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**PURIFICAÇÃO, CARACTERIZAÇÃO DE UMA LECTINA E COMPOSTOS
FENÓLICOS DA ENTRECASCA DE *Sebastiania jacobinensis* (MULL. ARG.):
EFEITO DE RADIAÇÃO GAMA SOBRE ESTRUTURA-ATIVIDADE PROTÉICA
E PROTEÇÃO DE FLAVONÓIDES ISOLADOS**

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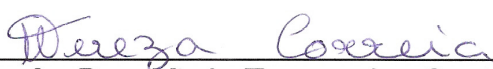
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“Purificação, caracterização de uma lectina e compostos fenólicos da entrecasca de *Sebastiania jacobinensis* (Muill. Arg.): efeito de radiação gama sobre estrutura-atividade protéica e proteção de flavonóides isolados”

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*"Uma rosa vermelha absorve todas as cores, menos a vermelha;
Vermelha, portanto, é a única cor que ela não é.
Essa Lei, Razão, Tempo e Espaço,
toda Limitação, cega-nos à Verdade
Tudo o que sabemos sobre o Homem, Natureza, Deus,
é apenas aquilo que eles não são;
é aquilo que rejeitam como repugnante."*

(Aleister Crowley, Liber CCCXXXIII, Cap. 40)

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Resumo

Uma lectina da entrecasca de *Sebastiania jacobinensis* (SejaBL) foi isolada por combinação de precipitação acetônica, fracionamento salino, cromatografia de troca iônica e gel filtração. A massa molecular de SejaBL foi determinada por SDS-PAGE e gel filtração, sendo de 52,0 kDa e 50,0 kDa, respectivamente. A lectina é uma glicoproteína com teor de carboidrato neutro de 6,94%, composta por duas subunidades com mesma massa molecular de 24 kDa. A lectina purificada hemaglutinou hemácias de coelho e humana. Glicoproteínas inibiram a hemaglutinação. SejaBL apresentou atividade máxima na faixa de pH 3,0-7,5, estabilidade térmica até 70 °C e é potencial inibidor da atividade da tripsina. Espectroscopia de fluorescência indicou a existência de superfícies hidrofóbicas. Lectina inibiu o crescimento micelial de *Fusarium moniliforme* e *Fusarium oxysporum* com um IC₅₀ de 123±0,5µg e 303±0,9µg, respectivamente. Organismos não alvos, larvas *Artemia salina* e embriões de *Biomphalaria glabrata*, não foram afetados, indicando baixa toxicidade ambiental. Como modelo experimental, a lectina purificada de *S. jacobinensis* (SejaBL) foi utilizada como prova de que mudanças no centro hidrofóbico, causados pela irradiação gama, alteram estruturalmente proteínas. Para elucidar o efeito dos radicais livres nas propriedades moleculares de SejaBL atividade, estrutura terciária (intrínseca e bis-ANS fluorescência), massa molecular e perfil cromatográfico foram examinados após γ-irradiação em várias doses (0,020 a 35kGy). Atividade hemaglutinante específica (AHE), na lectina irradiada, mostrou perda significativa ($p<0,05$) acima de 12,5kGy e aumento significativo ($p<0,05$) em 0,1, 0,8 e 1kGy. Espectroscopia de fluorescência indicou supressão da emissão de fluorescência quando excitado a 280 e 295nm, com elevada ligação do bis-ANS, após altas doses. SDS-PAGE e HPLC-RP mostraram fragmentação polipeptídica. Mudança significativa na superfície hidrofóbica indicou captação de hidrogênio radical em aminoácidos, causando desorganização estrutural da lectina. Flavonóides são produtos secundários de plantas com capacidade para capturar radicais livres. A lectina da entrecasca de *S. jacobinensis* (SejaBL), proteína capaz de ligar a carboidrato, tem sua superfície hidrofóbica modificada por radicais livres produzidos pela água radiólise. No presente estudo, o efeito protetor sobre a proteína pela estabilidade antioxidante de flavonóides da entrecasca de *S. jacobinensis* (SejaBF), após irradiação gama, foi investigada. Dois flavonóides foram isolados com rápida recuperação da capacidade para capturar radical mensurado pelo 2,2-difenil-1-picrilhidracilo radical (DPPH) a 0,5 mg/mL, após alta dose de irradiação. Os danos causados por irradiação na AH foi minimizado em 55% na presença dos flavonóides. Os resultados mostraram radioestabilidade da capacidade antioxidante de SejaBF, protegendo a lectina de radicais livres após irradiação.

Palavras-chave: lectina; bis-ANS; fluorescência; atividade antifúngica; toxicidade ambiental; radiação gama; flavonóides; capacidade de capturar radical DPPH

Abstract

A lectin from *Sebastiania jacobinensis* bark (SejaBL) was isolated using a combination of acetone precipitation, fractionation by ammonium sulphate, ion exchange and gel filtration chromatographies. The molecular mass of SejaBL was determined as approximately 52.0kDa by SDS-PAGE and 50.0kDa by gel filtration. The lectin is a glycoprotein with a neutral carbohydrate content of 6.94%, composed of two subunits with the same molecular mass of 24kDa. The purified lectin hemagglutinated rabbit and human erythrocytes. Glycoproteins inhibited agglutination of rabbit erythrocytes. SejaBL showed maximum activity over the pH range 3.0-7.5, heat stability up to 70°C and a potential trypsin inhibitory activity. Fluorescence spectroscopy indicated the existence of hydrophobic surface. Lectin inhibited the mycelial growth of *Fusarium moniliforme* and *Fusarium oxysporum* with an IC_{50} value of $123 \pm 0.5 \mu\text{g}$ and $303 \pm 0.9 \mu\text{g}$, respectively. Non-target organisms, *Artemia salina* Leach and embryo of *Biomphalaria glabrata*, were not affected, indicating low environmental toxicity. As an experimental model, the lectin purified from *S. jacobinensis* bark (SejaBL) was used as probe that changes on the hydrophobic center, by gamma radiation, structurally alter proteins. To elucidate the effect of free radicals in the molecular properties of SejaBL activity, tertiary structure (intrinsic and bis-ANS fluorescence), molecular weights and reversed phase chromatography were examined after γ -irradiation at various doses (0.020 to 35kGy). Specific Hemagglutinating Activity (SHA) in irradiated lectin showed significant loss ($p < 0.05$), above 12.5kGy, and a significant increase ($p < 0.05$) in 0.1, 0.8 and 1kGy. Fluorescence spectroscopy indicated suppression of emission intensity when excited at 280 and 295 nm, with high binding of bis-ANS, after high doses. SDS-PAGE and RP-HPLC showed polypeptide fragmentation. Significant change in the hydrophobic surface indicated hydrogen abstraction in amino acids, causing structural disorganization from the lectin. Flavonoids are ubiquitous plant secondary products with radical scavenger ability. The *S. jacobinensis* Bark Lectin (SejaBL), protein capable of binding to carbohydrates, has their hydrophobic surface modified by free radicals produced by water radiolysis. The present study shows the protective effect on protein and the antioxidant stability of *S. jacobinensis* bark flavonoids (SejaBF), after gamma irradiation. Two flavonoids isolated were present with fast scavenger ability measured by 2,2-diphenyl-1-picrylhydrazyl radical (DPPH) to 0.5mg/mL, after a high dose gamma radiation. The damage caused by irradiation on protein activity was reduced by 55% in the presence of the flavonoids. The results showed the radiostability of the antioxidant capacity of SejaBF, to protect the lectin from free radicals after irradiation.

Keywords: lectin; bis-ANS; fluorescence; antifungal activity; environmental toxicity, gamma radiation; flavonoids; DPPH radical scavenger capacity

LISTA DE ABREVIATURAS

AH: Atividade hemaglutinante

AHE: Atividade hemaglutinante específica

ALL: Lectina de musgo *Armillaria luteo-virens*

BSA: Albumina sérica bovina

Bis-ANS: 4,4'-Bis (1-anilidonaphthalene 8-sulfonate)

DTT: Dithiothreitol

Gal: Galactose

GlcNAc: N-acetilglicosamina

GalNAc: N-acetilgalactosamina

HA: Hemagglutinating activity

JCA: Lectina de *Artocarpus integrifolia* (Jacalina)

RIP: Proteína inativadora de ribossomos - tipo 2

HPLC-RP: Cromatografia líquida de fase reversa

SDS: Sulfato sódico de dodecila

SDS -PAGE: Eletroforese em gel de poliacrilamida contendo sulfato sódico de dodecila

SejaBL: Lectina da entrecasca de *Sebastiania jacobinensis*

SejaBF: Flavonóides da entrecasca de *Sebastiania jacobinensis*

SHA: Specific HA

TGL: Lectina da *tulipa gesneriana* L

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1. INTRODUÇÃO

A biodiversidade do Brasil abriga hoje entre 15% e 20% das espécies vegetais, animais e microrganismos do mundo. Apesar do crescente interesse internacional pela megabiodiversidade brasileira, sabemos que ela, por si só, não é garantia de crescimento econômico, tampouco de desenvolvimento sustentável. Mas espera-se que o País, além de importante exportador de matéria-prima, se converta no protagonista de uma nova economia mundial, baseado no uso sustentado da biodiversidade e dos seus recursos derivados, agregando valor para os diferentes setores produtivos (Ramalho, 1990).

Diferentes produtos utilizados pela sociedade possuem suas fontes na biodiversidade. São alimentos, fibras, itens farmacêuticos e químicos, óleos naturais e essenciais, entre outros. E sabemos que a geração de produtos de alto valor agregado a partir da biodiversidade está diretamente relacionada ao uso intensivo do conhecimento e do alto nível tecnológico. Este fato indica que a utilização de biotecnologia constitui uma opção viável e uma ferramenta fundamental para o uso sustentável e agregação de valor. O uso da biotecnologia implica na coleta de alguns poucos organismos ou parte deles, como fonte de matéria-prima para a prospecção e produção de novas biomoléculas, sendo assim uma estratégia adicional e efetiva de conservação da diversidade genética do meio ambiente (Wilson, 1988).

O controle de informações estratégicas, bem como das "tecno-ciências" que permitem agregar valor a essas informações – ao agregarem valor aos novos produtos e processos a partir daí gerados –, passa então a ocupar um dos centros de disputa e de conflito no jogo de forças políticas e econômicas internacionais. Tal controle pode ser exercido tanto com o domínio do acesso aos recursos da biodiversidade, quanto por intermédio de instrumentos de proteção de direitos à propriedade intelectual, seja sobre as modernas biotecnologias, seja sobre os conhecimentos tradicionais de populações locais (Sarita, 1998).

