Topic – Aneuploidy

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Aneuploidy

Variations in chromosome number may occur that do not involve whole sets of chromosomes, but only parts of a set. The term aneuploidy is given to variations of this nature.

Aneuploidy describes an numerical change in part of the genome, usually a change in the dosage of a single chromosome. Individuals that have an extra chromosome, are missing a chromosome, or have a combination of these anomalies are aneuploid. This definition also includes pieces of chromosomes. Thus, an individual in which a chromosome arm has been deleted is also considered to be aneuploid.

A loss or gain of one or few chromosomes as compared to the somatic chromosome number of a species is known as aneuploidy. The deviation is from a diploid setup (2n). The change in chromosome number do not involve the whole genome; they involve only one or few chromosomes of the genome.

Aneuploidy means presence of a chromosome number which is different than an exact multiple of a basic chromosome number. For instance, if 7 is the basic chromosome number, where the somatic chromosome number is 2n=14, then chromosome numbers like 2n=15 or 2n=13 would be aneuploids. Thus aneuploidy can be either due to gain of one or more chromosomes or due to loss of one or more chromosomes from a complete somatic or diploid chromosome complement.

Classification of Aneuploids

A. Hyperploidy- Aneuploidy which is due to gain of one or more chromosomes from a complete somatic or diploid chromosome complement.

1. Trisomy(2n+1) - Diploids that have one extra chromosome are represented by the chromosomal formula 2n+1. Trisomic individuals possess one extra chromosome than 2n. Trisomics are grouped into the following categories depending on the constitution of the additional chromosome.

- Primary trisomic- The additional chromosome is an unaltered member of the haploid chromosome complement of the species. Therefore, there can be n different primary trisomics of a species. e.g. 7 in barley 10 in maize, 12 in Datura etc.
- Secondary trisomic- The extra chromosome present is an isochromosome(a metacentric chromosome in which the two arms are identical; produced through a transverse division of the centromere of a chromosome. Therefore, 2n different types of secondary trisomics are possible in a species. e.g. 1.2 would produce 1.1 and 2.2 secondary trisomics.
- Tertiary trisomic- Trisomics where the extra chromosome present is a segment from non-homologous chromosome. It has a translocated chromosome in addition to the normal somatic complement. A large number of different types of tertiary trisomics are possible in a species, e.g. chromosome 1.2 and 3.4 may give rise to the following tertiary trisomics 1.3, 1.4, 2.3, 2.4, 3.1, 3.2, 4.1 and 4.2.
2. Tetrasomy (2n+2) - When one chromosome of an otherwise diploid organism is present in quadruplicate, this is expressed as 2n+2. Tetrasomic individuals have one additional chromosome pair. Complete one homologous pair is added.

3. B. Hypoploidy - Aneuploidy which is due to loss of one or more chromosomes from a complete somatic or diploid chromosome complement.
1. Monosomy (2n-1) - Diploid organisms that are missing one chromosome of a single pair are monosomics with the genomic formula 2n-1. There is deficit of a chromosome from the somatic number of a species. Thus, in this any one chromosome is missing from a homologous pair.

2. Nullisomy (2n-2) – An organism that has lost a chromosome pair is a nullisomic. In this
complete one homologous pair is lost. The result is usually lethal to diploids.

**Origin and Production of Aneuploids**

- Due to abnormal distribution of chromosomes during anaphase of meiosis, because of irregular chromosome distribution at the poles one daughter cell receives one or more extra chromosome and the counterpart lacks one or more chromosomes. This type of abnormal situation arises when one or more pair of chromosomes (bivalents) fail to separate or disjoin and move to one pole as such. This is called disjunction. So gametes having extra chromosome when fuse with normal haploid gametes results in hyperploidy and those containing one or more chromosomes less than a haploid no. if unite with normal haploid gamete produces hypoploidy.
- Loss of individual chromosome in meiosis leading to the formation of nuclei with hypoploid chromosome number.
- Irregularities in the segregation of chromosomes during meiosis in polyploids causes development of aneuploids.

