## 4. Chi-square statistics and Probability

For determining chi-square or goodness of fit, the size of the population must be considered. Suppose in one cross of tall and dwarf plants, out of the 100 plants of $F 2,70$ are tall and 30 dwarf instead of 75 and 25 as expected from a 3: 1 ratio. Obviously, there is a deviation of 5 from the normal. In a second cross where 1000 plants of $F 2$ progeny showed the same numerical deviation of 5 , there were 745 tall plants and 255 dwarf ones. In the third cross, the 1000 plants of $F 2$ generation appeared in the same proportion as in the first cross (70:30 or 7: 3 ) so that 700 plants were tall and 300 dwarf. The chi-square formula given below will show whether the differences in the observed results of the three crosses are significantly different from the 3: 1 ratio or not (See Table)

Thus, chi square is defined as the sum of the quotients obtained from division of squares of deviations of observed frequencies from the expected frequencies of various classes with the respective expected frequencies. This may have expressed as follows:

$$
X^{2}=\sum \frac{(0-e)^{2}}{e}
$$

Where, $\Sigma=$ summation, $\mathrm{O}=$ observed frequency of a class, and $\mathrm{E}=$ Expected frequency of the class.

| No.of plants in F2 generation | Cross no. 1 <br> $\mathbf{1 0 0}$ |  | Cross no. 2 <br> $\mathbf{1 0 0 0}$ |  | Cross no. 3 <br> $\mathbf{1 0 0 0}$ |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- |
|  | Tall | dwarf | Tall | dwarf | Tall | dwarf |
| Observed | 70 | 30 | 745 | 255 | 700 | 300 |
| Expected $(e)$ | 75 | 25 | 750 | 250 | 750 | 250 |
| Deviation $(d)$ | 5 | 5 | 5 | 5 | 50 | 50 |
| $d^{2}$ | 25 | 25 | 25 | 25 | 2500 | 2500 |
| $d^{2} / e$ | 0.33 | 1 | 0.033 | 1 | 3.3 | 10 |
| $\chi^{\alpha^{2}=\Sigma\left(d^{2} / e\right)}$ | $1+0.33=1.33$ | $0.033+1=0.133$ | $3.3+10=13.3$ |  |  |  |

The smaller the chi-square value the more likely it is that deviation has occurred due to chance. Now to find out whether the differences between expected and observed results are due to chance alone or not, we must be familiar with two more concepts namely

## degree of freedom and level of significance.

Degrees of Freedom: The number of degrees of freedom is calculated as the number of classes whose value is required to describe the outcome from all classes. The concept of degrees of freedom is important in experiments and genetic ratios because one must consider the total number of observed individuals in the experiment as a fixed or given quantity. This fixed quantity is composed of one or more classes some of which are variable. In the experiment between tall and dwarf pea plants there are only two classes, tall and dwarf. As soon as the number of one class is set, the other can be determined. Thus, when two classes are scored, there is one degree of freedom. In an experiment where three classes are scored, there are two degrees of freedom, and so on. The rule states that for the kind of genetic experiments described, the degrees of freedom are equal to one less than the number of classes.

Level of Significance: In the experiment described the actual ratio departs from that Which is expected. We must now determine how significant this discrepancy is so that we can decide to accept or reject the results. Small discrepancies are not significant; large discrepancies are significant and lead to rejection of a result or hypothesis. Therefore, values are assigned to these two kinds of discrepancies - the large discrepancies are the largest $5 \%$ and small discrepancies are remaining $95 \%$. On this basis if the discrepancy lies in the large class it is significant and the result may be discarded. The $5 \%$ frequency value that enables us to reject the result is called the $5 \%$ level of significance. The level of significance can be changed. If 5\% is too high we can decide on a low level of significance say $1 \%$. In this case it is not so easy to reject a result. Contrarily, if we decide on a high level of significance say $10 \%$, it is easier to reject a result. Usually the accepted level of significance is between the two extremes, that is $5 \%$.

