

An Overview on the Crystal Toxins from *Bacillus thuringiensis*

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ABSTRACT

Strains of *Bacillus thuringiensis* (*Bt*) are known to produce crystalline proteins (δ -endotoxins) concomitantly with sporulation during their stationary phase of growth, which are demonstrated as lethal to lepidopterous, coleopterous and dipterous insects in addition to mites, nematodes, protozoa and flukes. Upon ingestion, the δ -nascent endotoxin is an inactive protoxin complex of (*Cry* alone or *Cry* and *Cyt* toxins together) high molecular mass, which is cleaved upon ingestion into the active component proteins at the high alkaline environments in the digestive tract of these agricultural pests. Conventionally, *Bt*-crystals are being produced employing submerged or liquid fermentation techniques in commercial media, but recently many workers have used solid-state fermentation strategy for the enhanced production of *Bt*-toxin at low cost. Apart from δ -endotoxin, some isolates of *Bt* produce another class of insecticidal small molecules called β -exotoxin (thuringiensin), which may be harmful to humans. Moreover, resistance to *Bt* developed in various target pest is yet another concern for *Bt*-industry. Following a brief introduction, this review addresses various toxins produced by various strains of *Bt*, *Bt* production media and media formulations with emphasis to solid-state fermentation, general structure of *Cry* toxin, its mode of action, target pests, bioassay, resistance to *Bt* toxins and resistance management. Briefly, this review would provide the readers an overview on the general aspects of *Bt* toxin, its general structure and mechanism of action.

Keywords: *Bacillus thuringiensis*; δ -Endotoxin; Resistance

1. Introduction

Biological pesticide is one of the most promising alternatives over conventional chemical pesticides, which offers less or no harm to the environments and biota. *Bacillus thuringiensis* (commonly known as *Bt*) is an insecticidal Gram-positive spore-forming bacterium producing crystalline proteins called delta-endotoxins (δ -endotoxin) during its stationary phase or senescence of its growth. *Bt* was originally discovered from diseased silkworm (*Bombyx mori*) by Shigetane Ishiwatari in 1902. But it was formally characterized by Ernst Berliner from diseased flour moth caterpillars (*Ephestia kuhniella*) in 1915 [1]. The first record of its application to control insects was in Hungary at the end of 1920, and in Yugoslavia at the beginning of 1930s, it was applied to control the European corn borer [2]. *Bt*, the leading biorational pesticide was initially characterized as an insect pathogen, and its insecticidal activity was ascribed largely or completely to the parasporal crystals. It is active against more than 150

species of insect pests. *Bt* is normally marketed (as a mixture of dried spores and toxin crystals) under various trade names worldwide for controlling many plant pests, mainly caterpillars belonging to Lepidoptera (represented by butterflies and moths), mosquito larvae and a few others including unconventional targets like mites. The share of *Bt* products in agrochemical (fungicide, herbicide and insecticide) market is about only 1%. The first commercial *Bt* product was produced in 1938 by Libec in France, but the product was used only for a very short time due to World War II, and then in the USA in the 1950s [3]. The toxicity of *Bt* culture lies in its ability to produce the crystalline protein, this observation led to the development of bioinsecticides based on *Bt* for the control of certain insect species among the orders Lepidoptera, Diptera, and Coleoptera [3-5]. Nowadays, *Bt* isolates are reported also active against certain nematodes, mites and protozoa [6]. It is already a useful alternative or supplement to synthetic chemical pesticide for applications in commercial agriculture, forest management, and mosquito control, and also a key source of genes for

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transgenic expression to transfer pest resistance in plants. Due to this economic interest, numerous approaches have been developed to enhance the production of *Bt* bioinsecticides. The insecticidal activity of *Bt* is known to depend not only on the activity of the bacterial culture itself, but also on abiotic factors, such as the medium composition and cultivation strategy.

