

Genes $\left\{ \begin{array}{l} \text{Coding - Exons} \rightarrow 1.5\% \text{ of Human genome} \\ \text{Non-Coding - INTRONS - } \end{array} \right.$



Regulation of genes.

i) Centromeres

ii) Telomeres

iii) Transposons (Jumping genes) (33%)

func - chromatin organization

iv) Non-Coding RNAs

a) NTicso RNA - miRNA - 21-30 oligonucleotide

- gene silencing RNA (post-transcription)

- 5% of human genome

b) siRNA (small interfering RNA)

↓
exogenous RNA (invitro)

↓
To study gene function. - knock down technique

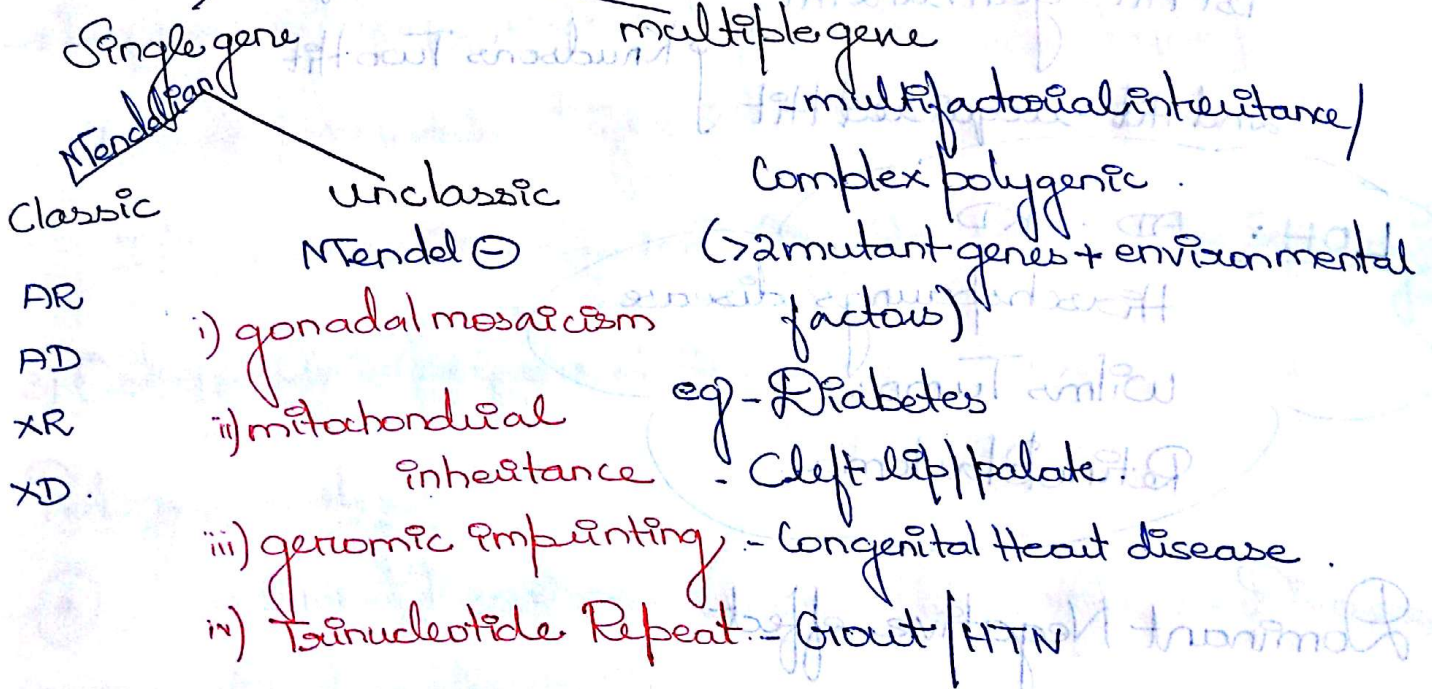
q LINC - RNA - Long intervening non coding
> 200 nucleotide

- associated with Paser-Body formation

Bau bodies

d) pi-RNA - m/c - Non coding RNA in Human genome
piwi - Interacting.

