



Review Article

Transgenic Pest Resistance

Beena M. R. *

Leibniz Institute DSMZ-Deutsche Sammlung von Mikroorganismen und Zellkulturen, Germany

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Abstract: After the first report of vegetative insecticidal proteins (VIP) from *Bacillus thuringiensis* there was an influx of scientific research to introduce insect resistance in crop plants using *Bt* or *Cry* genes. In this article most of the work happened on insect resistance including other insecticidal genes other than *Cry* is reviewed. However, transgenic plants including RNAi technology is not included.

Introduction

Legumes are used for their chemicals, timber, as cooking fuel, forage crops, pasture crops, cover crops, green manures, for feed and, most importantly food for vegetarian people in the world. They are grown in the Semi Arid Tropics particularly as rain fed crops and hence are subject to biotic and abiotic stress factors.

Biotic stress is mainly due to insect herbivores, bacteria, viruses and fungi. In the tropics the key factor limiting the yield of pulse crop is the insect pest since decades (Singh and Emden, 1979). The most economically important pests are those, which attack the plant at flowering and fruiting stages. Nevertheless, the pests that attack during the young stages of the plants and during storage are also equally important. Most plants have inherent defense system against most of them that has evolved over a period of time. But this is hardly of any use as the pathogens and the insect herbivores co-evolve for their survival. Plant breeders have been trying hard to develop the crops with higher level of resistance against the pathogens and insect pests since decades through hybridization, but

without significant success. Carnivore natural enemies that suppress herbivore populations can protect plants ecologically but at the same time thwart opportunities for selection of herbivore resistance. In other words the effectiveness of defense external to the plant in suppressing herbivores can prevent the evolution of internal resistance to herbivory (Strong and Larsson, 1992).

The pests on legumes are leaf hoppers (*Empoasca*), distributed widely in tropics and sub tropics that attack the legume plants at the seedling stage; aphids (*Aphis*) that deplete the assimilates by the removal of sap, and also act as vectors for viruses; bean flies (*Melanagromyza*) a widely distributed seedling pest; a large number of beetles - foliage beetle (*Ootheca*), a vector for yellow mosaic virus, striped foliage beetle (*Luperodes*), blister beetles (*Mylabris* and *Coryna*) that damage flowers and foliage, bean beetle (*Epilachna*, *Lagria* and *Chrysotragus*); weevils - striped bean weevil (*Alcidodes*), pod weevil (*Piezotrachelus* and *Apion*); thrips (*Megalurothrips*) and pod sucking bugs (*Nezara*, *Anoplocnemis*, *Riptorus* and *Acanthomia*) have been identified world wide causing damage mainly to the foliage and flowers resulting in stunted growth, late

*Corresponding author

E-mail: beena.ravindran@julius-kuehn.de



flowering and fruiting (Singh and Emden, 1979). However, the most widely distributed and most important foliage, flower and pod borers are those belonging to the order *Lepidoptera*. These have attracted a great deal of attention in recent years because of their wide host range, the severity of damage they cause to crops and their resistance to commonly used chemical insecticides. *Helicoverpa armigera*, *Maruca testulalis*, *Cydia tychora*, *Matsumuraeses phaseoli*, *Etiella zinckenella*, and *Spodoptera littoralis* (Singh and Emden, 1979) are few of them of which *Helicoverpa* and *Spodoptera* spp. are the most notorious ones.

With the advent of genetic transformation techniques, it has become possible to insert genes that confer resistance to insects into the host plant genome. However, protocols for regeneration and transformation are not routinely available for grain legumes. Biotechnology in association with the conventional crop improvement programs can make a drastic difference in the food production (Ortiz, 1998). The major difference between conventional breeding and biotechnology lies in the speed, precision, reliability and scope rather than in goals or process (Sharma and Ortiz, 2000). Among the biological pesticides, bacteria such as *Bacillus* species have been the most successful for use in pest control on a commercial scale. Insecticidal and anti-feedant genes like *Bt* genes, trypsin, protease and amylase inhibitors (enzyme inhibitors), enzymes, lectins, ribosome inactivating proteins, secondary plant metabolites, vegetative insecticidal proteins could be used singly or in combination for efficient pest control. Synthesised scorpion insect toxin AaHIT gene was also used for resistance to cottonbollworm (Wu et al., 2008) but in this review the subject

would be restricted to the above mentioned genes.

It is not surprising that insect resistant crops were the first to hit the market among the transgenic plants, because of the booming price of insecticides and pesticides and the intensity of crop loss due to insects and concern over the pollution caused to the environment. Among the insect resistant transgenics, the plants/crops harboring Crygenes are in fairly advance stage in Plant Biotechnology.

Genes from *Bacillus thuringiensis*

Bacillus thuringiensis is a gram-positive spore forming soil bacterium, which forms parasporal crystals during sporulation with unique insecticidal property. The parasporal crystals consist of one or more or truncated δ-endotoxin (pro-toxin sub units) or Insecticidal Crystal Proteins (ICP) of approximately 138 kDa and the active toxin that shows effective insecticidal properties is about 25 to 135 kDa. Ingestion of the insecticidal crystal protein by the target insect is essential for the toxin to be effective. Inside the gut of the insect, the pro-toxin breaks down to active form in presence of the high pH and gut enzymes, that binds to the gut membrane where it creates ion channels or 'pores' leading to leakage of mid-gut content, paralysis and death. In some insects, it leads to a loss of appetite and stops feeding ultimately leading to death. The high specificity of the protein is due to high-affinity binding sites in the brush border membrane of target insect mid-gut (Hofmann et al., 1998) that is lacking in human beings or ruminates.

