

## Cysteine Proteinases and Cystatins

**Adeliana S. Oliveira<sup>1</sup>, José Xavier-Filho<sup>2</sup> and Maurício P. Sales<sup>1\*</sup>**

<sup>1</sup> Laboratório de Química e Função de Proteínas – LQFP; Departamento de Bioquímica; Universidade Federal do Rio Grande do Norte; Natal - RN - Brazil. <sup>2</sup> Universidade Estadual do Norte Fluminense; Centro de Biociências e Biotecnologia; Laboratório de Química e Função de Proteínas e Peptídeos; msales@cb.ufrn.br; Campos dos Goytacazes - RJ - Brazil

### ABSTRACT

*This review describes the definition, localization, functions and examples of cysteine proteinases and their protein inhibitors in vertebrate, non-vertebrate animals and plants. These inhibitors are related with defense mechanisms of plant against pests. It also describes the factors involved in the specific cysteine proteinase-cystatin interaction and high degree of affinity and large specificity in this interaction which are not only represented by the compatibility between amino acid residues of the active site involved in catalysis, but also of all amino acid residues that participate in the enzyme-inhibitor interaction.*

**Key words:** Cysteine proteinases, Cystatin, Phytocystatins

### INTRODUCTION

Enzymes constitute a specialized and diverse group of proteins that have several roles in many physiological processes. Proteolytic enzymes such as proteinases which are involved in digestive processes, proenzymes activation, liberation of physiologically active peptides, complement activation, inflammation processes and others are part of this protein group (Neurath, 1984). Proteinases are grouped into four categories according to the essential amino acid residue at their active sites, the optimum pH range of activity, amino acid sequences similarities, similarity to inhibitors. Proteinases are classified as serine-, cysteine-, aspartic and metallo-proteinases (Bode and Huber 1992). Examples of these have been identified in plants, insects, microorganisms and are all like to those found in mammalian (Table 1).

Inhibitor proteins have been found for each of the four mechanistic classes of proteinases and a large number of proteinases inhibitors are directed towards serine- and cysteine proteinases (Barrett et al., 1986; Turk and Bode, 1991). In contrast, only a few of those inhibitors are known for aspartic- and metallo-proteinases (Ryan, 1990; Jouanin et al., 1998). Studies on plant protein inhibitors are important due their involvement in defense mechanism and in protection of seeds reserves from premature hydrolysis (Ryan, 1989, 1990; Domoney et al., 1993). This review describes the relationship between cysteine proteinases of insects and cysteine proteinase inhibitors of plants.

### CYSTEINE PROTEINASES

In vertebrate, the cysteine proteinases are involved in lysosomal protein degradation systems but also play a extracellular role such in some metabolic

\* Author for correspondence

disorders (Trysellius and Hultmark, 1997). In non-vertebrate such as nematodes and mite, cysteine proteinases are among the digestive enzymes (Rawlings and Barrett, 1994; Pernas et al., 1998). In arthropods such as lobsters, they play a digestive role, but are also related in the nervous system (Rawlings and Barrett, 1994; Le Boulay et al., 1995). In insects, they are utilized in the digestive processes (Rawlings and Barrett, 1994) but are found in several other tissues, indicating

that they may also play other roles (Yamamoto et al. 1994; Matsumoto et al., 1995, 1997). Studies on the pH dependence of cysteine proteinase activity in crude extract of insect larvae have indicated that activity was generally in the alkaline pH range (Bode and Huber, 1992), which was the optima pH of the cysteine proteinases activities. (Ryan, 1990).

**Table 1 - Group of proteinases**

PROTEINASES	AMINO ACID IN ACTIVE SITE	pH OPTIMUM (RANGE)	PROTEINS
Serine proteinase	Ser; His	7 - 9	Trypsin, Chymotrypsin, Elastase, Cathepsin(+) G
Cysteine proteinase	Cys	4 - 7	Papain*, Ficin, Bromelain, Ananain, Cathepsins(+) B, C, H, K, L, O, S and W
Aspartic proteinase	(2) Asp, Try	below 5	Cathepsin(+) D and E, Renin, Pepsin
Metallo-proteinase	Metal ion	7 - 9	Carboxipeptidases A and B, aminopeptidases

\* Papain isolated from the latex and fruit of *Carica papaya* have been used to identify cysteine proteinase inhibitor of the plant origin; (+) The term "Cathepsin" is generally used for the lysosomal cysteine protease (Cyglar and Morb, 1997; Matsumoto et al., 1997).

