

# Stability of Cry1Ab protein during long-term storage for standardization of insect bioassays

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The reliable use of purified Cry1Ab protein standards is a prerequisite for ecological studies and resistance monitoring programs of Cry1Ab-expressing transgenic corn. In this study the stability and activity of different Cry1Ab protein batches expressed in and purified from *Escherichia coli* were determined during two-year storage at different temperature conditions (4 °C, –20 °C, and –80 °C). SDS-Polyacrylamide gel electrophoresis showed degradation of the protein stored at 4 °C over four months, whereas no difference in the band intensity of the Cry1Ab proteins stored at –20 °C and –80 °C was observed. Bioassays with neonate larvae of *Ostrinia nubilalis* indicated that the biological activity of Cry1Ab varied from batch to batch, depending on the production process. Cry1Ab protein stored at 4 °C for four months showed a significantly decreasing activity measured as median lethal concentration (LC<sub>50</sub>), whereas the protein activity declined less than 11-fold after two years storage at –20 °C. When stored at –80 °C the toxin activity remained relatively stable for at least 30 months, as indicated by low LC<sub>50</sub> values of 7–10 ng Cry1Ab per cm<sup>2</sup> diet. These experiments demonstrate that appropriate long-term storage conditions of Cry1Ab protein standards are crucial for resistance monitoring programs of *Bt* corn, and storage at –80 °C is recommended.

**Keywords:** *Bt* corn / *Ostrinia nubilalis* / resistance monitoring / bioassay

## INTRODUCTION

Since the 1990s, transgenic crops expressing different Cry proteins of *Bacillus thuringiensis* (*Bt*) have become a new approach for insect pest control in agriculture. Cry1Ab is the primary protein used in *Bt* corn to protect the crop from damage by lepidopteran insect pests, particularly the European corn borer (ECB), *Ostrinia nubilalis* (Hübner). In 2006, different varieties of *Bt* corn were planted on more than 23.9 million hectares worldwide (Brookes and Barfoot, 2008). However, the evolution of resistance of target pest populations to transgenically expressed Cry1Ab is considered as a severe drawback of this new control strategy. In laboratory experiments using ECB populations it was demonstrated that the Cry1Ab susceptibility significantly decreased after only seven generations when selected with commercial *Bt* formulations (Huang et al., 1997). Field monitoring data in transgenic cotton revealed resistance of some field populations of *Helicoverpa zea* to Cry1Ac protein, though field-evolved resistance in other five major lepidopteran pests (*O. nubilalis*, *Sesamia nonagrioides*, *H. armigera*, *Heliiothis virescens*, *Pectinophora gossypiella*)

could not be demonstrated (Tabashnik et al., 2008). Hence, effective insect resistance management strategies appear to be essential to retain the efficacy of *Bt* crops. The development of reliable insect resistance monitoring methods allowing early detection of changes of the susceptibility of target insects are an important cornerstone of any resistance monitoring scheme (EPA, 1998; Gould, 1998). For *Bt* corn, the Cry1Ab susceptibilities of ECB field populations have been monitored in different countries, e.g. the US, Spain, France, Germany and others (Chaufaux et al., 2001; González-Núñez et al., 2000; Marçon et al., 1999; Martens et al., 1995; Saeglitz et al., 2006; Siegfried et al., 1995). Field studies on two ECB populations in Spain did not reveal any difference in their Cry1Ab susceptibilities (González-Núñez et al., 2000), whereas small variations in susceptibility to Cry1Ab protein among different geographical ECB populations were observed in the USA and in Germany (Marçon et al., 1999; Saeglitz et al., 2006). These variations, however, could be caused by (i) an existing natural variability among geographical populations; (ii) by different activities of Cry1Ab proteins used in the different tests; or (iii) by an intrinsic variability of different bioassays when using the same insect colony and the same Cry1Ab protein batch (Farinós et al., 2004; Marçon et al., 2000; Meise

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**Table 1.** Production and storage parameters of different Cry1Ab batches.

