Cloning and Expression of a Cotesia plutellae Bracovirus Gene, ELP

Keewoo Lee¹, Sungwhan Cho², Hyuksoo Lee¹ and Yonggyun Kim²

¹Department of Bioelectronic Engineering, College of Engineering, Andong National University ²School of Bioresource Sciences, College of Natural Sciences, Andong National University

A genome project has been launched and aims to sequence total genome of *Cotesia plutellae* bracovirus. On this process, several open reading frames (ORFs) have been identified. This study was intended to clone and express an EP1-like protein (ELP) gene, called 'ELP'. ELP ORF consists of 729 bp encoded 242 amino acid sequence. Its estimated protein size is 30,000 Da. A reverse transcriptase-polymerase chain reaction (RT-PCR) produced the corresponding product from cDNA of parasitized diamondback moth, *Plutella xylostella*. This PCR project was cloned into pBAD-TOPO (Invitrogen®) and over-expressed in response to an inducer, L-arabinose. The bacterial cell extract was subjected to Ni⁺-affinity column. The purified protein was confirmed by an immunoblotting with V5 antibody and its apparent molecular weight was about 32 kDa. ELP expression in parasitized *Plutella xylostella* was measured by real-time RT-PCR and indicate late expression pattern during parasitization.

Key words: *Plutella xylostella*, *Cotesia plutellae*, CpBV, Polydnavirus, RT-PCR, expression, real-time RT-PCR