# Genotypes, Phenotypes and Hardy Weinberg Equilibrium 

Biostatistics 666
Lecture II

## Previously: Refresher on Genetics

- DNA sequence
- Human Genome
- Inheritance of genetic information
- Sequence variation
- VNTRs, microsatellites and SNPs
- Common Types of Genetic Study


## Gregor Johann Mendel

- Discovered basic principles of genetics
- "The Father of Genetics"
- Monk, lived 1822 - 1884
- Crosses between strains of peas
- Garden pea (Pisum sativum)
- Each strain has particular characters
- Height, flower color, seed shape ...


## Mendel's Experiment

- Crossed different truebreeding strains
- Identical results for reciprocal crosses
- $\mathrm{F}_{1}$ resembled one of the parental strains
- In $F_{2}$ generation, the other parental trait


Pure-breeding green line

## Mendel's Numbers

- Seeds: Yellow vs. Green
- $F_{1}$ : All yellow
- $F_{2}$ : 6022 yellow, 2001 green
- 75.1\% yellow, 24.9\% green
- Seeds: Smooth vs. Wrinkled
- $F_{1}$ : All smooth
- $F_{2}$ : 5474 smooth, 1850 wrinkled
- 74.7\% yellow, 25.3\% wrinkled


## Phenotype vs. Genotype

- Genotype
- Underlying genetic constitution
- Phenotype
- Observed manifestation of a phenotype
- The yellow peas in the parental and $F_{1}$ generations are not the same


## Mendel's Interpretation

- Each trait determined by "particulate factors" (genes)
- E.g.: Seed colour
- Alternative forms for each factor (alleles)
- E.g.: Yellow seeds or green seeds
- Each plant carried two alleles
- Identical for true breeding parental strains
- Different for $F_{1}$ generation


## The Principle of Segregation

- Mendel's First Law
- The two alleles of a gene pair segregate from each other in the formation of gametes
(gametes are reproductive cells that fuse to form a new organism in sexual reproduction)


## Genotypes

- Each individual carries two alleles
- If there are $\mathbf{n}$ alternative alleles ...
- ... there will be $\mathbf{n}(\mathbf{n + 1}) / \mathbf{2}$ possible genotypes
- Homozygotes
- The two alleles are the same
- E.g.: Green/Green or Yellow/Yellow
- Heterozygotes
- The two alleles are different
- E.g.: Green/Yellow


## Penetrance

- Describes the relationship between phenotypes and genotypes
- Complete Penetrance
- Each genotype corresponds to only one phenotype
- Incomplete Penetrance
- Link between phenotype and genotype is only probabilistic


## The ABO blood group

- Important for blood transfusions
- Determined by alleles of the ABO gene
- 3 alternative alleles
- A, B and O
- 6 possible genotypes, $n(n+1) / 2$ - A/A, A/B, A/O, B/B, B/O, O/O


## ABO Blood Group II

| Phenotype | Antigen |  | Antibody |  |
| :--- | :--- | :--- | :--- | :--- |
|  | A | B | A | B |
| A | + | - | - | + |
| B | - | + | + | - |
| O | - | - | + | + |
| AB | + | + | - | - |

## Relationships between alleles

- Relation between alleles
- A and B are dominant over O
- $O$ is recessive in relation to $A$ and $B$
- $A$ and $B$ are codominant
- In this case all genotypes

| Genotype | Phenotype |
| :--- | :--- |
| A/A | A |
| A/B | AB |
| A/O | A |
| B/B | B |
| B/O | B |
| O/O | O | are fully penetrant

## BRCA1 and Breast Cancer

- BRCA1 mutations predispose to breast cancer
- About $0.1 \%$ of the population carries mutations in the BRCA1 gene
- Disease Risk
- Age
- Carriers 40\%
- Non-carriers 0.4\%

40
60
70\%
3\%

80
80\%
8\%

## Alleles, Genotypes and Phenotypes

- Classifying genotypes
- Homozygous
- Heterozygous
- Penetrance
- Relationships between alleles
- Dominant, Recessive, Co-Dominant


## Genes in Populations

- Genotype Frequencies
- Haplotype Frequencies
- Allele Frequencies
- Penetrance Function
- Derived measures of marker informativeness


## Notation

- $p_{i j}$
- frequency of genotype $\mathrm{i} / \mathrm{j}$ in the population
- $\mathrm{n}(\mathrm{n}+1) / 2$ of these
- $p_{i}$
- frequency of allele i in the gene pool
- n of these
- Write allele frequencies as function of genotype frequencies


## Hardy-Weinberg Equilibrium

- Random union of games
- Relationship discovered it in 1908
- Hardy, British mathematician
- Weinberg, German physician
- Shows $\mathbf{n}$ allele frequencies determine $\mathbf{n}(\mathbf{n + 1}) / \mathbf{2}$ genotype frequencies
- Large populations


## Required Assumptions

- Diploid, sexual organism
- Non-overlapping generations
- Autosome
- Large population
- Random mating
- Equal genotype frequencies among sexes
- Selection