Genetic disorders -



- m/c pattern of inheritance - Autosomal dominant (Vertical)

- m/c pattern in inheritance of metabolism - AR - (Horizontal)

AR
Aa - Carrier
aa - disease

AD
Aa - disease
aa - disease
(die in utero)

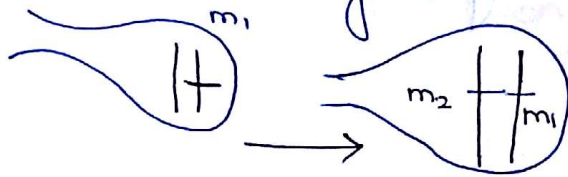
AD:

AR:

i) Reduced Penetrance

eg - Retinoblastoma

ch - 13q14



Rb - Two Hit Hypothesis - Homozygous
Loss of Heterozygosity - LHO. LOH

1st Hit - germline Hit

2nd Hit - acquired Hit

Knudson's Two Hit

LOH: AD - PKD

Hirschsprung's disease

Wilms Tumor

Retinoblastoma

ii) Dominant Negative effect.

eg Mutant - protein/gene

↓
affecting N genes/proteins.

M/c inherited Connective tissue disorder - Marfan's Syndrome

Fibrillinogen - C15q

Revised Ghent's Criteria:

i) Family H/o Marfan

ii) Cardinal features

iii) Fibrillinogen - 1 mutation

m/c - cause of death - aortic dissection

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1) Ehler's Danlos Syndrome - Rubber man Syndrome

Hyperextensibility of Joints + Skin Hyperextensibility

Collagen m/c collagen - III

m/c Subtype of EDS - Type - III

→ Rupture of colon/arteries - Type IV [Vascular type]

Worse prognosis - Type - IV

I/III - Scarring affected. Quagette paper like Scarring

3) Osteogenesis imperfecta - Collagen - I

Pedigree analysis -

● - x-linked Recessive Carrier

○ — || — □ - Divorce

□ - Proband / index Case

□ — | — ○ - Infertility

↗ - deceased / dead

○ --- □ - emotional attachment

◇ - unknown gender / Hermaphrodite

extra marital affair

△ - miscarriage / abortion

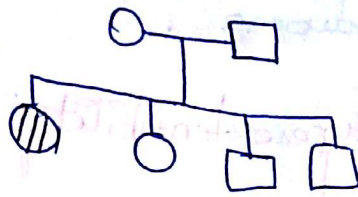
□ [○] - Adopted child

⑤ ③ - No. of siblings

△ - monozygotic

○ □ - Dizygotic

Pedigree analysis



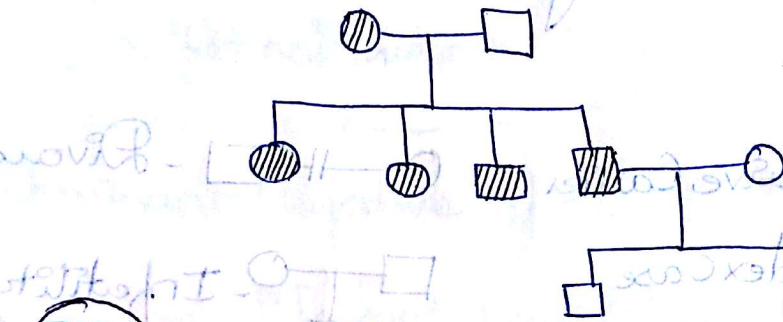
- Both parents normal,
one/variable child affected
undeclassic inheritance

① Gonadal mosaicism - variant of Autosomal dominant

→ mutation occur in zygote level

eg- Osteogenesis Imperfecta, Tuberous Sclerosis
Achondroplasia

② Mitochondrial Inheritance (Maternal Inheritance)



ovum - Rich in mitochondria

mtDNA

Normal

mutant

→ Heteroplasmy

when mutant allele > normal → disease

Sperm → also contain mtDNA - selectively degraded
during fertilization

↓
ubiquitin proteasome
pathway

→ Leber's Hereditary optic neuropathy - Prototype

→ NARP. - Neurogenic weakness
Ataxia.