**Table 2 - Cysteine proteinases from insects**

INSECT SPECIES	ORDER	D/ND	ENZYME TYPE	REFERENCE
<i>Sitophilus zeamais</i>	Coleopteran	D	Cathepsin L	Matsumoto et al., 1997
<i>Sarcophaga peregrina</i>	Diptera	ND	Cathepsin L	Homma et al., 1994
		ND	Cathepsin B	Takahashi et al., 1993
		ND	Bovine Trypsin-like	Nakajima et al., 1997
<i>Leptinotarsa decemlineata</i>	Coleoptera	D	Cathepsin L	Michaud et al., 1996
	Chrysomelidae		Cathepsin B, H and D*	Thie and Houseman, 1990 (*)
<i>Drosophila melanogaster</i>	Diptera	ND	Cathepsin L	Matsumoto et al., 1995
		D	Cathepsin L	
<i>Manduca Sexta</i> (tobacco hornworm)	Lepidoptera	D	Trypsin-like	Schukle and Murdock, 1983
	Sphigidae			
<i>Acanthoscelides obtectus</i>	Coleoptera	D	Cathepsin B	Wieman and Nielsen, 1988
	Bruchidae			
<i>Rhodnius prolixus</i> Stal	Hemiptera	D	Cathepsin B	Houseman and Downe, 1982, 1983
	Reduviidae	D	Cathepsin D	
<i>Zabrotes subfasciatus</i> (mexican bean weevil)	Coleoptera	D	CP	Lemos et al., 1990
	Bruchidae	D	AP	
<i>Tribolium castaneum</i> (red flour beetle)	Coleoptera	D	Catepsin D	Blanco-Labra et al., 1996
		D	Catepsin B	Murdock et al., 1987
		D	SP	Birk et al., 1960
		D	CP	Liang et al., 1991
<i>Callosobruchus maculatus</i> (cowpea weevil)	Coleoptera	D	Cathepsin B	Gatehouse et al., 1985
	Bruchidae	D	AP	Kitch and Murdock, 1986
		D	SP	Silva and Xavier-Filho, 1991
				Xavier-Filho and Coelho, 1980
<i>Spodoptera exigua</i> (beet army worm)	Lepidoptera	D	CP	Jongsma et al., 1995, 1996
	Noctuidae	D	SP	
<i>Mimosestes mimosae</i> (algaroba beetle)	Coleoptera	D	Trypsin-like	Pereira, 2000
	Bruchidae	D	CP, AP	

D, digestive enzymes; ND, non-digestive enzymes; CP, cysteine proteinase; AP, aspartic proteinase; SP, cysteine proteinase with similarity no related.

Experiments made with insect larvae showed that insects as well as some animals used a mixture of digestive enzymes to hydrolyze ingested proteins (Applebaum, 1985). The type of mixture could be the result of the type of ingested food (Ryan, 1990; Blanco-Labra et al., 1996) (Table 2).

Plant cysteine proteinases are found particularly in the vacuoles and are responsible for the mobilization of endosperm storage proteins during seeds germination. This provides nitrogenous nutrients to support the growth of young seedlings. Cysteine proteinases are also found in the

extracellular medium such as those from papaya and fig (Rawlings and Barrett, 1994) (Table 3).

The cysteine proteinases require processing such as removal of the amino-terminal fragment to produce the active enzyme. The amino-terminal region, also called proregion, plays important roles not only as inhibitors of enzymatic activity but for the correct folding of the newly synthesized protein and to protect it against denaturing effects in sudden changes in the pH conditions. In terms of the proregion, Cathepsin L-like and Cathepsin B-like enzymes are subfamilies of the papain family.

**Table 3** - Plant cysteine proteinases

SOURCES	SIMILARITY AND REFERENCE
Oryzains $\alpha$ and $\beta$	Papain
Oryzain $\gamma$ (Rice seeds)	Cathepsin H (Abe et al., 1991)
Aleurains (barley)	Cathepsin H (Mikkonen et al., 1996)
EP-C1 ( <i>Phaseolus vulgaris</i> )	Cathepsin L (Tanaka et al., 1993)
EP-B (barley)	Cathepsin L (Koehler et al., 1990)
SH-EP ( <i>Vigna unguiculata</i> )	Cathepsin L (Mitsuhashi et al., 1986)

**Table 4** - Alignment of the aminoacid sequences of the several cysteine proteinases

Pp	ExxxRxxxFxxNxxxIxxxN Karrer et al., 1993	(-)
Vm	ExxxRxxxFxxNxxxVxxxN Karrer et al., 1993	(-)
O- $\alpha$	ExxxRxxxFxxNxxxIxxxN Watanabe et al., 1991	57-75 aa residues
O- $\beta$	ExxxRxxxFxxNxxxVxxxN Watanabe et al., 1991	69-88 aa residues
EPB-1	ExxxRxxxFxxNxxxIxxxN Mikkonen et al., 1996	51-70 aa residues
EPB-2	ExxxRxxxFxxNxxxIxxxN Mikkonen et al., 1996	51-70 aa residues
SzL	ExxxRxxxFxxNxxxVxxxS Matsumoto et al., 1997	43-62 aa residues
mcL	ExxxRxxxWxxNxxxIxxxN Tryselius and Hultmark, 1997	(-)
mcK	DxxxRxxxWxxNxxxIxxxN Tryselius and Hultmark, 1997	(-)

