INSECT RESISTANCE IN TRANSGENIC PLANTS EXPRESSING Bacillus thuringiensis TOXIN GENS¹

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ABSTRACT

^Acrystal protein gene (ht2) has been cioned from plasmid DNA of *Bacillus thuringiensis* (*B.t.*) berliner 1715 and directs the synthesis of a 130 kd protein (Bt2) in *E. coli* which is toxic to larvae of *Pieris brassicae* and *Manduca sexta*. Treatment of the Bt2 protein with trypsin or chymotrypsin yields a 60 kd protease resistant fragment which is fully toxic towards insect larvae "in vivo" and insect cell lines "in vitro". The minimal portion of the Bt2 protein required for toxicity has been mapped by deletion analysis and coincides with the 60 kd protease resistant Bt2.-fraqment. Tobacco piant celis have been transformed with chimeric toxin genes using a Ti plasmid vector. Transformed plants express a functional toxin and exhibit resistance against insect larvas.

iNTRODUCTION

Bacillus thuringiensis $(B. t.)$ is a gram positive bacterium which produces ondogenous crystais upon sporulation. The crys tais are composed of protein and are specificaliy toxic against

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certain insect larvae, mainly lepidooteran and dipteran species(DiJLMAGE *et ai.,* 19801. Upon ingestion by larvae the crys tais disolve in the alkaline conditions of the insect midgut and release proteins of molecular weight (M.W.) 130 - 160 kd (HUBER *et ai.,* 19811. These high M.W. protoxins are proteolytically processed by midgut proteases to yield smaller toxic
fragments [LILLEY et al., 1980]. Most crystal protein genes fragments [LILLEY *et al.*, 1980]. Most crystal protein have been localised on large plasmids [GONZALEZ *et ai.,* 1982; KRONSTAD *et ai.,* 1983]. Some genes have recenity been cloned and functional toxin was expressed in *E. coli* [SCHNEPF *et al.*, 1981; KLIER *et ai.,* 1982; SHIBANO *et ai.,* 1985; ADANG *et ai.,* 1985].

B.t. toxins are highly specific in their activity and the refore represent a perfectly safe insecticide and an interesting alternativ for chemical control agents. Commercial prepa rations based on *B. t.* have already been used for several years as a biological insecticide, however only with limited success, mainly due to high production costs and instability of *B*, *t*. in the field. We have used the *Agrobacterium* vector system to the field. We have used the *Agrobacterium* vector transfer and express the *B.t.* toxin gene in tobacco plants.We present data on the resistance of those plants against insect larvae.

RESULTS

Cloning of a B. t. gene encoding a 130 kd'crystal protein exhibiting insect toxicity.

Since most crystal protein genes in *B. thuringiensis* strains are plasmid-born [KRONSTAD *et al.*, 1983], we used a purified plasmid preparation from strain *B. thuringiensis* berlineer 1715 for the cloning of crystal protein genes. A library was constructed by clining size fractionated Sau 3A digested plasmid DNA into an *E. coli* expression vector. Using a colony immunobiot assay with rabbit anti-crystal protein serum, 4 clones containing overlapping DNA fragments were isolated. A 7.5 kb BamHI-PstI fragment containing the toxin gene *was* subcloned into pUC8, generating plasmid pGI502.

Total ceil extract of *E. coii* K514 (pGI502) was anaiysed on SDS-PAGE. An intense protein band, with apparant M.W. 130 kd, was visible and was not present in K514 containing the pUC8 plasmid without insert. This protein, termed Bt2, comigrated in SDS-PAGE with one of the major crystal proteins of B, t , berliner, Bt2 protein represented between 5 and 10% of *B.t.* berliner. Bt2 protein represented between 5 and 10% the total protein content in K514 (pGI502) and was present as a precipitate in *E. coZi.*

The reiationship between Bt2 and *B.t.* crystal proteinswas confirmed by its imniunological properties and biologicai acti vity. In Western blotting, Bt2 reacted strongly with a rabbit

anti-Bt berliner crystal serum and in ELISA, 8 out of 16 monoclonal antibodies, generated against $B.t.$ berliner crystal proteins, reacted with the purified Bt2. The toxic activity proteins, reacted with the purified Bt2. The toxic of Bt2 protein was assayed on *Pieris brassicae* and *Manduca sex te* larvae. Purified preparations of Bt2 showed toxicity leveis comparable to those of solubilized crystals from *B.t.* berliner (Tabie 1).

TABLE 1 - Insect toxicity of B.T. crystal proteins

Nucleotide sequence of the gene and N-terminal amino acid sequence of the crystal protein

The *bt2* toxin gene was localized by deletion mapping on a 4343 bp HpaI-PstI fraqment. The sequence of this fragment showed one large open reading frame encoding a protein of 1155 amino acids with a predicted molecular weight of 130533 d.This
agrees well with the molecular weight of Bt2 determined in agrees well with the molecular weight of Bt2 determined in
SDS-PAGE, The amino acid sequence of the N-terminal end of SDS-PAGE. The amino acid sequence of the N-terminal end this purified protein was determined by gas-phase sequencing to be $X - Asp - Asn - Asn - Pro - Asn - Ile - Asn - Glu - X -$ Ile - Pro - Tyr - Asn - Leu - X - Asn - Pro. This sequence was identical to the N-terminal amino acid sequence deduced from the nucleotid sequence.

