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Expression, purification and bioassay of Cry55Aa protein against tomato root knot nematode, *Meloidogyne incognita*

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Abstract

A nematicidal Cry55Aa protein was expressed and purified from *E. coli* cells. Higher level of expression was observed in culture induced with 0.1mM of IPTG after four hours of induction. The Cry55Aa protein was purified by resuspending the crude protein in solubilizing buffer at pH 12 for 120 min. When *Meloidogyne incognita* was tested against different concentration of purified protein, a clear dose response lethality was observed with an LC₅₀ value of 31.4 µg/ml.

Keywords: Cry55Aa protein, expression, purification, bioassay, *M. incognita*

Introduction

Bacillus thuringiensis (*Bt*), a gram-positive soil bacterium, produces crystalline inclusions during sporulation, having insecticidal proteins called δ-endotoxins. Different *Bt* strains produce a large number of insecticidal crystal proteins, which are encoded by *cry* genes and these strains are used as bio-pesticides for several decades^[1]. Crystal proteins are known to be toxic against different insect orders such as Lepidoptera (butterflies and moths), Coleoptera (beetles) and Diptera^[2], as well as nematodes^[3]. Cry proteins such as Cry5, Cry6, Cry12, Cry13, Cry14, Cry21 and Cry55 are reported as nematicidal^[4] and found to be toxic against several nematode genera including *Criconebella*, *Globodera*, *Ditylenchus*, *Heterodera*, *Meloidiogyne*, *Helicotylenchus*, *Pratylenchus*, *Rotylenchus*, *Radopholus*, *Tylenchus* and *Bursaphalenchus*^[5]. *B. thuringiensis* strains produce more than one type of toxic proteins (*cry*, *vip*, *cyt* etc.) and to confirm the toxicity of a particular Cry protein, the respective *cry* gene is cloned and expressed in a heterologous bacteria like *E. coli* and used in bioassay studies^[6]. *E. coli* is one of the most attractive hosts^[7] for expression of heterologous protein with the advantages of fast growth in an inexpensive medium and the availability of a large number of cloning vectors^[8]. In this study, an attempt was made to optimize heterologous protein expression in *E. coli*.

The plasmid pET29a containing *cry55Aa* gene under the control of a T7 promoter was used for heterologous expression of Cry55Aa protein. In order to increase the yield of the Cry55Aa protein, different concentrations of IPTG were used with different duration of induction. Expressed Cry55Aa protein was purified by using solubilizing buffer with different pH levels and incubation time. The purified Cry55Aa protein was tested against tomato root knot nematode, *M. incognita*.

Materials and Methods**Gene construct**

The recombinant *E. coli* strain BL21 harbouring pET29a-*cry55Aa* gene (Accession no: HG764207.1) which was originally cloned from indigenous *Bacillus thuringiensis* isolate, T44 of *Bt* collections of CPMB & B, TNAU.

Expression of Cry55Aa in *E. coli* BL21 cell

Two *E. coli* cultures *viz.* BL21 (DE3) harbouring vector pET29a with *cry55Aa* insert and BL21 carrying pET29a vector without insert were grown in LB broth (5 ml) containing kanamycin 50 mg/l overnight at 37 °C. About 500 µl of overnight-grown culture was inoculated in 25 ml of fresh LB broth with kanamycin 50 mg/l and incubated at 37°C in orbital shaker (180 rpm) till OD₆₀₀ value reached ~0.6. Then different concentration of IPTG *viz.*, 0.01, 0.05, 0.10, 0.50 and 1.0 mM was added to induce the protein expression and the cells were further grown at 37 °C for different durations (1, 2, 3, 4, 5 and 6 hrs) with a view to

optimize the IPTG concentration and duration of induction. The cultures were harvested by centrifugation and washed with 1X TE buffer (10.0 mM Tris-Cl, 1.0 mM EDTA, pH 8). The pellet was dissolved in 5 ml of TE buffer containing 1 mM phenyl methyl sulfonyl fluoride and sonicated by using ultra-sonic liquid processor (Sonics and Material Inc., USA). Sonication was done with off pulsar mode for 4 min at 20 amplitude (4x 1 min with a time interval of 1 min.). The broken cells were pelleted by centrifugation at 7000 rpm for 15 min. The pellet was dissolved in the TE buffer and washed twice in the same buffer. The final product was suspended in 200 μ l of sterile double distilled water containing 2mM PMSF and an aliquot of 5 μ l was used for separation in SDS-PAGE.