**Trisomy**- Although trisomics may occur spontaneously in nature, they are rare.

**Tetrasomy**- In diploids tetrasomics are rarely obtained, but in polyploids these can be obtained by
selfing progenies of trisomics.

**Monosomy** - Spontaneous production of monosomics have been reported in wheat, tobacco, cotton and oats. Irradiation treatment of the inflorescence leading to non-disjunction of a normal bivalent produces monosomics in cotton. In wheat a haploid as female when crossed to normal hexaploid male produced monosomics.

**Nullisomy** - Selfing of monosomics give rise to nullisomics.

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**Cytological behaviour of Aneuploids**

**Trisomy** - One of the pairs of the chromosomes has an extra member, so that a trivalent structure may be formed during meiotic metaphase. If two chromosomes of the trivalent go to one pole and the third goes to the opposite pole, then gametes will be \((n+1)\) and \((n)\).

In primary and secondary trisomics, the three homologues of the concerned chromosome may pair with each other to form a trivalent of different configurations at metaphase1, or two of the chromosomes may form a bivalent leaving the additional chromosome as a univalent.

In a tertiary trisomic a dumb-bell shaped figure containing five chromosomes at metaphase1 is seen. Thus secondary and tertiary trisomics may be distinguished from primary trisomics due to the presence of a ring of three chromosomes and a dumb-bell shaped figure of five chromosomes respectively.

**Tetrasomy** – A quadrivalent may form for this particular chromosome during meiosis. Meiotic chromosome pairing usually produces quadrivalents (four synapsing chromosomes) that can
produce genetically balanced gametes if disjunction is by twos, that is, two chromosomes of the quadrivalent going to one pole and the other two to the opposite pole. If disjunction is not stabilised in this fashion for all quadrivalents, the gametes will be genetically unbalanced. Sterility will be expressed in proportion to the production of unbalanced gametes.

**Monosomy**- The single chromosome without a pairing partner may go to either pole during meiosis, but more frequently will lag at anaphase and fails to be included in either nucleus. Monosomics can thus form two kind of gametes, (n) and (n-1). In plants, the n-1 gametes seldom function. In animals, loss of one whole chromosome often results in genetic unbalance, which is manifested by high mortality or reduced fertility. Since in a monosomic one chromosome of the haploid complement does not have a pair, it remains unpaired during meiosis and is present as a univalent at metaphase1. During anaphase1, the bivalents disjoin normally, while the single univalent may follow any one of the following courses:  
- move to one of the two poles.  
- Lag behind at the metaphase plate and be ultimately lost.  
- Divide into two chromatids which move to the opposite poles.

**Nullisomy**- Chromosome pairing and the disjoining of bivalents to produce gametes is normal. But in certain nullisomics regular chromosome pairing is disturbed and multivalent associations are present at metaphase1.

**Uses of Aneuploids**  
1. They have been used to determine phenotypic effects of loss or gain of different chromosomes.  
2. Aneuploids have been used to produce chromosome substitution lines which yield information on the effects of different chromosomes of a variety on various plant characteristics, when they are placed singly in the same genetic background.  
3. Aneuploid analysis permits the location of a gene as well as linkage of a group, onto a specific chromosome.  
4. Aneuploids are useful in the identification of the chromosomes involved in translocations.  
   - Use of trisomics in chromosome mapping- trisomics can be utilized for locating genes on specific chromosomes and for finding out distances of these genes from centromere, a technique called chromosome mapping. If linkage groups are already established in an organism, trisomics can be effectively used for assigning these linkage groups to specific chromosomes.  
   - Tetrasomics are used to produce compensating nullisomic tetrasomic lines suggesting homeologous relationships.  
   - Monosomic analysis is used for locating genes on chromosomes in polyploid crops.  
   - Use of nullisomics in locating genes on chromosomes- The genes which are dominant can often be located by observing the absence of the trait in the nullisomics. For instance, Chinese Spring wheat has red seed, but nulli-3D has white seeds suggesting that the gene for red grain is located on 3D.
References