There are also vegetative insecticidal proteins (VIP) isolated from the *B. thuringiensis* and do not belong to δ-endotoxin family. The protein is expressed at the vegetative phase of the growth starting at mid-log phase unlike Cry



proteins. A list of Vip proteins can be retrieved from the web site <http://bttnomenclature.info/> (Crickmore *et al.*, 2016). A number of VIPs proteins have been identified and characterized against insects. As Cry and Cyt (cytolytic) proteins, VIPs are also very specific to the order of insects; Vip3A was found to be effective against Lepidopteran insects such as *Spodoptera exigua*, *Spodoptera litura* and *Helicoverpa armigera* (Estruch *et al.*, 1996; Chen *et al.*, 2002); Vip3V has been cloned and characterized by Doss *et al.* (2002) was found to work against Lepidopteran insects and it did not show any adverse effect on silk worms and mosquitoes. Vip1 and 2 have activity against insects belonging to order Coleoptera (Pitkin *et al.*, 2005). Yu *et al.* (1997) explained that the basic mode of action of Vip proteins lies in the lysis of mid gut epithelium of the susceptible insect larvae.

There are 341 *cry* genes so far identified of which, only a few have so far found application in technological plant breeding. The Cry and Cyt proteins are classified according to protein structural homologies. Revision of nomenclature of proteins was done by Crickmore *et al.* (1998) and the list of new names of all the insecticidal *cry* genes can be found at the website <http://bttnomenclature.info/> (Crickmore *et al.*, 2016). These δ-endotoxins are designated as *Cry I*, *Cry II*, *Cry III*, *Cry IV*, which are fatal to Lepidopteran, Dipteran and Lepidopteran, Coleopteran and Dipteran insects respectively and *cyt* genes and Cyt proteins, which are significantly different in structure and function from the *cry* genes. For more information on the *Cry* and *Cyt* genes with respect to the types and their insecticidal properties, a reference can be made to Hofte and Whiteley (1989), Schnepf *et al.* (1998). The host range of some of Cry and Cyt proteins are given in **Table 1**. Other than

those mentioned in the table, fusion genes were also developed for better expression, toxicity and host range in plants based on these genes; for example: *Cry1B-Cry1Ab* (Bohorova *et al.*, 2001), *Cry I E-C* (Singh *et al.*, 2004), *Cry2A** (Chen *et al.*, 2005).

Starting with the first successful demonstration of the power of Cry proteins in transgenic plants inducing insect resistance by Vaeck *et al.* (1987) and Barton *et al.* (1987) in tobacco trailed by Perlak *et al.* (1991), Carozzi *et al.* (1992), Koziel *et al.* (1993) in tobacco and maize, there are umpteen number of successful examples of transgenic plants for insect resistance in many crop plant species across monocots and dicots. However, insect resistant legume transgenic plants carrying Cry proteins are not too many in number. The usage of *Cry* genes in legumes started in 1996 with the introduction of *Cry IC* in alfalfa for *Spodoptera* resistance by Strizhov and group followed by Ozias Akins and co workers (Singsit *et al.*, 1997), who have developed peanut transgenic plants carrying *Cry I ac* for resistance against lesser corn stalk borer. Lesser corn stalk borer enhances the invasion of aflatoxin producing fungal species, *Aspergillus niger* on peanut seed. This has resulted in complete to partial protection in peanut. Subsequently, S. K. Sen's group demonstrated resistance to pod borer, *Heliothis* in transgenic chickpea carrying *Cry I ac* (Kar *et al.*, 1997). However, this pace was not sustained in legumes.

Transgenic plants expressing *Cry* genes produce the protein continuously; through all the phases of plant growth, though to varying extents, giving them protection and the problem of narrow specificity can be overcome by expressing them properly singly or in combination other insect resistance conferring proteins having with different specificities,



gene stacking/ pyramiding (de Maagd *et al.*, 1999; Estruch *et al.*, 1997; Broderick *et al.*, 2000; Sharma *et al.*, 2000; Datta *et al.*, 2002; Zhao *et al.*, 2003; Moar, 2003; Hilder, 2003; Beena *et al.*, 2008; Arvinth *et al.*, 2010). Stacking also helps in delaying development of resistance towards the toxins (Zhao *et al.*, 2003). Abdeen *et al.* (2005) suggested that using two dissimilar genes rendered stronger resistance to transgenic plants. Cao *et al.*, (2002a) demonstrated that two *Cry* genes (*Cry1Ac* and *Cry 1C*) pyramided in transgenic broccoli plants imparted resistance against diamond back moth that is resistant to *Cry1A* and *Cry1C* individually in transgenic plants. Use of more than two genes for resistance against yellow stem borer and sap sucking insects has been established by Ramesh *et al.*, (2004) by introducing *Cry IAb* + *CryIAc* + *GNA* into indica rice using super binary vectors of *Agrobacterium*. Transgenic rice plants exhibited significant levels of protection against the target insects. Potentiality of using other genes along with *Cry* genes has been demonstrated (Liang *et al.*, 1994; Fan *et al.*, 1999; Down *et al.*, 2001; Lawrence and Novak, 2001; Gujar *et al.*, 2004; Zhang *et al.*, 2004; Loc *et al.*, 2002; Mehlo *et al.*, 2005; Beena *et al.*, 2008). While Abdeen *et al.*, (2005) could not achieve 100% kill of the insect pest when transformed tomato plants with two different classes of Proteinase inhibitors (Potato protease inhibitor (*PI-II*) and Carboxypeptidase inhibitor (*CPI*)), Wang *et al.* (2005a) produced insect resistant *Brassica napus* by transforming it with chitinase (*Chi*) and scorpion Insect Toxin gene (*Bmk IT*), and Maqbool *et al.* (2001) combined *Cry1Ac* and *Cry2A* with *GNA* to develop resistance against three pests in rice viz., rice leaf folder, brown hopper and yellow stem borer. Datta *et al.* (2002) combined three entirely different genes conferring resistance to bacterial blight, yellow stem borer and sheath blight (*Xa2I* + *Bt*

+ chitinase), and produced stable transgenic rice lines resistant to all the three. This shows that stacking genes in a transgenic crop plant is technologically viable and sound for achieving stronger effect in one trait or satisfactory effect in multiple traits depending on the genes deployed. There are also reports on combining QTLs with *Cry1Ac* for better and sustainable lepidopteran insect resistance (Walker *et al.*, 2002). Haidi *et al.* (1996), Chen *et al.* (1998) and Campbell *et al.* (2000) have recorded multiple gene integrations (up to 13) with different plasmid vectors using biolistics method of transformation

