THE IMPACT OF INSERTING RNAI SUPPRESSOR GENE ON BACULOVIRUS PATHOGENICITY.

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B. Sc. Agric. Sc. (Genetics), Ain Shams University, 2004.

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ABSTRACT

Shimaa Mohamed Mohamed El-Sayed El-Gamal: The Impact of Inserting RNAi Suppressor Gene on Baculovirus Pathogenicity. Unpublished M.Sc. Thesis, Department of Genetics, Faculty of Agriculture, Ain Shams University, 2012.

Baculoviruses are natural enemies of pest insects but they have narrow host range. Recently, RNA interference (RNAi) confirmed to function as a natural defense mechanism against viruses. Too many plant and mammalian viruses encode RNAi suppressors to counteract this antiviral mechanism and enhance viral infectivity in host cells. Few studies have been done to illustrate the role of RNAi as an antiviral mechanism against DNA viruses. The influence of RNAi on baculovirus pathogenicity or replication has not been intensively studied. In the present study, the B2 gene of the flock house virus (FHV), which known as RNAi suppressor was synthesized and a recombinant baculovirus expressing it under polyhedrin promoter was produced. The mortality of this recombinant virus to S. littoralis larvae was compared with the mortality of a control virus. The data showed significant differences between both viruses in killing larvae, but it need three to four weeks post infection. Moreover, using the end point dilution method showed that the B2 enhanced the production of AcMNPV in Sf9 cells. The effect of B2 on viral replication needs to be studied in details.

Key Words:

Baculovirus, RNA interference (RNAi), RNAi suppressor, B2, Flock house virus (FHV), *Spodoptera littoralis*.

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LIST OF ABBREVIATION

AaIT Androctonus australis insect selective toxin

a.a. amino acid

AcMNPV *Autographa californica* nucleopolyhedrosis virus AGERI Agriculture genetic engineering research institute

Ago2 Argonaute
Anti antisense

ATP Adenosine triphosphate

B.P.B. Bromophenol blue

BEVS baculovirus expression vector system

Bt Bacillus thuringiensis

BV Budded virus

C. elegance Caenorhabditis elegansCrPV cricket paralysis virusCTV Citrus tristeza virusdsRNA double stranded RNA

DTT Dithiothreitol

E. coli Escherichia coli

EDTA Ethylenediaminetetraacetic acid EGFP enhanced green fluorescent Protein

egt ecdsteroid UDB-glucosyltransferase gene

Fig. Figure

FHV Flock house virus
GP64 Glycoprotein 64
GVs Granuloviruses

h.p.i. Hour post infection

HaSNPV Helicoverpa armigera nuclear polyhedrosis virus

Hc-pro helper component protein

HzSNPV Heliothis zea nuclear polyhedrosis virus

IE-0 immediate early - 0

IPTG Isopropyl-beta-thio galactopyranoside

LacZ Lactose Z gene

LB Luria Broth media

miRNAs microRNAs

MOI multiplicity of infection

NoV Nodavirus

NPVs nucleopolyhedrosis viruses

nt nucleotide

NTC no template control

O.D. Optical density

ODV occluded derived virus PCR polymerase chain reaction

PD proportionate distance

Pfu taq proof reading Taq DNA polymerase enzyme

Pfu/ml plaque forming unite/ml.
pIBs polyhedral inclusion bodies

Polh Polyhedrin

Pre-miRNAs precursor microRNAs

PTGS post transcriptional gene silencing

Rec. recombinant

RISC RNA induced silencing complex

RNAi RNA interference rpm round per minute S. littoralis Spodoptera littoralis

Schneider line 2

SDS sodium dodecyl sulfate

Sf9/Sf21 Spodoptera frugiperda 9/21

siRNAs small interfering RNAs

ss single stranded

TAE Tris acetate EDTA solution
TCID₅₀ tissue culture infectious dose

TEV Tobacco etch virus WhNV Wuhan Nodavirus

X-gal 5-bromo-4-chloro-indolyl- β -D-galactopyranoside

I. INTRODUCTION

Baculoviruses consider one of the most prominent natural enemies of pest insects. They are host specific and have narrow host range. Moreover, they are safe to human, plants, animals and environment. However, the major impediments on using them as biopesticides include low infectivity in some cases and slow killing speed. Recombinant viruses were the key to solve these problems. Too many recombinant viruses are produced using insect specific toxins to speed up their killing time (Stwart et al., 1991 and Tomalski and Miller 1991). Other genes have been inserted to or deleted from baculovirus genome to improve baculoviruses as biopesticides (O'Reilly and Miller 1991and Inceoglu et al., 2001).

One of the methods that has not been examined before is whether or not RNA Interference (RNAi) has effect on baculovirus replication or infectivity.

RNAi machinery (Fire et al., 1998) occurs in most of eukaryotic organisms and has been described as Post-transcriptional gene silencing (PTGS) in plants and quelling in fungi (Pandit and Russo, 1992 and Romano and Macino, 1992). This mechanism can degrade viral and transgene RNAs in a sequence specific manner. RNAi machinery is triggered by the presence of double stranded RNAs (dsRNAs) in most of the eukaryotic cells. This mechanism functions as a natural defense mechanism against viruses; therefore too many plant and mammalian viruses encode RNAi suppressors to counteract this antiviral mechanism in host cells (Delgadillo et al., 2004). The researchers found that some of these suppressors have other functions during virus infection such as B2 suppressor of Nodamura virus, which can increase viral RNA accumulation in both vertebrate and invertebrate cell lines (Johnson et al., 2004). In addition, B2 protein of Flock house virus enables virus infection in Drosophila cells (Li et al., 2002). Moreover, nonstructural

protein NSs of tomato spotted wilt virus enhances the Baculovirus replication in permissive and semipermissive insect cells. (Oliveira et al., 2011).

It is important to study the impact of RNAi suppressor protein isolated from insect virus on DNA virus pathogenicity, which may uncover the Role of RNAi machinery against DNA viruses as well as RNA viruses. In addition, more understanding for virus host interaction can be obtained from this study.

The main aim of this study is to synthesize the B2 gene of the FHV and express it under polyhedrin promoter to study if RNAi plays role in baculovirus infectivity.