Pp (papaya papain); Vm (*Vigna mungo*); O- $\alpha$  and O- $\beta$  (cysteine proteinases of the rice seeds, orizains); EPB-1 and EPB-2 (cystine proteinases of the barley seeds); SzL (Cathepsin L-like digestive cysteine proteinases of the *Sitophilus zeamais*); mcL and mcK (mammalian Cathepsins L, K,) and (-) data not found.

They form two group distinct of enzymes: the proregion of Cathepsin B is significantly shorter (~60 amino acid residues) than the proregion of Cathepsin L, with approximately 100 amino acid

residues. In three-dimensional structures of members of the papain family two distinct domains are commonly observed, the amino-terminal domain is mostly  $\alpha$ -helical while the

carboxyl-terminal domain contains predominantly  $\beta$ -sheets. The active site consists of Cys 25 and His 159 (papain numbering) and is supplemented by Asn 175 hydrogen bonded to His 159, and Gln 19. Active site of the Cathepsin B differ of others Cathepsin due to the presence of the so-called occluding loop, which can be responsible for the exopeptidase activity this protein. Several members of the papain family show preferences for bulky hydrophobic residues occupying the S2 subsite due to the presence of the Ser 205 residue in this subsite, whereas Cathepsin B prefers arginine (Karrer et al., 1993). In the papain family enzymes with different activities are found, including endopeptidases (such as papain and Cathepsins B, H, L and C), aminopeptidase (Cathepsin H and C) and carboxipeptidase activity (Cathepsin C). There are also differences in the

proteolytic activities of the Cathepsins in relationship to pI value and pH optimum: the Cathepsin B (pI = 4.5 - 5.5; pH 6.0), Cathepsin H (pI = 6.0 - 7.1; pH 6.8) and Cathepsin L (pI = 5.5 - 6.1; pH 5.5) (Rawlings and Barrett, 1994; Cygler and Morb, 1997; Turk et al. 1997).

Karrer et al. (1993) observed that all the cysteine proteinases had similarities to mammalian H and L Cathepsins containing the ERFNIN sequence, (e.g. ExxxRxxxFxxNxxxIxxxN; where x represented amino acid variants). However, Cathepsin B-like protein lacks or contains only part of this sequence, as result of the shorter proregions of the Cathepsin B-like protein (Table 4). These autores also suggest that ERFNIN sequence serves to inhibit protein activity and that its removal convert the protein to the enzymatically active form (Table 5).

**Table 5** - The alignment of the sequences of the proregions of mammalian Cathepsins L and B

mcL	...MNEEGWRRRAVWEKNMKMIELHNQEYR (amino acids: 25-50..)
mcB	HDKPSFHPLSDDM I NY I NKQN (amino acids: 1-21.....)

Difference exist between the two protein sequences (ERFNIN) and short pro-region of Cathepsin B in comparison with that of the Cathepsin L. The data were on references of Cygle and Morb, 1997; Trysellius and Hultmark, 1997.

## CYSTATINS

The term cystatin refers to proteins that specifically inhibit the activity of papain and related cysteine proteinases (Cathepsin B, H and L, ficin, Bromelain). Their presence in microorganism, animal and plant species may be ubiquitous. These proteins are all related by structure and function to an inhibitor of cysteine proteinase, which was first described in egg white and was later called chicken egg white cystatin (Colella et al., 1989). Cystatins have been found to be evolutionarily related, forming the "cystatin superfamily". Members of the superfamily may be divided into three groups (or families) of proteins more closely related which comprise the animal cystatins (Barrett et al., 1986; Barrett, 1987) and one family from plant cystatins. The classification in families is based on primary sequence similarities, molecular masses, number of disulfide bonds and subcellular localization. Others families are suggested with bases in these aspects.

## FOUR FAMILIES OF CYSTATINS

Family-1 cystatins, commonly known as stefins family, contain no disulfide bonds or carbohydrate groups with molecular mass of around 11 kDa, the molecules consist of about 100 amino acid residues and are cytosolic proteinases (Machleidt et al., 1983; Sato et al., 1990).

Family-2 cystatin, or only cystatin family, are proteins that contain two intra-chain disulfide bonds near the carboxyl-terminus, are glycosylated with molecular mass about 13-24 kDa with approximately 115 amino acid residues. Family-2 cystatins are known to be secreted and contain a signal sequence (Abranhanson et al., 1987).

