

**Session 1**  
**The Twentieth Bawden**  
**Lecture**

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## ABSTRACT

In this article I explore the benefits to mankind of various kinds of technology used to introduce new characteristics in living organisms, and consider the issues arising from the creation of new life forms. Bioengineering in its broadest definition includes engineering required for various methods to synthesise animal, plant and microbial products, and also includes devices to assist in the functioning of the human body. Recombinant DNA technology - genetic engineering - is the ultimate engineering, however, allowing mankind both to explore the very processes of life and to exploit the intellectual products. The subject matter encompasses a vast literature, replete with jargon and complex concepts. It is also one of the most rapidly advancing, innovative areas of science, worthy of substantial investment by private industry and by government alike. We live in the age of the biologist.

## INTRODUCTION

Last year, Professor C. R. W. Spedding presented the nineteenth Bawder Lecture, choosing as his topic the role and impact of technology, legislation and public opinion on modern agriculture. He noted that effective demand has the dominant impact on modern agriculture, and that technology is essentially an enabling procedure constrained by legislation, need and economics.

This paper focuses on technology of a special kind - variously referred to as bioengineering, genetic engineering, recombinant (r)DNA technology, genetic manipulation, genetic modification, depending on the sort of image that is meant to be conveyed. When applied to genetics, "engineering" has a strongly negative image in the minds of the public, many politicians and pressure groups; it indicates unnatural procedures with potentially calamitous risk. In my view, bureaucratic obfuscation, euphemisms, undue sensitivity, secrecy and dithering are no substitute for an open clinical approach to risk assessment, and recognition of the sheer brilliance and potential of the technology. It cannot be de-invented; it must be exploited responsibly. The need is pressing.

Genetic engineering currently has its greatest applications in pharmaceuticals and drug design. Studies on catalytic and other antibodies, human gene therapy, the Human Genome Project and drugs whilst being anthropocentric nonetheless mutually interact with related studies on plants, fungi and microorganisms to advance knowledge in the whole of genetic engineering. I shall concentrate on crop plants, the underlying theme of this Conference, avoiding as much as possible specialist vocabulary.

Agriculture in Developed Countries

From the perspective of most members of the public in the western world, agriculture is not viewed as a priority topic. In fact, its importance is overstated. Most references to economic analyses in agriculture and horticulture relate simply to gross traded values of commodities, hectares sown, yield estimates, major categories of land use, imprecise crop and livestock categories, unit prices, direct employment and percentage origin of Gross Domestic Product. Data relating to (i) processing values for the food and non-food sectors; (ii) development and maintenance of rural, industrial and marketing infrastructures; (iii) amenity and tourism; (iv) influence of subsidies; (v) costs of import substitution; (vi) assessments of trends in and prospects for exports and trading; (vii) relative social values to the national and local economies; (viii) cultivar performance and market share; and (ix) indirect employment, are usually ignored because they are often unreliable, incomplete, out of date, anecdotal, disputed or subject to commercial secrecy. International comparisons are made especially difficult by variations in the methods and dates of sampling, unspecified types of analyses and fluctuating currency exchange values. As a general point, it is germane to note that arable and horticultural crops produced in the UK had an annual value of £6,527 billion in 1990 (1.4% of Gross Domestic Product) before industrial processing and value-added contributions. During the past three decades there have been dramatic, research-driven, improvements in commodity processing, crop productivity and efficiency throughout the world.

Within Europe, public and political opinions of the effects of the Common Agricultural Policy of the European Communities tend to relate to unwelcome changes to the countryside, expensive food surpluses/stockpiles, subsidies, quotas, set-aside land, extensification, restrictions on pesticides and nitrogenous fertilizers, undignified animal welfare, and reappraisals of R&D priorities. All in all, opinions are not favourable towards agriculture, although the products are a basic need and civilisation is agriculture-dependent.