When transgenic plants and other strategies were compared for the potential ecological and human health consequences including effects on non-target organisms, food safety, and the development of resistant insect populations in insect management, the risks were definitely lower and beneficial than any other strategies deployed (Betz *et al.*, 2000; Shelton *et al.*, 2002, Romeis *et al.*, 2006) and can be successfully utilized in IPM. There are other reports that clearly demonstrate safety in using *Cry* and other genes for insect resistance in crop plants (Girard *et al.*, 1998; Wraight *et al.*, 2000; Gatehouse *et al.*, 2002; Hellmich *et al.*, 2001; Sears *et al.*, 2001; Tabashnik *et al.*, 2003; Dutton *et al.*, 2003; Cowgill and Atkinson, 2003; Hanley *et al.*, 2003; Wei-Xiang *et al.*, 2004; Schuler *et al.*, 2004; Bashir *et al.*, 2004; Huang *et al.*, 2005; Turlings *et al.*, 2005). Carriere *et al.* (2003) have shown in a ten-year study on *Bt* cotton in Arizona region that these transgenic plants helped long-term suppression of the pests and reduced the use of insecticide sprays considerably. Apart from this, there was no resistance developed by the insect against the introduced transgene encoded protein. There are only two insect species, which developed



resistance towards *Bt* insecticide sprays under commercial situations. As Shelton (2002) suggested, it is necessary to have optimal and consistent expression of the Cry and other proteins to achieve desirable effects in the transgenic plants, and delay the development of resistance in the insect and it is also likely that there are fewer genes in insects for resistance than estimated and recessive genes may be involved which might need more generations to develop resistance. Liu *et al.* (2001) observed that there is no cross resistance towards other insecticidal genes in diamond black moth that had evolved resistance against *Cry 1C*.

Since the list of transgenic plants carrying the different *Cry* genes singly or in combination with other genes is exhaustive, only some significant examples in other crops along with legume transgenic plants have been listed in Table 2.

Enzymes and Enzyme Inhibitors Conferring Insect resistance

The enzymes and enzyme inhibitoros that confer insect resistance would include protease inhibitors, chitinase, cholesterol oxidase, polyphenol oxidase, ehancin etc.

Plant proteinase inhibitors are mostly polypeptides or proteins, which occur in a variety of plants that have been evolved as a defense mechanism against herbivores. They inhibit the synthesis of proteins thereby affecting proteolysis of protein into amino acids, which in turn affects the growth of the insect larvae. At the same time, they enhance pernicious hyper production of proteinase activity that again leads to scarcity of amino acid supply for insect growth and development (Gujar *et al.*, 2000). According to Koiwa *et al.* (1997), protease inhibitors are an important element of the plant defense

response to insect predation and may also act to restrict infection by some nematodes. They are generally classified according to the class of proteases they inhibit. Four types of proteases have been identified as serine, cysteine, aspartic, or metallo-proteases (Koiwa *et al.*, 1997; Lawrence and Koundal, 2002). Some plant serine protease inhibitors are bifunctional possessing trypsin and α -amylase inhibitors activity. These proteinase inhibitors are insecticidal at high concentrations. Phytocystatins are cysteine proteinase inhibitors of plant origin that occur in a variety of higher plants including rice (Abe *et al.*, 1987; Kondo *et al.*, 1989, 1990) and corn (Abe *et al.*, 1992). Production of these inhibitors is highly regulated by a signal transduction pathway that is initiated by predation and transduced as a wound response. For more detailed information on enzyme inhibitors, please refer to Ryan (1990), Ussuf *et al.* (2001), De Leo *et al.* (2002), Oleveira *et al.* (2003), Valueva and Mosolov (2004).

Chitin is a linear homopolymer of β (1-4) linked 2-deoxy-2-acetamido-D-glucopyranosyl residues and is a major component of the exoskeleton and alimentary canal of insects. Together with proteinases in the moulting fluid, endo-splitting chitinases and exo-splitting β -N-acetylglucosaminidases break down the unsclerotized layers of the old cuticle prior to shedding of the sclerotized portion or exuvium. Transcripts of moulting fluid chitinase are seen only in epidermal and gut tissue of 5th instar larvae between 5 and 7 days after moulting and just prior to pupal transformation (Ding *et al.*, 1998a). This tight developmental regulation suggests that this chitinase could be detrimental to insect growth if presented in the diet at an inappropriate time. Transgenic plants expressing chitinase show enhanced protection against insects. Van



der Westhuizen *et al.* (1998) analyzed the differential induction of apoplastic peroxidase and chitinase activities in susceptible and resistant wheat cultivars by Russian wheat aphid infestation and suggested that peroxidase and chitinase have crucial roles in insect resistance. There are also reports showing that chitin synthesis inhibitors (CSI) applied externally also gives 100% mortality in pests (Cohen, 1993; Wilson and Cryan, 1997; Kostyukovsky and Trostanetsky, 2006).

Cholesterol oxidase is a monomeric bacterial flavoenzyme that catalyzes the oxidation and isomerization of cholesterol to ketosteroids and hydrogen peroxide and is widely used in clinical assays for determining levels of total serum cholesterol. This enzyme is produced by phylogenetically diverse group of microorganisms and usually occurs as a secreted protein. Cholesterol oxidases have been isolated from several sources including members of the genera *Streptomyces*, *Pseudomonas*, *Schizophyllum* and *Rhodococcus*. It has potent insecticidal activity against cotton boll weevil, *Anthonomus grandis* (Corbin *et al.* 2001). Upon ingestion, this enzyme causes developmental arrest and death of the larvae (Purcell *et al.*, 1993) and also marked decrease in the adult female fecundity (Greenplate *et al.*, 1995). It shows moderate mortality when used against some of the lepidopteran insects (Greenplate *et al.*, 1997). It was Purcell *et al.* (1993, 1994), who found that the filtrate of *Streptomyces* cultures killed the boll weevil larvae and this has been attributed to the oxidation of cholesterol in the mid gut epithelial membrane resulting in physical and functional disruption of the membrane; and that the active component that caused the mortality was cholesterol oxidase. Shen *et al.* (1997) reported that mild toxicity of cholesterol oxidase on Lepidopteran insects was due to

general susceptibility to cholesterol oxidase. Santos *et al.* (2002) showed that the emergence and viability of cotton boll weevil larvae (*Anthonomus grandis*) was acutely affected when fed with artificial diet containing cholesterol oxidase.

