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A Review

Exploring Plant Proteinase Inhibitors

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Abstract Proteinase Inhibitors (PIs) are small, natural antagonists of proteinases and present in all life forms. PIs are widely present in plants and often found in storage organs. They are known to be inducible in plants by injuries, such as insect damage. PIs have enormous diversity of function through regulation of target proteinases. Various plant sources have been explored for isolating PIs and broad-spectrum of biological activities have been elucidated. A range of strategies have been attempted to improve effectiveness of proteinase inhibitors as antimetabolites towards insects, bacteria and fungi. Much emphasis is yet to be given to address the health benefits of the PIs and implementing it in the most available forms throughout.

Keywords Serine proteinase inhibitor; Trypsin inhibitor; Insect pest; Cloning

Plants are continuously exposed to insect pest and pathogen attacks during their life cycle. Like animals, plants cannot move away from an endangering environment nor do they possess a characteristic immune system. Each plant species or cultivars have developed diverse defense mechanism, which includes both specific and general defense responses. Herbivorous insects, mites and nematodes are major contributors to yield loss either directly through consumption of plant biomass or indirectly as vectors/facilitators of pathogen infection. Consequently, introgression of insect-pest resistance into crops has become one of the major priorities for plant breeders. Resistant cultivars reduce dependence on insecticides and need for crop rotation, provide economic benefits without compromising the environmental stewardship. Studies on host plant resistance reveal that plants have evolved natural defense strategies against pests, including the production of compounds that contribute directly or indirectly, protection against herbivore invasion. The best-known plant substances supposedly involved in defense mechanisms against phytophagous insects are ribosome-inactivating proteins (RIPs), protease inhibitors (PIs), amylase

inhibitors (áAl-I) and lectins. Insects have developed several strategies to overcome plant defense barriers, allowing them to feed, grow and reproduce on their host plants. However, insects possess a powerful assemblage of enzymes that constitute their defense against chemical toxicant (Ryan, 1990; Smith and Boyko, 2006). The major digestive proteases present in herbivorous insect gut are serine, cysteine, and aspartyl proteases. Inhibitors of these proteinases have been identified in plants that inhibit gut proteolytic activity, and adversely affect the growth and development of insect pests. Studies reveal that PIs are induced under various stress conditions such as insect attack, mechanical wounding, pathogen attack and exposure to UV. Role of PIs in plants as natural defense has greatly stimulated thinking towards enhancing the plant defense against insect pests through genetic engineering. The strategy to improve plant defenses against insect pests is being tried using serine proteinase inhibitors. Evolutionarily during plant-insect interactions, insects have adapted to the PIs of their host plants. Therefore, non-host plants were also evaluated and subsequently used for transferring these traits into commercial cultivars.





PIs have broad-spectrum of biological activity, which includes suppression of pathogenic nematodes (Williamson and Hussey, 1996), inhibition of spore germination and mycelium growth (Dunaevskii et al., 1997) and hampering the growth of pathogenic fungi (Joshi et al., 1998; Bhattacharjee and Prasad, 2005). In addition, PIs also contribute to better and enhanced nutritional quality of grains, as they are rich in cysteine and lysine (Ryan, 1989, Bhattacharjee et al., 2006). PIs are specific for each of four mechanistic classes of proteolytic enzymes, classified as serine, cysteine, aspartic and metallo-protease based on the active amino acid in their reaction center. In general, PIs are competitive inhibitors, bind to the active site of the enzyme. For instance serine PIs bind to serine proteases, which include trypsin, chymotrypsin, elastase, subtilisin and thrombin. Serpins (serine protease inhibitors or classified inhibitor family I4) are the largest and most broadly distributed superfamily of protease inhibitors (Rawlings et al., 2004). Serpin-like genes have been identified in animals, plants, bacteria, and some viruses (Gettins, 2002). Most serpins are irreversible inhibitors of serine proteases of the chymotrypsin family, although some have evolved to inhibit other types of serine proteases, and a few are also able to inhibit cysteine protease (Schick et al., 1998, McGowan et al., 2006, Vercammen et al., 2006, Ong et al., 2007, Roberts and Hejgaard, 2008). Furthermore, some serpins have the ability to form complexes with very divergent proteases (Huntington, 2006). Serpins are involved in a number of fundamental biological processes, and a role in the protection of storage tissue against insects and pathogens has been proposed for plant serpins (Dahl et al., 1996; Rasmussen et al., 1996).