Population Pressures

A billion extra people are projected to be added over the next ten years to the world population which presently exceeds 5.4 billion (see annual reports from the World Commission in Environment and Development; FAO, World Resources Institute and Worldwatch Institute). Inordinate strains will be placed on the less-developed countries for food, water, shelter, fuel, education and welfare. Large parts of Africa, especially, face dismal prospects. Access to the media and advanced medicine, though, will ensure that all citizens will demand improving quality of life regardless of local economic situations. Low-grade grazing systems coupled to poor, unsustainable agricultural systems will inexorably lead to the acceleration of deforestation, soil erosion, desertification and the rapid loss of natural and managed ecosystems, destroying genetic and environmental diversity. Social instability, emigration and trade disruption seem inevitable for the poorer of the less-developed countries.

Largely as a result of major if unsung technological successes in the recent past, especially in plant breeding and pathology, agriculture and the related life sciences are universally assumed to be able to adapt without major investment to meet the challenges of population growth. At the same

time, it is expected that agriculture should not adversely affect the natural flora and fauna, nor exacerbate any potentially undesirable effects ("climate change") of the changing gaseous composition of the atmosphere since the advent of the Industrial Revolution.

It seems remarkable that there is so little publicity given to the loss of cultivated land throughout the world. Soil erosion, pollution, buildings, roads, airports, and recreation facilities account for the main loss of productive land. Modern monocultural agricultural systems can cause problems e.g. use of xenobiotics, soil compaction and erosion, salinity effects and changes in the soil flora and fauna, but traditional methods (e.g. slash-and-burn, uncontrolled grazing) can be even more erosive without even being productive. All too frequently, third-world agriculture can incorporate many of the bad practices of high-input agriculture now being phased out in the western world. About 85% of the growth in population occurs in developing countries where the numbers of malnourished people have increased by 35% since 1980. In the tropical zones, the area of cultivated land per capita has declined from 0.28 ha in 1971 to 0.22 ha in 1986; this figure masks urbanisation, fragmentation of farms, and expansion of cultivation into virgin lands unsuitable for arable farming in the medium term.

Given that the area of land under cultivation is a limited resource and difficult to increase without massive migrations of people and devastation of forests, that pests and diseases have a phenomenal ability to circumvent control measures, and that research and development demand long-term commitment, the global picture is far from bright until the demand of the world population matches sustainable resources.

One possible or probable scenario is that the industrially underdeveloped - or low income - world will become the major source of manufactured goods, effectively reversing the trend in trade established since the Industrial Revolution (Carruthers, 1993). The economies of most of the countries of the Pacific rim are buoyant, and in Asia there are several countries with sophisticated, urbanised workforces able to operate efficiently and compliantly with relatively low incomes. Scientific intercommunication, multinational trading, and improving education in the low-income world ensure that invention, intellectual property and service industries will not be the preserve of the present developed world. Moreover, agriculture in developing countries is no longer regarded as the engine of economic growth - witness the pressures on the Consultative Group on International Agricultural Research. Thus it is likely that most of the world's food production would take place in the temperate zones. Whether or not there would be the means to pay for the food is a moot point.

One global feature is the growing divide between stable or expanding urban populations and their rural counterparts. By way of example, in the UK 91.5% of the population is urban, with relatively low mobility, a birth rate per 1,000 population of half the world average, and a population doubling time in excess of 100 years. The overall population density is high (235 persons km<sup>-2</sup>) revealing the extent of crowding in the urban area. No wonder misunderstanding of the rural economy is becoming so pronounced.

Conventional management of terrestrial and aquatic resources will not meet future demands. Over 90% of the world's population depend on just 15 plant and 7 animal species for food (Hillman, 1992); a tiny genetic reservoir to combat the ravages and vagaries of pests, diseases and inclement conditions. To this must be added the fact that as the only animal to cook

and thereby broaden the range of acceptable food species and types, which is faced in the arid and semi-arid regions of the developing world with a shortage of fuel for cooking.

