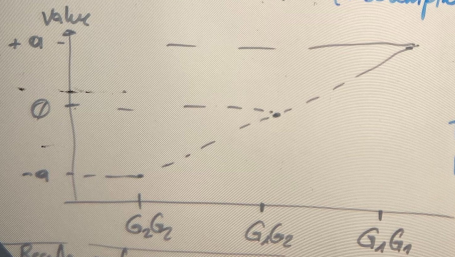


$$\begin{aligned}
 \Delta_1 BV_{11} - BV_{12} &= 2q\alpha - [(q-p)\alpha] \\
 &= 2q\alpha - q\alpha + p\alpha \\
 &= q\alpha + p\alpha = \underbrace{(p+q)}_{=1}\alpha = \alpha
 \end{aligned}$$

$$\begin{aligned}
 \Delta_2 - BV_{12} - BV_{22} &= (q-p)\alpha - [2p\alpha] \\
 &= q\alpha - p\alpha + 2p\alpha \\
 &= q\alpha + p\alpha = (p+q)\alpha = \alpha
 \end{aligned}$$

- The effect of substituting a  $G_2$ -allele by a  $G_1$ -allele on the breeding value is a constant and corresponds to  $\alpha$ . Hence,  $\alpha$  is called **Allele Substitution Effect**

- Special case for  $\alpha = a + (q-p)d$  occurs when  $d=0 \Rightarrow \alpha=a$  (Assumption is used in Genomic Selection in Swiss Dairy Cattle Breeding)



The effect of a single locus on a quantitative trait is determined by just one parameter  $q$ .

Results so far:

Genotype

BV