

**EFFECTS OF SOME PLANTS SEED EXTRACTS ON  
*HELICOVERPA ARMIGERA* HÜBNER (LEPIDOPTERA:  
NOCTUIDAE) MIDGUT PROTEASE ACTIVITY**

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**ABSTRACT:** *Helicoverpa armigera* is one of the most important pests of crops, such as cotton, cereals and vegetables. Using plant derived enzyme inhibitors in transgenic plants is one of the safe methods in IPM programs. In this study protease inhibitory activity of some plants seed extracts from poaceae and fabaceae family were studied. Insects reared in controlled condition and the last larval instars alimentary canal were used in enzymatic assays. The crude seed extracts were subjected for ammonium sulfate precipitation. The seed extracts of six plants were fractionated into four fractions (0-30 %, 30-50, 50-70 and 70-80%). The percentage of inhibition to cotton bollworm midgut protease activity obtained in crude and each protein fraction of ammonium sulfate. Proteolytic activity of midgut enzyme extracts was evaluated using the azocasein as substrate. The results revealed that seed extracts of *Phaseolus vulgaris* and *Cicer arietinum* are potentially effective in inhibiting the proteolytic activity of cotton bollworm (54.5 and 53.2% respectively). Also total extracts of *Triticum aestivum*, *Hordeum vulgare*, *Zea mays*, and *sophora alopecuroides* inhibited HGP activity by 17.7, 18.74, 20.62 and 29.31% respectively. Results revealed that the F1 fraction protein of all studied plants showed less than 20% inhibitory activity against HGP, and the F2 and F3 fraction exhibited the same inhibitory activity in the range of 10-20% in poaceae species. The legume plants especially in F1, F2 and F3 fractions exhibited near 15-50% inhibitory activity on HGP. In over all, among studied plants, *Z. mays*, *P. vulgaris* and *C. arietinum* have strong inhibitory activity in compare with others.

**KEY WORDS:** Inhibitory activity, poacea, fabacea, azocasein, ammonium sulfate, cotton bollworm

Cotton bollworm, *Helicoverpa armigera* (Hübner), is one of the most important pests worldwide. It is polyphagous insect with a wide range of host plants including cultivated and wild plants such as cotton, mays, chickpea, tomato, vegetables, and other crops (Harsulkar et al., 1999; Nair et al., 2013). The wide application of chemical insecticides has been the main strategy for the control of *H. armigera* in different parts of the world. High levels of resistance to conventional insecticides also harmful effects on the environment and human health were developed due to improper use of insecticides (Giri et al., 1998; Parde et al., 2010).

Pests such as cotton bollworm rely on a proteinases enzymes present in their guts to digest protease present in seeds, leaves and flowers of host plants (Shukla et al., 2005; Mohammadi et al., 2010). Insect pests depend on peptides obtained from proteolytic digestion for growth and development. Disruption in an insect's ability to digest protein by transgenic plants expressing proteinase inhibitors, seems to be an alternative approach to conventional insecticides (Fan & Wu, 2005).

Proteinase inhibitors are generally small proteins (less than 20 kDa) that mainly have been identified in storage tissues, such as tubers, seeds and aerial parts of plants such as flowers and leaves (Grover et al., 2014; Ryan, 2006). They

are also inducible in plants in response to attack by herbivores (Ryan, 2006). Protease inhibitors have been isolated and characterized from a large number of organisms, including plants, animals, and microorganisms (Christeller, 2005; Nair et al., 2013). A useful strategy for enhancing plant defense systems is to identify PIs with high activity against the target pests.

Protease inhibitors (PIs) are compounds that form complexes with proteases and inhibit their proteolytic activity and suppression of the normal assimilation of food proteins (Ryan, 1990; Fan & Wu, 2005). Plants utilize proteinase inhibitors in order to moderate the adverse effects from attacking herbivores.

So many studies have been carried on protease inhibitors, which active against different insect species, both in *in vitro* and in *in vivo* (Dorrah, 2004; Shukla, 2005).