2. *Bt* Toxins

Bt produces one or more types of parasporal crystalline proteins (called δ -endotoxins) concomitantly with sporulation. Crystalline (*Cry*) or cytolytic (*Cyt*) proteins singly or in their combination constitute the δ -endotoxins [7]. *Cry* proteins are parasporal crystalline inclusions produced by *Bt* that exhibit experimentally verifiable toxic effect to a target organism or have significant sequence similarity to a known *Cry* protein. *Cyt* proteins are also parasporal inclusions exhibiting hemolytic (cytolytic) activity with obvious sequence similarity to a known *Cyt* protein. These toxins are highly specific to their target insect, but innocuous to humans, vertebrates and plants, and are completely biodegradable [8]. These crystalline proteins are mainly encoded by extra-chromosomal genes located on the plasmids. The parasporal crystalline proteins produced during the stationary (senescence) phase of its growth cycle account for 20% - 30% of the dry weight of the cells of this phase [9]. Expression of most *Cry* genes (e.g., *cry1Aa*, *cry 2A*, *cry 4A*, etc.) are well regulated in the sporulation phase of growth. Studies have shown that several *Cry* proteins—when expressed in either *E. coli* or *B. subtilis*—expressed as 130 to 140 kDa protoxin complex molecules that retain their biological activity. More than 200 types of endotoxin gene have been cloned from various strains of *Bt*, and sequenced so far. The plasmid profiles of most *Bt* strains are rather complex, with molecular weight varying from 2 to 200 kb and the number of plasmids ranging from 1 to 10 in most strains [10]. The self-assembly of these 130 kDa proteins is spontaneous, mediated primarily by the C-terminus of the protein. Their cysteine-rich carboxyl terminus is highly conserved among lepidopteran-specific *Cry* proteins, which generates a number of disulfide bridges that allow good crystal packing and also protects the toxin from the attack of various proteases. Commercial insecticides derived from *Bt* have a long history of successful use in the biocontrol of insect pests [11,12]. Many studies examined the composition and methods of preparation of nutrient media for entomopathogenic bacteria [13,14]. Chromosomal insertion of *Cry* gene may enhance the production of δ -endotoxins in *Bt* strains [15]. Erythromycin resistance may affect the sporulation processes in *Bt* and *B. subtilis* [16,17]. Most *Bacillus* strains produce a mixture of structurally different insecticidal crystal proteins (*Cry* proteins), which are encoded by different *Cry* genes

which target different insect orders (Table 1). Each of these proteins may contribute to the insecticidal spectrum of a strain that makes it selectively toxic to a wide variety of insects belonging to the Lepidoptera, Coleoptera, Diptera, Hymenoptera and Mallophaga, as well as to other invertebrates [11,18-20]. *Bt* strains are able to produce exoenzymes, such as proteases and α -amylases [21]. Apart from δ -endotoxin, some isolates of *Bt* produce another class of insecticidal small molecules called β -exotoxin, the common name for which is thuringiensis [22].

Table 1. Endotoxin producing Bacilli and target organisms.

<i>Bacillus</i> spp.	Target pest	Reference
<i>B. laterosporus</i> & <i>Brevibacillus laterosporus</i>	<i>Musca domestica</i> and <i>Aedes aegypti</i>	[25]
<i>B. sphaericus</i> & <i>Bt israelensis</i>	<i>Culex quinquefasciatus</i> , <i>C. pipiens</i> , <i>C. tarsalis</i>	[26-28]
<i>Bt</i>	<i>Acyrtosiphon pisum</i> , <i>Aedes aegypti</i> , <i>Autographa californica</i> , <i>Cacyreus marshalli</i> , <i>Epinotia aporema</i> , <i>Lobesia botrana</i> , <i>Manduca sexta</i> , <i>Manduca sexta</i> , <i>Pectinophora gossypiella</i> , <i>Pieris brassicae</i> , <i>Plutella xylostella</i> , <i>Rhizoglyphus robini</i> , <i>Spodoptera exigua</i> , <i>Spodoptera frugiperda</i>	[29-43]
<i>Bt berliner</i>	<i>Lygus hesperus</i>	[44]
<i>Bt BRL 43</i>	First instar larvae of cotton leaf worm, cotton boll worm and black cut worm	[45]
<i>Bt finitimus</i> B-1166 VKPM	-	[46]
<i>Bt galleriae</i>	<i>Galleria mellonella</i> L.	[47]
<i>Bt</i> H14	<i>Leishmania. major</i>	[48]
<i>Bt</i> IPS 78/11	<i>Manduca sexta</i>	[49]
<i>Bt israelensis</i> IPS78/11	<i>Lucilia cuprina</i> , <i>L. sericata</i> , and <i>Calliphora stygia</i>	[50]
<i>Bt sotto</i>	Cabbage butterfly	[51]
<i>Bt kurstaki</i>	<i>Helicoverpa zea</i> ; <i>Scrobipalpula absoluta</i> ; <i>Malacosoma neustria</i> and <i>Lymantria dispar</i> larvae	[52,53]
<i>Btk</i> (serotype H3a, 3b, 3c) strain BNS3	<i>Prays oleae</i>	[54]
<i>Btk</i> HD1 & <i>Btk</i> HD73	<i>Manduca sexta</i> <i>Heliothis irescens</i>	[55]
<i>Bt tolworthi</i>	<i>Spodoptera frugiperda</i> , <i>Ostrinia nubilalis</i> and <i>Plutella xylostella</i> , <i>S. exigua</i>	[56,57]
<i>Bt. tenebrionis</i>	synanthropic mites	[58]