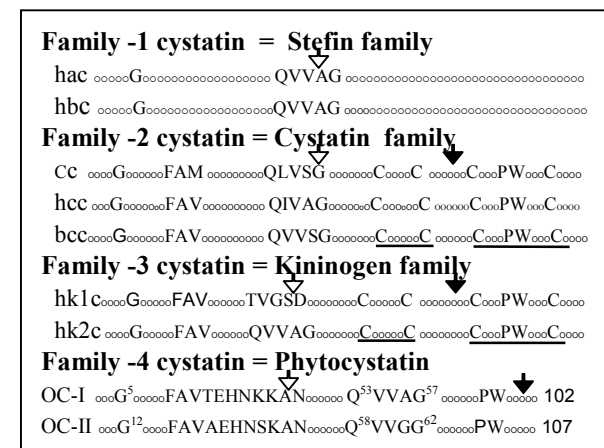
Family-3 cystatin or Kininogen family, comprise the blood plasm kininogens which are larger than the members of the two other families and are the most complex cystatin molecules with high molecular mass (60-120 kDa). These proteins are known to have domains in tandem that resulted from two duplications of genetic material of family-2 cystatins. Their composite sequences comprising about 355 amino acid residues that contain the bradykinin moiety, with additional

dissulfide bonds, are glycosylated molecules and are also secreted proteins. The Kininogens play an important role in blood coagulation process (Salvesen et al., 1986). Some family-2 and family-3 cystatins were reported to exhibit sequence homology with the active site sequence of Bowman-Birk type-trypsin inhibitor (Saitoh et al., 1991). Other cysteine proteinase inhibitors have their primary structure similar to those of the Kunitz-type soybean trypsin inhibitor family (Brzin et al., 1988). All the cystatins of which the primary structure have so far been elucidated contain the pentapeptide Gln-Val-Val-Ala-Gly (in most family-1 and -3 cystatins) and homologous sequences (e.g. Gln-X-Val-Y-Gly) are found in other cysteine proteinase inhibitors in family-2 cystatins (Barrett et al., 1986). The sequences, Phe-Ala-Val, localized near the carboxyl-terminus and the dipeptide Phe-Try, near the amino-terminal are conserved among family-2 but generally are absent in family-1 and family-3 cystatins. It has been speculated that this pentapeptide and dipeptide regions could be important for the binding to cysteine proteinase (Machleidt et al., 1989; Turk et al., 1997).

Family-4 cystatin, the phytocystatins (Turk and Bode, 1991; Turk et al., 1997), including almost all plant cysteine proteinase inhibitors. This family of proteins are known to have characteristics that are found commonly in most family-1 and -2 cystatins. Among plant cystatins, the oryzacystatin from rice seeds was the first inhibitor of plant origin considered one member of the cystatin superfamily that is related by structure and function to chicken egg white cystatin (Barrett et al., 1986). Cysteine proteinase inhibitors of the plant origin can be classified in two groups, one constituted by phytocystatins which possess single domain, comprised the majority of the phytocystatins (Abe et al., 1987a,b; Pernas et al., 1998) and the second only possess multiple domains, for example, multicystatin isolated from potato tubers (Walsh and Strickland, 1993) and cystatin of the tomato leaves (Bolter, 1993). Some researches showed that phytocystatins are inhibitory proteins that show non-competitive type inhibition Kinetic towards papain. These are corn cystatin I and oryzacystatin-I (Abe et al., 1994), soybean, L1 and R1 (Zhao et al., 1996), chestnut seed cystatin (Pernas et al., 1998), *Prosopis juliflora* seeds (Oliveira., 2000). Although, Barrett (1987) has related that cystatins

are reversible and competitive inhibitors of papain (Figure 1).

Phytocystatins have been identified in a variety of higher plants in both monocot and dicot species, such as rice (Abe et al., 1987a,b), maize (Abe et al., 1992), soybean (Hines et al., 1991; Botella et al., 1996), Wisteria (Hirashiki et al., 1990;), potato (Gruden et al., 1997), seedlings from sorghum (Li et al., 1996), leaves of the *Dianthus caryophyllus* (Kim et al., 1999), and apple fruit (Ryan et al., 1998). The molecular mass of the phytocystatins purified showed a range between 5 to approximately 87 kDa and stability to temperature and pH extreme (Abe et al., 1987a,b, 1992; Oliva et al., 1987; Abe and Whitaker, 1988; Pernas et al., 1998). The Table 6 shows several phytocystatins



investigated and their characteristics.

**Figure 1** - An alignment of the amino acid sequences of the four cystatin family. The sequences are those of human cystatin A (hac), human cystatin B (hab), chicken cystatin (cc), human cystatin C (hcc), beef colostrum cystatin (bcc); human Kininogen segment 1 (hk1c) and segment 2 (hk2c), oryzacystatin-I (OC-I) and oryzacystatin-II (OC-II). Positions of disulfide bonds in black line and the white dart and black dart indicated of the first and second intron position. The data are based on reference Barrett, 1987; Kondo et al., 1991; Mikkonen et al., 1996.