Reduction in fecundity and fertility, reducing larval growth and development and delay in pupation period after feeding of *C. annum* leaf extracts to *H. armigera* larvae through artificial diet has reported by Tamhane et al. (2005). Also they observed that about 91-98% of protease activity of *H. armigera* gut protease was inhibited by extracts of *C. annum*.

Protein proteinase inhibitors extracted from the seeds of *Momordica charantia* L. were identified as effective inhibitor of cotton bollworm gut proteinases (Telong et al., 2003).

Five plants including *Arachis hypogaea*, *Vigna sinensis*, *Dolichos lablab*, *Phaseolus aureus* and *Cassia siamea* reported inhibitory active plants with 22.91 to 58.33 % inhibition against *H. armigera* protease activity (Padul, 2012).

Serine protease activities of *S. littoralis* midgut were inhibited by soybean trypsin inhibitor *in vitro*. Consumption of SBI by the larvae, causes variable effects such as reduction in weight gain and survival of the larvae (Dorrah, 2004).

Partially purified inhibitor from soybean seeds extracts inhibiting cotton bollworm total protease activity by 91%. While inhibition of trypsin and chymotrypsin like proteases were found near 65 and 40% respectively (Ghodke, 2013).

In a study aimed to test the efficacy of pigeonpea genotypes against *H. armigera* development were observed that, insects fed with diet containing seed powder exhibited larval and pupal weight reduction. also certain abnormalities such as larval-pupal intermediates were reported (Grover, 2014). Grover et al. (2014) reported that cotton bollworm fed with diet containing pigeon pea seed powder exhibited larval and pupal weight reduction and certain abnormalities.

Some studies about screening of host and non-host plant-derived inhibitors has resulted in effective proteins that demonstrated high levels of inhibitory and biological effects against various insects. Some researchers reported that non-host plant PIs showed more inhibitory activity than host plants (Gruden et al., 1998; Harsulkar et al., 1999; Jamal et al., 2013). Parde et al. (2010) in their studies reported that, *in vivo* studies indicated that non-host plant PIs were good candidates as inhibitors of the HaGPs. The PIs from the non-host plants can be expressed in transgenic plants to confer resistance to insect pests.

Most of plants proteinase inhibitors that have been characterized are from the poaceae, fabaceae, and solanaceae families.

Usually seed extracts of plants showed more inhibitory activity than leaf and flower tissues and this may be due to higher accumulation of proteins in seeds than leaves and flowers (Qutchkourov et al., 2003; Harrison et al., 2012; Chougule et al., 2005). Currently, the main emphasis of plant proteinase inhibitor studies is on identifying potential inhibitors of digestive proteinases of the target insects and present study was conducted to evaluate the *in vitro* assays of the some poaceae and fabaceae family crude seed extracts and partial purified fractions proteins against the cotton bollworm gut proteases activity.

## MATERIALS AND METHODS

### *Insect rearing*

Cotton bollworm larvae were provided by a colony in plant protection department of Tabriz University. Larvae were reared on artificial diets based on cowpea (Shorey & Hall, 1965) in controlled condition of  $26\pm 2^{\circ}\text{C}$ ,  $50\pm 5\%$  relative humidity and a photoperiod of 16:8 (L: D) h.

### *Enzyme preparation*

One day old last-larval instars of cotton bollworm were selected for gut extraction. The individuals were chilled and dissected in cold petri dishes. Each gut with lumen contents was moved to 1.5 ml micro tubes containing 1 ml cold Glycine-NaOH, pH 10, buffer. The collected guts were then homogenized using Ultra turrax T8 homogenizer then centrifuged for 10 minutes at 10000 rpm and  $4^{\circ}\text{C}$ . The supernatant was used as enzyme solution (Mohammadi et al., 2015).

