

Dr. Rajesh Verma, Assistant professor and Head, U.G. Department of zoology, D.K. College Dumeron Notes for B.sc part 3rd, paper V11.

∴ Write Notes on Genetic Equilibrium

DEFINITION:- In population genetics, the Hardy Weinberg principle, also known as the Hardy - Weinberg equilibrium, model, theorem, or law, states that allele and genotype frequencies in a population will remain constant from generation to generation in the absence of other evolutionary influences.

Genetic equilibrium :-

Genetic equilibrium is the condition of an allele or genotype in a gene pool (such as a population) where the frequency does not change from generation to generation. Genetic equilibrium describes a theoretical state that is the basis for determining whether and in what ways populations may deviate from it. Hardy-

Weinberg equilibrium is one theoretical framework for studying genetic equilibrium. It is commonly studied using models that take as their assumptions those of Hardy-Weinberg, meaning,

- No gene mutations occurring at that locus or the loci associated with the trait.
- A large population size.
- Limited - to - no immigration, or emigration genetic flows.
- No natural selection on that locus or trait.
- Random mating (panmixis).

It can describe other types of equilibrium as well, especially in modeling contexts. In particular, many models use a variation of the Hardy-Weinberg characters being

present, these instead assume a balance between the diversifying effects of genetic drift and the homogenizing effects of migration between populations. A population not at equilibrium suggests that one of the assumptions of the model in question has been violated.

Theoretical models of genetic equilibrium :-

The Hardy-Weinberg principle provides the mathematical framework for genetic equilibrium. Genetic equilibrium itself, whether Hardy-Weinberg or otherwise, provides the groundwork for a number of applications, including population genetics, conservation and evolutionary biology. With the rapid increase in whole genome sequences available, as well as the proliferation of anonymous markers, models have been used to extend the initial theory to all manner of biological contexts. Using data from genetic

markers such as ISSR and RAPDs as well as the predictive potential of statistics. studies have developed models to infer what processes drove the lack of equilibrium. This includes local adaptation, range contraction and expansion and lack of gene flow due to geographic or behavioral barriers, although equilibrium geographic or behavioral barriers, although equilibrium modeling has been applied to a wide range of topics and questions.

Equilibrium modeling have led to developments in the field. Because allelic dominance can disrupt prediction of equilibrium, some models have moved away from using genetic equilibrium as an assumption. Instead of the traditional F-statistics, they make use of Bayesian estimates. Holsinger et al. develop

an analog to FST, called theta. studies I have found Bayesian estimates to be better predictors of the patterns observed.

Biological study systems :-

Genetic equilibrium has been studied in a number of taxa. Some marine species in particular have been used as study systems. The life history of marine organisms like sea urchins appear to fulfill the requirements of genetic equilibrium modeling better than terrestrial species. They exist in large, panmictic populations that don't appear to be strongly affected by geographic barriers. In spite of this, some studies have found considerable differentiation across the range of a species. This indicates that genetic equilibrium may be rare or difficult to identify in the wild, due to considerable local demographic changes on shorter time scales.