Oryzacystatin was the first well characterized plant cystatin, which could be classified as member of the family-1 and -2 cystatins. The characteristics common between oryzacystatin and the family-1 cystatins comprise: absence of the disulfide bond, pentapeptide constituted by Gln-Val-Val-Ala-Gly (residues 57-61) and molecular mass of approximately 11.5 kDa with 102 amino

acid residue. The comparison of the amino acid sequence of oryzacystatins showed approximately 30% identity with the family-2 cystatins and the presence of the regions constituted by Phe-Ala-Val (residues 29-31) and Phe-Try (residues 83-84), suggesting that it could be classified as a member this family (Abe et al., 1987 a, b; Abe and Arai, 1991; Arai et al., 1991). Not all cysteine proteinases inhibitors show sequence homology. Several cystatin have been isolated and sequenced, and some showed homology with serine proteinase inhibitor differing also in their effectiveness of inhibition of various cysteine proteinases (Brzin et al., 1988; Ishikawa et al., 1994). In potato tubers was found family of cysteine proteinase inhibitor, which represented a distinct group of proteins that could be belong to Kunitz-type soybean trypsin inhibitor family. They did not possess the conserved region (Gln-X-Val-Y-Gly) that usually

characterized the cystatin superfamily (Stubbs et al., 1990; Hatano et al., 1996, 1998). Bromelain inhibitor isoform isolated from pineapple stem by Lenarcic et al. (1992) showed to be a modest inhibitor for some cysteine proteinases including papain and bromelain. Their complete structure differed to those of other cystatins investigated due to the lack of the conserved region (Gln-X-Val-Y-Gly) that was generally present in the majority of the cystatins. Recently, eight bromelain inhibitor isoforms were isolated and their primary structures were not similar to cystatin superfamily. These inhibitor exhibited structural similarity to the Bowman-Birk trypsin/chymotrypsin inhibitor from soybeans seeds. These facts suggested that proteinase inhibitors might have a common ancestor (Hatano et al., 1996, 1998).

**Table 6** - Phytocystatins investigated and their characteristics.

SOURCE and REFERENCE	N	N. aa.	M.m. (kDa)	pI
<i>Helianthus annuus</i>	Sca	83	~9	5.6
Kouzuma et al., 1996	Scb	101	~11	9.6
<i>Persea americana</i>	(-)	100	~11	(-)
Kimura et al., 1995				
<i>Daucus carrot</i>	EIP18	133	18	(-)
Ojima et al., 1997				
<i>Oryza sativa</i>	OC-I	102	~12	5.3
Abe et al., 1987a,b				
<i>Vigna unguiculata</i>	(-)	97	10.7	(-)
Fernandes et al., 1993				
<i>Solanum tuberosum</i>	PMC	757	~87	(=)
Waldron et al., 1993				
(pineapple stem)	(-)	52	~5	(-)
Lenarcic et al., 1992				
<i>Glycine max</i>	(-)	245	26	(-)
Misaka et al., 1996				
corn seeds	(-)	135	18	(-)
Abe et al., 1992				
<i>Castanea sativa</i>	CsC	(-)	11.2	6.9
Pernas et al., 1998				
<i>Wisteria floribunda</i>	WCPI-3	(-)	~15.7	5.7
Hirashiki et al., 1990				
<i>Prosopis juliflora</i>	PjC	(-)	~20.5	(-)
Oliveira, 2000				

N, nomination; N.aa, number of amino acids; M.m., molecular mass; (-), data non-found; (=), domain-dependent. The molecular mass calculated by SDS-PAGE and/or filtration gel.

Several interaction model between the cystatins and their target cysteine proteinases were proposed that suggested three regions of the interactions: Gly residue near the amino-terminal, Gln-Val-Val-Ala-Gly central region conserved, and second hairpin loop near the carboxyl-terminal region. The data results of the X-ray crystallography did

not show the involvement functional of the disulfide bonds in the interaction (Barrett, 1987; Bode et al., 1988; Machleidt et al., 1983, 1989; Salmia et al., 1980; Stubbs et al., 1990). In order to determine which regions was responsible for the papain-inhibitory activity, Abe et al. (1988) and Arai et al. (1991) evaluated the effects of the wild

oryzacystatin and oryzacystatin truncated forms against papain activity. They observed that the primary region of interaction of the oryzacystatin responsible for the papain-inhibitory activity was the Gln-Val-Val-Ala-gly sequence and Pro-Try region could act also as a co-factor in the interaction, but the possibility that some other regions also contributed to inhibitory activity still remained (Table 7).