Cry1Ab-batch	Protein amount (mg)	Concentration at starting point (mg.mL <sup>-1</sup> )	Concentration after two years storage (mg.mL <sup>-1</sup> ), (SD) <i>n</i> = 10	Purity (%)	Treatment	Storage temperature
J6B	18	1.3	n.d.	80	1 × T	4 °C
H3	243	1.0	1.0 (0.15)	93	2 × T, UF	-20 °C, -80 °C
H4	200	0.8	0.8 (0.15)	84	1 × T, UF	-20 °C, -80 °C

SD = standard deviation, n.d.: not determined due to protein precipitation, T: trypsinization, UF: ultrafiltration.

and Langenbruch, 2007; Nguyen et al., 2004; Saeglitz et al., 2006). Hence, the standardization of the bioassay methodology as well as the application of standardized Cry1Ab proteins is of greatest importance in order to obtain meaningful data. Recently, different standardized methods for quantifying Cry1Ab protein from independent sources were compared and validated (Crespo et al., 2008). It was concluded that standardization of production and quantification of Cry1Ab batches may improve data consistency in monitoring efforts, and will be essential for the identification of changes of the susceptibility of target pests to Cry1Ab protein.

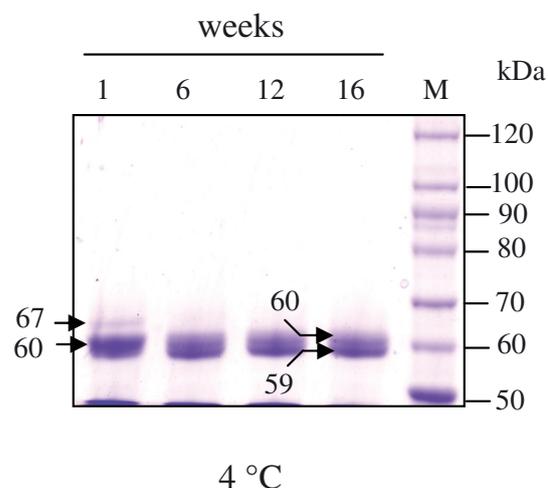
In our study we addressed the question how long a defined Cry1Ab protein standard keeps its bioactivity and what are the best storage conditions. Here, we present the results of storage experiments of different Cry1Ab batches, which were evaluated for their biochemical and biological stability during two year storage at different temperatures.

## RESULTS

### Production and stability of Cry1Ab protein batches

The production parameters of three trypsinized Cry1Ab batches J6B, H3 and H4, including protein concentration, purity, treatment during production and storage temperature are listed in Table 1. Batch J6B was stored for four months at 4 °C, whereas the batches H3 and H4 were stored for 24 months in duplicate at -20 °C and -80 °C. To compare the protein concentration before and after storage, Bradford dye assay was conducted. No decrease in the concentration of batches H3 and H4 was observed after two years storage at -20 °C (Tab. 1). Their protein concentration was still 1.0 and 0.8 mg.mL<sup>-1</sup>, respectively. In contrast, batch J6B was not measurable after 16 weeks storage due to protein precipitation.

The quality and stability of batch J6B was examined by SDS polyacrylamide gel electrophoresis (PAGE). After one week a minor band of 67 and a major band of 60 kDa were visible (Fig. 1). After 1.5–4 months the 67 kDa protein band had disappeared and two bands of