Woody perennial species present one of the stiffest challenges for crop management. New initiatives are desperately needed for breeding, selection, propagation and health of trees and shrubs.

#### Plant Breeding

Central to the ability of the bulk of the population to move from food cultivation and harvesting to engage in social and technological advancement is the provision of improved crop plants throughout the ages. Plant breeders have always been involved in genetic engineering. Characters including yield, performance, resistance or tolerance to pests and diseases, quality components, uniformity and lack of prolonged dormancy periods represent the main selection criteria. Together with advances in automation, storage and processing there is a complacent view that plant breeding will perpetually answer basic nutritional needs for burgeoning populations. This is acceptable.

Plant breeding programmes are protracted, expensive and are rarely allowed to proceed without interference. Basic to the needs of such programmes is access to genetic resources for parental material. Unfortunately, there has been severe attrition of genetic diversity by losses of diverse wild habitats, traditional farming areas, valuable collections and obsolete landraces. Breeders need to screen vast numbers of clones over many years, carry out regional trials, multiply stocks, access statutory trials and be involved in marketing. Other problems faced include imprecise predictions of genotype by environmental interactions, incompatibility systems between and within species affecting the ability to cross-breed, juvenility or ripeness-flower phases, seasonal growth patterns, changing disease virulence patterns and disease vector distributions, and complex breeding objectives involving polygenic characters. Conventional plant breeding is well-established but needs to be supplemented by bioengineering technology to allow access to new sources of genetic variability, to speed up the process, to unravel the complexities of genomes (genetic constitutions), to understand the processes involved in breeding, and to improve the prediction of performance of products arising from the breeding programmes.

#### BIOENGINEERING PLANTS

Modern technologies are reducing the reliance on a combination of serendipity and bulk selections for plant breeding and food processing facilities.

Modern biotechnology has its beginnings with the well-known early studies of deoxyribonucleic acid (DNA - actually a base!) molecules of which act as carriers of genetic information. In 1973 recombinant DNA techniques were discovered at Stanford University in California. Shortly thereafter, animal hybridomas were created in the Laboratory of Molecular Biology, Cambridge, a discovery which initiated the monoclonal antibody diagnostics industry. During the 1980s, the inherent similarity of the genetic language in the major groups of organisms was demonstrated by the insertion of genes or sequences into the DNA of recipient (transgenic, genetically manipulated or modified, recombinant) organisms and sub-cellular entities containing nucleic acid.

These insertions (constructs) lead to the induction of new traits in the genetically modified organisms (GMOs).

In this period, the speciality of protein engineering came to the fore with an aim of producing customised, biologically active (eg enzymes, antibodies) and structural (eg collagen) proteins for a wide range of purposes. Proteins are genetically coded amino-acid polymers with a molecular shape that helps determine function. Studies in protein engineering have been particularly facilitated by site-directed mutagenesis of the genetic code.

Both DNA and ribonucleic acid (RNA) consist of long sequences of nucleotide bases that are attached to a backbone of alternating sugar and phosphate groups. It is the sequence of bases that constitutes the genetic code. A gene is a portion of nucleic acid that carries the code for synthesising a specific protein or part of a protein. Each nucleic acid molecule is comprised of a linear series of genes. Between and within genes (the coding sequences or exons), though, there tend to be the so-called introns, or intervening sequences of nucleotide bases that apparently do not code for proteins. Functional RNA molecules have the introns removed and "spliced out" by cutting out and joining (ligating) the cut ends together.

DNA acts as the blueprint whereas RNA of different types act in a wide range of functions, including the role of messenger for conveying the blueprint code to the various sites in the cell where protein synthesis takes place.

### Splicing and Ribozymes

Splicing is one of the key control points for cell metabolism, development and differentiation generally (the "splice of life") and the mechanisms of intron recognition and splice-site selection are crucial to understanding regulated gene expression, especially in transgenic plants. Certain RNA molecules have the ability to act as catalysts - the ribozymes, which have led to the application of the so-called gene shears for selective cleaving of RNA molecules without the presence of any protein; they may also be used to inactivate or destroy RNA viruses.