2. REVISÃO DA LITERATURA

2.1 Lectinas

No final do século XIX, começaram a acumular evidências sobre a existência na natureza de proteínas com habilidade para aglutinar eritrócitos. Tais proteínas foram chamadas fitoaglutininas, porque eram originalmente encontradas em extratos de plantas. A primeira citação a respeito se deu em 1888, quando Peter Hermann Stillmark estudando a toxicidade de extratos de *Ricinus communis* (mamona) observou a capacidade de aglutinar eritrócitos devido à presença de uma proteína extraída, a ricina. Pouco tempo depois, James B. Sumner isolou, da semente de *Canavalia ensiformis*, uma proteína cristalina denominada Concanavalina A (Sharon & Lis, 2004). No entanto, o estudo sobre essas proteínas só começou a ganhar foco em 1960, abrindo vasta área de aplicação (Gabor et al., 2004).

O termo lectina, originado do latim *legere* que significa escolhido, refere-se à habilidade dessas proteínas para reconhecerem e se ligarem reversivelmente a carboidratos ou glicoconjugados (Sharon & Lis, 1972). As lectinas diferem entre si quanto à seqüência de aminoácidos, requerimentos de metais, peso molecular e estrutura tridimensional e, ao contrário dos anticorpos, não são oriundas de resposta imune.

Lectinas estão amplamente distribuídas no meio ambiente, sendo encontradas em seres unicelulares (Sulagna et al., 2004; Lee et al., 2008) e animais pluricelulares (Dunphy et al., 2002; Haijie et al., 2006). Nos vegetais, os isolamentos são freqüentes, principalmente em sementes e entrecascas (Branco et al., 2004; Qiaojuan et al., 2005; Feng et al., 2006; Lan et al., 2008).

2.1.1 Peculiaridades

As lectinas apresentam grande diversidade estrutural, e o aspecto comum entre elas é a presença de, ao menos, um sítio específico de ligação a carboidrato, denominado "domínio de reconhecimento de carboidrato". Em sua plenitude, liga-se a carboidratos ou glicoconjugados em solução ou que estejam conectadas ao envoltório celular (William & Drickamer, 1996). As lectinas podem ser monoméricas como a arcelina de *Phaseolus vulgaris* L (Lorís, 1998) e, quando apresentam duas subunidades polipeptídicas, são

denominadas diméricas, com cadeias semelhantes ou diferentes entre si, ganhando, respectivamente, a denominação de lectinas homodiméricas ou heterodiméricas. *Erythrina speciosa* Andr. é uma lectina homodimérica com 27,6kDa por subunidade (Konozy et al., 2003). A lectina presente em *Ricinus communis* L é heterodimérica, com uma subunidade de 32kDa e outra de 34kDa (Frigerio & Roberts, 1998). Ocorrem outros tipos estruturais, como as triméricas, pentaméricas, hexaméricas ou poliméricas, como a lectina de *Artocarpus heterophyllus*, também denominada jacalina, com 65kDa (Kabir, 1998).

A presença de lectina na amostra pode ser facilmente detectada a partir de ensaios de hemaglutinação, as quais interagem com carboidratos da superfície celular, através de seus sítios de ligação, formando diversas ligações reversíveis entre células opostas (Santos et al., 2005). Para assegurar que o agente aglutinante é uma lectina, são necessários ensaios subsequentes de inibição da atividade hemaglutinante (AH), utilizando solução de distintos carboidratos (Kawagishi et al., 2001; Jayati et al., 2005).

Em relação às cadeias polipeptídicas, as lectinas são caracteristicamente ricas em aminoácidos ácidos e hidroxilados, associadas por interações hidrofóbicas, pontes de hidrogênio e, em alguns casos, pontes dissulfetos (Kennedy et al., 1995). As especificidades e afinidades dos sítios são alcançadas principalmente por pontes de hidrogênio, com o auxílio de forças de van der Waals e interações hidrofóbicas com resíduos de aminoácidos aromáticos que estão próximos às porções hidrofóbicas de monossacarídeos (Sharon & Lis, 2002), contribuindo para a estabilidade dos complexos formados.

A maioria das lectinas apresenta um complexo glicídico, ligado covalentemente à sua estrutura, perfazendo, em geral, 4-10% da massa total da molécula. A porção glicídica da lectina liga-se apenas à asparagina, serina ou treonina. Na asparagina pode ocorrer a ligação com GlcNAc em animais e vegetais (Helenius & Aebi, 2004).

Muitas lectinas requerem íons metálicos bivalentes em sítios específicos, que estimulam o reconhecimento do carboidrato pelo qual têm afinidade ao estabilizar a ligação por fixar as posições dos aminoácidos que interagem com o carboidrato. Ficou demonstrado, por exemplo, que a lectina de *Erythrina speciosa* Andr. necessita de Mg^{++} e Ca^{++} para capturar o carboidrato para o qual tem afinidade (Konozy et al., 2003).

Em 51 lectinas investigadas, há uma região conservada rica em aminoácidos hidrofóbicos, que forma uma cavidade na estrutura nativa. Esse sítio hidrofóbico possibilita interação entre as lectinas e grande número de compostos químicos distintos dos carboidratos como, por exemplo, a interação de compostos fenólicos, ligantes naturais dessa superfície hidrofóbica (Barre et al., 2002).

2.1.2 Classificação

2.1.2.1 Subdivisões das lectinas de acordo com a ocorrência e especificidade

De acordo com Sharon (Sharon & Lis, 2004), lectinas de plantas e animais demonstram características estruturais, composição da estrutura primária e nível de glicosilação distintos. Entretanto, ambas podem ser reunidas e classificadas com base na especificidade para carboidratos, estrutura molecular e ocorrência (Tabela 1).

Tabela 1 - Famílias de lectinas de plantas e animais

<i>Superfamília</i>	<i>Família</i>	<i>Especificidade</i>	<i>Lectina</i>	<i>Referência</i>
Lectina de vegetais	Lectina de leguminosas	Glicanos diversos	<i>Cratylia mollis</i> (Cramell 1)	Correia e Coelho, 1995
	Lectina de floema curcubitaceae	(GlcNAc) _n	<i>Urtica dioica</i>	Van Damme et al., 1998
	Jacalina relacionadas	Glicanos diversos	<i>Artocarpus integrifolia</i> (jacalina)	Van Damme et al., 1998
	Ligadoras de quitina (heveína, RIP tipo-2 e monocotiledôneas)	(GlcNAc) _n e Manose	Heveína do látex de <i>Hevea brasiliensis</i> e RCA.	Van Damme et al., 1998
	Amarantáceas	GalNAc	ACA	Van Damme et al., 1998
Lectinas tipo C	Selectinas	Sialyl-Lewis ^x	P-Selectina	Moore et al., 1992
	Colectinas	Ligante de Colagene	-----	Kawasaki et al., 1978
	Lectinas endocíticas	Asialoglicoproteína	-----	Sharon and Lis, 2004; Drickamer, 1988
	Pentraxinas			
Lectinas tipo P	Receptor manose-6P	Manose-6P	-----	Hoflack e Kornfeld, 1985
Lectinas tipo S	Galectina	β-galactosídeo	Galectin-1	Barondes et al., 1994
Lectinas tipo N	-----	Glicanos complexos	IL-2	Sherblom et al., 1989
Ig	Siglectinas*	Glicanos complexos	Sialoadesina (siglec-4)	Crocker et al., 1994

GalNAc, N-acetilgalactosamina; GlcNAc, N-acetilglicosamina; ^x NeuAcα(2-3)Galβ(1-4)[Fucα(1-3)]GlcNAc;

* Esta família de lectina Ig conjugadas ligantes de ácido siálico, um membro da Superfamília Ig.

2.1.2.2 - Subdivisões de lectinas vegetais de acordo com a estrutura

Segundo Van Damme et al (1996), as lectinas vegetais podem ser classificadas de acordo com a característica estrutural em:

- merolectinas - são monovalentes e, por isso, não precipitam glicoconjugados ou aglutinam células. São representadas pela família da heveína;
- hololectinas - possuem dois ou mais sítios de ligação a carboidratos que são idênticos ou muito semelhantes; esse grupo compreende as lectinas que são capazes de aglutinar células e/ou precipitar glicoconjugados;
- quimerolectinas - são proteínas com um ou mais sítios de ligação a carboidratos e um domínio não relacionado. Esse domínio diferente pode ter atividade enzimática bem definida, ou outra atividade biológica, mas age independentemente do domínio ligante a carboidrato. Assim representadas pelos grupos RIPs tipo 2 e quitinase de plantas tipo I;
- superlectinas - são proteínas com dois domínios de ligação a carboidratos. Esse pode ser considerado um grupo especial de quimerolectinas, consistindo de dois domínios estrutural e funcionalmente diferentes de ligação a carboidratos, como a lectina TGL da *tulipa gesneriana L.*, que são formados por dois domínios de ligação a carboidrato, que reconhecem D-manose e L-fucose, respectivamente.

2.1.3 Lectinas de plantas (famílias)

2.1.3.1 Lectinas de leguminosa

Esta família é encontrada exclusivamente em leguminosa (Sharon & Lis, 2002). Todas são homólogas no nível molecular, mas mostram variação na especificidade ao carboidrato. Essa ampla especificidade certamente contribui para o sucesso dessa família de lectinas como ferramenta biotecnológica. Algumas lectinas estão envolvidas na simbiose entre legumes e a fixação de nitrogênio da bactéria do gênero *Rhizobium* (Van Damme et al., 1998).

2.1.3.2 Lectinas do floema de Cucurbitaceae

Muitas espécies de Cucurbitaceae contêm alta concentração de lectinas que ligam a oligômeros de GlcNAc. As lectinas do floema de Cucurbitaceae são proteínas diméricas, compostas por duas subunidades idênticas de 24kDa. Todas as lectinas conhecidas dos floemas de Cucurbitaceae mostram grau de similaridade seqüencial, mas não têm qualquer similaridade com outras famílias de lectinas de plantas (Peumans & Van Damme, 1998).

2.1.3.3 Lectinas relacionadas à jacalina

São lectinas específicas à galactose de sementes da fruta jaca (*Artocarpus integrifolia*). Lectinas similares têm sido encontrados em poucas espécies de *Artocarpus* e em *Maclura pomifera* (Young et al., 1989).

2.1.3.4 Lectinas ligadoras de quitina

A proteína ligante de quitina (polímero de N-acetil-glicosamina) contém o chamado domínio(s) heveína(s). O termo “heveína” refere-se à pequena proteína de 43 resíduos de aminoácidos, encontrados no látex de árvores de seringueira, *Hevea brasiliensis* (Waljuno et al., 1975). As quitinases classe I, por exemplo, são quimerolectinas compostas por um único domínio heveína, ligado a um domínio catalítico com atividade quitinásica (Sharon & Lis, 2004). Proteínas inativadoras de ribossomo (RIP), tipo 2, são quimerolectinas compostas de um polinucleotídeo constituído por um domínio adenosina glicosidase (cadeia A), ligado ao domínio ligante a carboidrato (cadeia B). A maioria das RIP, tipo 2, liga-se preferencialmente a Gal ou GalNAc. A RIP, tipo 2, tem sido encontrada em plantas das famílias Euphorbiaceae, Fabaceae, entre outras (Peumans & Van Damme, 1998).