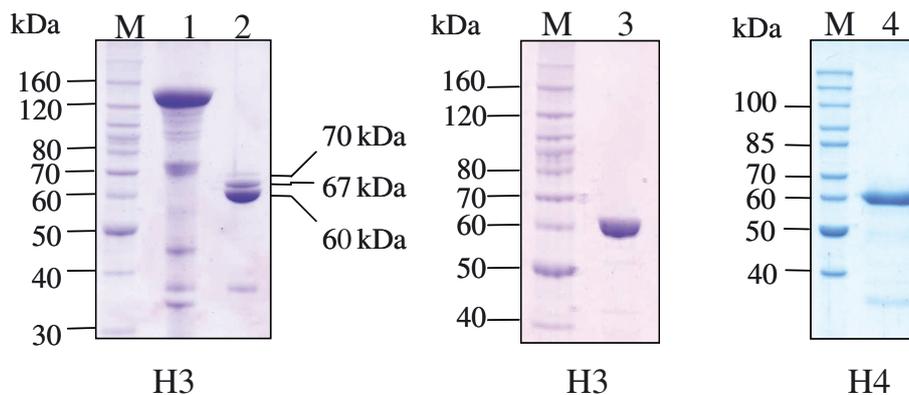


**Figure 1.** SDS polyacrylamide gel electrophoresis (7% gel) of Cry1Ab batch J6B after 1-, 6-, 12- and 16-week storage at 4 °C. (M = BenchMark™ protein ladder (GibcoBRL) with molecular masses (kDa) given to the right.)

60 and 59 kDa were observed. Due to rapid degradation of batch J6B at 4 °C, the other batches H3 and H4 were stored at -20 °C and -80 °C and their stability was examined.

Trypsinization of 130 kDa Cry1Ab protoxin of batch H3 resulted in three different bands with a molecular weight of 70, 67 and 60 kDa (Fig. 2, lane 2). Since it was assumed that the 70 and 67 kDa bands were due to incomplete trypsinization, a second trypsinization was performed. By adjusting the pH of the protein solution to 9.5 and repeating trypsinization at 22 °C overnight, a single Cry1Ab protein band of 60 kDa was obtained (Fig. 2, lane 3). Trypsinization of the batch H4 immediately resulted in a single 60 kDa band (Fig. 2, lane 4).

Batches H3, and H4 were stored for 24 months at -20 °C and -80 °C and aliquots were sampled after 6, 12, and 24 months. As shown in Figure 3, no change in number and intensity of the Cry1Ab band of batch H3 was observed at both storage temperatures. The same result



**Figure 2.** SDS polyacrylamide gel electrophoresis (7–10% gel) of Cry1Ab batches H3 and H4 after production and trypsinization. (Lane 1 = batch H3 full-length Cry1Ab protoxin 130 kDa, lane 2 = batch H3 (1× trypsinized), lane 3 = batch H3 (2× trypsinized), lane 4 = batch H4, M = PageRuler™ protein ladder (Fermentas) with molecular masses (kDa) given to the left.)

was obtained for the batch H4, indicating that storage at  $-20^{\circ}\text{C}$  or lower rendered stable Cry1Ab samples.

#### Stability of the biological activity of Cry1Ab batches

Bioassays with Cry1Ab batch J6B, which was stored at  $4^{\circ}\text{C}$ , were performed from one week to four months after production (Tab. 2). The  $\text{LC}_{50}$  dramatically increased from  $77\text{ ng}\cdot\text{cm}^{-2}$  (1 week) to  $1717\text{ ng}\cdot\text{cm}^{-2}$  (6 weeks) and  $1243\text{ ng}\cdot\text{cm}^{-2}$  (4 months), indicating a significant reduction of the insecticidal activity of this batch.