### Antisense Technology

Gene expression in living organisms can be prevented by synthesising relatively short RNA or DNA molecules (oligonucleotide primers or oligomers) which bind specifically and selectively to complementary sequences on the target RNA or DNA molecules, switching off genes such as those controlling ethene (ethylene) biosynthesis. Both ribozymes and antisense technology can be harnessed to combat viruses and control developmental functions in plants.

### DNA Fingerprinting or Profiling

When DNA is cleaved by highly specific restriction enzymes, the lengths of the resultant fragments produce an exceptionally reproducible pattern (restriction fragment length polymorphisms, RFLP) in electrophoretic separations. The patterns are inherited and can be used for diagnostic purposes. RFLPs (Bottstein *et al.* 1980) are used for tagging genes with tightly linked markers for selection in plant breeding programmes. They are also used for map-based gene cloning, assessment of genetic variability, and also for comparative genome mapping to study relationships between organisms. Locating genes with respect to DNA markers on an RFLP-based map provides

starting point for cloning genes by "chromosome walking" down overlapping large pieces of the chromosome from the RFLP tag to the gene of interest.

### Polymerase Chain Reaction (PCR)

A major advance has been the ability to produce simply, consistently, automatically and cheaply, identical copies of specific DNA sequences. In essence, and there are many variants, PCR involves a series of copying cycles in which double-stranded DNA fragments are firstly denatured to provide templates for the "annealing" or binding by base pairing of two synthetic primers (short DNA molecules of known sequence that flank either side of the target region) to complementary sequences on the DNA template strands. Thereafter, the bound primers are elongated enzymically by the addition of nucleotides from the reaction mixture. The resultant new DNA strands are complementary to the template strands. One PCR cycle duplicates one DNA fragment and hence produces two copies. This amplification system is employed to produce adequate amounts of single or multiple genes for both fundamental and applied studies. PCR has also been used for demonstrating, mainly in animal genome mapping, high levels of polymorphism in the repeat number for multiple sequence tandem repeats or microsatellites which are potentially ideal genetic markers.

### 2 DNA Approach

From the foregoing, a basic shift in philosophy has taken place in the last few decades, from the inference of the genotype from a study of phenotype morphology as was once carried out by conventional plant breeders, to a direct analysis of DNA sequence information i.e. a change in emphasis from Mendelian genomic genetics.

### Selectable Marker Genes

The incorporation of dominant selectable marker genes (marker genes or markers) together with the DNA construct aids in the identification and selection of transformed cells from a background of non-transformed cells. Marker genes also assist in confirming the identity of transgenic plants for legal purposes. Marker genes may encode a protein or enzyme that modifies a toxic substance to render it harmless, thereby allowing the transgenic cell to grow in the presence of the toxic substance. Other encoded proteins may act with compounds to produce chromogenic compounds or emit light. Yet others enable the degradation of organic compounds, the utilisation of sugars, the conversion of heavy metal components into their metallic form. Some are even silent, producing specific, amplifiable DNA fragments e.g. microsatellite sequences.

Plant pathogenic microorganisms and viruses have been a major source of DNA fragments for constructs which contain the marker gene. Bacterial transformation processes (e.g. *Agrobacterium*, *Escherichia*), chemically induced gene transfer, electroporation, liposomes, injection and particle bombardment, co-cultivation and incubation are the most common procedures for introducing the constructs.

In addition to normal biochemical assays, the marker gene protein or enzyme can be isolated and detected by Western blotting, in which total proteins are extracted, separated and reacted with antibodies with specificity towards the gene product protein. The specific complex formed between the



antibody and the protein product can then be detected using as little as  $5 \times 10^{-9}$  g by a secondary reaction directed at the complexed antibody.