2.1.3.5 Lectinas de Amaranthaceae

Clonagem molecular e análise de difração de raios X revelaram que amarantina, uma lectina de semente de *Amaranthus caudatus* não se assemelha a qualquer outra lectina

de planta com relação à sequência de aminoácidos e à estrutura tridimensional (Transue et al., 1997). Com base nesses dados, a amarantina é agora considerada como um protótipo da família de lectina amarantina, uma lectina ligante específica de N-acetilgalactosamina (Van Damme et al., 1998).

2.1.4 Papel fisiológico e biotecnológico

Hipóteses sobre o papel fisiológico das lectinas foram formuladas devido à ocorrência dessa classe de proteínas em microrganismos, animais e plantas, mas admite-se que a função primária das lectinas seja a de reconhecimento de carboidratos. Lectinas vegetais desempenham importantes papéis tais como proteínas de reserva (Peumans & Van Damme, 1995), no mecanismo de defesa contra micro-organismos (Sá et al., 2008a), insetos, animais herbívoros (Sá et al., 2008b) e como mediadoras da simbiose planta-micro-organismo (Limpens & Bisseling, 2003).

Diferentemente, lectinas animais estão envolvidas com controle de qualidade, no processo de N-glicosilação e na rota de secreção de glicoproteínas (Helenius & Aebi, 2004), como também nos processos específicos no citoplasma e/ou compartimentos nucleares (Levebvre, 2001). Lectinas de animais atuam como receptores e desempenham um papel no tráfego extracelular de glicoproteínas, adesão de celular, interações celulares e imunidade inata (Leffler, 2004; Hsu & Liu, 2004). Quanto aos micro-organismos, as lectinas desempenham papel de simbiose, sendo responsáveis pela associação específica entre bactérias fixadoras de nitrogênio e leguminosas (Bohloul & Schmidt, 1974), assim como entre bactérias da flora intestinal e mucosa (Sulagna et al., 2004).

É grande a versatilidade biotecnológica de lectinas. A lectina de *Arum maculatum* que previne eficientemente a infecção de culturas de fibroblastos por *Chlamydia pneumoniae* tem perspectivas de aplicação clínica (Mladenov et al., 2002); a propriedade de ligação a células tumorais torna as lectinas potenciais ferramentas bioadesivas na entrega de drogas (Gabor et al., 2004; Ponchel & Irache, 1998). A lectina de *Cratylia mollis* imobilizada em Sepharose CL 4B, foi capaz de isolar a enzima lecitina colesterol aciltransferase (Lima et al., 1997). Certas lectinas são capazes de atuar sobre linfócitos, fazendo com que essas células passem de um estado típico para um estado de crescimento e

proliferação. A lectina de semente de *Cratylia mollis* (Maciel et al., 2004) é exemplo de lectina com atividade mitogênica.

Em ensaios histoquímicos e imunoistoquímicos, as lectinas têm sido utilizadas para detecção de resíduos glicosilados em superfícies teciduais (Paessens et al., 2007). Essas proteínas têm sido aplicadas na diferenciação de tecidos normais, displásicos e neoplásicos, fornecendo contribuições importantes para o diagnóstico e distinção entre câncer de próstata e a hiperplasia benigna (Basu et al., 2003). Atuações larvicida e inseticida têm sido citadas. A lectina do Cerne de *Myracrodruon urundeuva* mostrou-se excelente na atuação sobre estágios larvais de *Aedes aegypti* (Sá et al., 2008c) e a lectina da folha de *Bauhinia monandra* delimitou ampla ação inseticida (Macedo et al., 2006).

2.2 Radioatividade

Foi descoberta casualmente por Antoine Henri Becquerel em 1896, quando o mesmo realizava experiências com um mineral do urânio. Becquerel observou que esse mineral $[K_2UO_2(SO_4)_2 \cdot 2H_2O]$ era capaz de impressionar filmes fotográficos. Esses raios foram inicialmente denominados "raios do urânio". Em 1897, Marie Sklodowska Curie concluiu que era a quantidade de urânio que determinava a intensidade da radiação e, conseqüentemente, as mesmas (radiações) dependeriam do elemento (urânio). Em seguida, foi evidenciado que esse fenômeno se tratava de fenômeno atômico, ou, na realidade, fenômeno nuclear (Knool, 1979).

2.2.1 Radiação

O termo radiação vem do latim *radiare*, que indica um fenômeno básico em que a energia se propaga através do espaço, ainda que interceptada pela matéria. O termo irradiação vem do latim *in* e *radiare*, empregado para indicar o tratamento da matéria pela energia radiante. As radiações são produzidas por processos de ajustes que ocorrem no núcleo ou nas camadas eletrônicas, ou pela interação de outras radiações ou partículas com o núcleo ou com o átomo (Tauhata et al., 2003). Distinguem-se, assim, dois tipos de radiação: as chamadas corpusculares, feitas por intermédio de elétrons (raios beta), núcleos

de hélio (raios alfa), núcleos de hidrogênio (prótons; p. ou H1) ou nêutrons (n ou n1); e as eletromagnéticas, constituídas pelos raios de comprimento de onda muito curto, os raios X e os raios gama (Pitorri, 1993). Enquanto os raios X são produzidos por geradores especiais, os raios gama emanam espontaneamente de substâncias radioativas como rádio, tório e cobalto (Magalhães & Tauhata, 1994).

2.2.2 Medidas de intensidade das radiações ionizantes

A intensidade das radiações ionizantes é medida na base do número de íons que elas produzem num certo volume padrão. A unidade padrão é denominada Roentgem (R). No organismo humano, isso corresponde a cerca de duas ionizações por micron cúbico (Knool, 1979). Em linhas gerais, as várias unidades de medida das radiações podem ser assim definidas:

- Roentgen (1 R), aplicado em tecidos moles, causa a absorção, por partes desses, de quantidade de energia igual a 93 ergs por grama. Um erg representa a energia desenvolvida pela massa de 1 grama, movendo-se com a velocidade de 1 centímetro por segundo;
- Rad, unidade empregada para raios alfa e beta, em tecidos moles, 1 R pode ser aceito como equivalente a 1 Rad;
- Gray (Gy), unidade empregada a qualquer radiação ionizante, quando submetida a qualquer material, representa a dose absorvida por um material no ponto P. 1 Gy equivale a 100 Rad.

2.2.3 Dose e taxa de dose

Dose é a quantidade total de radiação emitida; taxa de dose é a maneira como essa dose é distribuída ao longo do tempo. Assim, uma mesma dose (digamos 100Gy), podendo ser aplicada durante diferentes períodos de tempo (1 minuto, 10 minutos, 100 minutos etc.) se apresentará com diferentes taxas (de 100Gy/min, de 10Gy/min, de 1Gy/min etc.), apesar de, em todos os casos, a dose final de radiação emitida seja a mesma (100Gy). Quanto maior a taxa e a dose, maior o dano (MS/S.N.V.S, 1992).

2.2.4 Efeitos biológicos gerais da radiação

É fácil compreender que a radiação ionizante pode agir sobre a célula e modificar a concentração de íons hidrogênio e o potencial de oxirredução de diferentes biomoléculas, alterando, profundamente, a funcionalidade molecular. As alterações químicas, decorrentes da radiação ionizante, são realizadas por dois mecanismos: a) por ação direta - na qual a molécula sofre alterações, tornando-se ionizada ou excitada pela passagem de elétrons ou ondas eletromagnéticas; b) por ação indireta - na qual a molécula não absorve energia, mas recebe, por transferência, energia de outra molécula (Kempner, 2001).

Os mecanismos pelos quais a radiação age sobre a célula são vários e, em parte, desconhecidos, no entanto os efeitos agudos possivelmente se devem à ionização da água. A água se decompõe e, como consequência, verifica-se a formação de compostos químicos ativos, que influenciam várias classes de biomoléculas (Tauhata et al., 2003). Os compostos que se formam são instáveis, de curta duração, mas seus efeitos podem ser profundos. Entre eles são citados ânions superóxidos, radicais peróxidos, radicais hidroxil e peroxinitritos. Essa oxidação afeta facilmente grupos protéicos, em especial o grupo sulfidril e sítios hidrofóbicos (Riley, 1994).

Em sistemas biológicos, os efeitos das radiações ionizantes diferem qualitativamente segundo a dose da radiação. Pequenas doses agem por ação indireta e produzem principalmente oxidações. Grandes doses agem por ação direta e indireta ao mesmo tempo. Vários fatores influenciam os efeitos radiobiológicos. Sendo especialmente importante: a) a intensidade da radiação (efeito reversível e irreversível); b) a temperatura de exposição c) a maneira da exposição, isto é, se simples, continuada ou fracionada; d) o tempo de exposição; e) presença de oxigênio (MS/S.N.V.S, 1992).

2.2.5 Aplicação biotecnológica da radiação gama

A aplicação de radiação gama para inativação de patógenos é evento crucial para iniciar a comercialização de produtos derivados de plasma humano em grande escala, sem o potencial risco de contaminação para receptores. Um método de inativação viral, usando irradiação gama em alta dose (45kGy), provoca a inviabilidade de todo vírus presente no

plasma e essa efetividade antiviral está diretamente relacionada ao dano no genoma viral (Miekka *et al.*, 1998; Reid, 1998; Hiemstra *et al.*, 1991). No entanto, a aplicação de radiação gama pode afetar a integridade estrutural e a atividade protéica. A irradiação foi inicialmente descartada como ferramenta para a inativação de patógenos no plasma e derivados, por causa da pobre recuperação protéica. Pesquisas, envolvendo a proteção de proteínas contra os efeitos das radiações ionizantes, têm sido desenvolvidas com uso de antioxidantes (Zbikowska *et al.*, 2006) e estratégicos modelos de exposição (Terry *et al.*, 2007).

2.3 Flavonóides

Sydler, em 1915, propôs o termo “farmacognosia” para o estudo de drogas e medicamentos de origem natural, a maioria deles de origem vegetal. Essa área da Farmacologia, como ciência, passou a ser subdividida em áreas mais específicas como, por exemplo, a farmacoquímica (fitoquímica) que estuda a origem, a síntese e as formas de extração dos compostos (Di Stasi, 1995).

Os compostos produzidos pelos vegetais são divididos em dois grupos: os metabólitos primários, tais como carboidratos, aminoácidos e lipídeos e os metabólitos secundários: compostos elaborados a partir dos metabólitos primários, tais como compostos fenólicos, terpenóides e alcalóides. São esses compostos os responsáveis pelos efeitos medicinais, ou tóxicos, das plantas, com grande importância ecológica, uma vez que podem atuar na atração de polinizadores ou representar defesa química contra estresse ambiental (Baladrin *et al.*, 1985).

