

Inheritance

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parents $\xrightarrow{\text{characters}}$ offspring

\downarrow
Inheritance

Heridity
Similar

\downarrow
Dissimilar Variation
Dissimilar

Genes: \rightarrow Inheritance unit
 \rightarrow Particular nucleotide

sequence of a.a.

\rightarrow Allele: gene partner

\rightarrow homologous chromosome

\downarrow
positions: Locus: \rightarrow ABO \rightarrow 9

Sickle cell anemia \rightarrow 11

Gene H \rightarrow 19

X Linked \rightarrow X

X Linked \rightarrow Y

\rightarrow Homologous:

\rightarrow similar gene pair

\rightarrow Heterozygote

\rightarrow Dissimilar

- \rightarrow Dominance
- \rightarrow Recessive
- \rightarrow Incomplete
- \rightarrow Over Dominance
- \rightarrow Co-dominance

\rightarrow Trait:

Any character \rightarrow Height, color, shape

\downarrow
Genotype

\downarrow
phenotype

\rightarrow genetic composition \rightarrow phenotype \rightarrow particular

\rightarrow TT / Tt & tt

\rightarrow short & long

\rightarrow YY / Yy / yy

\rightarrow yellow & Green

\rightarrow RR / Rr / rr

\rightarrow Round & Wrinkled

Genotype:

↳ Total genotype of an individual.

Gene pool: breeding population

→ Total genes:

↳ Gene frequency ^{i.e.}

Eye color: $G-1, B-6, Bl-10$
 ↳ in specific population

Mendel's Work:

1866 → plant hybridization:

→ Pea plant → *Pisum sativum*

- Easily available
- Easy to grow
- Hermaphrodite → self & cross pollination
- Short life cycle
- Large progeny
- Distinct characters

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• 7 - Traits :

Seed shape	Round	Wrinkle.
Pod shape	Smooth	Constricted.
Seed colour	Yellow	green.
Pod colour	Green	Yellow
Flower Position	Axial	Terminal
Flower colour	Purple	White
Height	Long	Short

• Hidden Advantage:

- 7 Pair → chromosome
- 7 character → 4 chromosome
- No linkage.

• Red's covering:

→ 1900 → Correns, De Vries, Tschermak.

Mendel's First Law

→ Segregation, Dominance & Gametes purity:
Criteria:

→ monohybrid → single trait
seed shape

Round ↔ Wrinkle

1st Cross:

P₁ Round × Wrinkled

F₁ Round

2nd Cross:

F₁ × F₁ (self cross)

Round × Round

F₂ Round → 3 (75%)

Wrinkled → 1 (25%)

Test cross: → to find unknown genotype

1st Round: unknown genotype × P₁ recessive

1st Round: × P₁ recessive

Round → 100%

	r	r
R	Rr	Rr
R	Rr	Rr

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2nd Round: × P₁ recessive

Round → 50%

	r	r
R	Rr	Rr
r	rr	rr

Genotype:

1 : 2 : 1

3rd Round: × P₁ recessive

Round → 50%

	r	r
R	Rr	Rr
r	rr	rr

Phenotype:

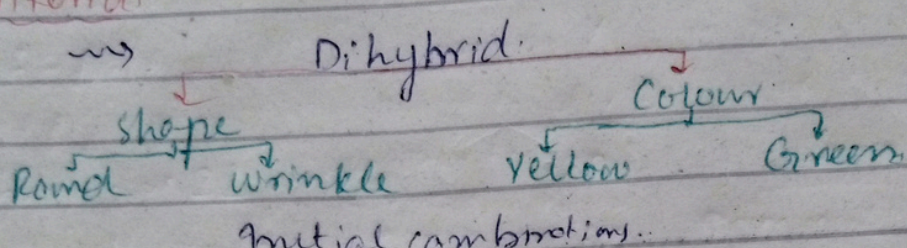
3 : 1

Heterozygous = 2
 Homozygous = 1

No. of gametes combination
 $RrYy = 2 \times 2 = 4$
 $RRYY = 2 \times 1 = 2$
 $RRYy = 2 \times 2 = 4$
 $RRyyPp = 2 \times 2 \times 2 = 8$
 $RRYYPp = 2 \times 1 \times 2 = 4$
 $RrYyPp = 2 \times 2 \times 2 = 8$

Mendel's 2nd Law:
 Law of Independent Assortment:

Criteria:



