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MOLECULAR MARKERS AND PLANT BIOTECHNOLOGY

change began after discovery of the structure of DNA by Watson and Crick. DNA is bimolecular part of living organism consists chemical language that determines the features and characteristics of them. The different traits and physical features of plants are encoded in the plant's genetic material, the deoxyribonucleic acid (DNA). Now a day, scientists can determine which parts of the DNA are responsible for certain traits.

Proteins and enzymes are useful parts of all living organisms to grow, metabolize energy, and become what their genetic code dictates. Genes determine traits by controlling the production of proteins, including enzymes. Every cell contains the entire genetic information in the form of code needed to create and develop the organism. The interaction of genetic makeup and environmental factors shapes the nature of all living things.

Applications of agricultural biotechnology have considerably shortened the time of the development of new crop varieties using molecular marker techniques. The method, which makes it easier and faster for scientists to select plant traits, is called marker-assisted selection (MAS). All of the plant's genes together make up its genome. Some traits like flower color, may be controlled by single gene called monogenic and complex characteristics, like crop yield, maybe influenced by several genes called polygenic. Traditionally, plant breeders have selected plants based on their visible or morphological traits. called the phenotype. But, this process takes time and influenced by the environmental conditions, and costly - not only in the development itself, but also for the economy, as farmers suffer crop losses. As a shortcut, plant breeders now use marker-assisted selection. Molecular markers help in the identification of specific genes. The molecular markers are located near the DNA sequence of the desired gene. Since the markers and the genes are close together on the same chromosome, they tend to stay together as each generation by generation. This is called genetic linkage. This linkage helps scientists to predict whether a plant will have the desired gene. If researchers find the marker for a particular gene, it is called linked or specific marker for that gene. If the marker is present in any generation of plant it means the gene itself is present in that. As scientists find the location of molecular marker on a chromosome, and how close it is to a desired gene, they can determined the position of the markers and create a map which helps in the identification of genes on specific chromosomes. Now Scientists can produce detailed maps in only one generation of plant breeding. Using detailed information regarding genetic maps and locations of molecular markers, researchers can analyze a tiny bit of tissue from a newly germinated seedling. This makes molecular marker approach faster in comparision to conventional breeding. Once the tissue is analyzed through molecular techniques; scientists know whether that seedling contains the appropriate gene. If it doesn't, they can quickly move on and concentrate analysis on another seedling, eventually working only with the plants which contain the desired trait. Currently, molecular marker-assisted selection is already a routine step in breeding of most crops where the gene and the markers for a desired trait are known. This technique is being used in the efficient introgression of important genes into rice such as increased beta carotene content, bacterial blight resistance and submergence tolerance, in maize opaque gene to enhance the level of essential amino acids. Molecular markers are also used to differentiation of a line or variety. Biotechnology is depending on bioinformatics for data analysis. Bioinformatics use computer programme to analyze the genetic relationship of different lines or varieties. The genetically diverse lines or varieties used for selection of parents for the production of hybrid seeds. The genetic information also provides detail on the parentage of the line, the possible traits, and the unique identity of the plant useful for germplasm collection database.

One of the most striking differences between conventional breeding and the molecular approach is that the source of genetic material need not come from the same species. These new biotechnology techniques have prompted considerable debate on the ethical and moral aspects of this branch of science. All living organisms share the same genetic language in the form of their genetic material made up of four nucleotides. In fact, person probably share about half of his or her genetic information with a plant. And the genetic information from that plant can function in a other plant. New techniques even allow scientists to decide in which part of the plant tissue a trait should be expressed, such as the pulp versus the skin of a tomato.

When working with plant based foods, scientists seek to improve foods for the benefit of producers, consumers, or the environment. Consumers may benefit from improved nutritional value or food quality of that. Producers should be able to grow crops under adverse conditions, such as drought. Some genetically improved plant foods require fewer chemical applications during their growth and therefore have less environmental impact. the first food plants produced using the tools of biotechnology was slow-ripening Flavr Savr tomato. When scientists isolated the gene responsible for the softening enzyme and inserted it backwards into the tomato's genetic code, the resulting tomato was able to maintain good eating quality for a longer time than regular tomatoes. This technique allowed better tasting tomatoes to be grown and shipped to distant markets.

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