Due to the lack of larvae at the beginning of the storage experiment of batch H3, the first determination of its bioactivity was performed after 5 months storage at  $-20^{\circ}\text{C}$  ( $\text{LC}_{50} = 240\text{ ng Cry1Ab}\cdot\text{cm}^{-2}$  diet) and 10 months storage at  $-80^{\circ}\text{C}$  ( $\text{LC}_{50} = 54\text{ ng Cry1Ab}\cdot\text{cm}^{-2}$  diet) (Tab. 2). Compared to the  $\text{LC}_{50}$  values of the batch H4 at comparable storage periods, the bioactivity of the double-trypsinized batch H3 appeared to be consistently lower. At  $-20^{\circ}\text{C}$  storage the  $\text{LC}_{50}$  values steadily increased until 25 months ( $\text{LC}_{50} = 1931\text{ ng Cry1Ab}\cdot\text{cm}^{-2}$  diet). After 25 months storage at  $-80^{\circ}\text{C}$  the  $\text{LC}_{50}$  value slightly increased ( $\text{LC}_{50} = 165\text{ ng Cry1Ab}\cdot\text{cm}^{-2}$  diet) compared to 15 months before (Tab. 2). Bioassays with batch H4 performed after 7 months storage at  $-20^{\circ}\text{C}$  and  $-80^{\circ}\text{C}$  gave similar  $\text{LC}_{50}$  values of  $3\text{ ng Cry1Ab}\cdot\text{cm}^{-2}$  and  $2\text{ ng Cry1Ab}\cdot\text{cm}^{-2}$  diet, respectively (Tab. 2). Until month 22, an 11-fold activity decrease could be observed at  $-20^{\circ}\text{C}$  storage, whereas no statistically significant difference was observed between the  $\text{LC}_{50}$  values after 7 and 13 months, and after 13 and 22 months storage at  $-80^{\circ}\text{C}$ . The results obtained with batches H3 and H4 indicated that the bioactivity of Cry1Ab remained more stable when stored at  $-80^{\circ}\text{C}$  compared to  $-20^{\circ}\text{C}$ .

#### DISCUSSION

The objective of this study was to investigate the stability of the purified Cry1Ab protein during long-term storage at different temperature conditions. Biochemical and biological methods were applied to control the quality of three different Cry1Ab protein batches JB6, H3, and H4. Bradford dye assays provided information about the total protein concentration of each batch after purification and demonstrated a stable concentration of Cry1Ab protein during long-term storage. The total protein concentration of Cry1Ab protein in different batches did not change after two years of storage at  $-20^{\circ}\text{C}$  (Tab. 1). Crespo et al. (2008) compared SDS-PAGE, ELISA and Bradford assays for estimating Cry1Ab concentrations in production batches and found that Bradford assays had the lowest coefficient of variation. However, this method did not allow determining whether a Cry1Ab protein in a production batch is really stable or degraded, because it is not possible to differentiate between active Cry1Ab proteins and its degradation products.

By SDS-PAGE, the molecular weight and the purity of Cry1Ab protein can be determined. SDS PAGE analysis of protein batch J6B during storage at  $4^{\circ}\text{C}$  allowed identifying the transition of this protein from 67 kDa and 60 kDa to 59 kDa (Fig. 2). This degradation could be the result of further proteolysis by the residual activities of trypsin, as it was not removed from the protein solution during production of batch J6B. Based on this result, the two other Cry1Ab batches H3 and H4 were ultra-filtrated and stored at  $-20^{\circ}\text{C}$  and  $-80^{\circ}\text{C}$ . SDS PAGE analyses did not show any degradation of these proteins during two years storage at both temperatures (Fig. 3). Thus, removing of trypsin by ultra-filtration and freezing kept Cry1Ab protein batches stable, without changing their biochemical properties.

**Table 2.** Bioactivity (median lethal concentration, LC<sub>50</sub>) of Cry1Ab batch J6B, H3 and H4 during storage at 4 °C, –20 °C and –80 °C.