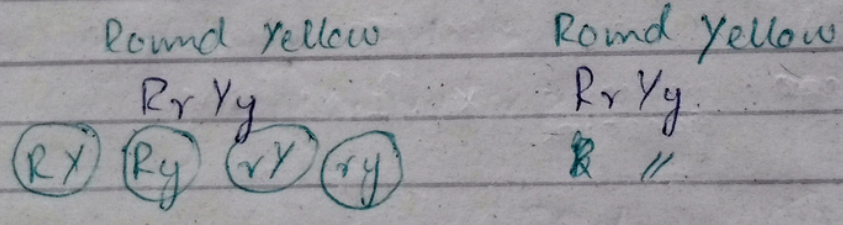
- Initial combinations:
- Round → Yellow
 - Wrinkled → green

→ Pure breed

1st Cross:

P₁ Round Yellow × Wrinkled green
 F₁ Round yellow - 100%

2nd Cross:



	RY	Ry	rY	ry
RY → yellow	RRYY*	RRYy	RrYY	(RrYy)
Y → green	RRYy*	RRyy*	(RrYy)	*Rryy
	RrYY	(RrYy)	*rrYY	rrYy
	(RrYy)	*RrYy	*rrYy	*rryy*

Homozygous dominant: 1
 Homozygous recessive: 1
 *Homozygous: 4

Genotypic ratio:
 1:2:1:2:4:2:1:2:1
 Total genotypes: → 9

Total phenotype No: 16

Total phenotype types: 4

- ↳ Round yellow = 9 $\frac{9}{16}$
- ↳ Round green = 3 $\frac{3}{16}$
- ↳ Wrinkled yellow = 3 $\frac{3}{16}$
- ↳ Wrinkled green = 1 $\frac{1}{16}$

→ Parental combinations:
 Round yellow: 9 } $\frac{10}{16}$
 Wrinkled green: 1 }
 → Recomb:
 Wrinkled yellow: 3 } $\frac{6}{16}$
 Round green: 3 }

Mendellian Traits:

Dihybrid: $F_2 \rightarrow 9:3:3:1$

Non Mendelian:

Deviate $\rightarrow 9:3:3:1$ in dihybrid.

\rightarrow Incomplete Dominance

\rightarrow Co-dominance

\rightarrow Over dominance

\rightarrow Linkage

\rightarrow multiple alleles

\rightarrow Pleiotropy

Linkage:

\rightarrow T.H. Morgan

B.W = broad wing

S.W = Short

R.A = Round abdom

P.A = Pointed

Dihybrid Drosophila

B.W & R.A

S.W & P.A

$\rightarrow 9:3:3:1 \times$

B.W & R.A $\rightarrow 100\%$

$\rightarrow 3:1 \checkmark$

$F_1 \times F_1 \rightarrow$ self cross

B.W & R.A \times B.W & R.A

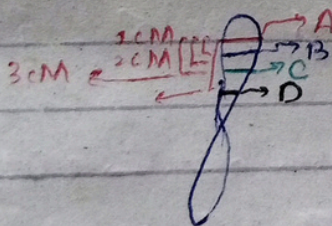
\rightarrow deviated

\rightarrow Linkage

\rightarrow Physical relationship b/w different genes

\rightarrow genes \rightarrow unlimited

\rightarrow Chromosome \rightarrow limited



\rightarrow transfer pattern \rightarrow enblock

Linkage $\propto \frac{1}{\text{distance b/w genes}}$

\rightarrow Ratio in F_2 :

\rightarrow All parental types

\rightarrow Complete linkage 3:1

\rightarrow More Parental & less recomb.

\rightarrow Incomplete

Parental & Recomb same.

\rightarrow No linkage. 1:1:1:1

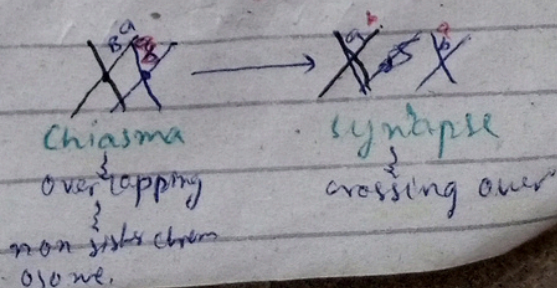
Crossing Over:

\rightarrow recomb linkage

\rightarrow prophase I

\rightarrow pachytene

\rightarrow homologous chromo \rightarrow among



chiasma overlapping non sister chromosome.

synapsis crossing over

(Linkage types)

Autosomal:

Autosomes

1-22 pair

♂ & ♀ equally

Chromosome 22

Sickle cell anemia

Leukemia

Albinism

Sex

Y-Linked

→ Chromosome Y

→ Direct transfer father to son

→ Holandric traits
↳ male specific

specific region on Y. **SRY gene**

↳ maleness

- (Hypertrichosis)
↳ hair on ear.