## FUNCTIONS ATTRIBUTED TO PHYTOCYSTATINS

Various biological functions have been attributed for phytocystatins: (a) Physiological functions of regulation of the endogenous proteinases activities in seeds during seeds maturation; (b) Participation of the inhibitors as important element in defence against attack by insects and nematodes that generally contain cysteine proteinases in their guts; and (c) Others functions not defined.

### (a) REGULATION OF ENDOGENOUS PROTEINASES

The phytocystatins participate in the control of endogenous cysteine proteinases during the maturation and germination of the seeds (Abe and Arai, 1991; Abe et al., 1992). Oryzacystatins are considered to be inhibitors of oryzains  $\alpha$ ,  $\beta$  and  $\gamma$ , cysteine proteinases that are produced during seed germination (Watanabe et al., 1991). Zeins and corn proteinases were inhibited by corn cystatin,

which suggested the involvement of the inhibitor in endogen defence mechanism (Steller, 1995). A proteinase prepared from germinating carrot seeds was inhibited by phytocystatin, named EICC, purified from seeds of carrot (Ojima et al., 1997). Two soybean cysteine proteinase inhibitors recombinants, named L1 and R1, were tested against plant cysteine proteinases from mungo bean cotyledons (Vicilin peptidohydrolase), and was observed that R1 had substantially more inhibitory activity than L1 (Botella et al., 1996). Phytocystatins and cysteine proteinases were also required to regulate the programmed cell death (PCD) (Solommon et al., 1999). This process is required for plant development and senescence. Plant programmed cell death or apoptosis has been implicated in several plant process such as xylogenesis (Cerós and Carbonell, 1993), some forms of senescence, in the pathogens attack response and stresses response (Greenberg, 1996; Lamb and Dixon, 1997). Recently cysteine proteinases was reported to regulate the expression of specific proteinase inhibitor genes in soybean cells (Groover et al., 1997) such as Sbcys, which was a specific soybean cysteine proteinase inhibitor (Abe et al., 1992).

The observations that endopeptidase activity of the Vicilin peptidohydrolase (cysteine endopeptidase from *Vigna radiata* cotyledons) occur in protein body and cytosolic localization of an inhibitory activity, has led to suggest to function in the cytoplasm protection against incidental rupture of protein body (Baumgartner and Chrispeels, 1976).

**Table 7** - Effects of the wild oryzacystatin and several oryzacystatin truncated forms against papain activity.

wOC-I (Wild oryzacystatin-I)	PPIA
Lacking 21 amino-terminal and 11 carboxyl-terminal amino acids	PPIA
Lacking 38 amino-terminal and 35 carboxyl-terminal amino acids	ACI
Substitution of Gln-Val-Val-Ala-Gly by <b>Pro</b> -Val-Val-Ala-Gly or <b>Leu</b> -Val-Val-Ala-Gly:	DR-PPIA
Substitution of Gln-Val-Val-Ala-Gly by Gln-Val- <b>Asp</b> -Ala-Gly	MPIA
Substitution of Gln-Val-Val-Ala-Gly by Gln-Val- <b>Gly</b> -Ala-Gly	E-PPIA

Wild oryzacystatin (wOC-I); PPIA, potent papain inhibitory activity; ACI, almost completely inactive; DR-, reduced drastically; MPIA, moderate papain inhibitory activity; E-PPIA, potent papain inhibitory activity equivalent to that of wOC-I. In boldface refer amino acid substituted by mutation. Data were obtained on references Abe et al., 1988; Arai et al., 1991.

## (b) PROTECTION AGAINST INSECTS AND NEMATODES

Several members of the Coleopteran order are tuber-, seed-, root- and leaf-eating insect. The presence of cysteine proteinase in the gut of these insects to hydrolyze ingested proteins, have implicated phytocystatin as proteins involved in the exogenous defense mechanisms (Abe et al., 1992; Gillikin et al., 1992; Waldron et al., 1993; Gruden et al., 1997). *In vitro* and *in vivo* inhibition of digestive proteinases from Coleopteran pests are summarized here that support this function.