Molecular assays, such as Southern or DNA blotting, detect the DNA sequence of the marker gene directly. Total DNA from the genetically modified plant is digested with restriction enzymes each of which cuts the DNA in a precisely defined manner. The resulting DNA fragments are separated and reacted with the DNA sequence of the marker gene that has been tagged with a radioactive or, increasingly commonly, a chemical label. Only the sequence of the marker gene will react with the labelled probe to give a complex that can easily be detected and quantified.

Where the DNA sequence of the marker gene is known from previous work, small fragments of it (oligonucleotide primers) can be synthesised. Under appropriate PCR conditions these primers when incubated with DNA from the genetically modified plant result in repeated and selective amplification of the marker gene with increases of several-million fold possible.

Concerns have been expressed about the use and safety of selectable markers in GMOs released into the environment and also incorporated in foodstuffs. For example, over 30 species of plant have been modified with genes encoding resistance for 9 different antibiotics used in human and veterinary medicine.

#### CONTAINMENT AND COST

Comprehensive regulations and procedures to deal with laboratory activities and the containment of GMOs in UK laboratories will shortly come into force (Ratledge, 1993). To a certain extent, the Regulations redefine the terms. Examples of the techniques which constitute genetic modification are as follows:

- (i) recombinant DNA technique consisting of the formation of new combinations of genetic material by the insertion of nucleic acid molecules produced by whatever means outside the cell, into any virus, bacterial plasmid (autonomously replicating DNA circle) or other vector system so as to allow their incorporation into a host organism in which they do not occur naturally but in which they are capable of continued propagation;
- (ii) techniques involving the direct introduction into an organism of heritable material prepared outside the organism, including micro-injection, macro-injection and micro-encapsulation;
- (iii) cell fusion (including protoplast fusion) or hybridization techniques where live cells with new combinations of heritable genetic material are formed through the fusion of two or more cells by means of methods that do not occur naturally.

For the sake of common sense and expedience, conjugation, transduction, transformation or any other natural process, polyploidy induction and in vitro fertilization do not constitute genetic modification if they do not involve the use of recombinant DNA molecules or GMOs. Likewise, the Regulations shall not apply to the following techniques of genetic modification if they do not involve the use of GMOs as recipients or parental organisms:

- (i) mutagenesis;
- (ii) the construction and use of somatic hybridoma cells (i.e. for the production of monoclonal antibodies);
- (iii) cell fusion (including protoplast fusion) of plant cells where the resulting organisms can also be produced by traditional breeding methods;

(iv) self-cloning of non-pathogenic, naturally occurring micro-organisms with proven histories of safe use and no known adverse consequences to the environment - the so-called Group I micro-organisms where it is necessary that the vectors and inserts should also be well-characterized and free from harmful sequences and should in themselves be poorly mobilizable;

(v) self-cloning and non-pathogenic, naturally occurring organisms other than micro-organisms. Such organisms (i.e. plants and animals) must be safe in the containment facility as any recipient or parental organism.

Three levels of containment are envisaged for Group II organisms (those that do not conform to the definition of Group I microorganisms) and certain procedures involving Group I, depending upon the evaluation of the risk assessment.

Individual scientists are legally obliged to carry out scale-dependent risk assessments of their work which will then be scrutinised and approved by a local safety committee. The Health and Safety Executive may be involved and will charge a fee for their services and consent. Attention will also have to be paid to transport, storage and destruction of GMOs.

Historically, the UK has a superb safety record of containing and monitoring GMOs. The new Regulations may convince the public that the most stringent conditions are applied. Rather like other legal and statutory aspects of health, safety and the environment, though, there is an exquisite balance to be struck between unfettered irresponsibility and high-cost restraint, demotivation and export of intellectual property elsewhere. Resource-strapped organisations will definitely be ineligible for involvement with GMOs unless they become associated with regional centres specifically designed to meet ever-changing standards imposed by the UK or European communities.

