# Bio 112: Hardy-Weinberg Equilibrium Examples General Info

- <u>Allele frequencies</u>: the frequency of each <u>allele</u> (R or r, for example) in the gene pool. The symbols p and q are used to represent these frequencies.
- <u>Genotype frequencies</u>: the frequency of each <u>genotype</u> (RR, Rr, rr for example) in the population. These are <u>always</u> equal to the number of individuals with a particular genotype divided by the total population size. They are <u>sometimes</u> equal to p<sup>2</sup>, 2pq, and q<sup>2</sup> only when the population is at HWE.

For a particular pair of allele frequencies ( $\mathbf{p}=0.2$  and  $\mathbf{q}=0.8$  for example), there are many possible sets of genotype frequencies that have the same allele frequencies (actually, infinitely many). This is illustrated by the 4 example populations below; all 4 of these populations have the same allele frequencies. However, given a pair of allele frequencies, there is only one set of these genotype frequencies that are at HWE. The predictions of HWE- the frequency of  $\mathbf{RR} = \mathbf{p}^2$  (0.04 in this example), the frequency of  $\mathbf{RR} = 2\mathbf{pq}$  (0.32 in this example), and the frequency of  $\mathbf{rr} = \mathbf{q}^2$  (0.64 in this example).

## Population 1

Observed #R's #r's contributed contributed Genotype Genotype # **frequency** to gene pool to gene pool RR 0 40 Rr Rr 60 totals

### 1.Allele frequencies

Freq. of R = p =Freq. of r = q =

2.Genotype frequencies

Freq. of RR = Freq. of Rr = Freq. of rr =

3. Compare: HWE or Not HWE

## Population 2

Genotype	<u>#</u>	Observed Genotype frequency	#R's contributed to gene pool	#r's contributed to gene pool
RR	400			
Rr	0			
rr	1600			
		<u>totals</u>		

#### 1. Allele frequencies

Freq. of R = p =

Freq. of r = q =

2.Genotype frequencies

Freq. of RR =

Freq. of Rr =

Freq. of rr =

3.Compare: HWE or Not HWE

## Population 3

Genotype	<u>#</u>	Genotype frequency	#R's contributed to gene pool	#r's contributed to gene pool
RR	4			
Rr	32			
rr	64			
		<u>totals</u>		

# 1.Allele frequencies

Freq. of R = p =

Freq. of r = q =

2.Genotype frequencies
Freq. of RR =

Freq. of Rr =

Freq. of rr =

3. Compare HWE or Not HWE

# Population 4

<u>Genotype</u>	<u>#</u>	Genotype frequency	#R's contributed to gene pool	#r's contributed to gene pool
RR	12			
Rr	96			
rr	192			
		totals		

# 1.Allele frequencies

Freq. of R = p =

Freq. of r = q =

2.Genotype frequencies

Freq. of RR =

Freq. of Rr =

Freq. of rr =

3. Compare: HWE or Not HWE