

## Evolution als Änderung der Genfrequenzen

Thema 4

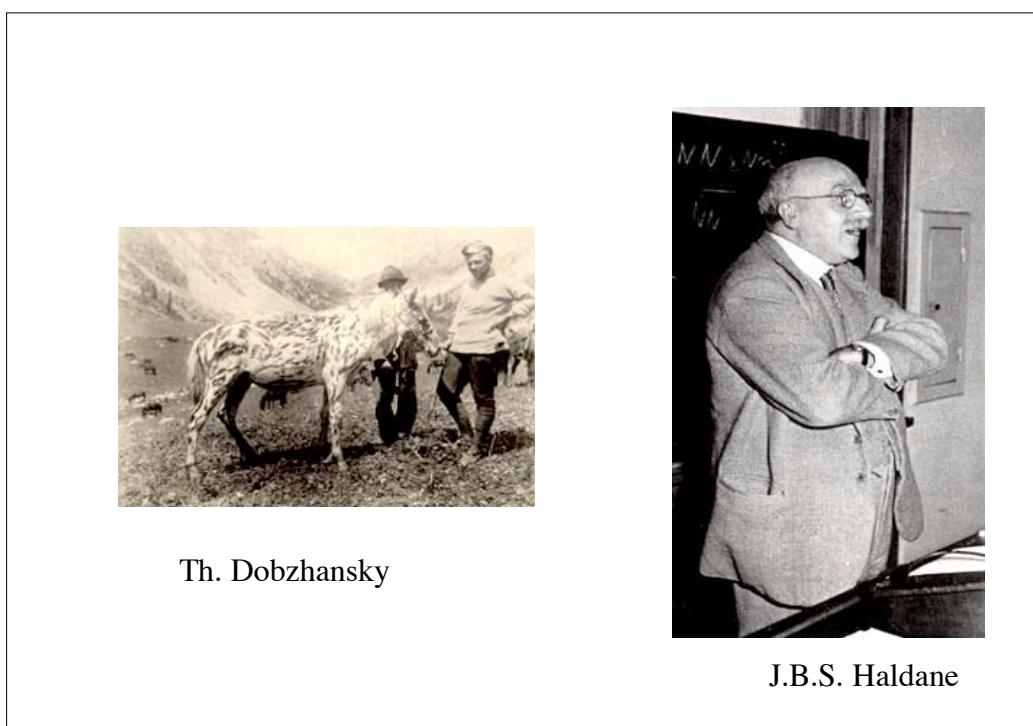
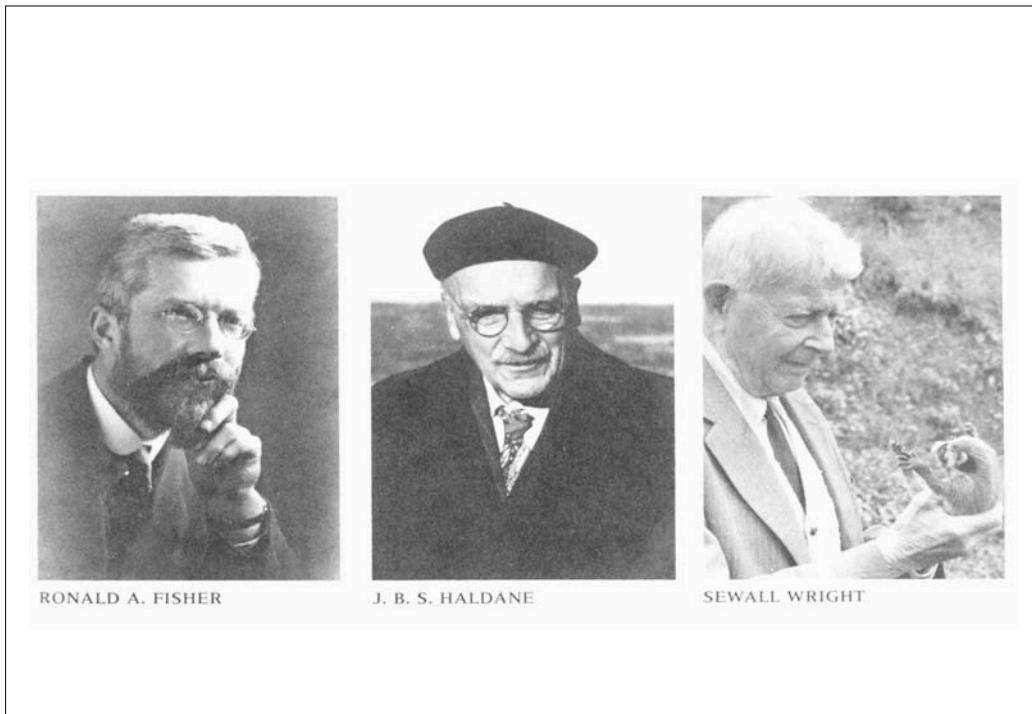