Batch	Storage temperature	Storage time (months)	N	LC <sub>50</sub> (FL)* [ng Cry1Ab.cm <sup>-2</sup> diet]	Slope ± SE	χ <sup>2</sup>	p**
J6B	4 °C	0.25	63	77 (46–121) <sup>a</sup>	1.0 ± 0.22	8.4	1
		0.5	221	111 (57–200) <sup>a</sup>	0.45 ± 0.08	1.3	0.69
		0.75	256	300 (150–630) <sup>b</sup>	0.36 ± 0.07	3.7	0.26
		1	383	541 (200–1800) <sup>bc</sup>	0.57 ± 0.13	8.8	0.14
		1.5	256	1700 (970–5400) <sup>c</sup>	0.61 ± 0.18	1.4	0.05
		4	256	1200 (680–3300) <sup>c</sup>	0.44 ± 0.09	0.4	0.06
H3	–20 °C	5	484	240 (60–1200) <sup>a</sup>	1.3 ± 0.29	16.4	1
		10	629	960 (470–2000) <sup>ab</sup>	1.2 ± 0.17	16.7	0.25
		19	637	1700 (410–20 000) <sup>b</sup>	1.6 ± 0.45	28.3	0.14
		25	639	1900 (1400–2600) <sup>b</sup>	1.2 ± 0.10	5.9	0.13
	–80 °C	10	768	54 (30–95) <sup>a</sup>	1.6 ± 0.21	14.6	1
		19	756	63 (20–159) <sup>ab</sup>	1.3 ± 0.24	32.0	0.86
H4	–20 °C	7	768	3 (2–4) <sup>a</sup>	1.8 ± 0.21	1.4	1
		13	511	4 (3–6) <sup>a</sup>	1.2 ± 0.11	1.2	0.75
		22	635	33 (19–65) <sup>b</sup>	1.4 ± 0.19	11.2	0.10
	–80 °C	7	767	2 (0–5) <sup>a</sup>	0.94 ± 0.21	14.0	1
		13	637	6 (2–14) <sup>ab</sup>	0.99 ± 0.17	12.2	0.33
		22	632	7 (4–11) <sup>b</sup>	1.2 ± 0.14	8.7	0.28

\* LC<sub>50</sub> values followed by the same letters within each batch of Cry1Ab protein were not statistically different ( $\alpha \leq 0.05$ ; Smith, 1997).

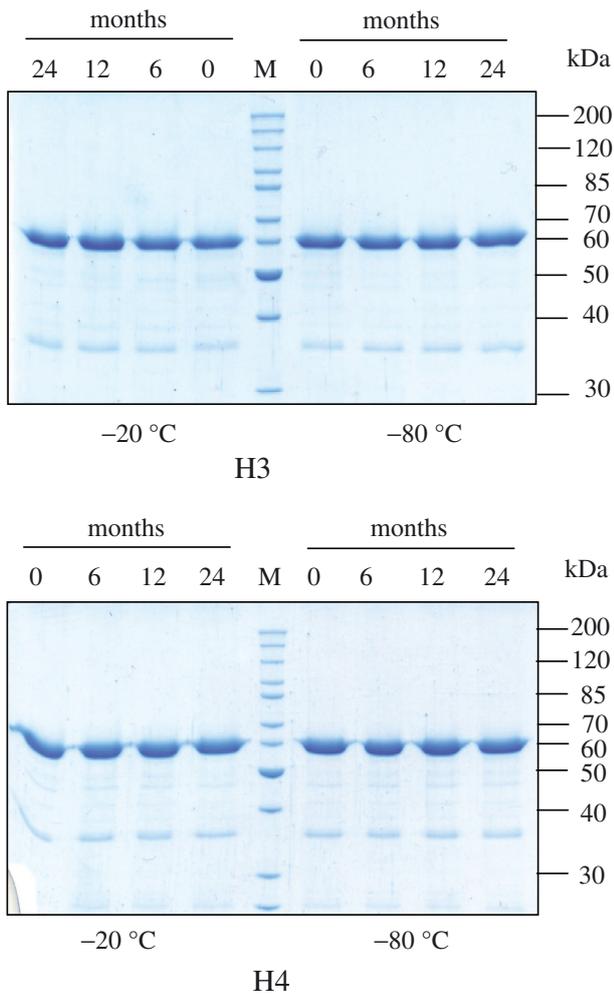
\*\* p: Potency of LC<sub>50</sub> from the first storage time point (T1) relative to that of LC<sub>50</sub> from the same batch after *n* months storage (T<sub>*n*</sub>) is estimated as:  $p_n = LC_{50(T1)}/LC_{50(Tn)}$ . N: number of tested larvae, FL = 95% fiducial limits, SE = standard error, χ<sup>2</sup>: Chi<sup>2</sup>.