Studies have shown that cysteine proteinases isolated from insect larvae can be inhibited by both non-proteinaceous and proteinaceous cysteine proteinase inhibitors. The effects of *in vitro* inhibition on the proteinases present in the guts from Coleopteran pests have been reported. Detection of the digestive proteinases present in the gut crude extract from *Acanthoscelides obtectus* were inhibited by 0.1mM E-64 [a specific non-proteinaceous cysteine proteinase inhibitor from *Aspergillus japonicus*, Trans-epoxisuccinyl-leucylagmatine], 20.0mM IAA (Iodoacetic acid) and 10.0mM NEM (N-ethylmaleimide) and 92%, 63% and about 7% inhibition of the cysteine proteinases present in the crude extract were obtained, respectively. The proteins partially purified from crude extract of these larvae were inhibited in 100, 100 and 20%, respectively in the presence of the same inhibitors (Wieman and Nielsen, 1988). Inhibitory activity against digestive proteinases from *Z. subfasciatus* larvae were measured with 0.028 mM E-64, 10mM NEM, 1.2mM IAA, 1.0mM DPT (dithiopyridine) and 0.075 mM cystatin that demonstrated 55%, 35, 29, 29 and 46% inhibition on crude extract midgut larvae, respectively. After partially purification of protein, the inhibition index obtained were 80, 75, 84, 43 and 57%, respectively (Lemos et al., 1990). Liang et al. (1991) analyzed the digestive cysteine proteinases present in the crude extract from *Sitophilus oryzae* and *T. castaneum* against E-64, PCMS (p-chloromercuriphenylsulfonic acid) and Antipain at  $10^{-3}$ M concentration that inhibited more than 95% from both crude extract. Luminal contents from *D. virgifera* were measured with the use of several inhibitors, which demonstrated that 0.02mM E-64, 1.0mM IAA and pCMB (p-chloromercuribenzoic acid), 0.01mM cystatin and 1.0mM  $\text{Cu}^{++}$  inhibited 30, 43, 39, 69 and 79% of

the luminal activities, respectively (Michaud et al., 1995). Orr et al. (1994) demonstrated maximal inhibition by E-64 to total activity in the gut extract from *D. undecimpunctata* and *D. virgifera* with 75 and 92% inhibition, respectively. Michaud et al. (1996) observed inhibition of the cysteine proteinases present in the *Otiorynchus sulcatus* crude extract by E-64 estimated at about 60% maximal inhibition. After elution of the crude extract in affinity chromatography used to remotion of oryzacystatin-sensitive proteinases using the GST/OCI fusion protein (GTS, glutathione S-transferase gene fusion system with oryzacystatin-I) preadsorbed onto GSH-agarose beads, the inhibitory activity decreased by 30%. The effects of the E-64 were available on crude extract from *L. decemlineata* and *O. sulcatus*. In order to determine the inhibitory activity 100 $\mu$ M E-64 was used to acquisition of 87 and 62% inhibitions against proteinases of the both insects, respectively (Michaud et al., 1996). The cysteine proteinases present in the crude extract from *D. virgifera* were inhibited in 25% by  $10^{-6}$ M E-64 and  $10^{-5}$ M E-64 inhibit 80% of the digestive proteinases from *L. decemlineata* (Zhao et al., 1996). The effects of E-64 were also analyzed against *T. castaneum* and *L. decemlineata* when 1.5 $\mu$ M of E-64 inhibited more than 50% of the digestive proteinases from both insects (Pernas et al., 1998). Studies on inhibition caused by oryzacystatin on the activity of the *S. oryzae* and *T. castaneum* showed that  $2 \times 10^{-5}$ M oryzacystatin caused 88 and 90% inhibition of the digestive proteinases, respectively (Liang et al., 1991). The crude proteinases extract of *L. decemlineata* and *O. sulcatus* were inhibited in 45 and 35% respectively by oryzacystatin-I. After application of the crude proteinases extract in affinity chromatography (GSH-GST/OCI) used to remove the oryzacystatin-insensitive proteinases, the inhibition dramatically decreased (Michaud et al., 1996). Irie et al. (1996) demonstrated that intrinsic oryzacystatin (genetically untransformed) inhibited 20% of the cysteine proteinases from *S. zeamais* insects. However, corn cystatin prepared from transgenic rice plants inhibited approximately 95%, when used in the same concentration. Orr et al. (1994) demonstrated the effects of PMC (multicystatin from potato tubers) on *D. virgifera* and *D. undecimpunctata*. Digestive enzymes from *T. castaneum* were inhibited by 1 $\mu$ M of the cystatin purified from



*Castanea sativa* seeds more than 50% inhibition, (Pernas et al., 1998). Recombinant soybean cystatin proteinase inhibitors, called L1, R1 and N2, showed substantial difference in inhibitory activities with about 10, 20 and 40% inhibition, respectively, to *L. decemlineata* when used in concentration of  $10^{-5}$ M. However, high inhibitory activities of the same proteins were observed to digestive proteinases from *D. virgifera* (30, 80 and 80%, respectively) used in concentrations of  $10^{-6}$ M. These results suggested that the differential inhibitory activities or specificity of the inhibitors could be due to motifs or residues that differed in the proteic structures (Zhao et al., 1996).