G. LEDYARD STEBBINS

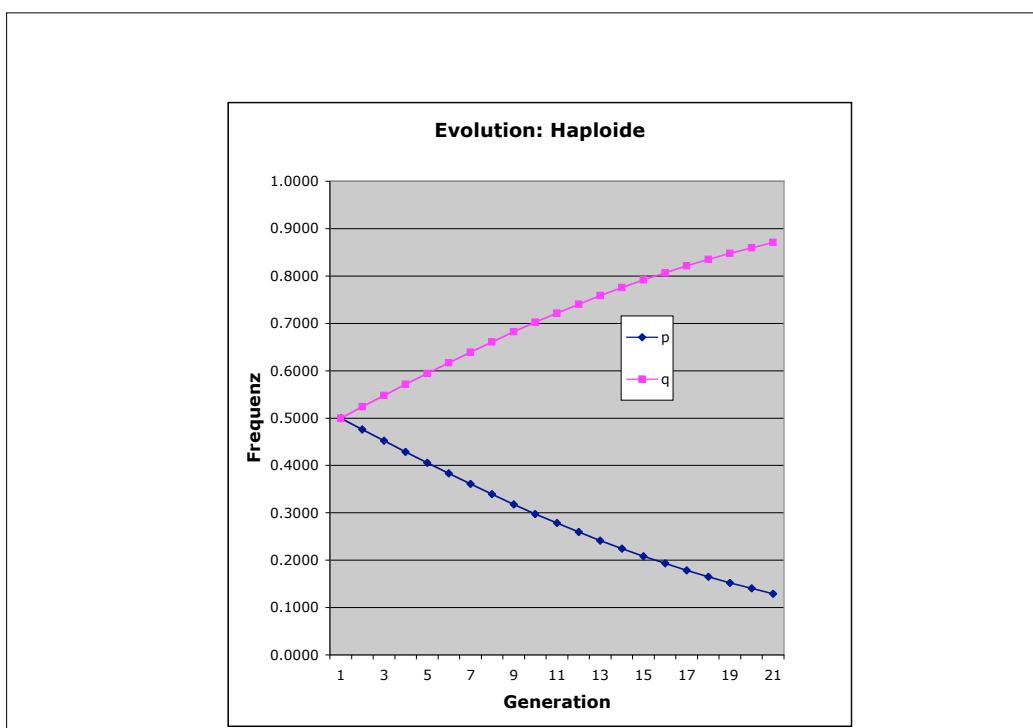
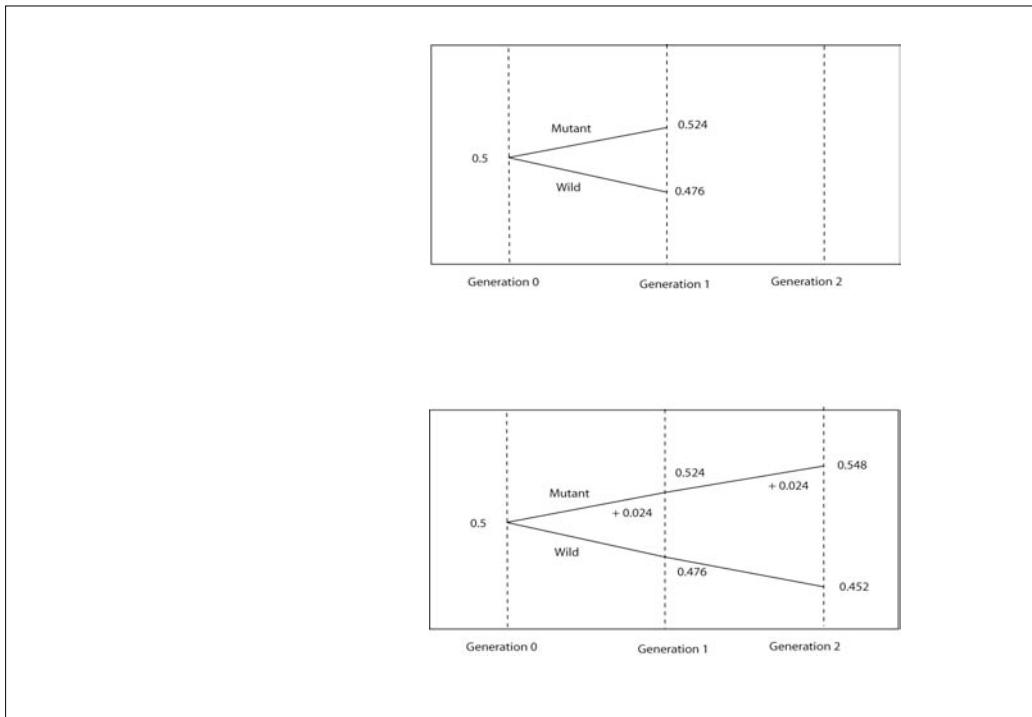
GEORGE GAYLORD SIMPSON