### *Crude extracts and protein fractions preparing of Inhibitors*

Plant seeds were prewashed with distilled water and dried in room temperature then ground using mortar and pestle. The prepared flour was soaked in Glycine-NaOH buffer, pH 10.0 for 90 minutes in  $6^{\circ}\text{C}$ . The homogenates were centrifuged at 10000 rpm for 30 minutes at  $4^{\circ}\text{C}$ . The proteins collected from the supernatant were used for protease inhibition assay (Baker, 1987; Melo et al., 1999).

For partial purification of proteins, crude extracts obtained from the seeds of studied plants were precipitated at 0-30, 30-50, 50-70 and 70-80% saturation with ammonium sulfate and four protein fractions (F1 – F4) were obtained (Mohammadi et al., 2010).

### *Determination of protein concentration*

Protein concentration was estimated by the method of Bradford (1976) using bovine serum albumin (BSA) as the standard.

### *Enzyme activity and inhibitory assays*

Total protease activity determined using azocaseinolytic assay. Azocasein at final concentration of 1% (w/v) was incubated with the enzyme fraction in Glycine-NaOH 200 mM buffer, pH 10, containing 5 mM  $\text{CaCl}_2$ , at  $37^{\circ}\text{C}$  for 60 min. The reaction was terminated by the addition of 300  $\mu\text{l}$  of TCA (10% v/v) and the sample was centrifuged for 10 min at 10000 rpm. The supernatant was added to 1M NaOH in equal volumes. And the absorbance of the supernatant was read at 450 nm. Protease and inhibitory activity was defined as the amount of enzyme that increased the absorbance by 1.0 OD under the given assay condition.

For the inhibitory assays, a suitable volume of seed extract was added to the gut proteinase extract and incubated at room temperature ( $27^{\circ}\text{C}$ ) for 15 min. The residual proteinase activity was then estimated for every assay.

### *Statistical analysis*

Statistical analyses were done using SPSS 15 software. The effects of PIs from plant materials on *H. armigera* last larval instars. enzyme activity analysed using one way-ANOVA. When a significant effect was found the Duncan's multiple range test was performed to compare the means ( $P=0.05$ ). All experiments carried out in three replications.

## RESULTS AND DISCUSSION

### *Efficiency of crude seed extracts against HGP inhibitory activity*

The crude extracts possess activities in the range between 18.11 to 55.33 % which are considered to have strong inhibitory activity (Fig. 1). The fabaceae species showed strong inhibitory activity against HGP (mean 45%) in while poaceae family species showed a moderate inhibitory activity (mean 19%).

All the species showed inhibitory activity *in vitro* against HGP activity while *P. vulgaris* crude seed extract showed a strong inhibitory activity of 55% against HGP among studied plant species.

#### *HGP inhibitory activity of different protein fractions of Poaceae species*

In different fractions of Poaceae species, inhibitory activity without significant differences was observed. Only *Z. mays* F4 fraction protein significantly affected HGP activity more than other fractions by 30% (Fig. 2).

#### *HGP inhibitory activity of different protein fractions of Fabaceae species*

Analysis of variances showed significant differences among F1 to F4 protein fractions in Fabaceae species. HGP activity strongly affected by *F. vulgaris* fractions and specially 70-80% fraction inhibited HGP activity more than 45%. About *C. arietinum* 70-80% protein fraction has more efficiency than other ones. The first fraction in all plants showed moderately to weak inhibitory activity in the range of 10-16%. The most inhibitory activity in *S. alopecuroides* was measured in 50-70% fraction by 26% (Fig. 3).

The use of conventional insecticides to control insect pests poses hazards to human health, non-target species, beneficial insects and environment. Indiscriminate use of chemical insecticides can also select insecticide resistance populations of pests (Harrison & Bonning, 2010). The digestive enzymes such as proteolytic and amylolytic enzymes are a target for insect pest management programs that are safe and environmentally friend method. Digestive enzymes play important roles in insect growth, development and reproduction; other functions including enzyme activation and detoxification are in relation with protease enzymes in insects digestive system (Christeller et al., 1992; Terra et al., 1996). Digestive systems of the lepidopteran larva contain proteases such as trypsin, chymotrypsin and elastase. Özgür et al. (2009) studies on *H. armigera* digestive protease showed that, serine proteases are dominant protease in the *H. armigera* midgut.