Os flavonóides seguem uma estrutura básica de C₆-C₃-C₆ (dois anéis fenil - A e B - ligados através de um anel pirano - C) (**Figura 1**). São bastante comuns na natureza, atuando na atração de polinizadores, e como copigmentos das antocianinas (Bruneton, 1995). Apresentam atividades anti-inflamatórias, antimicrobianas e antitumorais (Barnes *et al.*, 2001), exercendo esses benefícios pelo seu poder antioxidante (Wollgast & Anklan, 2000).

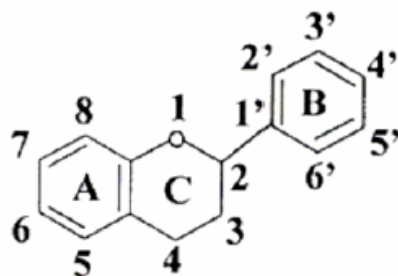


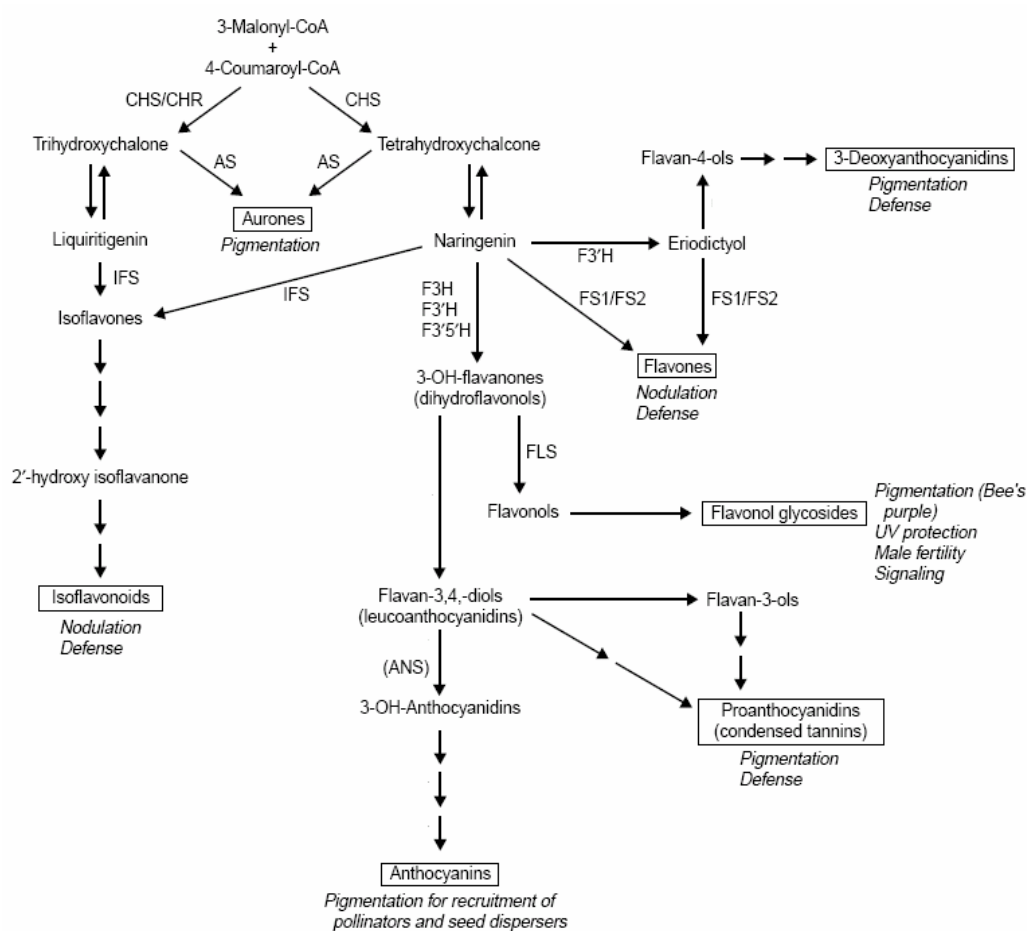
Figura 1 – Estrutura básica dos flavonóides (Bruneton, 1995)

Esses compostos têm origem biossintética mista, com parte da molécula proveniente da rota do ácido chiquímico e parte do ácido mevalônico, com estrutura química baseada no esqueleto 2-fenilcromano (Bruneton, 1995). Podem ser encontrados na forma livre ou conjugados a açúcares e, dependendo da substituição e do nível de oxidação no C₄, os flavonóides podem ser divididos em 14 classes (**Figura 2a**), tais como auronas, antocianidinas, flavonas, flavonóis, isoflavonóides e proantocianidinas (Harborne, 1984).

Flavonóides	Estrutura Básica
Auronas	
Antocianidinas	
Flavonas	
Flavonois	
Isoflavonóides	
Proantocianidinas	

Figura 2a - Diferentes classes de flavonóides segundo Bravo (1998)

Na biossíntese dos flavonóides, a primeira substância formada é uma chalcona, que está em equilíbrio enzimático com sua flavanona correspondente (**Figura 2b**). O par chalcona/flavanona é precursor das subclasses dos flavonóides. Diretamente deles, derivam as flavonas, os isoflavonóides e os dihidroflavonóides. Esses últimos são intermediários da síntese de catequinas, antocianidinas e flavonóis. As auronas derivam diretamente das chalconas (Winkel-Shirley, 2002). Os últimos passos na biossíntese dos flavonóides estão associados aos processos de O- e C- glicosilações, acilação dos açúcares e alquilações (Saleh, 1979). Estudos têm demonstrado a influência de alguns fatores ambientais na regulação gênica de flavonóides (Dooner, 1983).



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Figura 2b Esquema da via biosintética de flavonóides (adaptado de Winkel-Shirley, 2002)

2.4 *Sebastiania jacobinensis*

Segundo descrito em Flora Brasiliensis (2001), *Sebastiania jacobinensis* é uma árvore nativa, inerte, latexcente, possui copa pequena, irregular, semipersistente, de coloração verde-escura (**Figura 3a**). O tronco é geralmente curto, com casca externa rugosa e cor marrom-acinzentada (**Figura 3b**). Suas folhas são simples, alternas, glabras, de consistência coriácea, medem de 2 a 7cm de comprimento por 1 a 3cm de largura (**Figura 3c**). Apresentam geralmente ápice agudo, base cuneada, margem levemente serrada a dentado-crenada e pecíolo de até 7mm de comprimento. As flores são pequenas, amarelas, dispostas em espigas terminais de 3 a 6cm de comprimento (**Figura 3d**). Os frutos são cápsulas globosas com seis ângulos, de coloração parda, medem de 8 a 10mm de diâmetro e possuem 3 a 6 sementes em seu interior (**Figura 3e**). No levantamento bibliográfico não foi encontrado nenhum relato de investigação científica sobre essa espécie, deixando, desde já, aqui registrado, apenas aplicações fitoterápicas contra infecções e processos de hipersensibilidade.

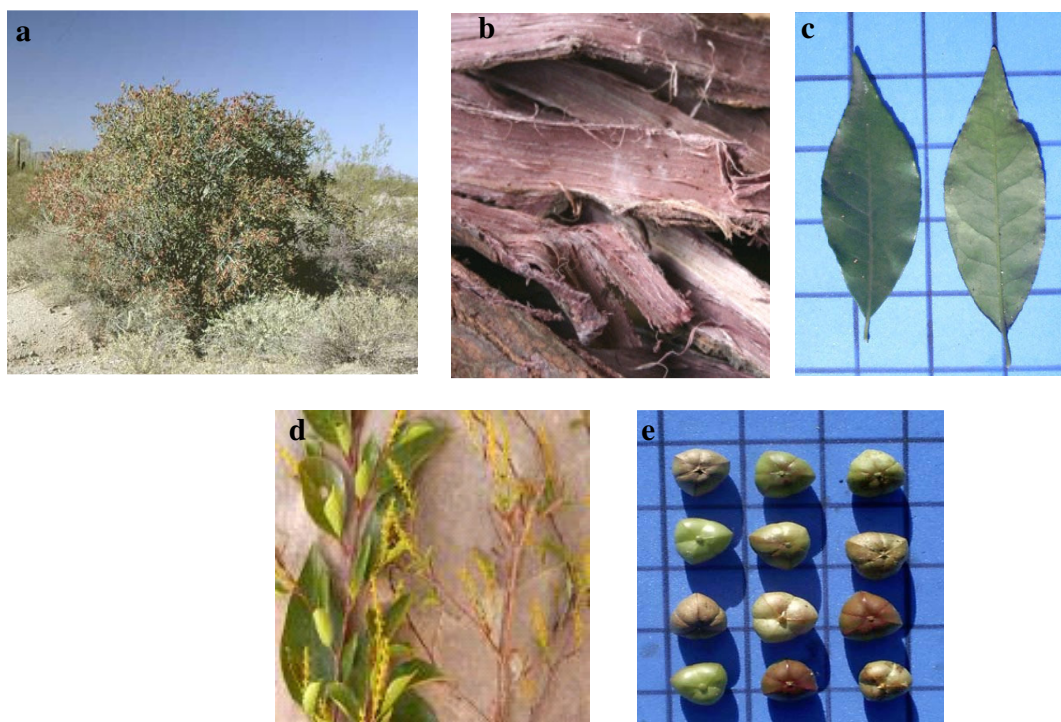


Figura 3 *Sebastiania jacobinensis*

3. OBJETIVOS

3.1 Objetivo Geral

Purificar, caracterizar uma lectina e compostos fenólicos da entrecasca de *sebastiania jacobinensis* (Muill. Arg.) e avaliar o efeito da radiação gama sobre a estrutura-atividade protéica sobre a proteção dos flavonóides isolados.

3.2 Objetivos Específicos

- Isolar a lectina, com homogeneizados da entrecasca de *S. Jacobinensis*, da melhor fração obtida pelo fracionamento utilizando cromatografia de troca iônica e filtração em gel;
- Caracterizar a lectina parcialmente purificada quanto à especificidade a carboidratos ou glicoproteínas, estabilidade térmica, especificidade para eritrócitos, estabilidade frente a variações de pH, influência de íons bivalentes e resistência a proteinases;
- Avaliar através de SDS-PAGE e Gel filtração a massa molecular, o padrão de pureza e o caráter glicoprotéico da lectina;
- Avaliar a atividade antifúngica e a seletividade ambiental “in vitro” da lectina purificada;
- Avaliar o efeito de radiação gama de alta taxa de dose na atividade e estrutura da lectina purificada;
- Isolar e caracterizar polifénóis da entrecasca de *S. jacobinensis* baseado na sua atividade antioxidante, no tempo de retenção cromatográfico e massa molecular;
- Avaliar a proteção da lectina purificada e a radioestabilidade contra radiação ionizante do antioxidante (i.e. flavonóides isolados).