1 Proteinase inhibitors in plants

Plants have developed defense systems to combat various pathogens throughout their life cycle, from the seed stage until senescence, and it is particularly important to keep embryo freefrom infection. There are several embryonic defense mechanisms detected, including the production of plant lectins and pathogen-relatedproteins, in response to pathogen or insect attack (Ye et al., 2001; Guiderdoni et al., 2002). Serine proteinase inhibitors are expressed in developing seeds and are thought to play an important rolein inhibiting trypsin andchymotrypsin of external origin. Two major serine class of proteinase inhibitor shave been studied extensivelyin plants: Kunitz inhibitors and Bowman-Birk inhibitors (Ryan, 1990). Proteinase inhibitors of high molecular weight with low cystein content are termed as Kunitz type (Odani and Ikeneka, 1973). Bowman-Birk inhibitors (BBIs) are cystein rich proteins of about 8 kD to 16 kD with disulfide bonds. Serine proteinase inhibitors are universal and most studied class of proteinase inhibitors in plant kingdom (Haq et al., 2004, Mello et al., 2002). Although they are present in lower concentration in vegetative tissues, are primarily localized to storage tissues such as seeds and tubers rich in storage proteins.

Plant cystatins or phytocystatins are the second most studied class of inhibitors from plants, viz, cowpea, potato, cabbage, ragweed, carrot, papaya, apple fruit, avocado, chestnut, and Job's tears. Seed cystatins have been reported from wide range of crops including sunflower, rice, wheat, maize, soybean, and sugarcane (Kuroda et al., 2001; Yozaura et al., 2002; Connors et al., 2002). Squash inhibitor, member of highly potent canonical serine proteinase inhibitors with typical knottin fold, was isolated and characterized (Chiche et al., 2004).

2 Families of proteinase inhibitors and their distribution

A comprehensive system of classification has been proposed for facilitating the exchange, storage and retrieval of information about this group of proteins (Rawlings et al., 2004). On the basis of three-dimensional structures, 31 families are assigned to 26 clans. The term "Clan" is to designate a single evolutionary line of inhibitors defined by single type of protein fold. Leo et al (2002) developed PLANT –PIs database to facilitate retrieval of information on plant protease inhibitors and related genes (Table 1). Christeller and Laing (2005) identified eight families of serine proteinase inhibitors (Table 2), which matched previously identified eight families in MEROPS.

Soybean trypsin inhibitor was the first PI to be isolated and characterized. Since then many PIs have been characterized that are widely distributed throughout the plant kingdom (Konarev et al., 2004). Most of the plant PIs characterized are from Gramineae, Poaceae, Leguminosae, Fabaceae, and Solanaceae families (Brzin and Kidric, 1995). PIs are usually found in storage organs, such as seeds and



1	5
Plant protease inhibitor family	PLANT-PIs code
Bowmann-Birk serine proteinase inhibitors	BBI
Cereal trypsin / a-amylase inhibitors	BRI
Cysteine proteinase inhibitors	CYS
Metallo carboxypeptidase inhibitors	MCI
Mustard trypsin inhibitors	MSI
Potato type I inhibitors	PI1
Potato type II proteinase inhibitors	PI2
Serpin	SP1
Soybean trypsin inhibitors (kunitz)	KNI
Squash inhibitors	SQI

Table 1 Plant proteinase inhibitor family in PLANT-Pis

However, their occurrence in aerial part of plants, as a consequence of several stimuli, has also been documented (De Leo et al., 2002). PIs accumulate to about 1 to 10% of the total soluble proteins of storage tissues. Number of PIs has been reported from non-storage tissues, such as leaves, flowers and roots (Brzin and Kidric, 1995; Xu et al., 2001; Sin and Chye, 2004). Trypsin inhibitor in mung bean is found localized to cytosol of cotyledonary cells (Chrispeels and Baumgartner, 1978). Soybean trypsin inhibitor (SBTI) was found localized in cotyledonary cell walls and embryonic cells, and to lesser extent in protein bodies, cytoplasm and nuclei. Soybean Bowman- Brik inhibitor (SBBI) was found to occur in protein bodies, nuclei, (Horisberger and cytoplasm and Tacchini-Vonlanthen, 1983). In tomatoes, serine proteinase inhibitors I and II selectively accumulated in endosperm and secretory cells of root cap (Narwaez-Vasquez et al., 1993). Xu et al (2004) reported the expression of phloem specific PIN2 protein in S. americanum stems, roots, and leaves.

3 Properties and regulation of plant PIs

Plants PIs are typical polypeptides composed of L-amino acids linked through peptide bonds, widespread in both monocots and dicot species (Ryan, 1990). Although the molecular size of PIs varies from 4 to 85 kD, majority of them are in the range of 8 kD to 20 kD (Hung et al., 2003). Plant PIs usually have high cysteine residues that form disulfide bridges contributing significantly to the stability of the inhibitors. Bowman-Birk type of Trypsin inhibitor from Brassica campestris seed (BCTI) of molecular size 8 kD, was found to be thermo stable, which is apparently related to the presence of the disulfide bridge (Hung et al., 2003). Plant PIs are low in methionione, histidine and tryptophan but are often



rich in aspartic acid, glutamic acid, serine, arginine and lysine. Glycosylated plant PIs similar to mammalian glycoprotein proteinase inhibitors have not been reported so far.