Beta-exotoxin and the other *Bacillus* toxins (δ -endotoxins) may contribute to the general insecticidal toxicity of the bacterium to lepidopteran, dipteran, and coleopteran insects. *Beta*-exotoxin is known to be toxic to humans and almost all other forms of life and, in fact, its presence is prohibited in *Bt* products [23]. Engineering of plants to contain and express only the genes for δ -endotoxins avoids the problem of assessing the risks posed by these other toxins that may be produced in microbial preparations [24].

3. General Structure of *Cry* Toxin

The major component of crystals toxic to lepidopteron larvae is a 130 kDa protein (protoxin), which upon cleavage in the insect yields the functional (insecticidal) proteins of lower molecular weight; very often the crystal formed is an assemblage of many proteins [59]. A *Bt* isolate (*Soil-47*) showed distinct bands of 32.1 and 34.6 kDa. The band corresponding to 32.1 kDa protein could arise from the type *Cry1* and/or *Cry 4* gene, while the other (34.6 kDa) protein is possibly encoded by the type *Cyt* gene [60]. An unexpected finding was that a 20 kb heterologous DNA fragment was found intimately associated with the crystals from *Btk* HD73. The DNA is not susceptible to nuclease attack unless the protoxin is removed or proteolyzed to toxin. The active toxin is not associated with DNA; however, evidence was obtained which indicated that the DNA was involved in the generation of toxin from the crystal protein. [61].

Structure determination of *Bt* toxins remains one of the most important tools in understanding and improving the utility of these proteins. Crystal structure of *Cry* III A has been published first [62] and several others are now available. Xia *et al.* [63] predicted the first theoretical model of the three dimensional (3D) structure of a *Cry* (*Cry* 5Ba) toxin by homology modeling on the structure of the *Cry1Aa* toxin, which is specific to Lepidopteran insects. The three-domain structure of *Cry*III A consisted of the following: an α -helical barrel (domain I) which shows some resemblance to membrane-active or spore-forming domains of other toxins [64]; a triangular prism of "Greek key" beta sheets (domain II); and a β -sheet jelly-roll fold (domain III) [62]. Members of this 3-domain *Cry* family are used worldwide for insect control, and their mode of action has been characterized in some details [8].

4. Mode of Action

Mode of action of δ -endotoxin involves several events that must be completed several hours after the ingestion in order to lead to insect death. Following ingestion of the inactive protoxin, the crystals are solubilized by the alkaline conditions in the insect midgut and are subse-

quently proteolytically converted into a toxic core fragment [65]. This activated toxin binds to receptors located on the apical microvillus membranes of epithelial midgut cells. For *Cry1A* toxins, at least four different binding sites have been described in different lepidopteran insects: a cadherin-like protein (CADR), a glycosylphosphatidylinositol (GPI)-anchored aminopeptidase-N (APN), a GPI-anchored alkaline phosphatase (ALP) and a 270 kDa glycoconjugate [66]. *Cry* toxins interact with specific receptors located on the host cell surface and are activated by host proteases following receptor binding, which would result in the formation of a pre-pore oligomeric structure that is insertion competent. In contrast, *Cyt* toxins directly interact with membrane lipids and insert into the membrane. Recent evidence suggests that *Cyt* synergizes or overcomes resistance (for instance, to mosquitoicidal-*Cry* proteins) by functioning as a *Cry*-membrane bound receptor [8].