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Manuscript

A nontoxic lectin from *Sebastiania jacobinensis* with antifungal activity

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Abstract

A lectin from *Sebastiania jacobinensis* bark (SejaBL) was isolated using a combination of precipitated acetone, ammonium sulphate fraction, ion exchange and gel filtration chromatographies. The molecular mass of SejaBL was 52.0 and 50.0 kDa by SDS-PAGE and gel filtration, respectively. The lectin was a glycoprotein with a neutral carbohydrate content of 6.94% composed of two identical subunits of 24 kDa. The purified lectin hemagglutinated both rabbit and human erythrocytes and this activity was not by monosaccharides. However, glycoproteins inhibited agglutination of rabbit erythrocytes. SejaBL showed maximum activity over the pH range 3.0-7.5, heat stability up to 70 °C and a potential trypsin inhibitor. Fluorescence spectroscopy indicated the existence of hydrophobic surface. Lectin inhibited the mycelial growth of *Fusarium moniliforme* and *Fusarium oxysporum* with an IC₅₀ value of 123 ± 0.5 and 303 ± 0.9 µg, respectively. Nontarget organisms, *Artemia salina* Leach and embryo of *Biomphalaria glabrata* were not affected indicating low environmental toxicity. Alternative viewpoints are presented that can hopefully help in future efforts to develop safer and more effective microbial control agents.

Keywords: Lectin; Bis-ANS; Fluorescence; Antifungal activity; Environmental toxicity

Introduction

Plant lectins are a heterogeneous group of proteins or glycoproteins that share the capacity to identify a specific carbohydrate. Their widespread distributions in the plant kingdom suggest a physiologically important function [1]. They have attracted great interest because of their various biological activities, as antiproliferative, antitumor, antifungal and antiviral properties [2]. Seeds, especially of leguminous species, are common sources of lectins, but they are also present in rice vegetative tissues, latex and bark of different species [3-6]. Intrinsic fluorescence and Bis-ANS have been used in folding, stability studies and as evidence of conformational change in proteins, by assessing hydrophobic regions [7,8].

The *Artemia salina* Leach is a salt-water crustacean used as food for fish. *A. Saline* is sensitive to the effect of active substances and its mortality is used to monitor toxicity because it is well sensitive to many chemical substances [9]. The *Biomphalaria glabrata* (Say, 1818) is the clam of the Planorbidae family with the widest distribution in Brazil. The antibiotic effect of drugs on the embryo development of *B. glabrata* and mortality of *A. Saline* have urged their use as a way to monitor the environmental impact and selectivity of microbiological control agents [10,11]. Antifungal proteins have been isolated from a large number of plants separated into many types comprising thaumatin-like proteins [12] chitinases and β -1.3-glucanases [13], thionins [14], plant defensins [15], ribosome-inactivating proteins [16], protease inhibitor-like proteins [17] and lectins [18].

Sebastiania jacobinensis Müll. Arg. (Euphorbiaceae family) is a common tree found in the tropical regions of Brazil. The bark of this plant is popularly used against infections and hypersensitivity processes. In view of the benefits that microbiological control agents with

low environmental toxicity provide to vegetable biotechnology, we describe the isolation and partial characterization of an antifungal lectin from the bark of *S. jacobinensis*, with ecotoxicological profile.

Material and methods

Purification lectin. *S. jacobinensis* bark was homogenized overnight at 4 °C in 10 mM Tris–HCl buffer (pH 8.5) with 0.2% (v/v) Triton X-100 detergent. The homogenate was centrifuged 5,000 g for 20 min (crude extract) followed by lyophilization. In brief, two volumes of cold acetone were added, mixed well and kept on ice for 10 min. The solution was centrifuged at 10,000 g for 20 min at 4 °C. The supernatant was concentrated under vacuum until total evaporation of the acetone. The residual aqueous solution obtained was precipitated with 60% (NH₄)₂SO₄. The precipitate was dialyzed against the Tris–HCl buffer. This fraction was applied to a CM-Cellulose column (2 x 20 cm) previously equilibrated with the Tris–HCl buffer into 150 mM of NaCl. Following removal of the unadsorbed proteins, the column was eluted with 1 M acetic acid. The largest fractions of absorbance eluted with acetic acid were pooled, dialyzed and concentrated by ultrafiltration (Amicon ultra-15, Mr 10.000 cut-off) and chromatographed on a Sephadex G-100 column (2 x 70 cm) equilibrated with 150 mM NaCl. The column was eluted with the same solution at a flow rate of 0.5 mL min⁻¹. The primary unadsorbed fraction (SejaBL), after concentration by ultrafiltration in 10 mM phosphate buffer (pH 7.0), was applied to a Superdex 75 HR 10/30 column coupled to an ÄKTA purifier system (Amersham Pharmacia Biotech). The column was preequilibrated and eluted with 300 mM NaCl, at a flow rate of 0.5 mL min⁻¹, monitored by absorbance at 215 nm, collected in 1-ml fractions. The lectin was submitted

to reverse-phase chromatography on a C-18 column performed on a HPLC system (Shimadzu) and monitored at 215 nm, as described below for chain separations. The total protein content from the crude extract and the purified lectin were determined by the Lowry *et al.* method [19], using BSA standard curve.

Hemagglutination activity and specificity sugar. Hemagglutinating activity (HA) was evaluated as described by Correia and Coelho [20] defined as the lowest sample dilution showing hemagglutination. Specific HA (SHA) corresponded to the relation between HA and protein concentration. The carbohydrate binding specificity of the lectin was determined by HA inhibition using several sugars (D-glucose, N-acetil-D-glucosamine, D-arabinose, D-mannose, L-fucose, L-raphinose, D-galactose, L-threulose, D-xylose, D-saccharose, L-rhamnose, L-cellobiose, D-lactose, methyl- α -D-mannopyranoside and methyl- α -D-glucopyranoside) and glycoproteins (bovine serum albumin, casein, tyroglobulin, ovalbumin, fetuin and asialofetuin) as described by Coelho and Silva [21].

Effect of pH, temperature and metal ions. The effect of pH on SejaBL HA was evaluated by incubating the lectin (0.5 mg mL^{-1}) at different pH values for 1 h at room temperature in selected buffers (10 mM citrate phosphate buffer, pH 2 – 7 and 10 mM tris-hydrochloric acid buffer, pH 8 - 12). The heat stability was determined by incubation of lectin solution at different temperatures (30-90 °C for 30 min and 100 °C, 30 to 90 min) and remaining SHA was determined. The effects of Mg^{2+} , Zn^{2+} and Ca^{2+} were determined by incubation at same volume of any metal ion (5, 10 and 20 mM) in 150 mM NaCl and lectin. An aliquot (50 μL) of the mixture was distributed in microtitre plate wells and the HA was proceeded as described for carbohydrate inhibition assay.

Molecular weight determination. Chain analyses were performed after disulfid bridges split by reduction and alkylation. Previously lyophilized SejaBL samples were reduced by the Friedman reaction [22] with some modifications as follows: lectin (1.5 mg) was dissolved in 250 μ L 50 mM Tris-HCl pH 8.6, 6 M urea, 10 mM EDTA, 179 mM DTT and incubated for 3 h at 37 °C, in the dark, before N₂ purging. The free sulphydryl groups were exposed to 100 μ L iodoacetate and the reaction was continued for another 2 h in the same initial conditions. The iodoacetate derivative chains were desalted and separated on a reverse-phase C-18 column (Vydac-protein peptide ultrasphere) performed on a HPLC system (Shimadzu LC-10AD-kyto, Japan) and monitored at 215 nm. The column was equilibrated with solvent A (0.1% TFA in H₂O) and eluted using solvent B (90% acetonitrile: 10% H₂O: 0.1% TFA) non-linear gradient in A - B = 5% at t = 5 min; B = 70% at t = 27 min; B = 80% at t = 60 min and B = 100% at t = 69 min. SDS-PAGE was carried out according to Laemmli [23] and acid gel electrophoresis for native proteins was performed with Davis' system [24]. SDS-PAGE on reducing conditions was made after Friedman reaction. The broad range standard marker proteins were purchased from Sigma Chemical Co. (St. Louis, MO, USA). Glycoproteins were detected by the periodic acid-schiffs (PAS) [25] and estimation of carbohydrate content from the protein samples was done by the phenol sulphuric acid method [26], with a curve of D-mannose as standard.

Fluorescence spectroscopy. Intrinsic fluorescence emission of lectin solution (0.2 mg ml⁻¹ in 10 mM phosphate buffer, pH 7.0) was measured at 25 °C using the spectrofluorimeter (JASCO FP-6300, Tokyo, Japan) in a cuvette (1-cm pathlength rectangular quartz). The excitation wavelength was 295 and 280 nm. The emission spectra were recorded at a range of 305 to 450 nm. The band passes were 5 nm. The hydrophobic surface was determined using the same condition as intrinsic fluorescence. The samples were transferred to the

cuvette and then mixed with 5 μ M bis-ANS (Molecular Probes Inc., USA) and the fluorescence measured. The fluorescence emission obtained was of 400 to 600 nm with excitation at 360 nm [27].

Resistance proteases assay and trypsin inhibitory activity. Sensitivity of SejaBL to protease was developed according to method of Rios et al. [28]. 5.0 μ g of α -chymotrypsin and trypsin at 0.1 M Tris-HCl buffer (pH 8.2) were preincubated for 3 h with 50 μ L of SejaBL (0.5 mg ml⁻¹). After, the reaction was stopped with phenylmethyl sulfonyl fluoride (Sigma). The sensitivity to protease was confirmed by lectin HA. The determination of trypsin inhibitory activity was carried out according to the method of Lee and Lin [29] by inhibition of hydrolysis of N-Benzoyl-L-arginine-4-nitroanilide (Sigma) at 0.1 M Tris-HCl buffer (pH 8.2) catalyzed by trypsin. Four determinations were averaged for trypsin inhibitory activity and expressed as μ g trypsin inhibited.

Antifungal activity. Assay of antifungal activity and quantitative assay to determine the IC₅₀ were performed as described by Wang and Ng [30]. The selected fungi, i.e.: *Trichoderma viride* (URM-3344), *Fusarium oxysporum* (URM-2489), *Colletotrichum gloeosporioides* (URM-4911), *Fusarium moniliforme* (URM-3226), *Aspergillus niger* (URM-5238) and *Candida albicans* (URM-4388), stock obtained from the Federal University of Pernambuco fungal cultures, were carried out in 60 mm – 15 mm petri plates containing 7 mL of potato dextrose agar (PDA). After the mycelial colony had developed, sterile blank paper disks (0.625 cm in diameter) were placed at a distance of 0.5 cm away from the border of the mycelial colony. A SejaBL aliquot (20 μ L with 300 μ g) was added to the disk. Positive control was Cercobim (2 μ g) and negative control (150 mM of NaCl). The plates were incubated at 23 °C for 72 h until mycelial growth had involved the disks containing the

negative control and had formed crescents of inhibition around disks containing samples with antifungal activity.