#### RELEASE OF GMOs AND THEIR PRODUCTS TO THE ENVIRONMENT AND FOR FOODSTUFFS

By and large there is relatively little opposition to using genetic engineering in most areas of health care, but as the technology is finding application in food production, storage and processing there is widespread consumer concern about safety. Some GMOs and their products will continue to be contained strictly within the laboratory environment, others will be monitored closely over a long period to examine their suitability for release into the environment and for consumption. A case-by-case analysis will be carried out in every instance.

Safety is not absolute; accidents, exceptional circumstances incidents and testing occur. Anything is possible. Without resorting to examples of food transport policy, holidays in Florida, and the consequences of oxidative processes in living organisms, the nub of the argument is one of setting the level of acceptability of risk. In this instance we are considering the risk of all products arising from genetic engineering, and the risk is typically classified according to effects on mankind.

By way of definition, hazard is a situation that may lead to harm or loss; risk is the chance in quantitative terms of a defined hazard occurring.

Biologists appreciate that there is natural transfer of genes controlling the formation of toxic principles in plants as well as desirable features.

For the most part, species are not static genetically. By natural selection for example, resistances to a range of adverse xenobiotics can be developed

One huge problem is that of forecasting performance "in the field". We are not knowledgeable about selection pressures operating on organisms, nor about unique recombination events likely to occur. Frankly, it is not feasible for any committee or individual to assess risk for all the possible combinations of genes. Assessment can only come from experimentation and monitoring, employing a battery of scientific disciplines. To date, release experiments have not been problematical. Even naturally occurring mobile DNA elements are limited in their natural hosts, and can be regarded as excessively promiscuous. In his recent article, Wilson (1993) cites that an analysis of 393 defined field trials of transgenic plants (25 species) between 1986-1991 (in 21 countries) reveals that 50 involved "virus-resistance" traits. Field releases have shown that coat-protein-mediated protection may not behave as predicted in laboratory and growth-chamber experiments; generally, there is greater susceptibility to virus challenge.

Selectable marker genes, especially relating to the potential impact of antibiotic resistance, raise questions about safety. The potential for uncontrolled gene transfer in the intestinal tract, soil or by cross-fertilization, or for example herbicide resistance leading to the creation of weeds have received the most attention. What little information is available would not indicate unacceptable risk. Obviously, a great deal of research is still required to quantify risk, if any, and to make recommendations on the use of marker genes. With time, their use will in any case decline, essential markers may need to be inactivated or eliminated prior to release and consumption of the transgenic organism.

Organisational structures to monitor GMO release are already in place in many countries. Much of the real work, though, is labour-intensive, from removal of flowers to stop breeding of transgenics with other plants (especially weeds), surveying experimental sites to eliminate propagules (seeds, tubers etc.) in the seasons following the experiment, and monitoring gene flow through ecosystems. It is always a good policy for any country to monitor its vegetation anyway. I am deeply suspicious about requirements for "analysis of benefit" prior to permission being given for the release and use of GMOs. Just who sets the criteria of benefit and performs the analysis? Central planning can be debilitating when prudence and responsible care are the objectives.

### Genetically Engineered Foods

In the UK, the assessment of the safety of foods which are themselves GMOs or which are produced in processes involving GMOs, is a part of the remit of the Advisory Committee on Novel Foods and Processes. A decision-tree scheme has been devised to pose a series of questions which indicate the type of information required for individual submissions to the Committee. Food components of food derived from GMOs must be as safe as, or safer than, their traditional counterparts. The Committee is refreshingly open, as are nearly all scientific committees, with regard to the advice it gives to Government and the reasoning behind that advice. Related committees include the Food Advisory Committee, the Committee on Toxicity and the Advisory Committees on Genetic Modifications and Releases to the Environment. Their deliberations range from food labelling to the nature of research. They function well and reinforce the confidence of the consumer that standards and proper control are in place in a democratic environment. I very much welcome the very recent

visible report of the Ethical Committee on Genetic Modification of Food, chaired by the Reverend Dr John Polkinhorne (1993).