Bioassays clearly demonstrated that different temperature conditions and storage periods can affect the biological stability of Cry1Ab protein. Activity of protein batch J6B significantly decreased 16-fold during one week to four months storage at 4 °C. This could be caused by further proteolysis of a partly incorrectly refolded protein with remaining trypsin after trypsinization of Cry1Ab protein, and is consistent with the SDS-PAGE analysis, which showed the degradation of a part of this protein from 60 to 59 kDa at 4 °C. The loss of activity of Cry1Ab protein was also observed by Höfte et al. (1986) and Martens et al. (1995) when a few amino acids from either the N-terminus or the C-terminus of the Cry1Ab trypsin-resistant core were removed. The potential impact of residual trypsin activity on batch J6B is further supported by a previous study, when no significant difference in the activity of an ultra-filtrated Cry1Ab protein batch could be observed during 4.5 months of storage at 4 °C (Nguyen et al., 2004).

The bioactivity of the Cry1Ab batches H3 and H4 decreased significantly by a factor of 8 to 11-fold when stored at –20 °C for two years (Tab. 2). In contrast, when stored at –80 °C they remained relatively stable during the same period of time. The observed difference in LC<sub>50</sub>

values was up to 3.5-fold. This could be caused by the variability in the response of the larvae and not by a decline of Cry1Ab bioactivity. An up to 3-fold variability of the LC<sub>50</sub> was observed when the same Cry1Ab batch had been used in repeated bioassays during one week (Nguyen et al., 2004).

The outcome of the bioassays clearly indicated that long-term storage of Cry1Ab protein at –80 °C can preserve the bioactivity better than storage at 4 °C and –20 °C. The reason why both Cry1Ab batches H3 and H4 lost some of their bioactivity during storage at –20 °C but remained stable at –80 °C is not fully clear. Normally, the protein must fold into compact structures with few exposed loops, in order to avoid excessive proteolysis in the insect gut (Bosch et al., 1994). We assume that storage at –20 °C and at –80 °C caused slightly differing folding structures of Cry1Ab proteins, which could not be detected by SDS-PAGE and Bradford assay but may have increased the susceptibility of Cry1Ab to protease digestion in the insect gut and thus resulted in a decreased activity in the bioassay. A considerable variation in the bioactivity of the two Cry1Ab batches H3 and H4 was observed. This is in agreement with previous studies, where different Cry1Ab sources resulted in a four- to eight-fold



**Figure 3.** SDS polyacrylamide gel electrophoresis (10% gel) of Cry1Ab batch H3 and batch H4 stored at  $-80^{\circ}\text{C}$  and  $-20^{\circ}\text{C}$  after 0, 6, 12, and 24 months. (M = PageRuler™ protein ladder (Fermentas) with molecular masses (kDa) given to the right.)

variation in their activity (Nguyen et al., 2004; Saeglitz et al., 2006). These batch-to-batch differences could be caused by different treatments during production of these batches and/or differences in the concentration of Cry1Ab depending on the protein determination methods. Crespo et al. (2008) noted that different protein batches should be produced and measured with standardized methods in order to eliminate the difference in their activity. When using the same standardized production and quantification methods, bioassay of three different Cry1Ab batches produced in *E. coli* and *B. thuringiensis* showed no significant difference in  $\text{LC}_{50}$  values (Crespo et al., 2008).