Systemic signals associated with translocation of wound response include systemin, abscisic acid (ABA), hydraulic (variation potentials) and electrical signals (Malone and Alarcon, 1995). In tomato, it has been shown that protease inhibitor initiation factor (PIIF) is wound inducible, triggers the cascade of events leading to the synthesis of serine proteinase inhibitor in the whole plant (Bryant et al., 1976; Miège et al., 1976). This suggests the existence of signal that moves from injured tissue to all parts of the plant leading to systematic induction of PI gene expression. Systemin regulates the activation of over 20 defensive genes in tomato, in response to herbivore and pathogen attack, by activating lipid based signal transduction pathway- linolenic acid released from plant membranes is converted into signaling molecule, jasmonic acid (JA). Systemin, cell surface receptor of molecular size~160 kD, regulates intercellular cascade by (1) depolarization of the plasma membrane and (2)opening of ion channels, thus increasing the intracellular Ca²⁺, which activates MAPK and phospholipase A. These rapid changes play vital role in intra cellular release of membrane linolenic acid and its conversion to JA, potent transcription activator of defense gene (Koiwa et al., 1997; Ryan, 2000). Plantinfected by pathogen trigger two possible signal pathways as its defense strategy- (1) recognition of penetration and colonization of pathogen through wound and (2) direct molecular recognition of the pathogen (Figure 1; Figure 2) (Cordero et al., 1994). Salicylic Acid (SA) and its methyl ester (MeSA) induce systemic acquired resistance in plants against pathogenic microorganisms (Hunt et al., 1996). Several jasmonic acid-dependent and independent wound signal transduction pathways have been identified and characterized. Components of these signaling pathways are similar to other signaling cascades that include reversible protein phosphorylation cascade, calcium/calmodulin- regulated events and production of reactive oxygen species (León et al., 2001). Stintzi et al (2001) demonstrated that in absence of JA, 1, 2-oxo- phytodienoic acid (OPDA), precursor of JA, elicits defense signal response. Induced rapid de novo synthesis of rice BBI was found in seedling/leaf





Serine proteinase inhibitor family	Class	Target protease
Kunitz Family	13	Trypsin, Chymotrypsin, Plasma Kallikrein
SERPIN Family	14	Trypsin, Chymotrypsin, Elastase
Proteinase Inhibitor 1 (PP1) family	113	Chymotrypsin, Elastase, Subtilisin, Trypsin
Ragi Seed trypsin / alpha –amylase inhibitor Family	16	Trypsin, Plasma kallikrein, Factor X11a
Squash Family	17	Trypsin
Bowman Birk Family	112	Trypsin, Chymotrypsin, CathepsinG, Matriptase
Mustard seed trypsin inhibitor family	118	Trypsin, Chymotrypsin
Proteinase inhibitor 2 (PP2) family	120	Trypsin, Chymotrypsin, Elastase, CathepsinG, Pronase, Subtilisin

in response to cut, exogenous application of jasmonic acid (JA), and protein phosphatase 2A (PP2A) inhibitors and completely inhibited by cycloheximide (Rakwal et al., 2001). Studies on herbivore-induced synthesis of glucosinolates, trypsin inhibitors, and resistance to herbivory in Brassica suggest that induced levels of trypsin inhibitors vary with genotypes (Cippolini et al., 2003). Changes in expression of jasmonic acid (JA)-and salicylic acid (SA)-dependent defense genes was observed in potato in response to potato and green peach aphids infestation (Martinez et al., 2003). Saedler and Baldwin (2004) demonstrated the potential of VIGS to manipulate and silence the expression of two jasmonate-induced genes that mediate the expression of proteinase inhibitor in Nicotiana attenuata roots and shoots. Hypothetical model for defense gene activation involving JA has been proposed which involves PI genes and jasmonate as critical signals in dicot's defense responses (Doares et al., 1995; Sivasankar et al., 2000). Farmer et al (1992) hypothesized that lipases may be synthesized or activated in response to wounding and that linolenic acid released from membranes may initiate the intracellular transduction pathway leading to proteinase inhibitor synthesis. Free linolenic acid could be converted to jasmonic acid, which may be a key signaling component that may be very close to the trans-element(s) that regulate proteinase inhibitor gene transcription (Figure 1).