To determine the IC_{50} value, three concentrations of SejaBL (75, 150, and 300 μg) were added, separately, to three aliquots each containing PDA (4 mL) at 45 °C, mixed quickly and poured into three separate small petri dishes. After the agar had cooled down, a small amount of mycelia (same amount to each plate) was added. Buffer only without SejaBL served as control. After incubation at 23 °C for 72 h, the area of the mycelial colony was measured and the inhibition of fungal growth determined.

Environmental Toxicity. *A. salina* (Brine Shrimp) encysted eggs (25 mg) were hatched in the becker filled with seawater under artificial light at 30 °C, pH 8–9 in constant aeration. After 28 h the nauplii were collected with a pasteur pipette macroscopically and counted in the stem of the pipette against a lighted background. Nauplii were transferred to test tubes containing the samples. The lectin concentration ranged from 10 to 1000 $\mu\text{g mL}^{-1}$ in vials containing 5 mL of seawater. Fifteen shrimp nauplii were added to each vial (45 shrimps per concentration). The plates were maintained under illumination. Survivors were counted after 24 h of incubation and the percentage of deaths at each dose and control was determined. The bioassay was performed as described previously by Meyer et al. [31] and developed with minor modifications. *B. glabrata* embryos (0 to 15 h after spawning) were exposed for 48 h to SejaBL (10 to 200 $\mu\text{g mL}^{-1}$) and the embryo development was followed for an additional 10 days in absence of exposure. The assay was carried out according to Oliveira-Filho and Paumgartten [32].

Statistical analysis. Data are represented as mean \pm S.E.M. The antifungal activity IC_{50} values and their 95% confidence intervals (CI 95%) were obtained by nonlinear regression using the GraphPrism® (GraphPad Software Inc., San Diego, CA, USA). LC_{50} on brine

shrimp and *B. glabrata* embryos was obtained from 24 and 48 h respectively; the percentage of deaths were estimated using the method described by Finney [33].

Results and Discussion

The lectin was best extracted with 0.2% Triton X-100 present in the solution. Only one peak bioactivity was detected per acetic acid elution at the CM-Cellulose column (Fig. 1A) and into Sephadex G-100 column (Fig. 1B) with SHA of 8192 (Table 1). Superdex-75 gel filtration (Fig. 1C) showed apparent molecular weight of 50 kDa. On the C-18 column (Fig. 1D) the lectin was eluted about 70% of acetonitrile. SejaBL is a single band of 52 kDa and under reducing conditions, analyzed by SDS-PAGE (Fig. 2A;B), showed dimeric protein composed of two subunits with the same molecular mass of 24 kDa and another fragment of less than 6 kDa. After desalting on the C-18 column, the two chains showed different mobility (Fig. 2C). This suggests that the chains are covalently connected by disulfide bonds and have no homology. In general, molecular mass of subunits back lectin is around 30 kDa, with the fully active lectin forming either dimers or tetramers. Molecular mass ranging from several kilodaltons to about 67 kDa has been reported for antifungal proteins [34-36].

The lectin is an acidic protein (Fig. 2D) with pH stability ranges of 5-8 (Fig. 3A). The lectin band was weakly stained by PAS, and positive phenol-sulphuric acid showed a glycoprotein with 6.94% covalently linked carbohydrate. The native active form was heat-stable up to 70 °C. A soft decrease in its activity was observed when temperature was raised from 70 to 100 °C (Fig. 3B). The loss of hemagglutinating activity with increasing temperature is evidently due to heat-induced denaturation of the lectin. The relative ease

with which SejaBL can be destroyed by heat enables its use as microbial control agents in cereal. SejaBL agglutinated erythrocytes from rabbits and humans of all blood groups except B. Fetuin, asialofetuin, bovine serum albumin and casein inhibited SejaBL induced rabbit erythrocyte agglutination, whereas monosaccharides did not inhibit its activity. Ions did not affect SejaBL HA and the lectin did not suffer apparent hydrolysis front proteases keeping HA. However, SejaBL had a dose-dependent inhibition against trypsin from the porcine pancreas (Table 2). Inhibitor-like proteins are generally found in vegetative tissues, believed to play a role of defence against predators [17].

The intrinsic protein fluorescence spectra of SejaBL (Fig. 4) revealed one major peak at 330 nm, indicating the presence of tryptophan in a highly hydrophobic region. Interestingly, bis-ANS fluorescence observed in SejaBL when compared only with buffer (i.e., the control) indicates hydrophobic surface turned to the solvent (Fig. 4). Prasad *et al.* [37] suggested that the immediate increase in the bis-ANS fluorescence intensity observed in proteins is due to the binding at primary hydrophobic site(s). This demonstrates the hydrophobic character of the protein confirmed by tryptophan and tyrosine intrinsic fluorescence. Steadman *et al.* [38] noted that a hydrophobic surface produced decreasing stability and increasing structural alteration in protein after chemical or physical stress. The lower agglutination activity, at the basic pH values and high temperature may be due to the hydrophobic character of SejaBL.

Among the fungi tested, SejaBL exerted antifungal action against *F. moniliforme* and *F. oxysporum*. The IC₅₀ values of its antifungal activity towards *F. moniliforme* and *F. oxysporum* was, respectively, 123 ± 0.5 and 303 ± 0.9 µg (Table 3), similar to those of previously reported for antifungal proteins [39,40], and some are linked to protease inhibitory activities [41,42]. The species of *Fusarium* sp are pathogens to banana, corn and

rice. Damage produced by these fungi cause loss of cereals with harmful effects on human health and agriculture.

The lectin showed percentage of *A. salina* deaths with LC_{50} of at $715.89 \pm 1 \mu\text{g ml}^{-1}$ (Table 3). Lectin did not present toxicity in blastulae and embryo development of *B. glabrata*. Lethality was not observed until $200 \mu\text{g ml}^{-1}$ (Table 3). This data suggests that the antifungal activity is unrelated to the lytic action or membrane instability. The inhibitory effect on embryo development was not observed since the first cleavage, and the cells presented a nuclear organization and homogeneous cytoplasm, with no DNA or protein synthesis in injury [43]. SejaBL, compared with the reference chemical fungicides, presents a higher degree of selectivity, does not affect the conservation and access to biological diversity and does not involve environmental contamination.

Conclusion

Some *Fusarium sp* species produce mycotoxins in cereal crops that can affect human and animal health [44,45]. The isolation of a plant lectin with inhibitory activity toward fungi and without environmental toxicity has important application in vegetable biotechnology as a promising biological control agent to genetically modified plants.

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Table 1

Table 1

Yields and hemagglutinating activities of various fractions (from 100 g meal of *S. jacobinensis* bark)

Sample	Yield (mg)	Specific hemagglutinating activity (units/mg)	Total hemagglutinating activity (units)	Recovery of hemagglutinating activity (%)	Folds of purification
Crude extract	14400	284.4	4.09×10^6	100	1
$(\text{NH}_4)_2\text{SO}_4$ fraction	1323.2	3951	5.22×10^6	127.67	13.9
CM-Cellulose	96	6827	6.55×10^5	16.01	24
Sephadex G-100	20	8192	1.63×10^5	4	28.8

Table 2

Table 2 Inhibition of purified SejaBL against trypsin

µg of SejaBL	0.5	1.25	2.5	5	7.5	12.5
% of inhibition	8.5	10	17	28	39	56.5

The assay for the inhibition of trypsin was carried out using 5µg porcine pancreas trypsin.

Table 3

Table 3 Linearity value for the antifungal activity and acute environmental toxicity

	Antifungal activity	IC ₅₀ ± S.D.	Regression equation	Coefficient of determination (r^2) ²
A	<i>F. moniliforme</i>	123 ± 0.5 µg	y = 0.1955x + 4.386	0.992
	<i>F. oxysporum</i>	303 ± 0.9 µg	y = 0.0741x	0.99
	Toxicity environmental	LC ₅₀ ± S.D.	Regression equation	Coefficient of determination (r^2) ²
B	Embryo of <i>B. glabrata</i>	nontoxic	—	—
	<i>A. salina</i> leach	715.89 ± 1 µg/mL	y = -0.1799x + 178.79	0.983

Figure

Figure captions

Fig. 1. (A) Purification by ion-exchange (40 mg of protein) was applied at a flow rate of 0.3 mL min⁻¹ and 2 mL fractions were collected. Arrows indicated elution with Tris-HCl buffer (1), followed by 1 M of acetic acid (2). Absorbance at 280 nm (- -); log of HA (-●-). (B) Gel filtration on Sephadex G-100 equilibrated and eluted with NaCl 150 mM at 0.5 mL min⁻¹ collected on 1-mL fractions. Absorbance at 280 nm (- -); log of HA (-●-). (C) Superdex 75 HR 10/30 column coupled to an ÄKTA purifier system equilibrated and eluted with NaCl 300 mM at 0.5 mL min⁻¹ collected on 1-mL fractions. (D) Reverse phase chromatography in C-18 column HPLC system. The column was equilibrated with solvent A (0.1% TFA in H₂O) and used a non-linear gradient elution (5-100%) with solvent B (90% acetonitrile: 10% H₂O: 0.1% TFA).

Fig. 2. (A) Molecular weight of SejaBL on a 10% SDS-PAGE. MW, molecular weight. (B) On reducing conditions. (C) Acid native protein. (D) Chains separations after desalting on the C-18 column.

Fig. 3. Effects of temperature and pH on SHA of SejaBL. SHA in NaCl 150 mM with rabbit erythrocytes was 3.01.

Fig. 4. Intrinsic fluorescence emission of SejaBL, excitation at 280 nm () and 295 nm (●). Bis-ANS fluorescence emission, excitation at 360 nm (■).