Once activated, the endotoxin binds to the gut epithelium and causes cell lysis by the formation of cation-selective channels, which leads to death. The activated region of the δ -endotoxin is composed of three distinct structural domains: an N-terminal helical bundle domain involved in membrane insertion and pore formation; a *beta*-sheet central domain involved in receptor binding; and a C-terminal beta-sandwich domain that interacts with the N-terminal domain to form a channel. After binding, toxin adopts a conformation suitable for allowing its insertion into the cell membrane. Subsequently, oligomerization occurs, and this oligomer forms a pore or ion channel induced by an increase in cationic permeability within the functional receptors contained on the brush borders membranes. This allows the free flux of ions and liquids, causing disruption of membrane transport and cell lysis leading to insect death [65,55]. The complete nature of this process is still elusive.

Differences in the extent of solubilization sometimes explain differences in the degree of toxicity among *Cry* proteins [67]. A reduction in solubility is speculated to be one potential mechanism for insect resistance [68]. *Cry3A* protein may be necessary for the solubilization of toxins in the midgut of insects [69]. Most recently, two models were proposed for the action of crystal proteins *i.e.*, the sequential binding model and signaling pathway model [70].

5. Target Pests

It is well documented that many insects are susceptible to the toxic activity of *Bt*; of them, lepidopterans have exceptionally been well studied, and many toxins have shown activity against them [66]. The order Lepidoptera encompasses majority of susceptible species belonging to agriculturally important families such as Cossidae, Ge-

lechiidae, Lymantriidae, Noctuidae, Pieridae, Pyralidae, Thaumetopoetidae, Tortricidae, and Yponomeutidae [71]. A novel crystal proteins exhibiting insecticidal activity against lepidopterans has been reported from *Bt* strains [72].

Dipterans are also important target pests, and many of them are highly susceptible to *Bt*. Discovery of novel strains of *Bt* containing parasporal crystal proteins having pesticidal properties against whiteflies, aphids, leaf hoppers, and possibly other sucking insects of agronomic importance extended the potential applications of this bacterium. However, the novel toxic activities found in these novel strains are not limited only to insects, as some of them produce crystals with activity against nematodes, protozoans, flukes, collembolans, mites and worms, among others [66].

6. Production Media and Media Formulations

Indeed, large quantities of spores with high insecticidal activity are required for practical applications. This means that while handling *Bt* as bioinsecticide, a high spore count is not sufficient to ensure toxicity, but it is necessary to reach high δ -endotoxin titers. One of the most underreported aspects of *Bt* is that of the production and formulation, although there are certain work existed in connection with *Bt* growth on several synthetic or complex media [73]. There are several formulations of media proposed by different authors. Our group explored the efficacy of various raw agricultural products as supplement to LB for enhancing the toxin production, and found potato flour as an efficient supplement to commercial Luria-Bertani (LB) medium [7]. To develop a cost-effective process for the production of *Bt*-based insecticide, it is imperative to cultivate the bacterial strain in a nutrient rich medium to obtain the highest yields of spore-crystal complexes. Conventionally, *Bt*-crystals are being produced employing submerged or liquid fermentation (SmF) techniques, but recently many workers use nutrient-rich waste water or sludge from various treatment plants as the medium for the production of *Bt*-toxin [74].

Solid-state fermentation: Solid-state fermentation (SSF) has been developed in eastern countries over many centuries, and has enjoyed broad application in these regions to date [75]. The term SSF denotes cultivation of microorganisms on solid, moist substrates in the absence of a free aqueous phase (water). There are several advantages for SSF; for example, high productivities, extended stability of products and low production costs, which say much about such an intensive biotechnological application. With increasing progress and application of rational methods in engineering, SSF will reach higher levels regarding standardization and reproducibility in future.

This can make SSF as the preferred technique in the special fields of application such as the productions of enzymes and secondary metabolites, especially foods and pharmaceuticals [76].