Early events of signaling cascade involve changes in protein phosphorylation pattern, which eventually regulate various cellular processes in eukaryotes (Hunter, 1995), including plant defense responses (Conrath et al., 1997). Phosphorylation of proteins is a transient process, can be regulated by using protein phosphatase inhibitors- cantharidin and endothall (Li et al., 1993; MacKintosh et al., 1994; Millward et al., 1999). JA is found highly effective in inducing defense-related and cellular defense proteins, including PRs (Agrawal et al., 2000; Rakwal et al., 1999). It is initiated with the interactions of local or systemic signal molecules (Abscisic acid, systemin, oligogalacturonic acid, chitosan, electrical, and hydraulic signals) and putative plasma membrane receptors (β-glucan-elicitor-binding protein and systemin-binding protein). *P*-chloromercuribenzene sulfonic acid (PCMBS) has been shown to inhibit phloem systemin translocation, and attenuate systemic induction of protease inhibitor (PI) gene expression. Wound signal is transduced by an unidentified lipase that facilitates the release of linolenic acid from membrane lipids, a process stimulated by ABA. Volicitin, which originates from oral secretion of insects, also function like linolenic acid (Alborn, 1997). Defl regulates the conversion of 13(S)- hydroperoxylinolenic acid (HPOTrE) to 12-oxo- phytodienoic acid (12-oxo-PDA). Defl tomato mutant sensitive to insect attack is found defective in octadecanoid pathway. Diethyldithiocarbamic acid (DIECA) reduces the conversion of HPOTrE to 13-hydroxylinolenic acid (HOTrE) and decreases the synthesis of jasmonic acid (Howe, 1996). Inhibitors of ethylene action (norbornardiene (NBD) and silver) and salicylic acid prevent iasmonic acid-stimulated induction of PI expression (Borella, 1996; O'Donnell, 1996). Okadaic acid, inhibitor of protein phosphatase1 and 2A, inhibits jasmonate-induced PI expression; suggest the involvement of these phosphatases downstream regulatory pathway of jasmonate. The protein kinase inhibitor, staurosporine, inhibits PI gene expression induced by ABA, specifically, but not jasmonate (Dammann, 1997) (Figure 1).





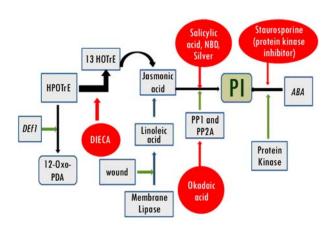


Figure 1 Regulation of proteinase inhibitor gene in plant Note: Green arrow: activation; red arrow: inhibition; black/blue arrow: direction of reaction

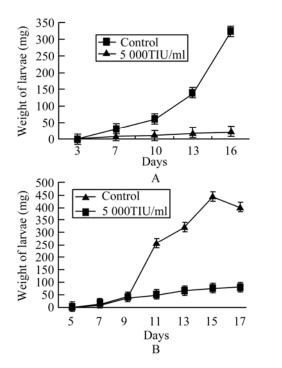


Figure 2 Larval growth curve of second (A) and third (B) instar Helicoverpa armigera larvae fed on artificial diet containing ChTI (5000 TIU/mL) and control on same diet without ChTI

Note: Each value represents the mean weight of 20 larvae from five independent replicates \pm SE

4 Role of proteinase inhibitors in plants

Plant PIs are known to play important role in plant's defense against insect-pest and pathogens as well as regulation of endogenous proteinases (Mosolov et al., 2001; Birk, 2003; Shewry, 2003). They are of interest in plant biology as source of (1) resistance against pests and pathogens, (2) drugs with antiviral properties and (3) markers for studying plant diversity

and evolution (Konarev et al., 2002; Lawrence and Koundal, 2002; Korsinczky et al., 2004).

4.1 In insect pest control

Serine proteinases have been identified in gut extracts of many lepidopteran insects (Houseman et al., 1989) and many of these enzymes are inhibited by proteinase inhibitors. The pH of lepidopteran guts are alkaline, ranging from 9.0-11.0 (Applebaum, 1985) wherein serine- and metallo-proteinases are most active. Hence, serine proteinase inhibitors show anti-nutritional effects against several lepidopteran insects (Shulke and Murdock, 1983; Applebaum, 1985). Nandeesha and Prasad (2001) have reported partially purified subabul trypsin inhibitor (STI) from Leucaena leucacephala with molecular weight of 15 kD that shows high-level thermo tolerance and pH stability. Bioassay reveals that STI is a strong inhibitor of H. armigera, extending larval growth period by 12 days and induces mortality by 40% TIU/mL concentrations. at 20,000 Dimorphandra mollis seed trypsin inhibitor (DMTI-II) showed 67% mortality among bruchid, (Callosobruchus maculates) fed at 1% level in artificial diet (Macedo et al., 2002). Giri et al (2003) reported at least 14 trypsin inhibitors from Psophocarpus tetragonolobus seeds. WBTI-1 (28 kD) was identified as a potent inhibitor of HGP activity (94%). WBTI-2 (24 kD) and WBTI-4 (20 kD) inhibited HGP activity up to 85%. WBTI-3-5-7 showed limited inhibition of HGP as compared with trypsin. Barley (Hordeum vulgare L.) malt contains all the four classes of endoproteinases as well as low molecular weight peptides that inhibit many of these endoproteinases. These are chloroform/ methanol soluble peptides, possibly play a significant role in controlling the activity of barley proteinases during germination, and protecting seed and young plant from pathogens/pests (Jones and Fontanini, 2003). Cotton boll weevil, Anthonomus grandis, feeds on fruits and buds causing severe crop loss. Trypsin/ chymotrypsin inhibitor (BTCI) purified from Vigna unguiculata seeds showed low inhibitory activity against trypsin-like proteinases of adult weevils. The bioassay results strongly suggest that BTCI has potential to engineer crop plants for resistance to the cotton boll weevil (Franco et al., 2003). Peltophorum dubium seed trypsin inhibitor (20 kD), thermo stable kunitz type inhibitor showed 56% mortality in Anagasta kuehniella at 1.6% level (Rodrigues et al., 2003). Chougule et al (2003) analyzed seeds of 53