THEODOSIUS DOBZHANSKY

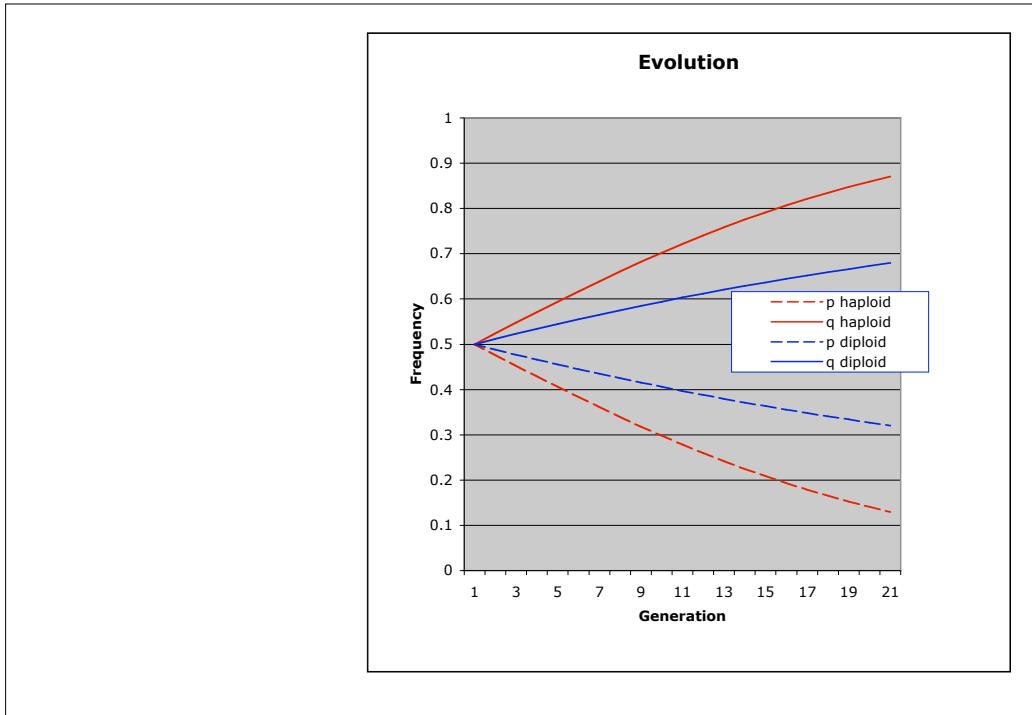
## Änderung von Genfrequenzen