X-Linked

→ On X chromosome

→ Affects male & female

→ Ziggzag pattern

Dominant

→ Affects more ♀

→ hypophosphatemic rickets (vitamin D resistant rickets)

Recessive

affects more ♂

→ haemophilia
→ color blindness
→ Gout

RR ↓
XX → affect } 66%
Rr Xx → affect }
rr xx → normal } 33%

R ↓
XY → affect } 50%
rY → normal } 50%

Hh ↓ 66%
XX → normal }
Hh Xh → normal }
hh xx → affect } 33%
Hh Y → normal } 50%
hh Y → affect } 50%

Haemophilia:

→ clotting factors → absent

→ blood fails to clot ^{excessive} _{bleeding} → death

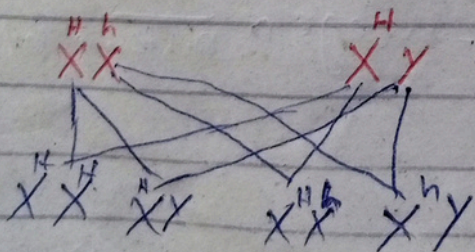
→ Royal disease → Queen Victoria

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Types:

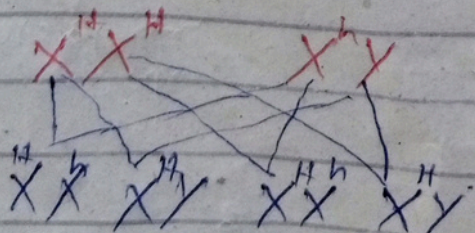
Absent clotting factor

Type	Chromosome	Linkage	Prevalence	Affected
H.A	VIII	X-linked	80%	♂ more
H.B	IX	X-linked	20%	♂ more
H.C	XI	4 th chromosome → autosomal	<1%	♂ & ♀



hemophiliac → 25%
Affected → 25%
Carrier → 25%
Healthy → 25%

50% sons are affected



Haemophiliac → 0%
Healthy 100%
Carrier 50%

Multiple Alleles:

Single trait controlled by many alleles
 → maximum alleles discovered → 300.

Population → all alleles

Individual → 2 alleles

↳ gamete → 1 allele.

Example:

A.B.O Blood Group.

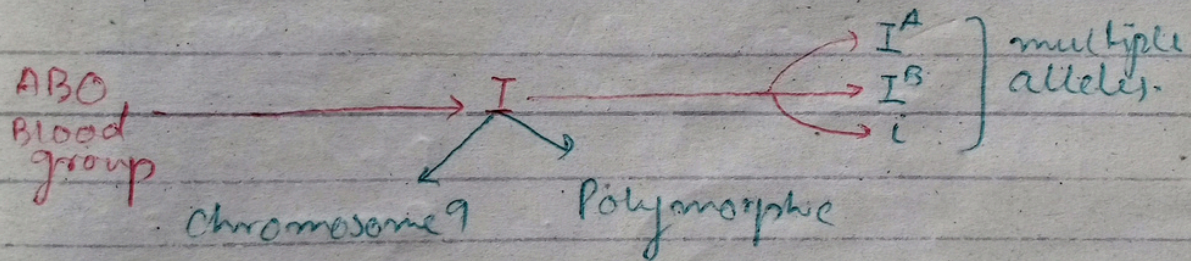
Discovered by:

Karl Landsteiner

Genetic bases:

Bernstein.

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→ I - express → Early embryonic stage.

Genotype	Antigen	Diagram	Blood group	Anti body	Donate to	Donate from.
$I^A I^A / I^A i$	A		A	B	A, AB	A, O
$I^B I^B / I^B i$	B		B	A	B, AB	B, O
$I^A I^B$	Both		AB	X	AB	A, AB, B, O
i	X		O	Both	O, A, B, AB	Only

→ Total phenotypes → 4

→ Total genotype → 6

→ Total antigen → 2

→ Total antibodies → 2

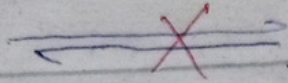
→ Universal Donor → O

→ Universal Receptor → AB

antigen 'A' + Anti body 'A' = agglutination

Children
Parents

AB



O

Parents
children

AB
no child
If any parent $I^A I^B$ will be O
If any parent O no child will be AB.

$I^A I^B \times ii$

	i	i	
I^A	$I^A i$	$I^A i$	50% A
I^B	$I^B i$	$I^B i$	50% B

$I^A i \times I^B i$

	I^B	i	
I^A	$I^A I^B$	$I^A i$	
i	$I^B i$	ii	

→ all 4 blood groups.

25% → each