pigeon pea cultivars and wild genotypes, resistant and susceptible to insect-pests and pathogens, for the presence of proteinase inhibitors against Helicoverpa armigera gut proteinases (HGPs). PIs from pigeonpea cultivars showed complete inhibition of trypsin and chymotrypsin, and moderately towards HGPs. PIs of wild relatives showed stronger inhibition with HGPs. Telang et al (2003) purified bitter gourd (Momordica charantia L.) seed proteinase inhibitors (BGPIs) that strongly inhibit HGPs. Electrophoretic analysis revealed the presence of two major proteins (BGPI-1 and-2) and two minor proteins (BGPI-3 and-4) having inhibitory activity against trypsin and HGPs. BGPIs inhibited proteolytic activity of larvae fed on different host plants, artificial diet with or without PIs supplementation and proteinases excreted in fecal matter from respective samples. BGPIs were found to retard growth and development of Helicoverpa armigera and Spodoptera litura. Reports indicate that BGPIs mediated inhibition of insect gut proteinases directly affect fertility and fecundity of H. armigera and S. litura. Patricia et al (2003) used bioassays to investigate the effect of Bowman Birk and kunitz -type soybean trypsin inhibitor on growth pattern of Diatraea saccharalis moth using two diets- Diet 1 was less nutritious, with low protein content; and reduced minerals and essential aminoacids (cysteine, lysine and methionine) content while Diet 2 was richer and more complete. Food intake and utilization; larval development and mortality were monitored. When PI supplemented, larval development was was significantly altered in larvae fed with diet1, with reduced trypsin-like activity of midgut enzymes. Diet 2 fed larvae also showed reduced level of trypsin-like activity but it was less marked than diet 1. Similar feeding experiment was done with subabul high and low molecular weight trypsin inhibitor (HSTI, LSTI) using artificial diet, chickpea seeds and leaves. Larvae fed with artificial diet showed reduction in larval weight up to 21% (HSTI) and 43% (LSTI). However, larvae fed on seeds showed significant reduction in weight, 52.4% (HSTI) and 60.3% (LSTI), suggesting the diet also play vital role on the effectiveness of the inhibitors on larval growth and development (Bhavani et al., 2007). Franco et al (2004) suggested that SKTI can be an effective in developing transgenic plants against the cotton boll weevil, Anthonomus grandis. Cotton boll weevil gut digestive system

contains serine proteinases. In vitro assay showed that SKTI inhibit these enzymes. Neonate larvae reared on an artificial diet containing SKTI showed reduction in larval weight of up to 64% and caused mortality and severe deformities of larvae, pupae and adult insects. Oryza sativa chymotrypsin inhibitor (OCPI1) transgenic positive plants showed higher grain yield and seed setting rate than the wild type and control under the severe drought stress conditions, whereas the potential yield of transgenic plants under normal growth conditions was not affected. Chymotrypsin- inhibitor activity assay from positive transgenic plants showed stronger inhibition. The decrease of total proteins in transgenic plants is less than the wild type under drought stress (Huang et al., 2007). The defensive role of PIs is based on their inhibitory activities towards proteolytic enzymes of insect gut and phytopathogens resulting either in a critical shortage of essential amino acids (Hilder et al., 1993; Jongsma and Bolter, 1997) or interfering with metabolic processes, such as the proteolytic activation of enzymes, molting of insects, or replication of viruses (Gutierrez-Campos et al., 1999). Direct evidence for the involvement of PIs in the plant defense system has come from studies on transgenic plants. Transgenic plants expressing PIs have been produced in the last two decades and tested for enhanced defense capacities, particularly against insect-pests (De Leo et al., 2002). Expression of cowpea trypsin inhibitor (CpTI) in transgenic tobacco was shown for the first time to confer resistance to feeding by tobacco budworm Heliothis virescens (Hilder et al., 1987). Plant proteinase inhibitors are known to confer natural protection against nematode attack (Atkinson et al., 2003; Cai et al., 2003; McPherson and Harrison, 2001). Nematode control with PIs expressed in transgenic tomato (Urwin et al., 1995), Arabidopsis thaliana (Urwin et al., 2000), and rice (Vain et al., 1998) has been well documented (Hepher and Atkinson, 1992).