Chi-square: After determining the degrees of freedom in an experiment and deciding on the level of significance, the actual size of the discrepancy between expected and
observed is found by chi-square. Statisticians have prepared tables that relate the number of degrees of freedom with the probability that particular groups of chi-square values will be found (Table).

We can now examine the results of the experiment described in Table 1. The chisquare values of the first two crosses are 1.33 and 0.133 . Both are acceptable discrepancies because these values are smaller than the chi-square value for one degree of freedom given as 3.84 in Table 2. The results of the first two crosses therefore may be considered to be consistent with Mendel's hypothesis, the difference between expected and observed being due to chance.

TABLE 2. Values of chi-square corresponding to $\mathbf{5 \%}$ and $\mathbf{1 \%}$ levels of significance and up to $\mathbf{1 0}$ degrees of freedom

| Degrees of freedom | Level of significance |  |
| :---: | :---: | :---: |
|  | $\mathbf{5 \%}$ | $\mathbf{1 \%}$ |
| 1 | 3.84 | 6.64 |
| 2 | 5.99 | 9.21 |
| 3 | 7.82 | 11.34 |
| 4 | 9.49 | 13.28 |
| 5 | 11.07 | 15.09 |
| 6 | 12.59 | 16.81 |
| 7 | 14.07 | 18.48 |
| 8 | 15.51 | 20.09 |
| 9 | 16.92 | 21.67 |
| 10 | 18.31 | 23.21 |

## Probability:

In the experiments described above the $F 2$ generation segregates in a particular ratio. The numbers of individuals in each ratio result from chance segregation of genes during gamete formation, and their chance combinations to form zygotes. Since these are chance events, accurate predictions about the results cannot be made. This is especially true in cases where the progeny is limited to a small number such as in experiments in animal breeding and even more so in human pedigree studies. It is hardly possible to predict the appearance of a certain phenotype or genotype. But what we can say is that there is a certain probability of occurrence of a given genetic event.

In a general way, we can say that the probability or chance that an event will occur can be defined as the proportion of times in which that event occurs in a very large number of trials. If there are $x$ trials and the event occurs on the average $y$ times during the $x$ trials, then the probability is expressed as $y / x$. The probability would be between zero and one. The closer the probability is to zero, the less chance is there for the event to occur. When probability is one, the event is certain to occur.

## Rules of probability:

The multiplication rule: While considering probabilities it is important to note that the inheritance of certain genes such as for height of pea plants (tall or dwarf), flower colour (red or white), seed texture (round or wrinkled) are independent events. If we consider each gene separately, the probability of any one $F 2$ plant being tall is $3 / 4$, and of its being dwarf $1 / 4$.

Similarly, there is a probability of $3 / 4$ that an $F 2$ plant will bear red flowers and $1 / 4$ that the flower will be white. Now what is the probability that an $F 2$ plant would be both tall and coloured? Assuming that the inheritance of each gene is an independent event, the probability that a plant be coloured and tall is equal to the product of their individual probabilities, i.e., $3 / 4 \times 3 / 4=9 / 16$. Thus there are 9 chances out of 16 that an $F 2$ plant be coloured and tall. Stated in a general way, when the probability of an event is independent of that of another event, and the occurrence of one does not influence the occurrence of the other, probability that both events will occur together is the product of their individual probabilities.
The Addition Rule: The probability that one of several mutually exclusive events will occur is the sum of their individual probabilities. This law is applicable when different types of events cannot occur together. If one occurs the other is excluded. When a coin is tossed there are two alternatives, either heads or tails will appear. If the probability for heads is $1 / 2$, for tails $1 / 2$, then the probability that either heads or tails will appear is $1 / 2+1 / 2=1$.

## DOMINANCE RELATIONSHIPS

## i) Complete dominance:

In case of complete dominance, the phenotype produced by heterozygotes is identical with that produced by homozygotes for the concerned dominant allele. The dominant allele in such a situation is known as completely or fully dominant. An example of

