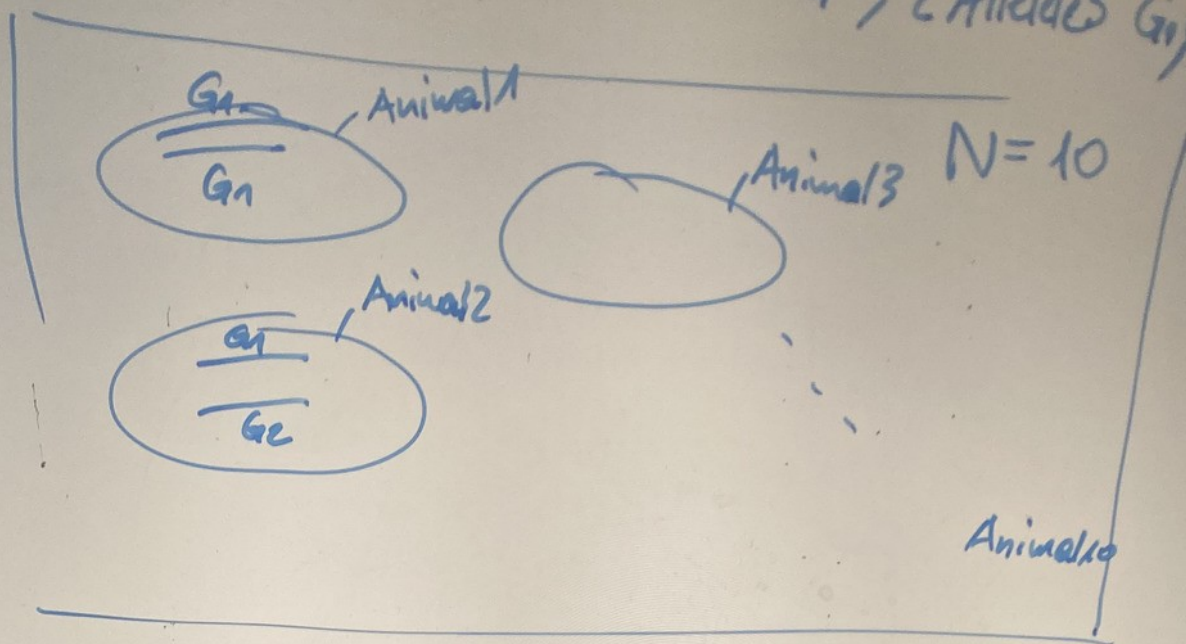


Population :

1 Locus : G , 2 Alleles G_1, G_2



- Locus G shows 2 different variants (G_1 and G_2) in the population
- Di-plotoid : 2 alleles mean 3 genotypes
 - G_1G_1
 - $G_1G_2 \Rightarrow$ heterozygous
 - G_2G_2
- Genotypes $G_1G_2 = G_2G_1$
(not the case with haplotypes)