Retinitis Pigmentosa.

Retinitis pigmentosa m/c pattern - of inheritance

60% Sporadic

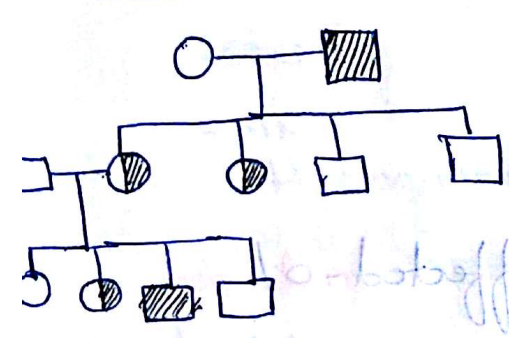
40% Hereditary

AR (60%)

30-40% AD

Dominant Inheritance

X-linked Recessive



→ X-chromosome of father transmitted to daughter only.

→ mostly males are affected.

→ Females also affected when carrier mother + affected father.

Y-chromosome - Y linked Disorders

- only males are affected. Retinitis pigment

- Hairy ears.

fragile X-Syndrome

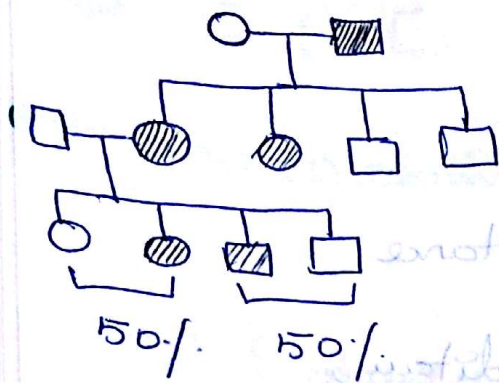
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→ Hemophilia - A/B - XR

Hemophilia - C - AR

X Linked dominant



Father
↓
all daughters

↓
Sons spared.

↓
from affected daughters
50% Sons 50% Daughters

All parts - 85% → X linked Dominant

Autosomal Recessive - m/c

Aa

AA

- Affected - 0%
Carrier - 50%

AA AA aA aA

Galactosemia → AR

Uniparental Disomy - one parent giving both genes.

Carrier - Normal



→ Functional gene inherited from 1 parent only
 no imprinting - epigenetics.

Chromosome-15q



Prader-Willi

- Obesity
- MR
- Hypogonadism

Ghrelin ↑ - orexogenic.

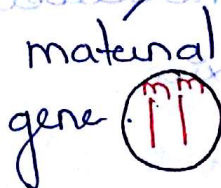


Angelman

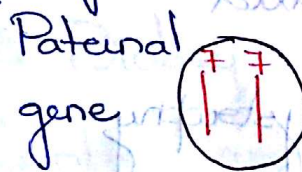
- Ataxia
- Laughter
- Happy puppet

m/c i) deletion

ii) uniparental disomy



↓
Prader-Willi



- Angelman

UP associations

- Angelman
- Prader-Willi
- Russel-Silver

iv) Bloom - Rare (m/c-AR)

v) Beckwith-wiedman (UPD of WT-1ch11)

trinucleotide Repeat

Codons - normally Triplets - Repeat normally

Times - 29 (6-55)

55 - 200 - **Premutation** - asymptomatic || - Carrier.

> 200 Repeats - **Disease**

∴ No. of Repeats correlate with severity of disorder.

No. of Repeats ∝ Severity.

Dynamic mutation - No. of Repeats - gain of mutation

Oogenesis
affecting Noncoding
areas -

eg - 1) Fragile - X - Syndrome

2) Friedreich's ataxia

3) Myotonic Dystrophy

Spermatogenesis
Coding areas
exons

1) Huntington's chorea

2) Spinocerebellar ataxia

Huntington's Disease Father → 300 Repeats.