Polyphenol oxidase oxidizes phenols by catalyzing the oxidation of the *o*-diphenols to *o*-diquinones, as well as hydroxylation of monophenols. This activity of the enzyme increases in response to different types of stress, both abiotic and biotic, mostly due to physiological injury (Rivero *et al.*, 2001). Felton *et al.* (1992) reported that polyphenol oxidase significantly reduced protein quality in *Spodoptera exigua* larvae, which influenced the larval growth rate. Polyphenol oxidase cDNAs have been cloned from sugarcane, apple, grape, potato etc. and upon characterization of this cDNA, Constabel *et al.* (1995) proposed the role of defense for these genes. Upon chewing and feeding, the interaction of polyphenol oxidase with phenolic substrates generates *o*-quinones and these highly reactive compounds covalently modify free amino and sulphydryl groups in dietary proteins within the insect. Constabel *et al.* (2000) have also cloned a polyphenol oxidase gene from poplar expressed in response to wounding and herbivory.

Enhancin gene from baculoviruses encodes baculovirus metalloproteases that can degrade mucin in the mid guts of insects (Lepore *et al.*, 1996; Wang and Granados, 1997). Disruption of peritrophic membrane increases viral infection and leads to faster larval mortality (Wang *et al.*, 1994). Hayakawa *et al.* (2000) and Cao *et al.* (2002b) have shown that it is effective against larvae of *Spodoptera exigua* and *Trichoplusia ni* in transgenic tobacco.

Protease and amylase inhibitors, chitinases and other enzymes deployed in raising transgenic plants including legumes, that resisted herbivores are listed below in Table 3 including the legume transgenic plants.



Other secondary metabolites with insecticidal activity

Oleoresins, lectins, phytoalexins (antimicrobial secondary metabolites) etc. are other secondary metabolites induced by wounding are shown to have increased effect against herbivore insects and are used in transgenic plants for developing insect resistance.

Oleoresin is a complex mixture of terpenoids, consisting of a turpentine (monoterpene and sesquiterpene) and rosin (diterpene) fraction. The turpentine fraction contains a range of insect and microbial toxins such as limonene and 3-carene and other biologically active agents that often act synergistically to discourage insect predation. Turpentine also acts as the solvent for transporting the higher molecular weight diterpenoid resin acids (rosin fraction) to the site of injury. Upon exposure to the atmosphere, the volatile turpentine evaporates leaving a semi crystalline mass of resin acids that oxidatively polymerize to form a hardened barrier that seals the wound, often trapping insect invaders and microbial pathogens in the matrix (Phillips and Croteau, 1999).

Host plants play a key role in the production and use of sex pheromones by herbivorous insects through larval or adult sequestration of chemically active compounds and pheromone precursors. Some of the plant semiochemicals/ volatiles have inhibitory or repellent effects that interrupt insect responses to pheromones and attract predators and parasitoids to the attacking species after herbivory injury (Reddy and Guerrero, 2004). The synergism between insect pheromones and plant odors can increase attraction of natural enemies, offering new strategies for IPM. Altering the plant volatile compounds will make it difficult for the phytophagous insects to locate their host plants for oviposition. In the same way nectarines in the nectar of the angiosperms are also found to have defense role against non-pollinating

insects and other air borne microbes (Carter and Thornburg, 2004).

Lectins act almost like *Cry* genes where they bind to the gut surfaces of the insect larvae causing lesions and death of the larvae. They show detrimental effects on homopteran insects and aphids (Wang *et al.*, 2005b). Lectins belong to a broad group of bioactive peptides called defensins that elicit a variety of responses including plant defence against a range of factors including environmental stress, predation by insects, and infestations by bacteria, fungi, and nematodes. Many agricultural plants lack lectin defenses, however, making them susceptible to yield losses from pest infestations. The insertion of specific lectin genes into crop plants gives protection against many pestilent pests attacking them. Most commonly used lectins in transgenic studies are GNA (*Galanthus nivalis* agglutinin) and Con A (Jack bean: *Canavalia ensiformis* lectin). Lectins are favored over other pesticidal plant compounds because of their low toxicity to humans and domesticated animals at levels that are effective against insect larvae, and GNA does not appear to damage other organisms. Chrispeels and Raikhel (1991) made a comprehensive review on lectins and their role in plant defense.

Avidin is a water-soluble tetrameric glycoprotein isolated originally from chicken egg white. It strongly binds to vitamin biotin, which is an essential compound for insects, which they acquire from dietary sources. Biotin is a co-factor of major carboxylases involved in gluconeogenesis, lipogenesis, fatty acid and amino acid catabolism. It causes mortality of a wide range of pests if mixed with artificial diets and fed (Morgan *et al.*, 1993; Markwick *et al.*, 2001 and Burgess *et al.*, 2002). Transgenic plants, developed for resistance against various insects using secondary metabolites are given in Table 4.



Each insecticidal gene is unique. It is important to assess the activity of the protein encoded by a gene against a particular insect and in a particular crop. Different genes for the proteins with insecticidal activity that have been characterized from different species are summarized in the Table 5.

Conclusions and Future

The efficacy of insect resistance engineered by deployment of the various insect resisting gene products, the proteins has been well documented and the level of cultivation of insect resistant transgenic crops cultivated on 16. 2 million hectares with Bt crops including soybean and 10. 1 million hectares on the stacked herbicide tolerance and insect resistance bears testimony to the concept of insect resistance through genetic engineering (James, 2005). The technology should be used judiciously in tackling insect problems. The most important insect(s) attacking a given legume crop need to be identified and the genes for controlling them need to be ascertained. There are about 341 *Cry* genes reported and it is imperative to study the relative efficiencies of the individual *Cry* proteins against target insects in legumes. A thorough study is needed in this direction for identifying effective *Cry* proteins for deploying the corresponding genes in legume transgenic plants for insect resistance. Compared to other crops, the legumes, barring soybean which has received attention of the various laboratories and Multinational Companies on a very large scale because of the importance world wide, have started receiving attention of late and the desired transformaiotn protocols for the recalcitrant legume are standardized. With this scenario, more insect resistant legume transgenic crops are expexted to be developed in the near furture.