Crystal protein purification

The crude Cry55Aa protein was collected by centrifugation and resuspended with solubilization buffer (8.7 mM tripotassium citrate, 43.4 mM citric acid and 10 mM dithiothreitol) at different pH levels (pH 9.0, 10.0, 11.0 and 12.0) and incubated at 37°C for different durations (30, 60, 90 and 120 minutes) to solubilise the Cry55Aa protein. The contents were spun out at 8,000 rpm for 10 min and protein was precipitated from the supernatant by adding 1 M tripotassium citrate to adjust pH to 5.0 and incubated in ice overnight. The Cry55Aa protein was harvested by centrifugation for 10 min at 12,000 rpm. The precipitated Cry55Aa protein was washed with ddH₂O, dissolved in 20 mM 4-(2-hydroxyethyl)-1-piperazine ethane sulfonic acid (HEPES) (pH 8.0) and resolved in SDS-PAGE. The sds-page was performed as per the protocol given by Laemmli (1970) [9].

Lethal activity assay against *M. incognita*

Initial *M. incognita* culture was collected from infested tomato fields from Coimbatore, Tamil Nadu and multiplied in potted tomato plants maintained in greenhouse condition at 25 \pm 2°C. The J2 stage of *M. incognita* were hatched on a filter paper which was saturated with water at 20-25 °C and used to test the toxicity of the Cry55Aa toxin and the assay was performed in 24 well plates. Each well with 400 μ l S medium [10] contained purified Cry55Aa protein at 10, 20, 30, 40 and 50 μ g/ml, about 100 J2 worms and 1 μ l of chloramphenicol (10 mg/ml). BSA (20 μ g/ml) was used as the control and each treatment was replicated four times. After incubation of 5 days at 25 °C, 50 μ l culture from each well was transferred onto glass slides and observation made on live and dead worms under a dissecting microscope. Observation were made on at least three samples of 50 μ l from each well.

Results and Discussion

Optimization of protein expression by IPTG induction

SDS-PAGE results showed that the protein expression increased gradually with increase in concentration of IPTG and maximum expression was observed with 0.1mM IPTG (Plate 1). The protein from recombinant *E. coli* BL21 (harbouring pET29a+cry55Aa gene) showed an expected prominent band of ~45 kDa on SDS-PAGE, which confirmed the cry55Aa gene expression, whereas no band was noticed in protein isolated from *E. coli* BL21 (harbouring pET29a vector alone). When IPTG (0.1 mM) induction was done for different durations (1, 2, 3, 4, 5 and 6 hrs) to check the level of expression, higher level of expression of protein was observed in cultures after four hours of induction (Plate 2). The present findings are comparable with previous report [11] suggesting that 0.1 mM IPTG was enough to induce the

protein expression [12]. An IPTG concentration of 0.1mM and an induction period for four hours were found to be optimum for Cry55Aa expression in *E. coli* BL21 cells. Conditions for recombinant protein expression appears to be protein dependent [13]. For instance, induction with IPTG 10 μ M was appropriate to induce the cHSPA6 expression which was 100 times less than generally used concentration and 5 h of post-induction incubation period was found to be better to produce folded cHSPA6 [14].

Purification of Cry55Aa protein by alkali solubilization

Solubility of crystal protein was found to have significant effect on toxicity [15]. Proteolytic activation of protoxin, is a crucial step in Cry protein toxicity mechanism, and found to be reliant on crystal protein dissolution in host intestine [16]. Some types of crystal protein from *B. thuringiensis* were found to be soluble at pH 12, making it nontoxic to insect with gut pH of 9.0 [17]. However, when crystal protein are presolubilized at high pH, these nontoxic crystal proteins exhibit significant toxicity to kill specific insect.

In this study, purification of Cry55Aa protein from recombinant *E. coli* (BL21) using solubilization buffer with different pH (9, 10, 11 and 12) and duration of induction (30, 60, 90 and 120 min) was attempted. Cry55Aa protein solubilization was maximum with pH 12, showing an expected prominent band of ~45 kDa on SDS-PAGE. When solubilization was done for different duration of incubation (30, 60, 90 and 120 minutes), solubilization was found to increase gradually from 30 to 120 minutes (Plate 3). The results are comparable with earlier reports on Cry5B (>pH 10), Cyt (pH 9.5), Cry4B (pH 12) [18, 19] and Cry6Aa2 (pH 9.5) [20].

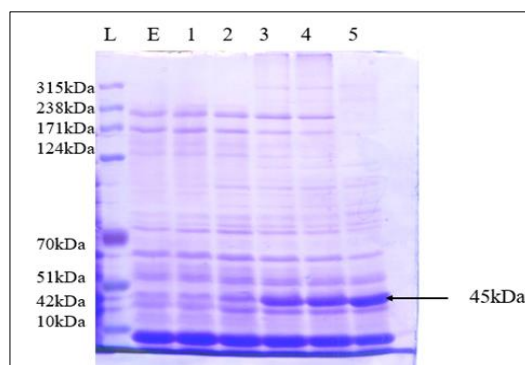


Plate 1: Standardization of IPTG concentration for Cry55Aa protein expression (L- Protein ladder (10-315kDa), E-empty vector (pET29a), 1 to 5- pET29a with Cry55Aa induced with IPTG 0.01, 0.05, 0.10, 0.50 and 1.0Mm)