Anticipation

↑ Severity,
years - earlier
onset

Son - ↑ Repeats 600 Repeats.
(earlier age); ↑ Severity.

next generation - earlier onset

→ X-Chromosome shows fragile sites, when grown on folate deficient medium



Large Testis (macroOrchidism) > NTR [Hallmark]

Large face
mandible <
everted ears

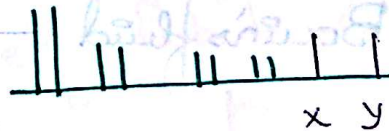
Small / Normal nose

KARYOTYPING

→ Study of chromosome

→ paired chromosome

↓ length



xy - last

Sample - i) Amniotic fluid - shedding of epithelium
(skin, GIT, GIT, R.T)

ii) Chorionic villi Biopsy

iii) Fetal tumor Sample - m/c - fetal tumor - Teratoma

iv) Cordocentesis - umbilical cord Blood

vi) Peripheral blood - lymphocytes

Other cells reversed - don't proliferate
in Test tube

Fixative of choice -

Karyotyping -

Carnoy's Fluid²

MTethanol + glacial acetic acid
3 parts + 1 part

Histopathology - 10% neutral buffered Formalin

Electron microscopy → glutaraldehyde → Best
Osmium Tetroxide

Cytology → 95% ethanol (cytology) - Pap Smear

Hematopoietic cells → mercuric chloride - Zenker's fluid

Penic acid - Bouin's fluid → Glycogen/Lipid storage disorders

Testicular biopsy → Bouin's fluid - ideal (preserves architecture)

Routine Karyotyping - neutral buffer formalin - universal
Cells

↓
Proliferate in Test

↓
metaphase arrest Colchicine

Resolution - 800 bands/Haploid set

To increase Resolution - prophase arrest

Resolution - 1500 bands/Haploid set

Routine Cytogenetic analysis → Light micro → Resolution of chromosome



Types of chromosomes

- i) NTelocentric - p-q - Rest all chromosomes metacentric
- ii) Sub-METACENTRIC - X-chromosome q > p
- iii) Acro-centric - Y-chromosome q >> p.

i) associated with pathogenesis

Down-Syndrome

- Robertsonian Translocation

t → 21 → 13
14
15

} All are Acrocentric chromosomes

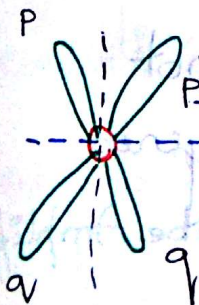
Acrocentric group -

21, 13, 14, 15, 22, Y

iv) Telocentric



- not seen in Humans



Normal division - vertical

↓ to normal axis - Horizontal plane



Isochromosome (i)

m/c isochromosome - Xq - x long arm

m/c in Human Cancer - 17q (ii)

gammaltomas of ovary & Testis
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i12p — m/c cytogenetics in Medulloblastoma
i17q — m/c cytogenetics in Medulloblastoma
↓
Poor prognosis

Uses of Karyotyping -

- i) Numerical disorders
 - ↳ euploid - $23 \times$
 - ↳ Aneuploidy - odd
- m/c Chromosomal disorder - Trisomy 16 (But die in utero)
- m/c chromosomal disorder in live birth } **Down Syndrome**

Limitations - i) microdeletion

- ii) Amplification
 - iii) Complex Translocation
- } Cannot be seen

→ FISH - can overcome this

Best for known gene loci - FISH - Limitation - Cannot detect Small genes.

∴ For point mutation - PCR Limitation - Cannot detect larger genes.

MLPA - Multiplex Ligation Associated Probe Amplification

- overcome limitations of both PCR & FISH
- Both Small & large genes
- Best for Cystic fibrosis detection

mech - i) Meiotic - Non dysjunction - Meiosis - I (in all)

95%

trisomies, except for Edward Syndrome - Meiosis II

Rock Bottom feet - more Edward Syndrome

extra chromosome in Down's → Maternal in origin

ii) Robertsonian - 4%

No Relation with advanced maternal age

iii) NTosauism - 1%

Robertsonian

NTosauism

Normal + abnormal no. of chromosomes.

→ + 21 → 13, 14, 15; 22.