Apart from the *Cry* genes, the bacterium produces other proteins, which are essentially

insecticidal in nature. Plants also try to defend themselves against the invading insects by producing different molecules and proteins that have a direct bearing on the insects feeding on them. It could be through various morphological, biochemical and or molecular mechanisms. The biochemical mechanisms against insects are of wide ranges which are dynamic. More and more such anti-insect proteins and molecules need to be identified for their utilization in genetic engineering. The larvae of insect pests have essentially large guts as they spend their whole life, till they pupate, feeding on the host plants. Most of the mechanisms described in this article targets the mid gut of the larvae and stops them from feeding. As the pro-toxins will be converted to active toxin only in the mid guts of the target larvae, even as the plant dies and degenerates, it will not be converted to active toxin; hence do not leave any toxin traces in the ecosystem to be biomagnified at a higher level. Occurrence of resistance alleles in the target insect populations should be assessed so that strategies such as pyramiding of two or more similar or dissimilar genes can be deployed to overcome the resistance. Over expression of a single gene also could be considered for which, lethal concentration of the protein should be measured.

Technology has changed and newer methods of engineering resistance have been discovered. Use of double stranded RNA in the forms of artificial micro RNA, small RNA, and tasi-RNA is taking over the normal gene stacking mechanisms against insects which are acting as pests and also as vectors for many devastating diseases, which are not elaborated in this review. Among the still existing intense debates whether or not transgenic plants itself and toxins and antibiotic resistance genes they carry are harmful in a long run to human kind as well as to other harmless useful insects, *Bt* still remains a very good candidate for crop improvement against insect pests.



Table 1. Host range of some Cry and Cyt proteins: (Please refer to Hofte and Whiteley (1989), Schnepf *et al.* (1998)

Cry protein	Acc. No.	Insect
CrylA(a)	M11250	Lepidoptera
CrylA(b)	M13898	Lepidoptera
CrylA©	M11068	Lepidoptera
CrylB(a)	X06711	Lepidoptera and Coleoptera
Cry1B©	AAQ52387	Lepidoptera
CrylC(a)	X07518	Lepidoptera
CrylD(a)	X54160	Lepidoptera
CrylE(a)	X53985	Lepidoptera
CrylF(b)	173895	Lepidoptera
Cry1G(a)	Y09326	Lepidoptera
Cry1I(a)	X62821	Lepidoptera
Cry1I(b)	U07642	Lepidoptera
Cry1I(d)	AF047579	Lepidoptera
Cry1K(a)	U35780	Lepidoptera
CryIIA(a)	M23723	Lepidoptera and Diptera
CryIIA(b)	M23724	Lepidoptera
CryIIA©	AAQ52385	Lepidoptera
CryIIIA(a)	Y00420	Coleoptera
CryIIIB(a)	X17123	Coleoptera
CryIIIC(a)	X59797	Coleoptera
CryIVA(a)	Y00423	Diptera
CryIVB(a)	X07423	Diptera
CryIVC	--	Diptera
CryIVD	--	Diptera
CryVA©	134543	Coleoptera and Lepidoptera
CryIXA(a)	X58120	Lepidoptera
CryIXB(a)	X75019	--
CryXA(a)	M12662	Diptera
CryXIA(a)	M22860	Diptera
CryXIB(a)	X86902	Diptera
CryXIB(b)	AF017416	Diptera
CryXVA(a)	M76442	Lepidoptera
Cry20A(a)	U82513	Diptera
Cry26A(a)	AF122897	--
Cry38A(a)	AAK64559	Siphonaptera and Coleoptera
CytIA(a)	X03182	Diptera
CytIA(b)	X98793	Diptera
CytIC(a)	AL731825	Diptera
CytIIA(a)	Z14147	Diptera
CytIIB(a)	U52043	Diptera
CytIIC(a)	AAK50455	