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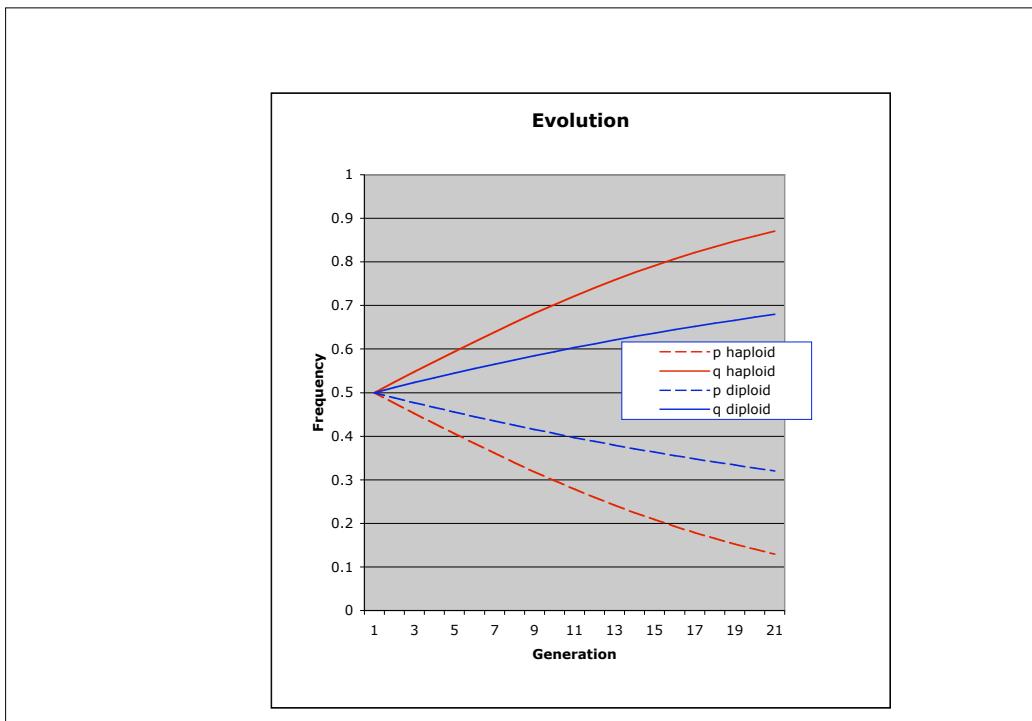
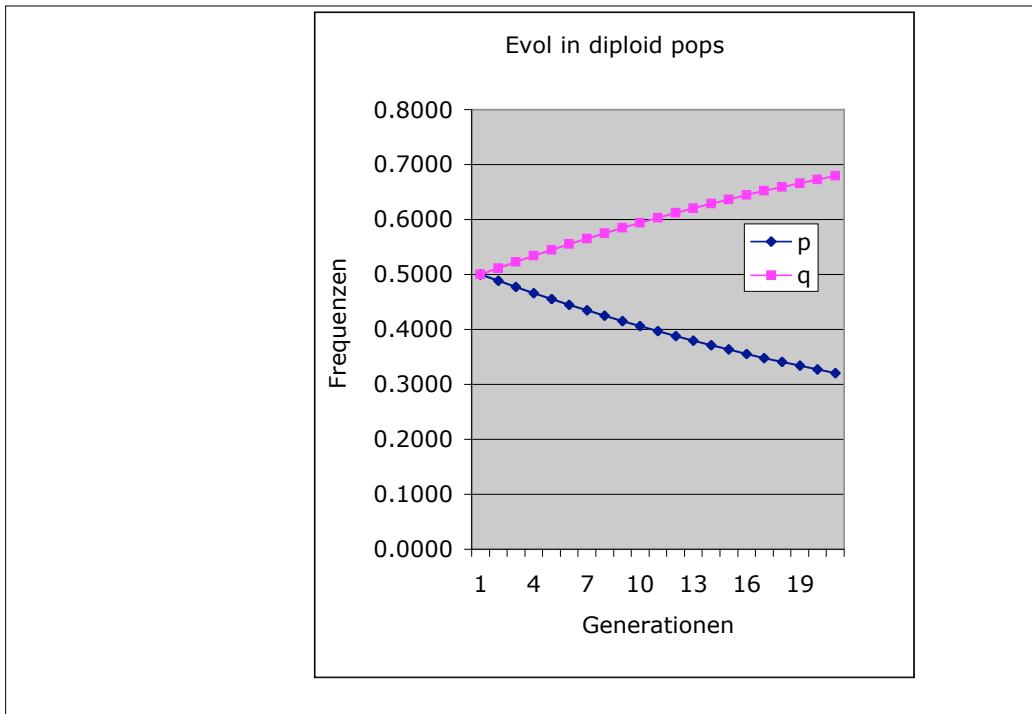