### Public Attitudes

Voters comprise heterogeneous groups who determine the political, industrial and economic climate of democratic countries. Their taxes and use of private companies support R&D programmes in the public sector. They are also consumers who should be free to exercise choice. Scientists should be providing them with factual basis for reaching informed decisions.

In contrast to healthcare, applications of modern biotechnology to food and the environment are greatly influenced by the level of education, perceived social and ethical issues, as well as reaction, frequent irrational, responses towards non-medical sciences.

There are also objections at a secondary level to the role of multinational companies carrying out genetic engineering and failing to take adequately into account the impact of their activities on the less-developed world or playing one economy off against another. Although not necessarily associated with religious organisations, there is also the oft-cited "unnatural" or "ungodliness" aspect of science replacing natural functions, creating chimaeric organisations, or fiddling with life for profit.

Ignorance of science and technology, as much as ignorance of business, adds to fear, anxiety and reluctance to fund research and development projects. Pressure groups of all kinds are formed. It seems that the public receive most of their limited understanding of science through the arts-influenced media, especially television, where all too often artistic license embellishes scientific observation with imaginative doom-laden claptrap. This tends to cause the scientific and advanced industrial communities a measure of respectation. Healthy scientific scepticism, questioning, sharp debate, experimentation and wide-ranging open-minded interpretations and conclusions are the stuff of science. So is presentation. There can be no room for ignorance nor ineptitude. Scientists, like the public, cover a spectrum of views and attitudes and are difficult to organise except into cliques. The public must realise that it is entirely technology-dependent. Scientists in turn accept justifiable control as much as the financial backing. I am worried that the Foucault pendulum is swinging towards harsher controls which will not be sustained in the longer term even though it is unfashionable to argue against any moves restraining science. Sometimes, pressure groups have tied political interests or social engineering at heart. Nonetheless, a balance-point must be reached, taking into account illogical fears, damage to the environment, healthcare, and the need for science. The need for genetic engineering is irrefutable. It is how we do it that we must get right.

At this juncture, there is a view that only by being aware of obvious benefits to the consumer (e.g. increased safety because of reduced natural toxins, lower costs, post- and disease-free produce, better and more consistent quality etc.) or to the environment (e.g. reduced pesticide inputs, remediation etc.) will there be general acceptance to genetic engineering. Fewer problems are experienced with plant-plant than with plant-microbe or transgenetics; plant-animal and animal-animal combinations, most notably where "man" genes are concerned can provoke virulent public and pressure group objections. The greatest level of acceptance will be for transgenic plants used for non-food purposes. Meanwhile, there will be a plethora of legislative barriers.

## CURRENT APPLICATIONS OF BIOENGINEERING

Aided by the rapid uptake of biotechnology in the higher education sector, and relatively crude but accurate assessments of its potential, decision-makers in government and private industry, genetic engineering of crop plants is a world-wide phenomenon. Selective herbicide resistance to a better crop management; introduction of plant-derived insecticidal genes (e.g. protease inhibitors); introduction of characteristics associated with resistance to pests, diseases, abiotic and biotic stresses; enhanced quality (e.g. amino acid composition); production of engineered oils, proteins, carbohydrates, enzymes etc. are examples of projects currently underway using several crops or related species.

The diagnosis and quantification of disease organisms are increasingly reliant on biotechnology, as are studies on the relationships between different races, pathotypes and virulence groups. To investigate the mode of action and effectiveness of control agents requires the new technologies.

Plant breeding is one of the leading beneficiaries of genetic engineering. All parts of normal breeding schedules are being revolutionised: form describing the genetic architecture of parental material, overcoming natural breeding barriers and selections, to propagation, prediction of performance and identifying more accurately the added-value properties of the progeny. Speed is of the essence, so is protection of intellectual property.