Figure

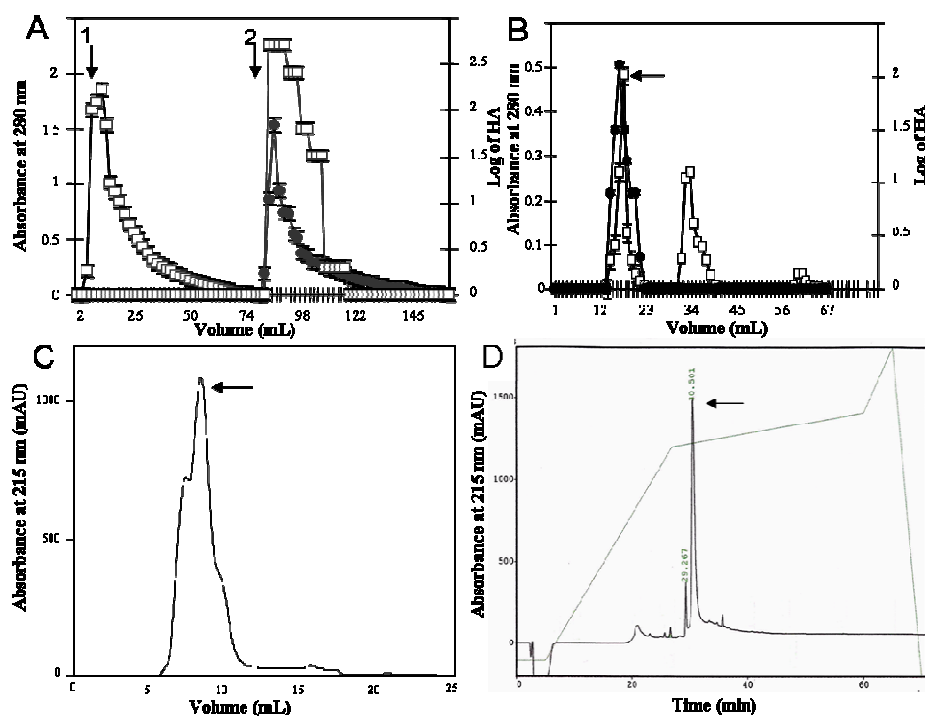


Figure 1

Figure

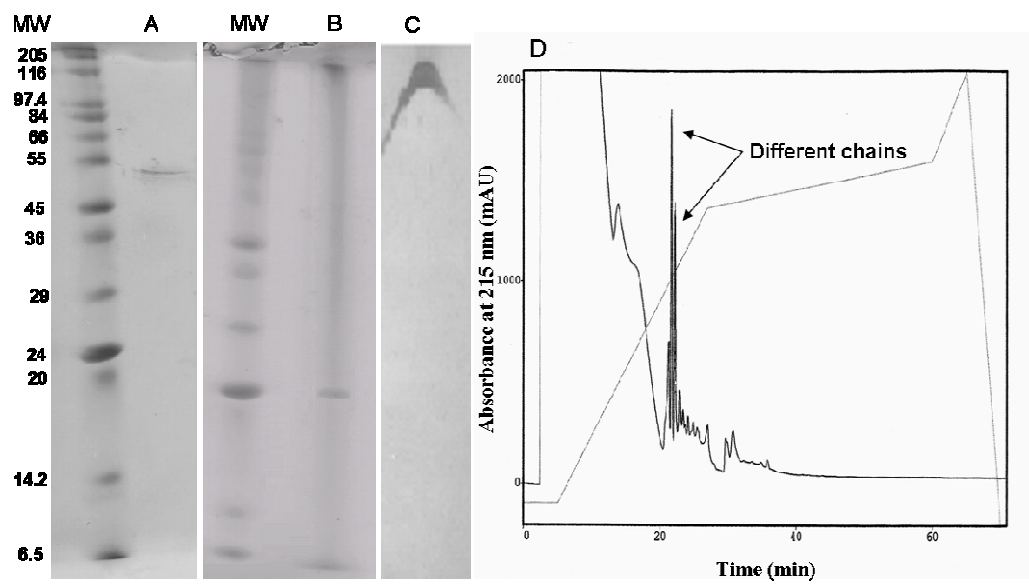


Figure 2

Figure

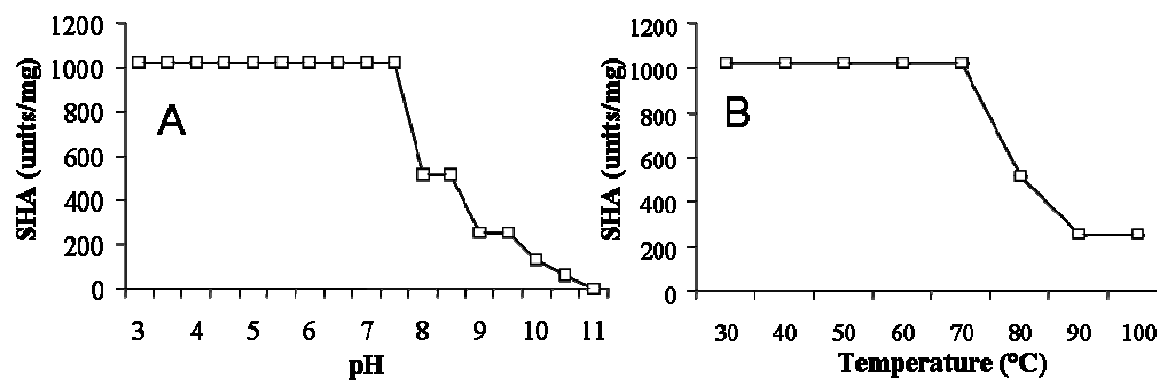


Figure 3

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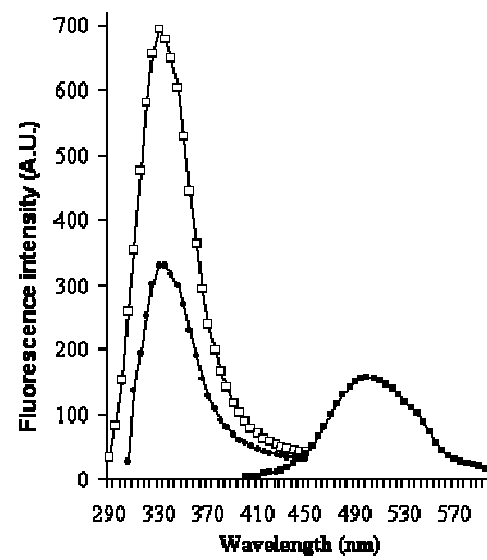


Figure 4

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Manuscript

Effect of gamma radiation on the molecular properties of *Sebastiania jacobinensis* bark lectin

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Abbreviations used: CM-cellulose, Carboxymethyl cellulose; SDS-PAGE, Sodium dodecyl sulphate polyacrylamide gel electrophoresis; SejaBL, *Sebastiania jacobinensis* bark Lectin; %SHA, percentage relative Specific Hemagglutinating Activity.

Abstract

Purified lectin from *Sebastiania jacobinensis* bark (SejaBL) is an effective antifungal agent of low environmental toxicity with hydrophobic surface that is displayed to free radicals produced by water radiolysis. In an experimental model, SejaBL was used as probe that changes on the hydrophobic center by irradiation, structurally altered proteins. To elucidate the effect of free radicals in molecular properties of SejaBL hemagglutinating activity (HA), tertiary structure (intrinsic and bis-ANS fluorescence), molecular weights and reversed phase chromatography were examined after γ -irradiation at various doses (0.020 to 35 kGy). Specific Hemagglutinating Activity (SHA) in irradiated lectin showed significant loss ($p < 0.05$), above 12.5 kGy, and a significant increase ($p < 0.05$) in 0.1, 0.8 and 1 kGy. Fluorescence spectroscopy indicated suppression of emission intensity when excited at 280 and 295 nm with high binding of bis-ANS after high doses. SDS-PAGE and reversed phase chromatography showed polypeptide fragmentation. Significant change in the hydrophobic surface indicated hydrogen abstraction in the amino acids causing structural disorganization from the lectin. Irradiation is a promising new safety technology that can eliminate pathogens in biomolecules, and to ratify their application it is important to clarify changes in various biological materials to start the application in large scale.

Keywords: Gamma radiation; Lectin; Intrinsic fluorescence; Bis-ANS

Introduction

Radiation treatments of biological materials have been applied in various processes as the sterilization of medical supplies and food. The applicable high dose of gamma radiation may affect the functional integrity of biopolymers. In proteins, structural damage is extensive abolishing biological activity [1-3]. Irradiation can inactivate proteins through two different mechanisms. First, it can split covalent bonds in protein target as a direct result of depositing photon energy [4]. Second, it can act indirectly, via water molecules, producing reactive oxygen species (ROS) responsible for the majority of protein damage [5]. These products interact with biological molecules, forming secondary organic radicals [6]. The primary damages are hydrogen abstraction in amino acid side chain, rings of aromatic residues, and reaction with sulfur [7,8].

Exposure of proteins to radiation has altered their physical and chemical structures with fragmentation, cross-linking, aggregation, unfolding and formation of new reactive groups, resulting in distortions of secondary and tertiary structures [9-11]. This amendment depends on several factors, such as protein concentration, presence of oxygen and quaternary structure [12]. Many conformation studies involved intrinsic and extrinsic fluorescence measurements of proteins. Denaturant proteins often exhibit changes in solvent accessibility to hydrophobic regions. One of the most important probes for study protein denaturation is bis-ANS [13]. Bis-ANS is non-fluorescent in aqueous mediums but shows fluorescence when bound to proteins. It has been used to probe the structure-function relationship in several proteins, such as tubulin [14] and myosin [15].

Lectins are proteins or glycoproteins of ubiquitous distribution in nature, which have one or several carbohydrate binding sites without catalytic function or immunological

characteristic [16]. They present a variety of chemical and biological strategies as insecticidal, fungicidal, antibacterial actives [17,18]. SejaBL is a purified lectin from *Sebastiania jacobinensis* bark with antifungal activity of low environmental toxicity which has a hydrophobic surface sensitive to the effect of free radicals. In this paper we describe the effect of gamma radiation on the activity and molecular properties of SejaBL, with focus on the changes in hydrophobic regions suffered by free radicals generated by the water radiolysis.

Material and methods

Purified lectin S. jacobinensis

S. jacobinensis bark was homogenized overnight at 4 °C in 10 mM Tris–HCl buffer (pH 8.5) with 0.2% (v/v) Triton X-100 detergent. The homogenate was centrifuged 5,000 g for 20 min (crude extract) followed by lyophilization. In brief, two volumes of cold acetone were added, mixed well and kept on ice for 10 min. The solution was centrifuged at 10,000 g for 20 min, at 4 °C. The supernatant was concentrated under vacuum until total evaporation of the acetone. The residual aqueous solution obtained was precipitated with 60% (NH₄)₂SO₄. The precipitate was dialyzed against the Tris–HCl buffer. This fraction was applied to a CM-Cellulose column (2 x 20 cm) previously equilibrated with the Tris–HCl buffer into 150 mM of NaCl. Following removal of the unadsorbed proteins, the column was eluted with 1 M acetic acid. The largest fractions of absorbance eluted with acetic acid were pooled, dialyzed and concentrated by ultrafiltration (Amicon ultra-15, Mr 10.000 cut-off) and chromatographed on a Sephadex G-100 column (2 x 70 cm) equilibrated with 150 mM NaCl. The column was eluted with the same solution at a flow

rate of 0.5 mL min^{-1} . The primary unadsorbed fraction (SejaBL) was concentrated by ultrafiltration in 10 mM phosphate buffer (pH 7.0) and stored. All other chemicals were of analytical grade and were purchased from Sigma (St. Louis, MO, USA). All solutions were made with water purified by the Milli-Q system.