$w = \text{mittlere Fitness der Population}$

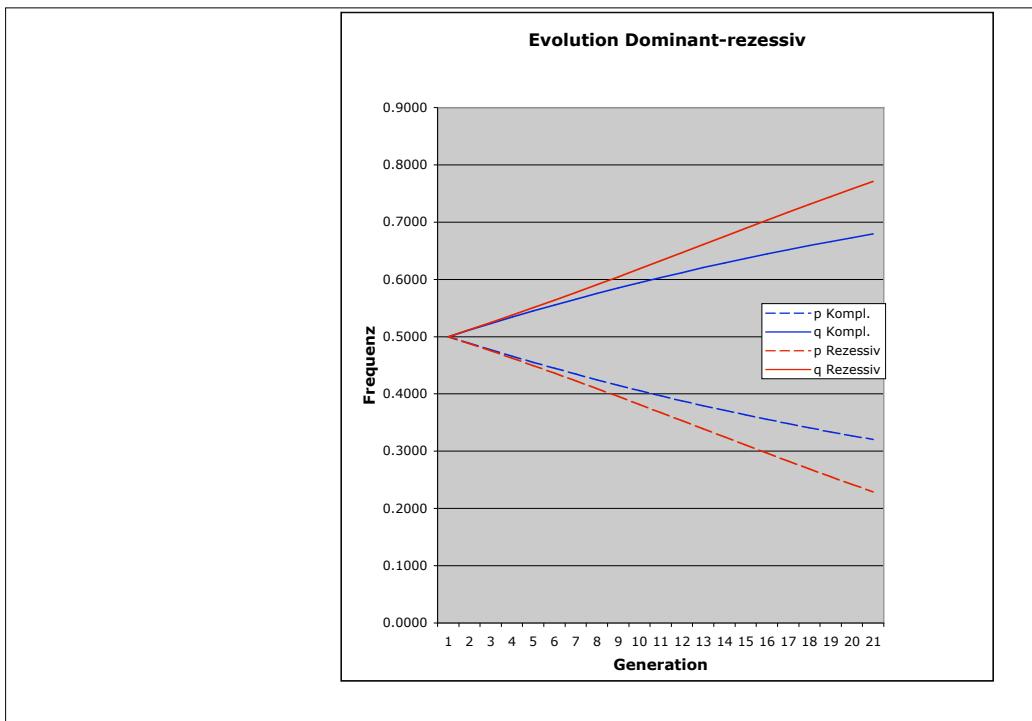
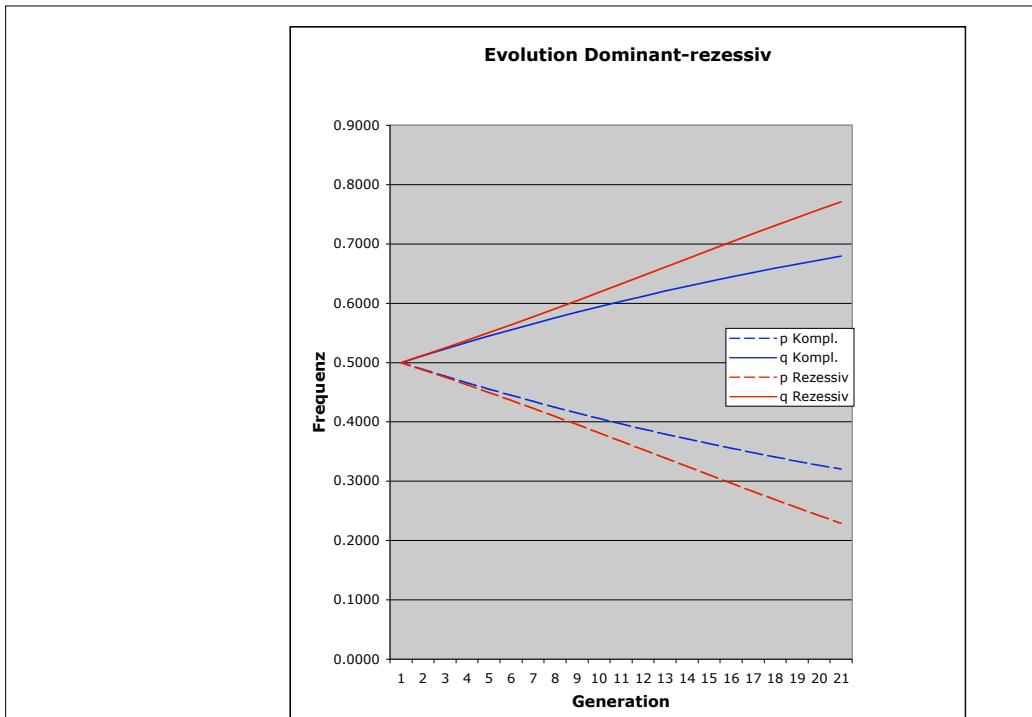
$d = \text{Dominanz } (0 \leq d \leq 1)$