## Random Mating: <br> Mating Type Frequencies

| Mating | Frequency |
| :---: | :---: |
| $\mathrm{A}_{1} \mathrm{~A}_{1}{ }^{*} \mathrm{~A}_{1} \mathrm{~A}_{1}$ | $\mathrm{p}_{11}{ }^{2}$ |
| $\mathrm{~A}_{1} \mathrm{~A}_{1}{ }^{*} \mathrm{~A}_{1} \mathrm{~A}_{2}$ | $2 \mathrm{p}_{11} \mathrm{p}_{12}$ |
| $\mathrm{~A}_{1} \mathrm{~A}_{1}{ }^{*} \mathrm{~A}_{2} \mathrm{~A}_{2}$ | $2 \mathrm{p}_{11} \mathrm{p}_{22}$ |
| $\mathrm{~A}_{1} \mathrm{~A}_{2}{ }^{*} \mathrm{~A}_{1} \mathrm{~A}_{2}$ | $\mathrm{p}_{12}{ }^{2}$ |
| $\mathrm{~A}_{1} \mathrm{~A}_{2}{ }^{*} \mathrm{~A}_{2} \mathrm{~A}_{2}$ | $2 \mathrm{p}_{12} \mathrm{p}_{22}$ |
| $\mathrm{~A}_{2} \mathrm{~A}_{2}{ }^{*} \mathrm{~A}_{2} \mathrm{~A}_{2}$ | $\mathrm{p}_{22}{ }^{2}$ |
| Total | 1.0 |

## Mendelian Segregation: Offspring Genotype Frequencies

## Offspring

$\begin{array}{ccc}\text { Mating } & \text { Frequency } & \mathbf{A}_{1} \mathbf{A}_{\mathbf{1}} \\ \mathrm{A}_{1} \mathrm{~A}_{1}{ }^{*} \mathrm{~A}_{1} \mathrm{~A}_{1} & \mathrm{p}_{11}{ }^{2} & \mathrm{p}_{11}{ }^{2}\end{array}$
$A_{1} A_{1}{ }^{*} A_{1} A_{2} \quad 2 p_{11} p_{12} \quad p_{11} p_{12} \quad p_{11} p_{12}$
$\mathrm{A}_{1} \mathrm{~A}_{1}{ }^{*} \mathrm{~A}_{2} \mathrm{~A}_{2} \quad 2 \mathrm{p}_{11} \mathrm{p}_{22}$
$\mathrm{A}_{1} \mathrm{~A}_{2}{ }^{*} \mathrm{~A}_{1} \mathrm{~A}_{2} \quad \mathrm{p}_{12}{ }^{2}$
$1 / 4 \mathrm{p}_{12}{ }^{2}$
$1 / 2 p_{12}{ }^{2}$
$1 / 4 \mathrm{p}_{12}{ }^{2}$
$A_{1} A_{2}{ }^{*} A_{2} A_{2} \quad 2 p_{12} p_{22}$
$\mathrm{p}_{12} \mathrm{p}_{22}$
$p_{12} p_{22}$
$\mathrm{A}_{2} \mathrm{~A}_{2}{ }^{*} \mathrm{~A}_{2} \mathrm{~A}_{2}$
$\mathrm{p}_{22}{ }^{2}$
$\mathrm{p}_{22}{ }^{2}$

## And now...

$$
\begin{aligned}
p_{11}^{\prime} & =p_{11}^{2}+p_{11} p_{12}+1 / 4 p_{12}^{2} \\
& =\left(p_{11}+1 / 2 p_{12}\right)^{2} \\
& =p_{1}^{2} \\
p_{22}^{\prime} & =p_{22}^{2}+p_{22} p_{12}+1 / 4 p_{12}^{2} \\
& =\left(p_{22}+1 / 2 p_{12}\right)^{2} \\
& =p_{2}^{2} \\
p_{12}^{\prime} & =2 p_{11} p_{22}+p_{11} p_{12}+p_{12} p_{22}+1 / 2 p_{12}^{2} \\
& =2\left(p_{11}+1 / 2 p_{12}\right)\left(p_{22}+1 / 2 p_{12}\right) \\
& =2 p_{1} p_{2}
\end{aligned}
$$

## Conclusion

- Genotype frequencies are function of allele frequencies
- Equilibrium reached in one generation
- Independent of initial genotype frequencies
- Random mating, etc. required
- Conform to binomial expansion
- $\left(p_{1}+p_{2}\right)^{2}=p_{1}^{2}+2 p_{1} p_{2}+p_{2}^{2}$


## A few more notes...

- Can be expanded to multiple alleles
- Expand $\left(p_{1}+p_{2}+p_{3}+\cdots+p_{k}\right)^{2}$
- Holds in almost all human populations
- Little inbreeding (typical F = ~0.005)
- Deviations can suggest:
- Problems with experimental assays
- Non-independence of observations
- Selection
- Disease locus


## Heterozigosity

- Probability that two alleles will differ

$$
H=1-\sum p_{i}^{2}
$$

- For $a$ equally frequent alleles

$$
H=1-\frac{1}{a}=\frac{a-1}{a}
$$

- Sometimes called "gene diversity"


## PIC

- Probability that alleles of parent can be distinguished in offspring
- Botstein et al, 1980.
- Markers that could track dominant alleles
- Probability that parent will heterozygous and informative in relation to spouse


## PIC - Definition

- In general:

$$
\text { PIC }=1-\sum_{i=1}^{n} p_{i}^{2}-\sum_{i=1}^{n} \sum_{j=i+1}^{n} 2\left(p_{i} p_{j}\right)^{2}
$$

- For $a$ equally frequent alleles

$$
\text { PIC }=\frac{a-1}{a}-\frac{a-1}{a^{3}}
$$

- PIC <= H


## Exercise

- ABO locus allele frequencies
- A - frequency 0.3
- B - frequency 0.1
- O - frequency 0.6
- Calculate genotype frequencies
- Calculate heterozygosity and PIC
- Calculate phenotype frequencies


## NOD2 and Bowel Disease

- Leu1007fs
- Frame shift mutation at position 1007
- Frequency of about 5\%
- Disrupts gene
- Penetrance
$\begin{array}{llll}\text { - Genotype } & +/+ & -/+ & -/- \\ \text { - } P(\text { Crohn's } \mid G) & 0.1 \% & 0.2 \% & 3 \%\end{array}$
- Calculate frequency of -/- genotype in population and among patients...