+ (21, 21) - 100% chance

of 2nd Baby

ALL - m/c leukemia associated with Down's.

AMLNT₇ - Acute NTegakaujocytic - m/c leukemia in children < 3 years.

> 40 years - Alzheimer's - 100%.

m/c Cardiac defect - endocardial cushion defect > VSD > ASD

Turner's Syndrome i) 45 XO m/c

ii) 46 X(r) (Ring chromosome)

iii) 46 Xp

~~46 Xq~~

Histology - Abnormality of ovarian Tissue

Histology - Breast Tissue Normal

m/c Cardiac - Bicuspid aortic valve

Tumors -

- i) Colon Ca
- ii) AML

AML - Down's
Turner

Klinefelter

Klinefelter Syndrome -

Trisomies - $\geq 2x$ and $\geq 1y$

m/c - $47XXY$ - meiotic nondisjunction (meiotic I)

→ Associated with both Advanced Maternal/Paternal age

→ Testosterone $\downarrow\downarrow$; $\uparrow$ FSH, oestrogen.

→ gynecomastia

m/c Cancer Breast Cancer

m/c germ cell tumor - Extragonadal > gonadal

Teratoma

m/c site - mediastinum

m/c Site EGCT in Brain - pineal gland

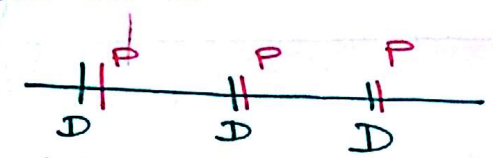
Genetic polymorphism

15-million Base pairs

- polymorphic genes / marker genes - can detect multifactorial / unknown gene mutation,

Polymorphic genes - (2 characters)

i) always dose to disease causing allele.



ii) disease

Polymorphic genes - always stably transmitted to next generation.

Linkage - Disequilibrium

major type of polymorphic gene - SNP - Single nucleotide Polymorphism
- 1 nucleotide difference / 1000 base pairs

ii) Repeat length polymorphism -

a) microsatellite Macro satellite

(2-6) Base Pair repeats

15-70 Base Pair repeats

Uses

Genetic

Single gene

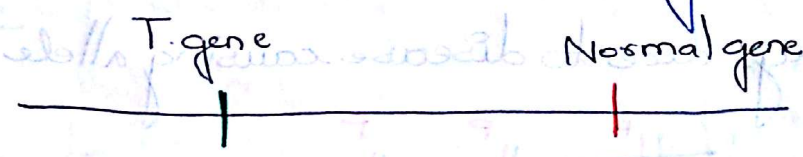
multifactorial

Direct Sequencing with PCR / FISH

Polymorphic genes - Indirect Sequencing

Large Population - **GWAS** (genome wide Analytical Study)

To detect Same chromosome - Tumor gene / Normal gene



Patient

Control (entire chromosome)
gene

Fluorescence

Red

Green

Hybridization

Fluorescence

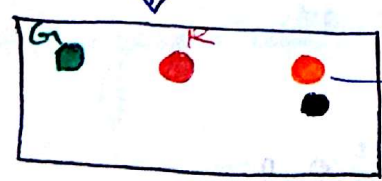
Scanner

Computer

Red Signal - Duplication (gain)

Green Signal - Deletion (loss)

Normal chromosome - yellow/Black



yellow/Black } NTA array Analysis

NTA array based Comparative, genomic Hybridization **CGH**

i) gene expression profiling eg

ERT+; PR+ve; Her. neu.

ii) molecular classification of Cancer.

→ Molecular classification of Breast Ca based on ^{gene} expression profiling

a) FISH Congenital malformation - metaphase

Oncological specimens - Interphase

2 variants-

i) Chromosomal painting (entire human genome on same chromosome)

fluorochrome painted on entire genome of one chromosome

a

2) Multicoloured FISH - entire genome on all gen chromosomes

Spectral Karyotyping
Spectral Karyotyping

simultaneously

fluorochromes Computer generalized.

Immunology

Immunity

Innate

Acquired

- Birth / 1st line of defence

- memory ⊕

- Non memory

- Specific ⊕

- Non-Specific