Different production media and media compositions can change either the relative toxicity against several target insects or the insecticidal potency of products obtained from the same *Bt* strains [77]. According to Farrera *et al.*, [78], media with different composition showed changes in crystal production, *i.e.* different amounts of *Cry* proteins produced per spore would vary. The ingredients in the media affect the rate and synthesis of the different δ -endotoxins and also the size of the crystals produced. Using barley as the carbon source, Amin [79] developed a cost-effectively protocol for the mass production of *Bt*.

Several media based on complex substrates such as corn steep liquor [80], peptones [81] blackstrap molasses and Great Northern White Bean concentrate [82], or LB supplemented with agricultural products [7] have been found efficient for *Bt* bioinsecticide production. Various investigators modified such commercial media by supplementing it with mineral nutrients or various salts, *i.e.*, enriched medium. Zouari *et al.* [73] showed that *Bt* subspecies *kurstaki* produced 1 g/L of δ -endotoxin in 4.5 g/L total dry biomass in a complex liquid medium, in which the sugar was replaced by gruel hydrolysate. A mixture of extracts from potato and Bengal gram or bird feather and de-oiled rice bran or wheat bran, chickpea husk and corncob was used to cultivate *Bt israelensis* and found that the mosquitocidal activity of the crude toxin was higher than that produced in the conventional medium [83]. Valicente *et al.* [84] used LB medium supplemented with various salts, and agricultural by-products like soybean flour (0.5%) and liquid swine manure (4%) to increase *Bt* biopesticide production by SmF, which resulted in 1.18 g/L dry cell mass. Zhuang *et al.* [85] also claimed that they have purified δ -endotoxin (up to 7.14 mg/g medium) by one step centrifugation from wastewater sludge-based medium, however they did not provide any physical evidence for the purified crystals. From these reports, it seems that maximum yield of *Bt* toxin could be attained is 3.6 g/L [84] in SmF or 7.14 g/Kg medium in SSF [85], where they did not provide the actual cost effect.

7. Bioassay

Well-designed studies under confined conditions required to understand the effect of *Bt* toxins on different organisms. It is considered that *Bt* toxins also to be toxic to lepidopterous, coleopterous and dipterous insects in addition to mites, nematodes, protozoa and flukes [18, 19,40]. These proteins are usually thought to act only on the actively feeding larvae of susceptible species by a

mechanism involving consumption and proteolytic processing of the protein followed by binding to, and the lysis of midgut epithelial cells. It was found that proteolytically activated insecticidal crystal proteins significantly reduced the lifespan of adult *Heliothis virescens* and *Spodoptera exigua* at concentrations of 500 µg/ml, but not 167 or 25 µg/ml at their assay conditions [86]. *Bt* crystal proteins showed *in vitro* cytotoxicity against human cancer cells and leukemic T cells [87]. Interestingly, Xu *et al.* [88] demonstrated that the *Bt* crystal proteins can protect plasmodium-infected mice from malaria. Moreover, non-conventional targets such as *Caenorhabditis elegans* (nematode) has been demonstrated for the first time [89].

Toxins of *Btk* strain HD1 have widely been used to control the forest pests such as gypsy moth, spruce bud worm, the pine processionary moth, the European pine shoot moth and the nun moth [90]. Direct feeding of crude pellet containing *Bt*-toxin [91], pollen diet formulation [92] are the normal mode of applications being practiced in entomotoxicity assays. A different feeding strategy was successfully used for the bioassay of *A. guerreronis*, in which the dried solid fermented powder directly brushed on the infested coconut buttons [7,21]. Many authors used surfactants like BIT (1,2-benzisothiazolin-3-one), one of the inert ingredients in Foray 48 B (a *Btk* formulation); the siloxane (organosilicone) Triton-X-100, Tween 20 and Latron CS-7 are some surfactants for *Btk* formulations [93].

The mortality rate of *Thaumetopoea solitaria* on the application of *Btk* toxin has been demonstrated by Er *et al.* [94]. Purified *Btk* toxin inhibited the growth of monarch larvae, but did not cause mortality [95]. The LC₅₀ value of *Btk* was found to be 398.1 µg/ml against caterpillars of *Arctornis submarginata* [96]. Toxicity of several formulations of *Btk* to beet armyworm (*Spodoptera exigua*) was determined using neonate larvae in a diet incorporation bioassay.