Our study indicates that the choice of the storage temperature is a crucial factor concerning the bioactivity of Cry1Ab protein in long-term monitoring the

susceptibility of a target insect. When using soluble protein batches, we recommend storing the Cry1Ab protein at  $-80^{\circ}\text{C}$ . On the other hand, when using the same batch of Cry1Ab, intrinsic variations of the  $\text{LC}_{50}$ s by a factor of 3–5 were observed in this and in a previous study (Nguyen et al., 2004). Due to this variability of insect response from bioassay to bioassay, a beginning shift towards resistance of a target ECB population measured as change of  $\text{LC}_{50}$  cannot be statistically substantiated, unless the susceptibility changed by a factor of 5 to 10. This natural variability of bioassays has also to be considered, when resistance testing is performed at a discriminative dose, e.g. at a  $\text{LC}_{95}$  or  $\text{LC}_{98}$  (Roush and Miller, 1986). Thus, initial changes in the susceptibility of an ECB population will be hardly detected. Nevertheless, the standardization of production and storage of Cry1Ab protein for long-term monitoring keep the variability of bioassays to a minimum and are a prerequisite of any successful resistance monitoring scheme.

## MATERIALS AND METHODS

### Preparation and storage of Cry1Ab protein

*E. coli* XL1-Blue containing the plasmid pBD140 carrying the *cry1Ab* gene was kindly provided by Ruud de Maagd (Wageningen, The Netherlands) (De Maagd et al., 1996). An overnight preculture was grown at  $37^{\circ}\text{C}$  with vigorous shaking (220 rpm) in 5 mL of TB medium (12 g bacto-tryptone, 24 g bacto-yeast extract, 4 mL glycerol, 2.31 g  $\text{KH}_2\text{PO}_4$ , and 12.54 g  $\text{K}_2\text{HPO}_4$  per liter) complemented with 2% glucose and ampicillin ( $100\ \mu\text{g}\cdot\text{mL}^{-1}$ ). This preculture was transferred to 500 mL of the TB medium containing ampicillin ( $100\ \mu\text{g}\cdot\text{mL}^{-1}$ ) in a 2-L Erlenmeyer flask and grown at  $28^{\circ}\text{C}$  with shaking (220 rpm). After 3 days of incubation, the cells were harvested by centrifugation at  $5000\times g$  for 10 min at  $4^{\circ}\text{C}$  using a SLA-1500 rotor (Sorvall RC5B Plus), stored at  $-20^{\circ}\text{C}$  and then purified as described in Höss et al. (2007). Briefly, inclusion bodies (IBs) were washed several times in washing buffer (Tris/HCl, NaCl, Triton X-100), then in phosphate buffer saline (PBS), and then in double-distilled water. Finally, the IBs were solubilized in 50 mM cyclohexyl aminopropene sulfonic acid (CAPS) buffer (pH 10.5). The solubilized Cry1Ab protoxin (130 kDa) was trypsinized with L-1-tosylamido-2-phenylethyl chloromethyl ketone (TPCK)-treated bovine pancreas trypsin ( $1\ \text{mg}\cdot\text{mL}^{-1}$ ) in order to obtain activated Cry1Ab protein ( $\sim 60\ \text{kDa}$ ). The trypsinized Cry1Ab protein samples were ultrafiltered by using polyethersulfone membrane (Millipore, NMWL: 50 000). In total, three batches of Cry1Ab protein, J6B, H3, and H4 were produced independently. Protein batch J6B was the first produced and stored at  $4^{\circ}\text{C}$ . In order to examine the stability