PIs have the potential to enhance the current Bt toxin technology because they target a broader range of pests, including nematodes and fungi (Hilder and Boulter, 1999). There is a major concern that the effectiveness of Bt will be negated if field-evolved Bt resistance (Tabashnik et al., 2009) becomes a more widespread problem. A proposed management strategy for delaying insects' development of





resistance to plant-protection transgenes, such as *Bt* toxins, is to deploy multiple insect-control genes (such as PIs) with different modes of action in a single plant (Manyangarirwa et al., 2006). There is evidence that the combination of PIs with a sublethal-dose *Bt* toxin has a strong effect on the growth and development of insects (Zhu et al., 2007). Dunse et al (2010) coexpressed *Nicotiana alata* proteinase inhibitor (NaPI) and II proteinase inhibitors and *Solanum tuberosum* potato type I inhibitor (StPin1A). The combined inhibitory effect of NaPI and StPin1A on *H. armigera* larval growth in the laboratory was reflected in the increased yield of cotton bolls in field trials of transgenic plants expressing both inhibitors.

In our own study; Bhattacharjee et al (2009) showed that a thermotolerant monomeric trypsin inhibitor from Cocculus hirsutus (ChTI) caused 74% and 59.53% inhibition of bovine trypsin and Helicoverpa gut proteases respectively. The second and third instar larvae of H. armigera fed with ChTI (5000 TIU/mL) resulted in 84.59 and 58.71% reduction in mean larval weight respectively. An increase in the larval growth period was observed in ChTI fed larvae at all instars and inhibitor fed larvae could not complete their life cycle (Figure 2; Figure 3). In a recent study by Alvarez-Alfageme et al (2011) it was observed that AtSerpin1 (Arbidopsis thaliana Serine proteinase inhibitor) inhibited proteases from all pest and non-target species assayed. AtSerpin1 supplied in the artificial diet or by transgenic plants reduced the growth of S. littoralis larvae by 65% and 38%, respectively, relative to controls. Macedo et al (2010) suggested that trypsin inhibitor (ApTI) purified from Adenanthera pavonina have a potential antimetabolic effect when ingested by Anagasta kuehniella, a polyphagous pest that feeds on a wide variety of stored products. Larval and pupal developmental time of larvae fed on ApTI diet at 1% was significantly longer; the larval period was extended by 5 days and pupal period was 10 days longer, therefore delaying by up to 20 days and resulting in a prolonged period of development from larva to adult. The percentage of surviving adults (%S) decreased to 62%. The fourth instar larvae reared on a diet containing 1% ApTI showed a decrease in tryptic activity of gut and that no novel proteolytic form resistant to ApTI was induced. da Silva et al (2012) showed that 0.1% trypsin inhibitor (ApTI) have great toxic potential on the

development of *Diatraea saccharalis*, a major sugarcane pest, causing damage to the stalks of sugarcane plants larvae. 0.1% ApTI produced approximately 67% and 50% decreases in weight and survival larval, respectively. The level of trypsin was significantly decreased (ca. 55%) in the midgut of larvae reared on a diet containing 0.05% ApTI and the trypsin activity in ApTI-fed larvae demonstrated sensitivity to ApTI.

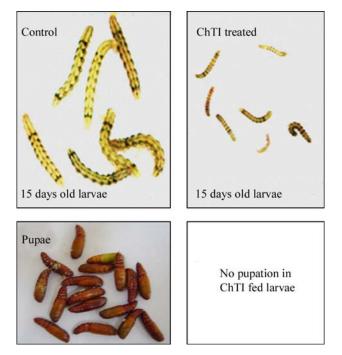


Figure 3 Effect of ChTI on growth of *Helicoverpa and* Spodoptera larvae

Note: The second star larvae were fed with artificial diet containing ChTI (5000 TIU) and control on same diet without ChTI

4.2 Proteinase inhibitor-antifungal and antibacterial property

Proteinase inhibitors in plants are able to suppress enzymatic activity of phytopathogenic microorganisms. Trypsin and chymotrypsin inhibitors of plant origin were shown to suppress activity of proteinases secreted by *Fusariumsolani* (Mosolov et al., 1976). Heat stable antimicrobial peptide, Potide G, completely suppress the proteolytic activity of trypsin, chymotrypsin and papain, in addition to growth inhibition of variety of bacterial and fungal strains (Kim et al., 2006). In tomato and potato tubers infected with the oomycete fungus, *P. infestans*, increase in trypsin and chymotrypsin inhibitor content was observed, which correlated with resistance to pathogen (Valueva et al., 1998, 2003). Mendieta et al (2004) reported that