Probit analysis (LC₅₀) has been used by many authors for ascertaining the efficacy of various *Bt* formulations. For instance, Yashodha and Kuppasamy [97] successfully used dipping method for testing the efficacy of *Btk* formulation in Tween 20 on Brinjal. Gobatto *et al.* [98] used various concentrations of spore suspension of *Bt* for estimating the probit value on mosquito and a moth. Payne *et al.* [40] employed artificial feeding assay for Two-spotted spider mite (*T. urticae*), a related mite to *E. orientalis* with different feeding regime. They fed the mite with 5 mg spray-dried powder of *Bt* broth (a mixture of pores, crystals, cellular debris) in 1 ml sucrose (10%) containing preservatives and surfactant.

Possible use of *Bt* preparation (Dipel 2X) as a substitute for chemical insecticides (Lannate and Hostathion) was evaluated against two major pests of potato crop,

Agrotis sp. and *Spodoptera exigua*. The toxicity studies of *Bt* to four instars larvae of diamondback moth, *Plutella xylostella* (L.) suggested that *Bt* could be an important agent for the control of larval instars of *Plutella xylostella*. [42]. The *Bt* diet suppressed the growth of the four mite species such as *Acarus siro* L., *Tyrophagus putrescentiae* (Schrank), *Dermatophagoides farinae* Hughes, and *Lepidoglyphus destructor* (Schrank) via feeding tests [58].

8. Resistance to *Bt* Toxins

Laboratory-selected strains: In the past, it was believed that insects would not develop resistance to *Bt* toxins, since *Bt* and insects have coevolved. Starting in the mid-1980s, however, a number of insect populations of several different species with different levels of resistance to *Bt* crystal proteins were obtained by laboratory selection experiments, using either laboratory-adapted insects or insects collected from wild populations [99,100]. Examples of laboratory-selected insects resistant to individual *Cry* toxins include the Indian mealmoth (*Plodia interpunctella*), the almond moth (*Cadra cautella*), the Colorado potato beetle (*Leptinotarsa decemlineata*), the cotton leafworm (*Spodoptera littoralis*), the beet armyworm (*S. exigua*), *etc.* [19]. Given the multiple steps in processing the crystal to an active toxin, it is not surprising that insect populations might develop various means of resisting intoxication. It is important, however, to keep in mind that selection in the laboratory may be very different from selection that occurs in the field. Insect populations maintained in the laboratory presumably have a considerably lower level of genetic diversity than field populations. Several laboratory experiments to select for *Bt* resistance in diamondback moths failed, although the diamondback moth is the only known insect reported so far to have developed resistance to *Bt* in the field [19]. It is possible that the genetic diversity of the starting populations was too narrow and thus did not include resistance alleles. In the laboratory, insect populations are genetically isolated; dilution of resistance by mating with susceptible insects—as observed in field populations—is excluded [19].

In addition, the natural environment may contain factors affecting the viability or fecundity of resistant insects, *i.e.*, factors excluded from the controlled environment of the laboratory. Resistance mechanisms can be associated with certain fitness costs that can be deleterious under natural conditions [101]. Natural enemies, such as predators and parasites can influence the development of resistance to *Bt* by preferring either the intoxicated, susceptible or the healthy resistant insects. In the former case, one would expect an increase in resistance development, while in the latter, natural enemies can help to retard resistance development to *Bt*. Never-

theless, selection experiments in the laboratory are valuable because they reveal possible resistance mechanisms and make genetic studies of resistance possible.

Field-selected strains: The first case of field-selected resistance to *Bt* was reported from Hawaii, where populations of diamondback moth showed different levels of susceptibility to a formulated *Bt* product (Dipel). Populations from heavily treated areas proved more resistant than those populations treated at lower levels, with the highest level of resistance at 30-fold [100].