**Table 2.** A curtailed list of transgenic plants hosting *Cry* genes; legume transgenes emphasized in bold letters

Bt gene	Plant	Insect	Reference
<i>Cry2AX1</i>	rice	Rice leaffolder (<i>Cnaphalocrosis medinalis</i>)	Manikandan <i>et al.</i> , 2016
<i>Cry1Ac</i>	Sugarcane	Sugarcane borer (<i>Diatraea saccharalis</i>)	Gao <i>et al.</i> , 2016
<i>Cry1la12</i>	Cotton	Fall armyworm (<i>Spodoptera frugiperda</i>) Cotton boll weevil (<i>Anthonomus grandis</i>)	de Oliveira <i>et al.</i> , 2016
<i>Cry1Ac + Cry2A</i>	Cotton	American boll worm <i>Heliothis</i> sp	Puspito <i>et al.</i> , 2015
<i>Cry1Ac and Cry2A</i>	Cotton	American boll worm <i>Heliothis</i> sp	Muzafar <i>et al.</i> , 2015
<i>Cry1Ac</i>	Tomato	<i>Helicoverpa armigera</i>	Koul <i>et al.</i> , 2014; 2015
<i>Cry1C</i>	Maize	<i>Ostrinia furnacalis</i>	Du <i>et al.</i> , 2014
<i>Cry1Aa3</i>	Sugarcane	Sugarcane borer	Kalunke <i>et al.</i> , 2009
<i>Cry 1 E-C</i>	Peanut	<i>Spodoptera litura</i>	Beena <i>et al.</i>, 2008
<i>Cry1Ac</i>	Sugarcane	Sugarcane borer	Weng <i>et al.</i> , 2006
<i>Cry 1 E-C</i>	Pigeon pea	<i>Spodoptera litura</i>	Surekha and Beena <i>et al.</i>, 2005
<i>Cry3Bb1</i>	Maize	Cut root-worm (<i>Diabrotica</i> spp.)	Vaughn <i>et al.</i> , 2005
<i>Cry1Ac</i>	Sorghum	Spotted stem borer (<i>Chilo partellus</i>)	Girijashankar <i>et al.</i> , 2005
<i>Cry2A*</i>	Indica rice	Lepidopteran pests	Chen <i>et al.</i> , 2005
<i>Cry 1 Ab protoxin</i>	Soybean	Velvet bean caterpillar	Dufourmantel <i>et al.</i>, 2005
<i>Cry 1 A(c)</i>	Chick pea	<i>Helicoverpa armigera</i>	Sanyal <i>et al.</i>, 2005
<i>Cry 1 A(b)</i>	Pusa Broccoli KTS-1	Diamond back moth (<i>Plutella xylostella</i>)	Viswakarma <i>et al.</i> , 2004
<i>Cry1 Aa10</i>	Oil seed rape	Diamond back moth (<i>Plutella xylostella</i>)	Huo <i>et al.</i> , 2003
<i>Cry 2Ab</i>	Cotton	Pink boll worm	Tabashnik <i>et al.</i> , 2002
<i>Cry 1 A(b)</i>	Cauliflower	Diamond back moth (<i>Plutella xylostella</i>)	Chakrabarty <i>et al.</i> , 2002
<i>Cry 1 A(b)</i>	Cabbage	Diamond back moth (<i>Plutella xylostella</i>)	Bhattacharya <i>et al.</i> , 2002
<i>Cry9Aa</i>	Tobacco, Potato, Cauliflower and Turnip	Diamond back moth (<i>Plutella xylostella</i>)	Kuvshinov <i>et al.</i> , 2001
<i>Cry V</i>	Potato	<i>Phthorimaea operculella</i> and <i>Symmetrischema tangolias</i>	Douches <i>et al.</i> , 2000; Madkour <i>et al.</i> , 2000; Lagnaoui <i>et al.</i> , 2000
<i>Cry1 Ab3</i>	Cabbage	Diamond back moth (<i>Plutella xylostella</i>)	Jin <i>et al.</i> , 2000
<i>Cry2A</i>	Rice	Yellow stem borer and rice leaf folder	Maqbool <i>et al.</i> , 1998
<i>CryIA(b)</i>	Sugarcane	Sugarcane borer (<i>Diatraea saccharalis</i> F.)	Arencibia <i>et al.</i> , 1997
<i>CryIIIB</i>	Eggplant	Colorado potato beetle (<i>Lepinotarsa decemlineata</i> Say)	Arpaia <i>et al.</i> , 1997



Cry 1 A(c)	Chick pea	Pod borer (<i>Heliothis armigera</i>)	Kar et al., 1997
Cry 1 A(c)	Peanut	Lesser Corn Stalk Borer (LCB) <i>Elasmopalpus lignosellus</i>	Singsit et al., 1997
Cry 1 C	Alfalfa and tobacco	<i>Spodoptera</i>	Strizhov et al., 1996
Cry 1Ac	Brassica napus	Diamondback moth (<i>Plutella xylostella</i> L.) and cabbage looper (<i>Tricholusia ni</i> Hübner)	Stewart et al., 1996a
Cry 1 Ac	Soybean	<i>Helicoverpa zea</i>, <i>Pseudoplusia includens</i>, <i>Heliothis virescens</i>, and <i>Anticarsia gemmatalis</i>	Stewart et al., 1996b
CryIIIA	Potato	<i>Leptinotarsa decemlineata</i>	Perlak et al., 1993
Cry1A(b)	Rice	stemborer (<i>Chilo suppressalis</i>) and leaffolder (<i>Cnaphalocrosis medinalis</i>)	Fujimoto et al., 1993
HD-1 (<i>cry1A(b)</i>) and HD-73 (<i>cry1A(c)</i>)	Cotton	<i>Trichoplusia ni</i> and <i>Heliothis zea</i>	Perlak et al., 1990
Bt protein	Tomato	<i>Helicoverpa zea</i>, <i>Manduca sexta</i>, <i>Keiferia hypersicella</i>	Delannay et al., 1989

Table 3. List of enzymes and enzyme inhibitors used for developing transgenic plants; legume transgenes highlighted in bold face

Plant	Gene	Insect	Order	Reference
Cotton	Pin I from <i>Solanum tuberosum</i> and Pin II inhibitor from <i>Nicotiana alata</i>	<i>Helicoverpa</i> spp.	Lepidoptera	Dunse et al., 2010
Tobacco	Proteinase inhibitor from <i>Solanum americanum</i>	<i>Helicoverpa armigera</i> and <i>Spodoptera litura</i>	Lepidoptera	Luo et al., 2009
Tomato	Potato protease inhibitor (PI-II) + Carboxypeptidase inhibitor (CPI)	Tomato fruit worm (<i>Heliothis 11bsolete</i>) and serpentine leafminer (<i>Liriomyza trifolii</i>)	Lepidoptera; Diptera	Abdeen et al., 2005
Chick pea	Bean α-amylase inhibitor 1	<i>Callosobruchus</i> spp.	Coleoptera	Sarmah et al., 2004
<i>Pisum sativum</i>	α-amylase inhibitor	Pea weevil (<i>Brochus pisorum</i>)	Coleoptera	de Sausa-Majer et al., 2004
Sugarcane	Soybean proteinase inhibitor	<i>Diatrea saccharalis</i>	Lepidoptera	Falco and Silvia-Filho., 2003
Rice	Trypsin inhibitor from Barley	<i>Sitophylus oryzae</i>	Coleoptera	Alfonso-Rubi et al., 2003
Tobacco	Enhancin (<i>Tn-En</i>) and (<i>Ha-En</i>)	<i>Trichoplusia ni</i>	Diptera	Cao et al., 2002b



