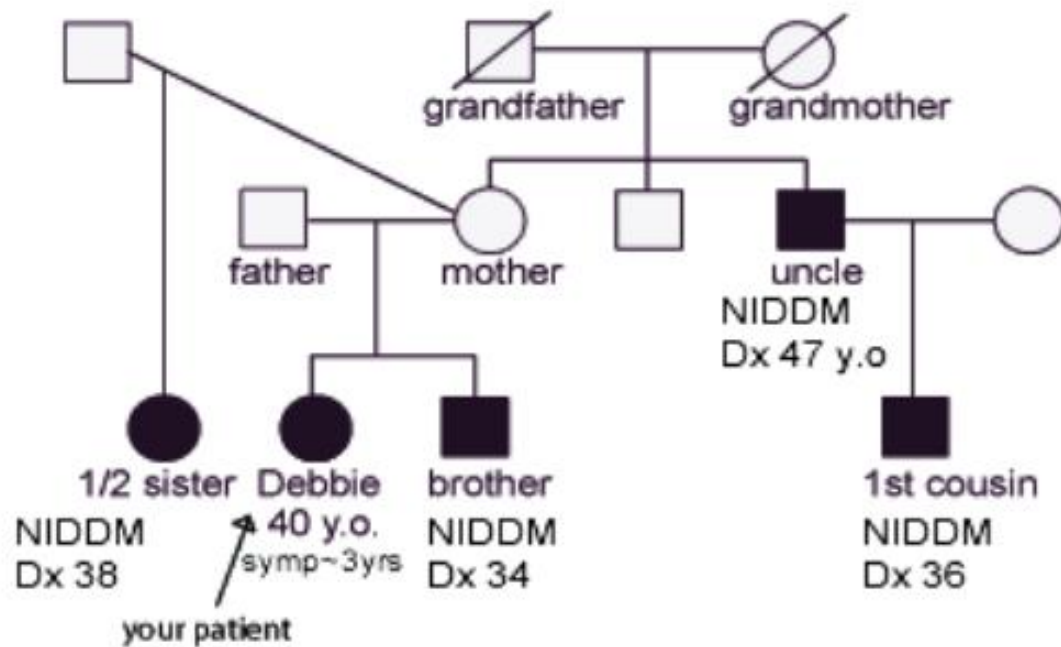


QUESTIONS: MONOGENIC DIABETES

1. Diabetes diagnosed before one year of age; or
2. Type 1 diabetes and a parent or child with type 1 diabetes; or
3. Diabetes other than Type 1 diagnosed at 30 years of age or less; or
4. Type 2 diabetes diagnosed by 45 years of age, not extremely overweight at diagnosis, and 2 or more relatives diagnosed with diabetes by 50 years of age; or
5. Diabetes along with other features, such as birth defects, intellectual disability, deafness or blindness; or
6. Lean and have or have had gestational diabetes; or
7. Diabetes suspected by your physician to be monogenic or unusual in some way

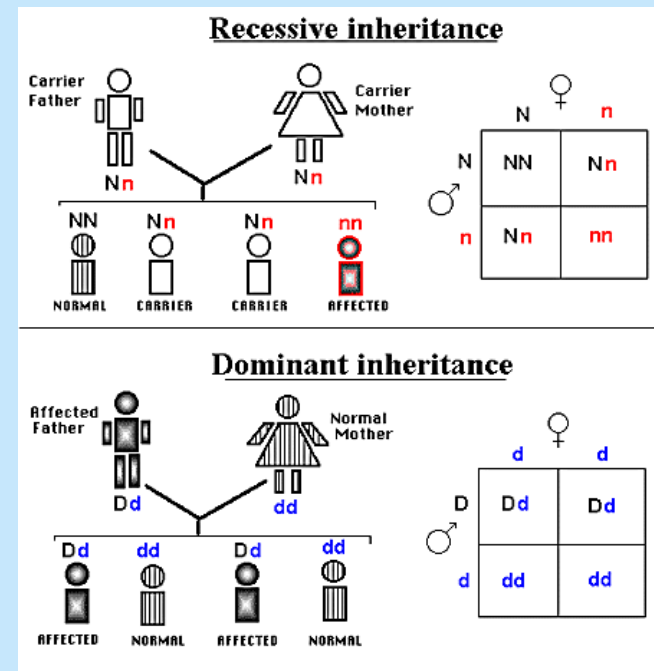
ENVIRONMENTAL CONSIDERATIONS:

- City and Country where born
- Culture
- Diet
- Lifestyle
- Occupation
- Stress
- Exposure to chemicals (pesticides, insecticides, paints, solvents, asbestos, hair dyes, etc...)
- Tobacco Use
- Alcohol Use



Does a clear pattern of inheritance emerge

- A. Autosomal Dominant Inheritance
- B. Autosomal Recessive Inheritance
- C. X-Linked Recessive Inheritance
- D. None of the above





Thank-You

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