Generation of a 60 kd toxic polypeptide through proteolytic degradation of the 130 kd Bt2 protein.

The delta-endotoxins of *B.t.* are generaily believed to be protoxins which, upon ingestion by insects, are degraded by insect gut proteases into smaller active toxin(s) [KLIER et. $a1.$, 1982]. We therefore investigated whether smaller toxic polypeptides could be generated from Bt2 by proteolytic cleavage. Purified Bt2 protein was digested with either trypsin
or chymotrypsin and at defined time intervals, aliquots were or chymotrypsin and at defined time intervals, aliquots analysed on SDS-PAGE. The results demonstrated that the 130 kd Bt2 protein is rapidiy degraded by trypsin or chymotrypsin (after 10 min. at 370C) yielding a predominant polypeptide of 60 kd.

This 60 kd polypeptide was relatively resistant to further degradation by both enzymes over a 2hperiod, indicating that it constitutes a protease resistant fragment within the Bt2 protein. The 60 kd poiypeptide generated by trypsin cieavage was purified by gel filtration and its insect toxicity was
determined. It was equally toxic to P. brassicae and M. sexta determined. It was equally toxic to P. *brassicae* and larvae as intact 130 kd Bt2 (Tabel 1). The 60 kd protein starts at the isoleucine at position 29 in the N-terminai haif of the moi ecu 1 e.

Using insect cell lines it was deduced that the 60 kd tryp sin fragment represent an active toxin molecule which is specifically toxic against the ceil lines derived from Lepidopte ra species and most toxic towards a Diptera cell line. In con trast, the intact 130 kd Bt2 protein was not toxic towards all four celi lines.

Genetic mapping of the minimal gene fragment ecoding an active toxin

Different deletions at the 5' or the 3' end of the $bt2$ gene were constructed and expressed in E. coli. We analysed the produced polypeptides to deiineate the minimal Bt2 fragment exhibiting insect toxicity (table 2).

Type of construct	$E.$ coli clone	Position 3' end Bt2 sequence		1/10
Deletions	pLB16	2170	100	100
	pLB820	2031	100	100
	pLB822	1911	100	100
	pLB828	1830	100	100
	pLB826	1827	100	100
	pLB884	1821	100	100
	pLB879	1821	98	50
	pLB834	1798	0	$\overline{2}$
	pLB950	1731	$\begin{array}{c} 6 \\ 2 \\ 0 \end{array}$	
	pLB876	1695		$\begin{smallmatrix}0\0\0\end{smallmatrix}$
	pLB12	1692		
$Bt2-IacZ$	pBZ12	1821	100	74
fusions	pBZ13	1798	4	Ω

TABLE 2 - Toxicity of celi extracts of NF1 strains containing different plasmids expressed as % mortality after 4 days; 50 3rd instar P. brassicae were used per dilution of extract.

To delineate the 5' end of the gene fragment encoding an active toxin, we constructed a 5' deletion in the Bt2 gene up to the ClaI site at the 110th bp, removing the first 36 codons of the *bt2* qene.

The experimental data show that the minimal toxic frag-
of the Bt2 protein is a 60 kd polypetide delineated by ment of the Bt2 protein is a 60 kd polypetide delineated by
positions 29 and 37 at the N-terminus and amino acid 601 and positions 29 and 37 at the N-terminus and amino acid 601 607 at the C-terrninus.

Trypsin cleavage site GLY GLU ARG ILE GLU THR GLY TYR THR PRO ILE ASP ILE ...'GGA'GAA'AGA'ATA'GAA'ACT'GGT'TAC'ACC'CCA'ATC'GAT'ATT'...
80 0 110 110 80 90 100 110 pRB2 10 600 605 TYR ILE ASP ARG ILE GLU PHE VAL PRO ALA GLU VAL 'TAT'ATA'GAT'CGA'ATT'GAA'TTT'GTT'CCG'GCA'GAA'GTA' 1800 1810
pLB834 pLB879 FIGURE 1. Mapping of the 5' and 3' end for the minimal toxic fragment.

Transfer and expression of chimeric B.t. genes in tobacco plants.