New plant varieties arising from traditional breeding methods are protected in many countries by plant variety (or breeder's) rights (PVR) without recourse to patent law. Patents are now being granted for the protection of recombinant methods for the production of transgenic plants and their resultant products. Attempts are being made to harmonise patent law and practice internationally. Ethical concerns are expressed about patenting life-forms and claiming ownership. There is freedom to research under both patent and PVR law, but freedom to commercialise is complex, and therefore plant breeders using modern technologies seek protection of both types of law.

Patent protection is unlikely to affect access to existing germplasm and traditional varieties. Genetic resources and diversity are internationally seen as the common heritage of humankind. Biotechnology adds to genetic diversity. Counterarguments centre on the farmer's privilege to save seed and produce subsequent crops without royalty payments to recoup R&D costs, abuse of monopoly provisions, "ordre public" and the public interest, and also the nature of more discovery. Unfortunately for all concerned, a patent on an invention does not guarantee a reward for the inventor; simply put, it gives an opportunity for the inventor or patent proprietor to profit from the invention where there is a profitable market for it. Secrecy in some cases is the best commercial protectant in the short term.

For transgenic plants to achieve a small portion of their potential, basic studies will need to expand on the factors (e.g. promoters etc.) regulating the expression of introduced genes in different organs and tissues at various phases of growth and differentiation. Industry would be assisted by studies on the biosynthesis and degradation of complex natural polymers such as lignin, cutin, suberin and cellulose, manifestations of cell differentiation, and the cellular components of industrial relevance. Single gene studies will give way to polygenic linked constructions.

Bioremediation could have a highly publicity profile for genetic engineering to assist in measures to reverse environmental degradation. Most recent research is initially concerned with the construction of organisms that can degrade, on command, oil and organic xenobiotics such as polychlorinated biphenyls. Future research is likely to include the construction of transgenic plants that can withstand abiotic "stresses" such as high salinity, heavy metal and radionuclide contamination, sewage and factory effluents.

Dinomania apart, the retrieval of nucleic acids from fossils and preserved tissues is of merit for evolutionary studies of all kinds, and for research on adaptation to climates, predation and disease.

#### Protection of Crop Plants - The Virus Example

Crop protection deserves conferences of its own. A daunting matrix of pests and diseases, vector systems, host types, mechanisms of infection and host adaptation, control measures, economics, impact assessments, and variability confronts every reviewer. Biotechnology finds its greatest sophistication in crop protection in the area of virology, for understandable reasons connected with the viral modus operandi.

Most plant species are naturally resistant to the majority of the 675 or so plant viruses currently identified. All crops, however, are prone to significant yield and quality losses caused by one or more viruses. Plant viral genomes are plastic and resistance-breaking virus strains are rapidly introduced in monocultural agricultural systems with intense selection pressures. This, in turn, causes difficulties for plant breeders attempting to introduce dominant and durable resistance genes: there may be true immunity, sublethal infection or symptomless tolerance to infection. No resistance genes per se have been characterised to date. For many years, there have been observations that infection with mild, symptomless or attenuated strains of viruses could "cross-protect" a range of field crops against closely related, but severely pathogenic virus strains. Virus-resistant crops have been created in many countries by genetically engineering them to express part of a viral genomic or virus-associated sequence. Cognate effects on virus control measures are aimed at the virus vectors (nematodes, insects and fungi) and the molecular features determining virus transmission and replication.

Transgenic plants expressing viral-derived sequences have been discussed as sites for hyper-evolution of pathogenic viruses through recombination events. There is no supporting evidence for this. Any long-term genetic or epidemiological effects would seem remote.

An exciting concept for future work is the protection against fungal, bacterial and viral diseases by expressing appropriate antibodies in transgenic plants. Eventually, we would like to be in a position to solve resistance gene construction and action.

#### CONCLUSION

Genetic engineering is here to stay. No doubt the introduction of the technology was foreseen by some to require road traffic legislation as a way to promote its development. The technology is clever and takes science and industry

into a new phase of opportunity. Our shared responsibility is to get it right.

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