The resistance trait is conferred largely by a single autosomal recessive locus [102]. This “Hawaii” resistance allele simultaneously confers cross-resistance to *Cry1Aa*, *Cry1Ab*, *Cry1Ac*, *Cry1Fa*, and *Cry1Ja* but not to *Cry1Ba*, *Cry1Bb*, *Cry1Ca*, *Cry1Da*, *Cry1Ia*, or *Cry2Aa* (369). At least one *Cry1A*-resistant diamondback moth strain has been shown to be very susceptible to *Cry9C* [31].

Resistance to *Btk* products and resulting failure in diamondback moth control has resulted in the extensive use of *Bt* subsp. *aizawai*-based insecticides in certain locations [19]. Insects in two colonies from Hawaii showed up to a 20-fold resistance to *Cry1Ca*, compared to several other colonies, including one obtained earlier from the same location, as well as moderately high resistance to *Cry1Ab* and *Btk*-based formulations [19].

A Malaysian strain simultaneously highly resistant to the *kurstaki* and the *aizawai* subspecies was apparently mutated in several loci [31]. A *Cry1Ab* resistance allele associated with reduced binding to brush border membrane vesicles receptors was partially responsible for resistance to both subspecies. Genetic determinants responsible for subspecies *kurstaki*-specific and subspecies *aizawai*-specific resistance segregated separately from each other and from the *Cry1Ab* resistance allele in genetic experiments [103].

After less than 2 decades of intensive use of *Btk* in crucifer agriculture, resistant insects have evolved in numerous geographically isolated regions of the world, and subspecies *aizawai* resistance is beginning to appear even more rapidly.

Defying the expectations of scientists monitoring transgenic crops such as corn and cotton that produce insecticidal proteins derived from *Bt*, target insect pests have developed little or no resistance to *Bt* crops thus far, according to US Department of Agriculture-funded scientists. These findings suggest that transgenic *Bt* crops could enjoy more extended, more profitable commercial life cycles and that the measures established to mitigate resistance before the crops were introduced are paying off [104].

Evolution of resistance in pests can reduce the effectiveness of insecticidal proteins from *Bt* produced by transgenic crops. Field outcomes support theoretical pre-

dictions that factors delaying resistance include recessive inheritance of resistance, low initial frequency of resistance alleles, abundant refuges of non-*Bt* host plants and two-toxin *Bt* crops deployed separately from one-toxin *Bt* crops. The results imply that proactive evaluation of the inheritance and initial frequency of resistance are useful for predicting the risk of resistance and improving strategies to sustain the effectiveness of *Bt* crops [105].

9. Resistance Management

Resistance management strategies try to prevent or diminish the selection of the rare individuals carrying resistance genes and hence to keep the frequency of resistance genes sufficiently low for insect control [106,107].

Proposed strategies include: the use of multiple toxins (stacking or pyramiding), crop rotation, high or ultrahigh dosages, and spatial or temporal refugia (toxin-free areas). Retrospective analysis of resistance development does support the use of refugia [99]. Experience with transgenic crops expressing cry genes grown under different agronomic conditions is essential to define the requirements of resistance management. In transgenic plants, selection pressure could be reduced by restricting the expression of the crystal protein genes to certain tissues of the crop (those most susceptible to pest damage) so that only certain parts of the plant are fully protected, the remainder providing a form of spatial refuge. It has been proposed that cotton lines in which *Cry* gene expression is limited to the young bolls may not suffer dramatic yield loss from *Heliothis* larvae feeding on other plant structures, since cotton plants can compensate for a high degree of pest damage [108].

Another management option is the rotation of plants or sprays of a particular *Bt* toxin with those having another toxin type that binds to a different receptor. A very attractive resistance management tactic is the combination of a high-dose strategy with the use of refugia [19].

10. Conclusion

Development of resistance to *Bt* toxin is one of the concerns of *Bt*-based agroindustry. It was expected that resistance would be developed in transgenic crops such as corn and, interestingly target insect pests have developed little or no resistance to these *Bt* crops. It suggests that transgenic *Bt* crops could enjoy more extended, more profitable commercial life cycles and the measures established to mitigate resistance before the crops were introduced are paying off. Nevertheless, making *Bt* toxin at low cost for the farmers, especially in the developing and underdeveloped countries remains one of the major challenges, wherein SSF offers great potentials. In fact, the ill effects of the exotoxin, thuringiensin from *Bt* on humans (and other animals too) are a growing concern.

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