at  $-20\text{ }^{\circ}\text{C}$  and  $-80\text{ }^{\circ}\text{C}$  storage, two protein batches, H3 and H4, were produced subsequently. The full-length protoxin of batch H3 was treated with trypsin twice in order to obtain the trypsin-resistant core protein. Cry1Ab protein batch H4 was produced using the optimized protocols obtained during production of batch H3. Each Cry1Ab batch was aliquoted and stored under the conditions described in the different experiments. The samples were aliquoted to avoid multiple freezing and thawing and to prevent protein degradation. To test stability, samples were diluted to  $400\text{ }\mu\text{g}\cdot\text{mL}^{-1}$ , aliquoted ( $30\text{ }\mu\text{L}$  each) and stored at  $4\text{ }^{\circ}\text{C}$ ,  $-20\text{ }^{\circ}\text{C}$  and  $-80\text{ }^{\circ}\text{C}$ . Every six months one aliquot of different protein batches was taken out and treated with 0.25 volumes of 4 X sample buffer containing 40 mM Tris/HCl, 4 mM EDTA, 8% (w/v) SDS, 40% (w/v) glycerol, 0.004% bromophenol-blue (w/v) and 0.1 volumes of  $\beta$ -mercaptoethanol. The mixture was heated to  $100\text{ }^{\circ}\text{C}$  for 10 min and stored again at  $-20\text{ }^{\circ}\text{C}$  or  $-80\text{ }^{\circ}\text{C}$ .

### Protein determination

Total protein concentration of each batch was determined using Bradford dye assay (Bio-Rad Laboratories GmbH, Munich, Germany) (Bradford, 1976). Bovine serum albumin (BSA, Fraktion V, Pierce) with concentrations of 100, 200, 400, 600, 800, 1000 and  $1200\text{ }\mu\text{g}\cdot\text{mL}^{-1}$  was used as quantification standard.

### SDS polyacrylamide gel electrophoresis (SDS PAGE)

SDS PAGE was performed following the procedure of Laemmli (Laemmli, 1970). Scan of the SDS gels were analysed using ImageMaster 1D Software (Pharmacia System) to estimate the purity level of the protein.

### Insect bioassays

An ECB laboratory strain was obtained from the Julius Kuehn Institute, Institute for Biological Control, Darmstadt (Germany), and has been reared at the DLR Rheinpfalz since 2005. Bioassays were performed with first-instar ECB larvae at an age of 3–24 h after hatching. A volume of 1 mL of artificial diet was dispersed in each well (surface area of  $1.77 \pm 0.08\text{ cm}^2$  per well) of the bioassay tray (Bio-ba-128, Color-Dec Italy). Cry1Ab protein was diluted in 50 mM CAPS buffer (pH 10.5) to six different concentrations and pipetted on the surface of the diet and air dried. The concentrations of Cry1Ab protein applied in the bioassay depended on the different batches, but generally ranged between 3–1000  $\text{ng}\cdot\text{cm}^{-2}$ .

For each dilution, 16 larvae were assayed individually in wells. In untreated controls, 32 larvae were exposed to 50 mM CAPS buffer (pH 10.5) instead of Cry1Ab protein. Each bioassay was conducted at  $22\text{ }^{\circ}\text{C}$  and the mortality was assessed after 7 days. The assays were independently repeated 3–5 times.

Bioactivity was measured during 16 weeks for batch J6B stored at  $4\text{ }^{\circ}\text{C}$  and approximately every half year for batches H3 and H4 stored at  $-20\text{ }^{\circ}\text{C}$  and  $-80\text{ }^{\circ}\text{C}$ , respectively. The detailed period between different bioassays is described in the results.

### Statistical analysis

After correction for control mortality (Abbott, 1925), probit analysis implemented in SAS (SAS Institute, version 9.1.3, 2007) was applied to estimate the median lethal concentrations ( $\text{LC}_{50}$ ) and the slopes of the concentration-mortality lines. Significance testing of the  $\text{LC}_{50}$  was done by pairwise comparisons using 95% fiducial limits (FLs) ( $\alpha \leq 0.05$ ) (Smith, 1997). Parallelism of slopes of two lines was also tested (Unkelbach and Wolf, 1985).

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