Several families of proteinase inhibitors has recognized among the animal and plant kingdom. Majority of proteinase inhibitors studied in plant kingdom originates from three main families namely Fabaceae, Solanaceae and Poaceae (Wee, 2000). Many investigators have isolated and characterized enzyme inhibitors from poaceae species such as barley, wheat and maize. Divya et al. (2014), reported that *Z. mays* contains PIs with trypsin and chymotrypsin inhibitory activity and these enzymes are abundant in cotton bollworm gut (Ozgur et al., 2009). Also Gourinath et al. (2000) reported that the members of Poaceae family have serine protease inhibitors. Odani et al. (1983) has reported that a large number of inhibitors in poaceae family have only  $\alpha$ -amylase-inhibitory activity; however inhibitors from barley, rye and tall fescue are active against trypsin.

Maize and ragi inhibitors showed dual activities and can inhibit serine proteinases as well as  $\alpha$ -amylase (Mahoney et al., 1984; Shivraj & Pattabiraman, 1981; Habib & Fazili, 2007). Boisen (1983), reported that, Inhibitors of trypsin, chymotrypsin and microbial proteases are the most common PIs in barley and are present mostly in seeds also trypsin and chymotrypsin inhibitory activity was detected in barley by Casaretto et al. (2004). Constantin et al. (2008) and Poerio et al. (2003) demonstrated that protease inhibitors are present in leaf and seeds of wheat. In present study protease inhibitory activity of different poaceae species has observed and all fractions with minor differences contains HGP inhibitory activity which is in agreement above mentioned reports. The present study concluded that all the species possess the potential to inhibitory *H. armigera* gut protease activity. Results revealed that the F1 fraction protein of all studied plants

showed less than 20% inhibitory activity against HGP, and the F2 and F3 fraction exhibited the same inhibitory activity in the range of 10-20% in Poaceae species.

Fabaceae species are rich of proteins and protease inhibitors and plenty of studies isolated and characterized different PIs from leaves, seeds and foliage of different Fabaceae species (Giri et al., 1998; Franco, 2003; Fan & Wu, 2005; Kansal, 2008; Abd El-latif, 2015). The PIs from the wild relatives of pigeonpea showed considerable potential against the HaGPs (Parde et al., 2012).

Our results were concurrent with those by Nair et al. (2013), *C. arietinum* (chickpea) seeds are known to contain, inhibitors of proteases. This study has shown that *P. vulgaris* and *C. arietinum* protease inhibitor caused a significant decrease in proteolytic activity in the gut of *H. armigera* compared to the other inhibitors. The legume plants especially in F1, F2 and F3 fractions exhibited near 15-50% inhibitory activity on HGP. In over all, among studied plants, *Z. mays*, *P. vulgaris* and *C. arietinum* have strong inhibitory activity in compare with other plants that are in agreement with above mentioned reports.

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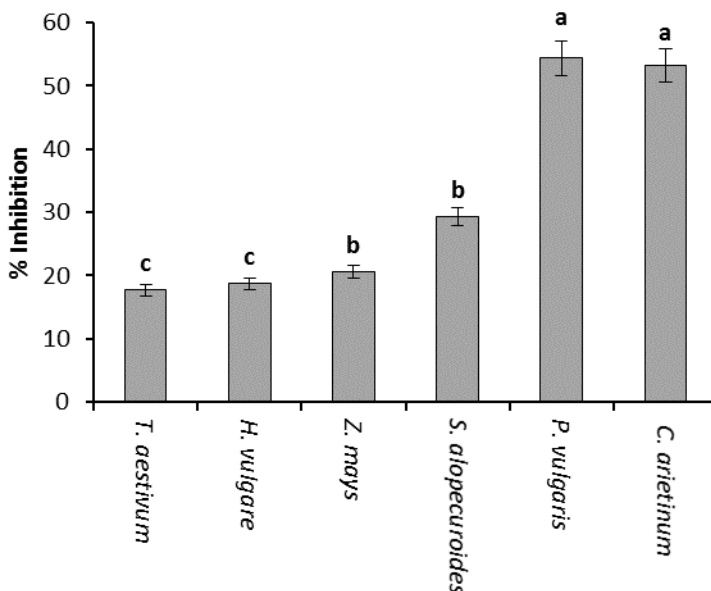


Figure 1. PI activity of different plant species crude seed extracts on *H. armigera* midgut protease activity (The means followed by different letters are significantly different,  $p=0.01$ ).