Potato	Trypsin inhibitor	<i>Lacanobia oleracea</i>	Hymenoptera	Bell <i>et al.</i> , 2001
White poplar	<i>Arabidopsis</i> proteinase inhibitor (<i>Atcys</i>)	Chrysomelid beetle (<i>Chrysomela populi</i>)	Coleoptera	Delledone <i>et al.</i> , 2001
Pea	α -amylase inhibitor 1	Pea weevil	Coleoptera	Morton <i>et al.</i> , 2000
Tobacco	<i>Autographa californica</i> nucleopolyhedrovirus Enhancin	<i>Spodoptera exigua</i>	Lepidoptera	Hayakawa <i>et al.</i> , 2000
Potato and Tobacco	Soybean Kunitz C-II and PI-IV serine protease inhibitors	<i>Spodoptera littoralis</i>	Lepidoptera	Marchetti <i>et al.</i> , 2000
Rice	Bean trypsin inhibitor (<i>WTI-1B</i>)	Rice stem borer (<i>Chilo suppressalis</i>)	Lepidoptera	Mochizuki <i>et al.</i> , 1999
Tobacco and Peas	<i>Nicotiana alata</i> protease inhibitor	<i>Helicoverpa armigera</i>	Lepidoptera	Charity <i>et al.</i> , 1999
Wheat	Barely trypsin inhibitor (<i>BTI-Cme</i>)	Angoumois grain moth (<i>Sitotroga cerealella</i>)	Lepidoptera	Altpeter <i>et al.</i> , 1999
Rice	Soybean Kunitz trypsin inhibitor	<i>Nilaparvata lugens</i> Stal	Hemiptera	Lee <i>et al.</i> , 1999
Tobacco	Insect (<i>Madhuca sexta</i>) chitinase (EC 3. 2. 1. 14)	Tobacco bud worm (<i>Heliothis virescens</i>)	Lepidoptera	Ding <i>et al.</i> , 1998a
Bean	Bean α -amylase inhibitor	Bean weevil	Coleoptera	Grossi de Sa <i>et al.</i> , 1997
Rice	Cowpea Trypsin inhibitor (<i>CpTi</i>)	Striped stem borer (<i>Chilo suppressalis</i>) and pink stem borer (<i>Sesamia inferens</i>)	Lepidoptera	Xu <i>et al.</i> , 1996
Azuki bean	Common bean α -amylase inhibitor	Bruchid (<i>Zabrotes subfasciatus</i>)	Coleoptera	Ishimoto <i>et al.</i> , 1996
Pea	Bean α -amylase inhibitor	Bean weevil (<i>Bruchus pisorum</i>)	Coleoptera	Schroeder <i>et al.</i> , 1995
Cotton	<i>Manduca sexta</i> Protease inhibitor	<i>Bemisia tabaci</i>	Hemiptera	Thomas <i>et al.</i> , 1995
Alfalfa	<i>Manduca sexta</i> Antielastase proteinase inhibitor	Thrips	Thysanoptera	Thomas <i>et al.</i> , 1994



Table 4. Transgenic plants developed using genes producing insecticidal secondary metabolites; Fabacean plant highlighted in bold

Plant	Gene	Insect	Order	Reference
Cotton	<i>Allium sativum</i> agglutinin	Jassid (<i>Amrasca devastans</i>) and Whitefly (<i>Bemisia tabaci</i>)	Hemiptera	Vajhala <i>et al.</i> , 2013
Rice	<i>Dioscorea batatas</i> tuber lectin 1	Brown plant hopper	Hemiptera	Yoshimura <i>et al.</i> , 2012
Rice	<i>Allium sativum</i> (<i>asal</i>) + <i>Galanthus nivalis</i> (<i>gna</i>)	<i>Nephrotetix virescens</i> , <i>Nilaparvata lugens</i> , <i>Sagotella furcifera</i>	Hemiptera	Bharathi <i>et al.</i> , 2011
Chickpea	<i>Allium sativum</i> (<i>asal</i>)	<i>Aphis craccivora</i>	Hemiptera	Chakraborti <i>et al.</i>, 2009
Rice	<i>Allium sativum</i> (<i>asal</i>)	Sap sucking insects	Hemiptera	Yarasi <i>et al.</i> , 2008
Cotton	<i>Amaranthus codatus</i> agglutinin	Aphid	Hemiptera	Wu <i>et al.</i> , 2006
Tobacco	<i>Allium sativum</i> lectin	Peach potato aphid (<i>Myzus persicae</i>)	Hemiptera	Dutta <i>et al.</i> , 2005
Maize	Snow-drop lectin (GNA)	Corn borer	Lepidoptera	Wang <i>et al.</i> , 2005c
Maize	Snow-drop lectin (GNA)	Corn leaf aphid (<i>Rhopalosiphum maidis</i>)	Hemiptera	Wang <i>et al.</i> , 2005b
Tepary bean	Arcelins (<i>arc5</i> or <i>arc1</i>)	Mexican bean weevil	Coleoptera	Zambre <i>et al.</i>, 2005
Populus	Polyphenol oxidase	Forest tent caterpillar	Lepidoptera	Wang and Constabel, 2004
Rice	Snow-drop lectin (GNA)	White backed plant hopper (<i>Sogatella furcifera</i>), brown plant hopper and green leaf hopper	Hemiptera	Nagadhara <i>et al.</i> , 2004
Rice	Snow-drop lectin (GNA)	Brown plant hopper and green leaf hopper	Hemiptera	Nagadhara <i>et al.</i> , 2003
Rapeseed	Pea lectin	Pollen beetle (<i>Meligethes aeneus</i>)	Coleoptera	Melander <i>et al.</i> , 2003
Sugarcane	Snow-drop lectin (GNA)	Mexican rice borer (<i>Eoreuma loftini</i>)	Lepidoptera	Wachtel <i>et al.</i> , 2003
Tobacco and apple	Avidin and Streptavidin	Potato tuber moth (<i>Phthorimaea operculella</i>) and light brown apple moth (<i>Epiphyas postvittana</i>)	Lepidoptera	Markwick <i>et al.</i> , 2003