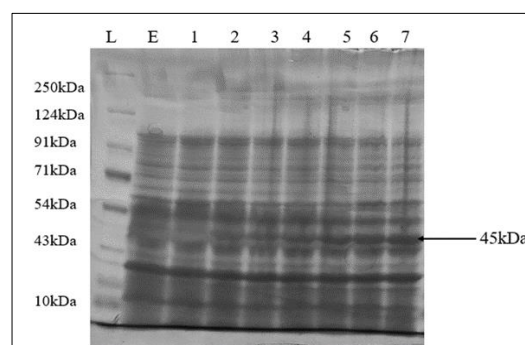


Plate 2: Standardization of induction period for Cry55Aa protein expression (L- Protein ladder (10-250kDa), E-empty vector (pET29a), 1 to 7- pET29a with Cry55Aa induced with 0.1mM IPTG for 0, 1, 2, 3, 4, 5 and 6 hrs)

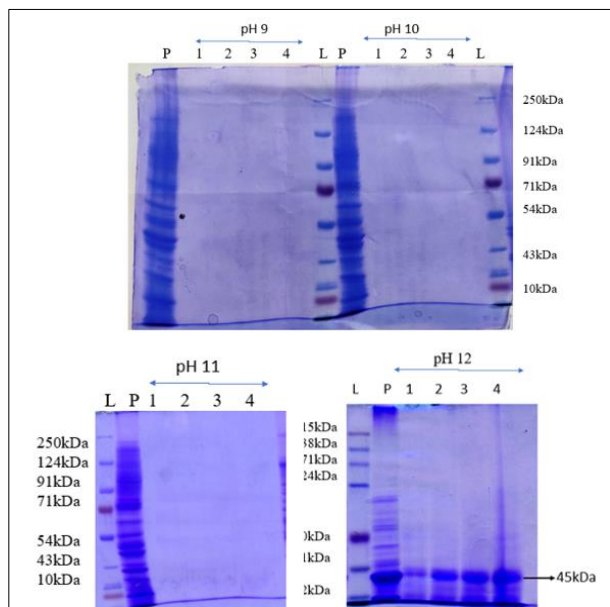


Plate 3: Purification of Cry55Aa protein from recombinant *E. coli* (L- Protein ladder, P- Protein without solubilization; 1 to 4- in each gel represent 30, 60, 90 and 120 minutes of incubation at 37 °C)

Lethal activity of Cry55Aa protein on *M. incognita*

The per cent corrected mortality of Cry55Aa protein against *M. incognita* was 14.01, 30.61, 40.94, 48.51 and 58.81 at concentrations of 10, 20, 30, 40 and 50 µg/ml respectively (Table 1). Calculated LC₅₀ value for Cry55Aa protein against second juvenile stage of *M. incognita* was 31.4 µg/ml (162.66–1109.62 µg/ml for 95% fiducial limits determined by profit analysis). The results are comparable with estimated LC₅₀ values of Cry55Aa1 protein which was 23.2 µg/ml [21] against *M. hapla* as. In addition, the combination of Cry6Aa and Cry55Aa was reported to have toxicity on *M. incognita* [22]. The LC₅₀ value of crude protein from *Bt* strain YBT-021 against *Tylenchorhynchus sp.*, *M. hapla*, *Pratylenchus scribneri*, *Ditylenchus destructor* and *Aphelenchoides sp.* was about 94.31 µg/ml, 35.62 µg/ml, 75.65 µg/ml, 215.21 µg/ml and 128.76 µg/ml respectively [23].

In the present study, have shown that the *cry55Aa* gene could be expressed in *E. coli* by inducing BL21-DE3 cells with 0.1 mM IPTG. The results presented here demonstrate that the Cry55Aa protein is toxic to *M. incognita*. This Cry55Aa is a potential nematocidal protein and could be included as a component of IPM, either as a formulated product or through transgenic means.

Table 1: Lethal activity assay of purified Cry55Aa protein against *M. incognita*

Treatment		Number of nematodes		Per cent mortality	Per cent corrected mortality	LC ₅₀	Fiducial limit LC ₉₅	
		Released	Dead				Lower	Upper
Cry55Aa protein (ug)	10	97.38	23.32	23.95 (29.29) ^c	14.01	31.4	162.66	1109.62
	20	89.94	34.64	38.62 (38.40) ^b	30.61			
	30	87.93	41.98	47.76 (43.71) ^b	40.94			
	40	88.52	48.28	54.46 (51.46) ^a	48.51			
	50	87.25	55.50	63.57 (52.98) ^a	58.81			
Control		91.96	10.64	11.55 (19.80) ^d				
CD (P=0.5)				7.20				

Mean of four replications. Values in parentheses are arcsine transformed. Means in a column followed by same superscripts are not significantly different at $P \leq 0.05$

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