Recent progresa in plants genetic engineering techniques made it possible to transfer foreign DNA into many plant species using vectors derived from the Ti-plasmid of Agrobacterium tumefaciens. Using such vectors, several genes of bacterial origin mostly antibiotic resitance genes have been expressed in transgenic plants using plant specific promotor se-
quences. The latter have been obtained from the Ti-plasmid quences. The latter have been obtained from the quences. The factor have been obtained from the 11 -prasmid
T-DNA genes [HERRERA-ESTRELLA *et al.*, 1983; VELTEN *et al.*, 1–DNA genes [nERRENA-ESTREDLA *et al.*, 1983; VELIEN *et al.*, 1984], from plant viruses [ODELL *et al.*, 1985] or from nu-1964], from plant viruses [ODELL ℓt $d\ell$, 1985] or from nu-
clear plant genes [HERRERA-ESTRELLA 1984; SIMPSON ℓt $\alpha \ell$., 1985; KREUZALLER *et al.*, 1986]. We have used Ti-plasmid de-
rives vectors (DEBLAERE *et al.*, 1984) to transfer *bt2* gene rives vectors (DEBLAERE *et al.*, 1984) to transfer *bt2* fragments into *N. tabacum* plants using a leaf disc transforma-. tion procedure (HORSCH. et al., 1985). NPTII was used as a selectable marker. Leaf discs have been grown for 2×3 weeks in a selective medium containing 50-200 mg/ ℓ of Km. Afterwards a selective medium containing 50-200 mg/ ℓ of Km. shoots were removed from the leaf disc, grown in non selecti-
ve conditions and retested for resistance in a callus inducve conditions and retested for resistance in a callus induc-
tion assay on different concentrations of Km (50-500 mg/l). tion assay on different concentrations of Km (50-500 Plants showing high Km resistance were retained for turther tests.

Transformed tobacco plants containing a chimeric *bt2* gene were analysed for production of Bt2 related polypeptides in an

ELISA assay usinq a rabbit anti-Bt crystal serum, as well as monoclonal antibodies generated against the Bt2 protein.

Variable leveis of Bt, ranging from 10 to 100 ng per g of plant tissue were recorded in independent transformants.

Insect resistance of transformed tobacco plants.

We evaluated the toxic activity exhibited by transformed tobacco plants on insect larvae. Twenty first instar larvae of M . sexta were placed on leaf discs of 4 cm diameter in petri-dishes. Fresh leaf material was qiven each day, growth ra te and mortality of the larvae feeding on the leaves were sco red over a 6 day period. Larvae feeding of Bt positive plants rapidly showed marked growth inhibition, which retarded their transition to the second larval stage. Most ímportantly, signi ficant mortality among the larvae became apparent at day 3 and reached 100% at day 6 for several of the independent transforrned plants tested (Tabie 3).

Table 3 - Mortality rates in M. sexta larvae feeding on transformed tobacco plants.

In contrast, larvae growing on untransformed control plants developed normally and their mortality rate was only 0-5%. Thus, the effect on insect larvae of B . t . toxin expressed in plant celis was comparable to the effect of *R.* i. toxin of bacterial origin, namely growth inhibition in the initial stage followed by death.

Independant transformante was also variable (Tabie 3). Ho wever for one and the same transformant the degree of insect resistance was highiy reproduciblo when tested in independant experiments over a period of time.

In a next set of experiments, we investigated whether the levels of $B. t.$ toxin produced in transformed plants would pro tect them from significant insect damage. Fifteen freshly hat ched larvae were placed on the leaves of 40 cm high tobacco ched larvae were placed on the leaves of 40 cm high plants in the greenhouse. On control plants, considerable de mage was obvious after 4-7 days and they were completey consume after 10-15 days. In contrast, all larvae died within 4 days
and they were completely consumed after 10-15 days. In conand they were completely consumed after 10-15 days. trast, ali iarvae died within 4 days when feeding on plants expressing high leveis of Bt2 protein, and leaf damage was res tricted to holes of a few mm2 caused by each insect.

Conclusion and prospects.

A functional analysis has been performed on Bt2, a clo ned Bacillus thuringiensis crystal protein. The Bt2 protein is highly toxic to larvae of Pieris brassicae and Manduca sexta. This 130 kd polypetide is a protoxin which is not toxic in an in vitro assay on insect cell lines. However, treatment of Bt2 with proteolytic enzymes results in a 60 kd protease resistant and fully toxic polypeptide.

We have precisely determined, through genetic mapping, the minimal fragment of Bt2 which still exhibits full toxicity. This fragment was found to coincide with the 60 kd polypeptide generated by proteolytic degradation and located in the NH₂-terminal half of the protein. According to our data, it is very unlikely that toxic fragments, smaller than this
60 kd can be obtained. A detailed analysis' to identify the 60 kd can be obtained. A detailed analysis to identify functional domains of Bt2 is presently in progress, using monocional antibodies specific for well defined regions on the molecuie.

The feasibility of using plants genetic engineering tech niques to obtain insect resistant plants is cieariy exemplified here. Fully normal plants were obtained, expressing suffi ciently high levels of $B. t$. toxin to give resistance against insect damage under greenhouse conditions.Further experiments will be directed to fully evaluate this new approach under na turai conditions of stress caused by insect pests and to investigate the inheritance of the newiy acquired defense mecha nisms.

A large number of natural B. t. strains have been isolated, covering a wide spectrum of insecticidal activities and directed against Lepidoptera, Coleoptera and Diptera species.

Analysis of the molecular basis of the specificity will aliow to design new types of toxins with the desired activity against agriculturally important insect pests. This technology, in combination with new deveiopments in the field of piant engineering will enable us to engineer genetically modified varieties of commercialiy important crops, expressing resistance agains major insect pests.

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