Stadium	Wild	Heterozygot	Mutant	TOTAL
Individuen Frequenz A1, A2	$A1A1$ $p$	$A1A2$ $2pq$	$A2A2$ $q^2$	100 $p + q = 1$
Frequenz Fitness Nachkommen	$p^2$ $\frac{1}{p^2}$	$1 + ds$ $2pq(1+ds)$	$1 + s$ $q^2(1+s)$	1.0 $w = p^2 + 2pq(1+ds) + q^2(1+s)$
Frequenz, neu	$\frac{p^2}{w}$ $p' = \frac{p(1 + sdq)}{w}$	$\frac{2pq(1 + ds)}{w}$	$\frac{q^2(1 + s)}{w}$ $q' = \frac{q(1 + s[q + dp])}{w}$	1.0 $p' + q' = 1$
	(Individuen, neu) = 100	$A1' = p' \cdot 100$	$A2' = q' \cdot 100$	100

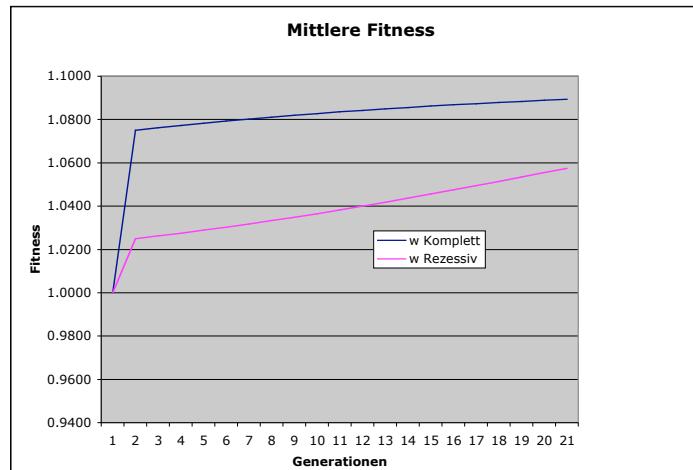
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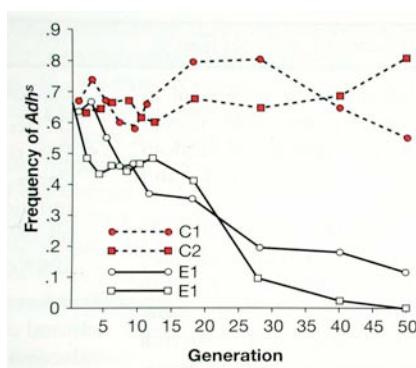
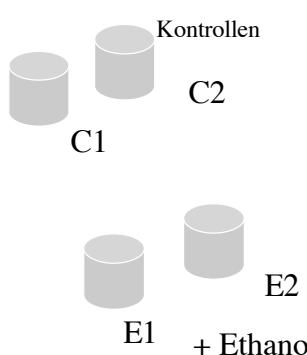
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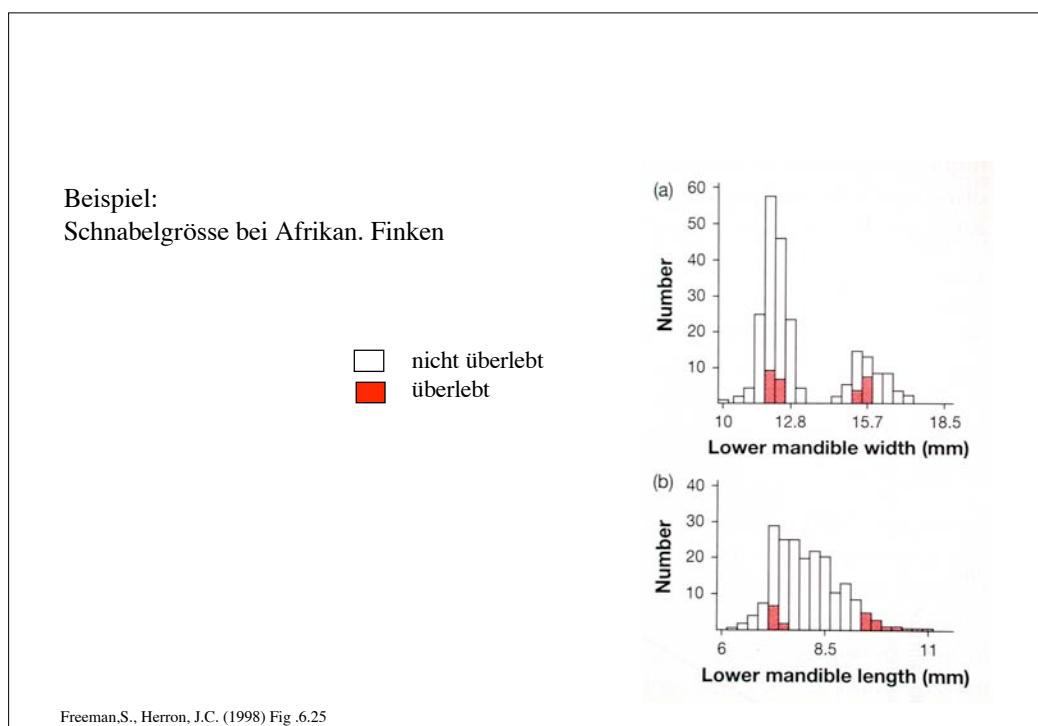
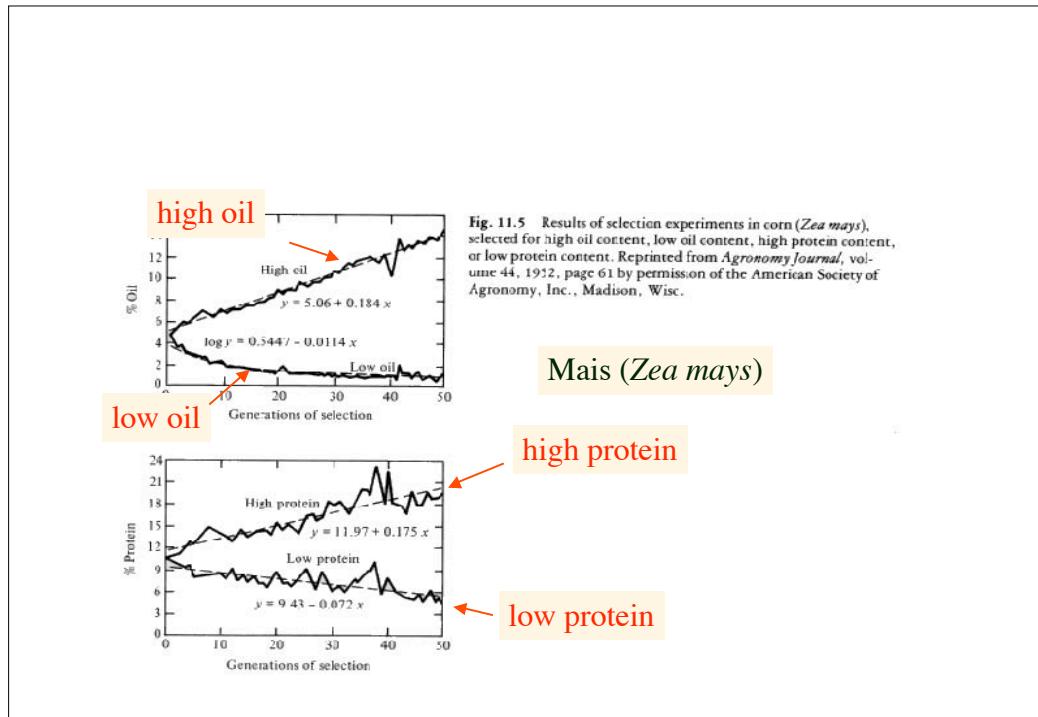
## Änderung von Genfrequenzen