trypsin inhibitor inhibited the spore germination of fungal pathogen Sclerotinia sclerotiorum. However, proteinase inhibitors induced in response to infection differed from inhibitors present in healthy plant (Geoffroy et al., 1990). Such induction in response to infection by pathogenic microorganisms is not limited to serine proteinase inhibitors, synthesis of cystatin like inhibitor was observed in chestnut leaves inresponse B. cinerea infection (Pernas et al., 2000). Kim et al (2005) purified potamin-1 (PT-1), a 5.6 kDa trypsin-chymotrypsin protease inhibitorfrom the tubers of the potato (Solanum tuberosum). PT-1 strongly inhibited pathogenic microbial strains, including Candida albicans, Rhizoctonia solani, and Clavibacter michiganense subsp. Michiganinse. This protease inhibitor, PT-1, was composed of polypeptide chains joined by disulfide bridge(s). Reduced PT-1 almost completely lost its activity against fungi and proteases indicating that disulfide bridge is essential for its protease inhibitory and antifungal activity. Wang et al (2006) purified 10 kDa proteinase inhibitor from mung bean (Phaseolus mungo) seeds. It exerted a potent inhibitory action toward a variety of fungal species including *Physalospora* piricola, *Mycosphaerella* arachidicola, **Botrytis** cinerea, Pythium aphanidermatum, Sclerotium rolfsii and Fusarium oxysporum, as well as an antibacterial action against Staphylococcus aureus. Our own study showed that ChTI exhibited strain specificity and inhibited growth and development of plant fungal pathogens. Bioassay studies on yeast strains indicated that Δ YNK and MNN1 are more sensitive to ChTI. The results suggest that phosphodiester linkage in cell wall components is likely to be the key determinants for binding of ChTI (Figure 4). It had no effect on bacteria (Bhattacharjee et al., 2009). Shilpa and Murugan (2010) purified a 14.3 kD proteinase inhibitor from Coccinia grandis. PI strongly inhibited pathogenic microbial strains, including Staphylococcus aureus, Klebsiella pneumoniae, Proteus vulgaris, Eschershia coli, Bacillus subtilis and pathogenic fungus Candida albicans, Mucor indicus, Penicillium notatum, Aspergillus flavus and Cryptococcus neoformans. Examination by bright field microscopy showed inhibition of mycelial growth and sporulation. Morphologically, PI treated fungus showed a significant shrinkage of hyphal tips. Reduced PI completely lost its activity indicating that disulfide bridge is essential for its protease inhibitory and antifungal activity.

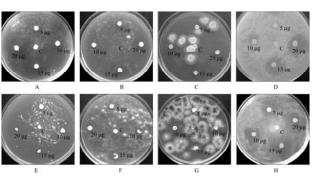


Figure 4 Antifungal assay using fungal isolates

Note: Zone inhibition assay of ChTI against Phytopathogens; A: Alternaria alternate; B: Aspergillus flavus; C: Colletotrichumcapsici; D: Fusarium oxysporum; E: Fusarium solani (crossandra isolated); F: Fusariumsolani (chick pea isolated); G: Rhizoctonia oryzea; H: Sclerotia sp.

4.3 Proteinase inhibitors in regulation of programmed cell death (PCD)

The compounds that were effective in controlling PCD in soybean- leupeptin, PMSF and AEBSF have inhibitory activity against proteases- leupeptin is an established inhibitor of cysteine proteases, whereas PMSF and AEBSF are inhibitors of serine proteases (Alonso et al., 1996). In plants, proteinase inhibitors are subject of regulation by intercellular signaling molecules, such as jasmonic acid (Farmer et al., 1992), salicylic acid (Doares et al., 1995) and systemin (Constabel et al., 1995). Salicylic acid was found to suppress the expression of cystatin (Doares et al., 1995). Salicylic acid also was shown to promote cell death induced by direct oxidative stress or pathogen attack leading to the possibility-(1) increased generation of H₂O₂ and (2) repressed cystatin expression (Shirasu et al., 1997). Controls of cellular fate through regulated expression of specific proteases in combination with the associated protease inhibitor genes provide additional plasticity to responses from outside stimuli. Plant cysteine proteases seem to play an important role in preventing PCD triggered by oxidative stress, wounding caused by insect chewing or during chilling-induced oxidative stress (Prasad et al., 1994; Solomon et al., 1999).