The effects of proteinase inhibitors on the digestive physiology of insect is an important area of study and their results are attractive for the biological control of insect pests. Studies relating the effects of proteinase inhibitors on insect diets have been developed long time ago (Bolter and Jongsma, 1995). Recently, several studies have reported the consequences of the incorporation of proteinase inhibitors in diets of insect. These proteinase inhibitors, when present in the diet, resulted in an increased mortality rate, decreased weight gain, reduced fecundity and in some cases led the death (Wolfson and Murdock, 1987; Murdock et al., 1988; Campos et al., 1989; Bolter, 1993).

*In vivo* deleterious effects of E-64 and proteinaceous cysteine proteinase inhibitors (phytocystatins) on growth and development in Coleopteran have been reported by various authors. E-64, when ingested by larvae of the *L. Decemlineata* (Wolfson and Murdock, 1987), *A. obtectus* Say (Wieman and Nielsen, 1988; Chen et al., 1992), *C. maculatus* (Campos et al., 1989; Kuroda et al., 1996; Murdock et al., 1988) showed adverse physiological effects on these organisms. Chen et al. (1992) and Kuroda et al. (1996) demonstrated the effects of the Oryzacystatin when ingested by *C. chinensis* and *T. castaneum*, respectively. Incorporation of PMC in diet of *D. virgifera* and *D. undecimpunctata* demonstrated that multicystatin from potato tubers (PMC), which incorporated in concentration of  $31.25 \mu\text{g}/\text{cm}^2$  diet for first instar southern corn rootworm produced maximal inhibition growth of approximately 70%, but increasing the concentration to  $125 \mu\text{g}/\text{cm}^2$  had no additional effect, suggesting that the larvae had a population of proteolytic enzymes not sensitive to

proteinaceous inhibitors that were able to generate sufficient amino acids to sustain limited growth (Orr et al., 1994). Recently Soyacystatin, cystatin from soybean, called soyacystatin N (scN) and L (scL), with 70% sequence identity, showed different insecticidal activities. The purified protein, scN, substantially retarded growth of first and second instar larval of cowpea weevil, *C. maculatus*, by 0.2% of inhibitor in bioassay, while scL affected neither insect growth nor mortality (Koiwa et al., 1998). Oryzacystatin-I (OC-I, wild-type) and variant form, named OC-IdeltaD86 in which residue Asp86 was deleted, both expressed and regenerated from tomato hairy roots, showed detrimental effect on growth and development of *Globodera pallida* (plant parasitic nematode) (Urwin et al., 1995). Both oryzacystatins were also expressed and regenerated in *Arabidopsis thaliana* showed effect on the size and fecundity of *Heterodera schachtii* and *Meloidogyne incognita* females (Urwin et al., 1997).

### (c) OTHER FUNCTIONS

The effects of the proteinaceous cysteine proteinase inhibitors has also been tested *in vitro* against serine proteinases and Cathepsins H, B and L too. Weak inhibitory effect from CSC was observed on trypsin and elastase (Pernas et al., 1998) and a cysteine proteinase inhibitory purified from cell cultures and seeds of carrot was weakly active against trypsin (Ojima et al., 1997). The results could be inferred to the fact that some family-2 and 3 cystatins exhibited some homology to the active site of the Bowman-Birk type-trypsin inhibitor (Saitoh et al., 1991; Watanabe et al., 1991), but homology was not found between proteins two classes.

Abe et al. (1991) observed that oryzains  $\alpha$  and  $\beta$  (papain-like enzymes) were preferentially inhibited by oryzacystatin-I (OC-I). However, oryzain  $\gamma$ , with similarity of cathepsin H-type, was natural target proteinase of the oryzacystatin-II (OC-II). On the other hand, Kondo et al. (1990) demonstrated that oryzacystatin-I (OC-I) showed significant inhibitory activity against papain and while oryzacystatin-II (OC-II) was more effective against cathepsin H, neither OC-I nor OC-II showed any effective inhibition against cathepsin B or cathepsin L.

## CONCLUSIONS

The informations reviewed here showed that the interactions between cysteine enzymes and cysteine inhibitor are an integral part of plant defence. This fact has opened up new researches with fundamental and practical applications that could be used in plant breeding to produce modified plant with proteinaceous inhibitors, especially with plant cystatins by DNA recombinant technology.

## RESUMO

Nesta revisão foram descritas definições, localizações, funções e exemplos de proteinases cisteínicas e suas proteínas inibidoras em animais vertebrados e invertebrados e plantas. Tratamos principalmente com aqueles inibidores que são relatados com o mecanismo de defesa da planta contra pestes. Em adição, comentamos sobre recentes trabalhos que contribuíram para uma melhor compreensão dos fatores envolvidos na interação específica proteinase cisteínica-cistatina. Por outro lado, chamamos atenção para o alto grau de afinidade e grande especificidade na interação que não são apenas representadas pela compatibilidade entre os resíduos de aminoácidos do sítio ativo envolvidos na catalise, mas também de todos os resíduos de aminoácidos que participam da interação enzima-inibidor.

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