Experiment mit *Drosophila melanogaster*  
2 Allele  $Adh^F$ ,  $Adh^S$

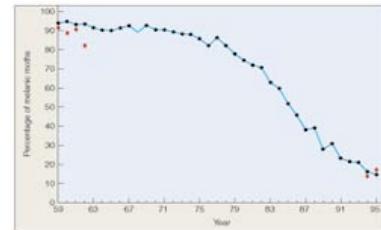


## Änderung von Genfrequenzen



Freeman,S., Herron, J.C. (1998) Fig .6.25

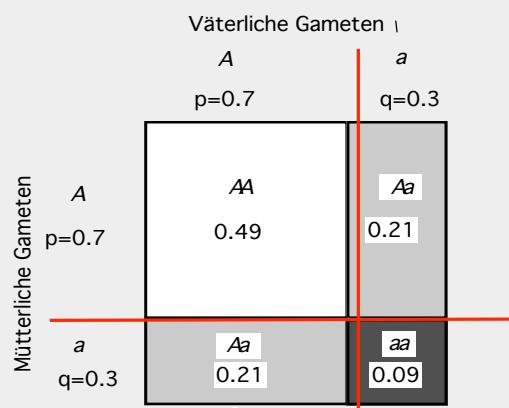
## Änderung von Genfrequenzen



Frequency of melanic form in England  
(1959-1995)

Melanic and wild-type *Biston betularia*

## HWG: Schema



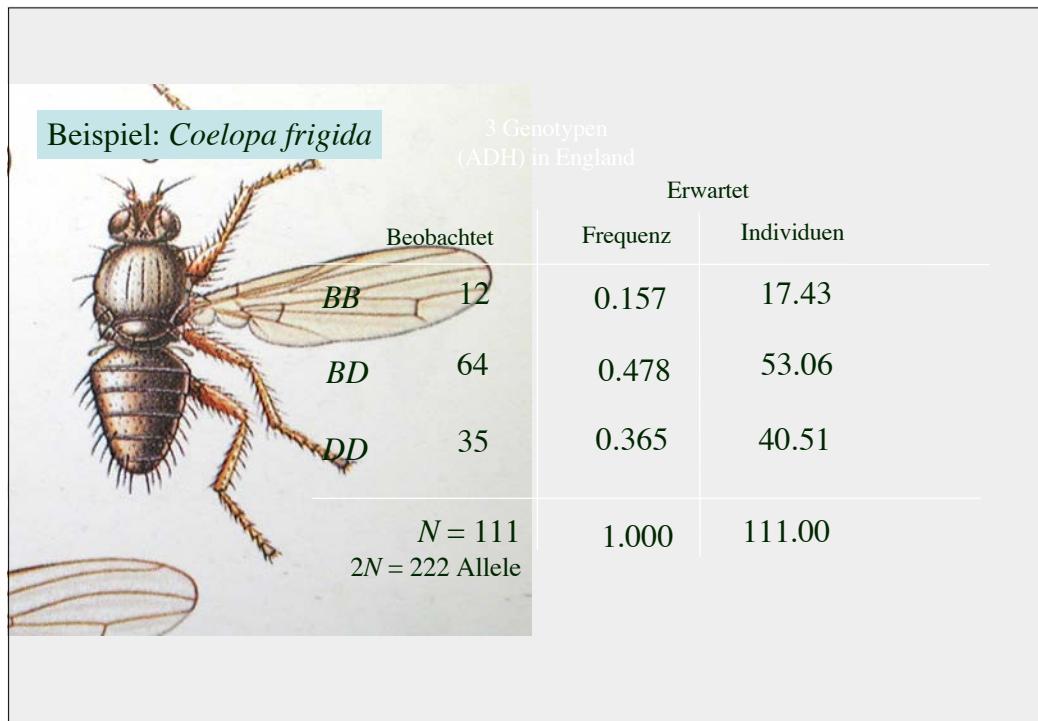
## Änderung von Genfrequenzen

Beispiel: Motte (*Panaxia dominula*)

$$\chi^2 = \sum (O-E)^2/E$$

FIGURE 5.4 Variation due to two alleles at a single locus in the moth *Panaxia dominula*. Type AA is most common genotype; AB is middle type heterozygote; BB is least common genotype. In AA, the central white spot on the forewing is reduced or absent and the amount of black on the hindwing is less than in AA. The heterozygote is intermediate. (After Paul 1971.)

	Beobachtet	Erwartet	$\chi^2$
AA	1469	$p_A^2 N =$ 1467.1	0.002
AB	138	$2p_A p_B N =$ 141.5	0.087
BB	5	$p_B^2 N =$ 3.4	0.753
$N = 1612$			



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Beispiel: Fliegen (*Coelopa frigida*)

		Erwartet	
	Beobachtet	Frequenz	Individuen
<i>BB</i>	12	0.157	17.43
<i>BD</i>	64	0.478	53.06
<i>DD</i>	35	0.365	40.51
$N = 111$ $2N = 222$ Allele		1.000	111.00