Tobacco	<i>Helianthus tuberosus</i> agglutinin	Peach-potato aphid (<i>Myzus persicae</i>)	Hemiptera	Chang <i>et al.</i> , 2003
Tobacco	<i>Pinellia ternata</i> agglutinin	Aphid	Hemiptera	Yao <i>et al.</i> , 2003
Mustard	Wheat germ agglutinin	Aphid	Hemiptera	Kanrar <i>et al.</i> , 2002
Sugarcane	Lectin (GNA)	Mexican rice borer (<i>Eoreuma loftini</i>) Sugarcane borer (<i>Diatraea saccharalis</i>)	Lepidoptera	Setamou <i>et al.</i> , 2002
Rice	Lectin (GNA)	Brown plant hopper(<i>Nilaparvata lugens</i>)	Hemiptera	Tang <i>et al.</i> , 2001
Tobacco	Cholesterol oxidase	Cotton boll weevil (<i>Anthonomus grandis grandis</i>)	Lepidoptera	Corbin <i>et al.</i> , 2001
Maize	Avidin	Storage pests	-	Kramer <i>et al.</i> , 2000
Potato	Lectin (Con A)	Tomato moth (<i>Lacanobia oleracea</i>)and peach potato aphid (<i>Myzus persicae</i>)	Hemiptera	Gatehouse <i>et al.</i> , 1999
Wheat	Lectin (GNA)	Grain aphid (<i>Sitobion avenae</i>)	Hemiptera	Stoger <i>et al.</i> , 1999
Rice	Lectin (GNA)	Brown plant hopper(<i>Nilaparvata lugens</i>)	Hemiptera	Rao <i>et al.</i> , 1998
Potato	Lectin (GNA)	Tomato moth (<i>Lacanobia oleracea</i>)	Hemiptera	Gatehouse <i>et al.</i> , 1997
Rice	Lectin (GNA)	(Brown plant hopper) <i>Nilaparvata lugens</i>	Hemiptera	Gatehouse <i>et al.</i> , 1996
Grand fir	Oleoresin	Bark beetle	Coleoptera	Steele, 1995

Table 5. Some genes characterized for resistance against pests

Origin	Gene	Insect	Order	Reference
Chemical/Soybean	Serine proteinase inhibitor	<i>Eurygaster integriceps</i>	Hemiptera	Saadati and Bandani, 2011
Corn	α -amylase inhibitor	<i>Sitophilus zeamais</i>	Coleoptera	Marsaro Junior <i>et al.</i> , 2005
Rice	<i>Bph 1</i>	Brown plant hopper (<i>Nilaparvata lugens</i>)	Hemiptera	Kim and Sohn., 2005
<i>Capsicum Annum</i>	Proteinase inhibitors (<i>CapA1</i> and <i>Cap A2</i>)	<i>Helicoverpa armigera</i>	Lepidoptera	Tamhane <i>et al.</i> , 2005
Chick pea	Trypsin inhibitor	<i>Helicoverpa armigera</i>	Lepidoptera	Srinivasan <i>et al.</i> , 2005
Jackbean, Snowdrop	<i>Con A</i> , GNA (lectins)	<i>Lacanobia oleracea</i>	Hymenoptera	Bell <i>et al.</i> , 2004



Jackbean, Snowdrop	Lectin (<i>Con A</i>)	Pea aphid (<i>Acyrthosiphon pisum</i>)	Hemiptera	Suavion <i>et al.</i> , 2004
<i>Poecilanthe parviflora</i> seeds	Trypsin inhibitor	<i>Diatraea saccharalis</i> , <i>Anagasta kuehniella</i> , <i>Spodoptera frugiperda</i> , and <i>Coryza cephalonica</i>	Lepidoptera	Garcia <i>et al.</i> , 2004
<i>Talisia esculenta</i>	Lectin (<i>TEL</i>)	<i>Callosobruchus maculatus</i>	Coleoptera	Macedo <i>et al.</i> , 2004
Tobacco	Cembratrienols (CBTols)	Aphids	Hemiptera	Wang <i>et al.</i> , 2004
<i>Vigna unguiculata</i>	Proteinase inhibitor (vicilins)	Sugar cane stalk borer (<i>Diatraea saccharalis</i>)	Lepidoptera	Mota <i>et al.</i> , 2003
Winged bean)	Trypsin inhibitor	<i>Helicoverpa armigera</i>	Lepidoptera	Giri <i>et al.</i> , 2003
<i>Pinellia ternata</i>	Lectin (<i>PTA</i>)	Peach potato aphids	Hemiptera	Yao <i>et al.</i> , 2003
<i>Koelreuteria paniculata</i>	Lectin	<i>Callosobruchus maculatus</i> and <i>Anagasta kuehniella</i>	Coleoptera Lepidoptera	Macedo <i>et al.</i> , 2003
<i>Streptomyces</i> sp.	Cholesterol oxidase (E. C1. 1. 3. 6)	Cotton boll weevil larvae	Lepidoptera	Santos <i>et al.</i> , 2002
Jackbean, Snowdrop	Lectins (<i>Con A, GNA</i>)	<i>Lacanobia oleracea</i>	Lepidoptera	Fitches <i>et al.</i> , 2001
Soybean	Proteinase inhibitor	Sugarcane borer (<i>Diatraea saccharalis</i>)	Lepidoptera	Pompermayer <i>et al.</i> , 2001
Snowdrop, Jack bean and <i>Psophocarpus tetragonolobus</i>	Lectins (<i>GNA, Con A, PTA</i>)	Plant hoppers (<i>Tarophagus proserpina</i> and <i>Nilaparvatha lugens</i>)	Hemiptera	Powell 2001
<i>Phaseolus vulgaris</i>	α -amylase inhibitor	Coffee berry borer (<i>Hypothenemus hampei</i>)	Coleoptera	Valencia <i>et al.</i> , 2000
<i>Phaseolus vulgaris</i>	α -amylase-like	Cowpea weevil (<i>Callosobruchus maculatus</i>), azuki bean weevil (<i>C. chinensis</i>)	Coleoptera	Ishimoto <i>et al.</i> , 1999
Kidney bean (<i>Phaseolus vulgaris</i>)	Arcelin-1 (lectin-like protein)	Bruchids	Coleoptera	Fabre <i>et al.</i> , 1998
<i>Griffonia simplicifolia</i>	N-acetylglucosamine-specific lectin	Cowpea weevil (<i>Callosobruchus maculatus</i>)	Coleoptera	Zhu <i>et al.</i> , 1996
<i>Streptomyces</i> sp.	Cholesterol oxidase (E. C1. 1. 3. 6)	Cotton boll weevil larvae	Coleoptera	Purcell <i>et al.</i> , 1993; 1994



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