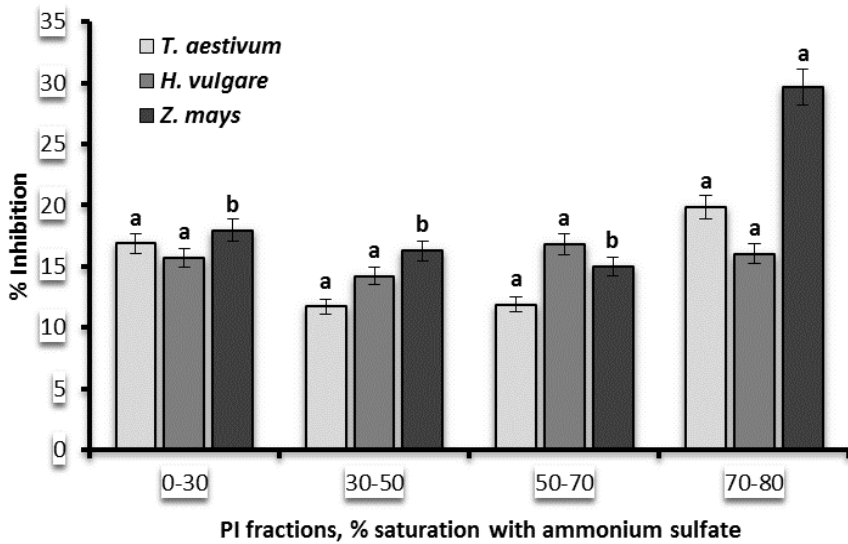


Figure 2. HGPI activity of different protein fractions of Poaceae family plants prepared with saturation in Ammonium sulfate (The means followed by different letters in each plant are significantly different,  $p=0.01$ ).

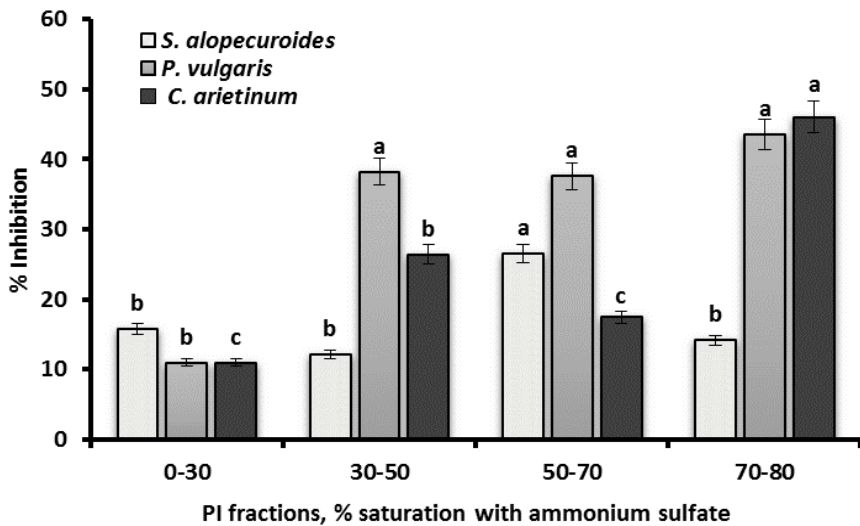


Figure 3. PI activity of different protein fractions of Fabaceae family plants prepared with saturation in Ammonium sulfate (The means followed by different letters in each plant are significantly different,  $p=0.01$ ).