4.4 Proteinase inhibitors in health and disease control-medical and industrial aspects

Proteolytic enzymes are involved in numerous physiological processes in humans including digestion of food, tissue remodeling, host defense, blood coagulation and activation of proenzymes and prohormones. Proteinase inhibitors also play an





important physiological role in the regulation of enzymes. There are many examples of pathological conditions in which uncontrolled proteolytic activity of host enzymes leads to irreversible tissue destruction (e.g., in inflammatory processes, including rheumatoid arthritis and periodontitis), or to tumor growth and metastasis. In addition to host proteinases, exogenous proteinases from infectious agents such as bacteria, viruses or protozoa play a role in the onset and perpetuation of infection, suggesting the potentially therapeutic application for proteinase inhibitors in combating diseases. Therapeutic possibilities of plant PIs in the treatment of a wide range of disorders, such as pancreatitis, shock, allergy and inflammation associated with enhanced proteolytic activities had resulted in several kallikrein trypsin inhibitor-based drugs (Richardson, 1977; Park and Ohba, 2004). Epidemiological studies indicate decreased occurrence of breast, colon, and prostates cancers in vegetarian populations suggested the role of plant PIs in preventing these cancers (Birk, 1993), which has eventually led to extensive studies of plant PIs as cancer chemo preventive agents (Troll and Kennedy, 1993). Plant PIs are found active in regulating human physiological processes, e.g. cell signaling/migration, digestion. fertilization. growth. differentiation. immunological defense, wound healing and apoptosis, have great potential in therapeutic applications (Abdel-Meguid et al., 2000; Leung et al., 2000). Purified homodimeric trypsin inhibitor of 54kD from Clausena lansium seed inhibited trypsin with no inhibitory activity on chymotrypsin and proteinase K. However, Clausena lansium trypsin inhibitor inhibited the uptake of MTT by human leukemia HL60 and hepatoma Hep G2 cells. The activity of HIV-1 reverse transcriptase was reduced in the presence of the inhibitor (Ng et al., 2003). Two molecular species of oryzacystatin (OC), OC-I and OC-II were investigated to haveantiviral action against herpes simplex virus type 1 (HSV-1) in vitro and in vivo. In the mouse modelwith HSV-1-induced keratitis and encephalopathy, topical administration of OC-I to produced significant cornea а decrease in virusproduction and improvementin survival rates (Aoki et al., 1995). Kunitz trypsin inhibitors act on platelet aggregation, blood coagulation, fibrinolysis and inflammation. Due to its ability to block enzymes,

plant Kunitz inhibitors are useful as tools in the study of biochemical processes of these phenomenon (Sampaio et al., 1996, Souza-Pinto et al., 1995). Serine proteinase inhibitor isolated from Leucaena leucocephala seeds has shown to inhibit plasmin, human plasma kallikrein, trypsin, chymotrypsin and Factor XIIa but not factor Xa (Oliva et al., 2000). Soybean derived KTI inhibits LPS-induced up-regulation of cytokine expression possibly through suppression of ERK1/2 and p38 kinase-mediated NFkappaB activation. These findings may identify anti-inflammatory properties of KTI at the level of gingival fibroblasts and suggest the use of KTI in modulating inflammation, including periodontal disease (Kobayashi et al., 2005). The Bowman-Birk trypsin-chymotrypsin inhibitor from soybean has been described as a potential chemopreventive agent of cancer. Effects of inhibitor were compared with two variants of pea seed protease inhibitors, rTI1B and rTI2B homologous to BBI but differing in inhibitory activity, on the growth of human colorectal adenocarcinoma HT29 cells. Significant and dose-dependent decrease in the growth of HT29 cells was observed using these proteinase inhibitors (Clemente et al., 2005).Squash family inhibitors are the smallest protein serine protease inhibitors, composed of 30 amino acid residues. Squash family inhibitors (from the seeds of bitter gourd, squash, gourd and luffa) were examined on serine proteases associated with blood coagulation. Five of them prolonged the activated thromboplastin time of human plasma. All inhibitors inhibited the amidolytic activities of factor XIIa, plasma kallikrein, factor Xa, but not to the same extent with factor XIa, IXa, VIIa and thrombin. The extension of activated partial thromboplastin time by inhibitors appeared to correspond to their inhibitory potencies for factor XIIa (Hayashi et al., 1994). Siritapetawee and Thammasirirak (2011) investigated biological effect of Artocarpus heterophyllus (jackfruit) latex on human blood coagulation. They demonstrated the serine protease inhibitory property of a heteromultimeric glycoprotein (HSGPL1) purified from jackfruit latex. This protein affects the intrinsic factors of human blood coagulation by prolonging the activated partial thrombin time (APTT) and inhibiting blood coagulation factors XIa and α -XIIa. In addition, this protein was provisionally identified as a





heat-shock/chaperone protein. These properties may be a medicinal benefit, e.g., in wound healing, blood coagulation and fibrinolysis.

Conclusion

The potential of PIs is quite evident and using PIs for developing transgenics can give durable resistance than using Bt genes alone. As such, the inhibitors can probably act not only as self-sufficient protective proteins, but can also protect other recombinant proteins from the deleterious effects of proteolytic enzymes. Apart from this, medicinal properties of proteinase inhibitors are being explored. Much importance and awareness are required to implement the same for health benefits in it's the most available forms throughout.

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