Sample irradiation

The lectin 0.5 mg mL^{-1} in phosphate buffer (pH 7.0) added in borosilicate glass vials (16-125 mm) were frozen and irradiated into atmospheric O_2 , using the ^{60}Co gamma ray irradiator Gammacell 220 Excel (Ontario, Canada); in doses of 0.020; 0.050; 0.075; 0.1; 0.2; 0.4; 0.6; 0.8; 1.0; 3.0; 6.0; 12.5; 25; 35 kGy. The dose rate was 12.5 kGy / h . Each sample was analyzed 1 h, 24 h and 7 days after exposition by the methods below.

Hemagglutination activity and Protein concentration

Hemagglutinating activity (HA) was evaluated as described by Correia and Coelho [19] defined as the lowest sample dilution showing hemagglutination. Specific HA (SHA) corresponded to the relation between HA and protein concentration measured according Lowry *et al.* [20], using the BSA standard curve. The percentage of remaining SHA ($\% \text{SHA}_{\text{REM}}$) was calculated according to the equation: $\% \text{SHA}_{\text{REM}} = [\text{SHA}]_{\text{GM}} / [\text{SHA}]_{\text{G0}} \times 100$, where G_M is the lectin SHA of each dose irradiated (0.02-35 kGy) and G_0 is the not irradiated lectin SHA (control).

Sodium dodecyl sulphate–polyacrylamide gel electrophoresis (SDS–PAGE)

SDS–PAGE was performed to Laemmli [21]. Protein samples were mixed with sample

buffer (60 mM Tris-HCl, 2% SDS, 25% glycerol, 0.1% bromophenol blue, pH 6.8), resolved on a 10% separation gel and stained with silver kit (Bio-Rad). The broad range standard marker proteins were purchased from Sigma Chemical Co. (St. Louis, MO, USA).

Reverse phase chromatography analysis

The fraction irradiated lectins (0.1, 3 and 35 kGy) were submitted to reverse phase chromatography on the C-18 column (Vydac-protein peptide ultrasphere), performed on the HPLC system (Shimadzu LC-10AD-kyto, Japan) and monitored at 215 nm. The column was equilibrated with solvent A (0.1% TFA in H₂O) at a flow rate of 0.7 mL min⁻¹ and a non-linear gradient elution was used with solvent B (90% acetonitrile: 10% H₂O: 0.1% TFA) in A, with B = 5% at t = 5 min; B = 70% at t = 27 min; B = 80% at t = 60 min and B = 100% at t = 69 min.

Fluorescence spectroscopy

Fluorescence emission intensity of the irradiated lectin solution was measured at 25 °C using the spectrofluorimeter (JASCO FP-6300, Tokyo, Japan), at 0.2 mg mL⁻¹ of SejaBL (phosphate buffer, pH 7.0) in cuvette (1-cm pathlength rectangular quartz). The excitation wavelength was 295 and 280 nm. The emission spectra were recorded at a range of 305 to 450 nm. The band passes were 5 nm. Light scattering was measured at 90° for aggregation assays, in lectin previously irradiated, and light scattering values at 320 nm were monitored at 300 to 340 nm. The spectra reported averages of 3 scans. The spectrum of the solution was subtracted from each spectrum.

Hydrophobic surface

The lectin hydrophobic surface was determined using the same condition to intrinsic fluorescence. The samples were transferred to the quartz cuvette and then mixed with 5 μ M bis-ANS (Molecular Probes Inc., USA) and the fluorescence measured in the JASCO spectrofluorimeter. The fluorescence emission was obtained at 400 to 600 nm with excitation at 360 nm [22].

Statistical analysis

Statistical analysis for paired data was carried out by Student's t test using the GraphPrism® (GraphPad Software Inc., San Diego, CA, USA). The significance accepted difference between the mean values of parameter was analyzed to the level of $p < 0.05$.

Results and Discussion

Effects of irradiation on SHA

Lectin showed significant loss ($p < 0.05$) with 12.5, 25 and 35 kGy due to direct action of radiation on structure in 1 h after exposure. In doses (0.02 to 6 kGy) no significant change was observed. The doses 0.1, 0.8 and 1 kGy showed a significant increase ($p < 0.05$) with 1 h and 24 h after radiation target analysis (Fig. 1). The amplification is associated with reduced dosage of protein, except for dose of 0.1 kGy, where there was an increase in HA. This reduction has been observed [23-25] and is a consequence of chemistry oxidation suffered by aromatical amino acids (mainly tyrosine and tryptophan residues) responsible for the formation of the measurable colorful complex based on the

Lowry et al. method [20]. These data are confirmed by intrinsic fluorescence emission reduction of tyrosine and tryptophan amino acids (Fig. 3). In relation to the increase of HA with 0.1 kGy, according to Macklis [26] and Shuzheng [27], low doses of radiation yield a rise of active molecules exerted into closed systems. This peculiar behavior, confirmed by increased intrinsic fluorescence, reveals changes in native tertiary structure with a possible modification of hydrogen bonding, van der Waals and hydrophobic interactions. Although there was a significant loss of HA, SejaBL showed high resistance compared to other previously analyzed proteins [3]. The concentration of proteins influenced the degree of structural change of the irradiated proteins. Thus, the damage is greater to low protein concentrations [12].

Structural analysis

None of these doses (0.02 to 0.075 kGy) changed the molecular weight pattern of SejaBL. The initial degradation of lectin (MW 52 kDa) was observed in doses (0.2 to 6 kGy) with aggregation tracked by light scattering (data not shown). Above 3 kGy, only a degradation of the main band was viewed. The reverse phase chromatography analysis revealed loss of the peak area with structural collapse (Fig. 2). Puchala & Schuessler [28] suggested that break points, caused by radiation on protein, are fragile bonds in the polypeptide chain. Proteins can be converted to higher molecular weight aggregates, due generation of inter-protein cross-linking reactions, hydrophobic and electrostatic interactions, as well as the formation of disulfide bonds [5,24,29].

Intrinsic fluorescence

The intrinsic fluorescence emission showed increase (0.02 to 0.1 kGy) and decreased (0.2 to 35 kGy) without change in λ_{max} at around 330 nm for tryptophan and tyrosine residues (Fig. 3). The hydroxyl radicals are generated by water radiolysis [30] and all aliphatic amino acids are potential targets of modification [11,31]. Aromatic amino acids residues become particularly sensitive, producing hydroxylated products. The formation of formylkynurenine from tryptophan, 3,4-dihydroxyphenylalanine (DOPA) from tyrosine, and o-tyrosine from phenylalanine as major oxidation products was recognized [32]. The partial aromatic amino acid substitution by hydroxylated products results in a lesser emission of fluorescence [33]. Such oxidative damage distorts the polarity of these residues modifying the hydrophobic surface and tertiary structure favoring the denaturation and aggregation.

Hydrophobicity

Bis-ANS fluorescence increased at all doses with maximum blue shifts to 490 nm above 0.8 kGy (Fig. 4). Bis-ANS fluorescence is observed in non-irradiated SejaBL (i.e., the control) and when compared only with the buffer it indicates that the hydrophobic surface turned to the solvent. In protein, hydrophobic surfaces are easily attacked by hydroxyl radicals and, in this case, the change in protein hydrophobicity is considered the preponderant factor for the structural collapse. Hydrogen abstraction of aromatic amino acids promotes greater exposure of the hydrophobic surface to the solvent, due to decrease of apolarity surface, leading to greater binding bis-ANS (maximum blue shift emission at 490 nm).

A survey of the literature showed that lectins bind their ligands most commonly through hydrogen bonds (some mediated by water), hydrophobic interactions, and that in rare cases electrostatic interactions (ion pairing) and coordination with metal ions also play a role [16]. The hydrophobic face of sugar rings on aromatic side chains of amino acids is a common feature of the complexes with carbohydrates of other lectins, for instance, of the galectins, and certain bacterial toxins, such as *Escherichia coli* lytic toxin, and also of non-lectin proteins [34]. Several independent properties of irradiated SejaBL revealed different responses. They indicate that the tertiary structure plays a critical role in HA, where the various modifications of hydrophobic amino acid residues via free radicals promote the collapse of hydrophobic regions, changing several kinetic properties in lectin. Irradiation at appropriate doses is a promising approach for producing safe and pathogen-free biomolecules. Therefore, it is important to clarify changes in various biological materials in order to ratify the application of gamma radiation, a crucial event to begin the production of bioderivatives in large scale.

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Figure

Figure captions

Fig. 1. γ -radiation effect on lectin activity. %SHA_{REM} (▲) 01 h; (O) 24 h; () 7 days after irradiation. The error in the determination of %SHA_{REM} for different doses is approximately $\pm 1\%$, smaller than the size of the symbols. *Significant difference ($p < 0.05$) in relation to those non-irradiated lectin (i.e., the control).

Fig. 2. Chromatography profile of SejaBL under gamma irradiation. (—) control non-irradiated; (—) 0.1 kGy; (—) 3 kGy; (—) 35 kGy. Reverse phase chromatography in C-18 column HPLC system (Shimadzu). The column was equilibrated with solvent A (0.1% TFA in H₂O) and a non-linear gradient elution was used with solvent B (90% acetonitrile: 10% H₂O: 0.1% TFA) in A, with B = 5% at t = 5 min; B = 70% at t = 27 min; B = 80% at t = 60 min and B = 100% at t = 69 min.

Fig. 3. SejaBL intrinsic fluorescence. (A) Total intrinsic fluorescence; relative intensity (%) with (▲) 01 h; (O) 24 h; () 7 days after irradiation lectin. (B) Tryptophan fluorescence. The error in the determination of fluorescence from the relative peak areas of integral intensities to different doses is approximately $\pm 0.5\%$, smaller than the size of the symbols. (C), (D) and (E) fluorescence with 1 h, 24 h and 7 days after irradiation, respectively. Excitation 295 nm. Emission 305-450 nm. (—) control non-irradiated; (---) 0.02 to 0.075 kGy; (—) 0.02 to 6 kGy; (...) 12.5 to 35 kGy.

Fig. 4. SejaBL fluorescence treated with bis-ANS. (A) bis-ANS Fluorescence; relative intensity (%) with (▲) 01 h; (O) 24 h; () 7 days after irradiation. (B) Center mass. The

error in the determination of the fluorescence from the relative peak areas of the integral intensities to different doses is approximately $\pm 1\%$, smaller than the size of the symbols. (C), (D) and (E) fluorescence with 1 h, 24 h and 7 days after irradiation, respectively. Excitation 360 nm. Emission 400–600 nm. (—) control non-irradiated; (---) 0.02 to 0.075 kGy; (—) 0.02 to 6 kGy; (...) 12.5 to 35 kGy.

Figure

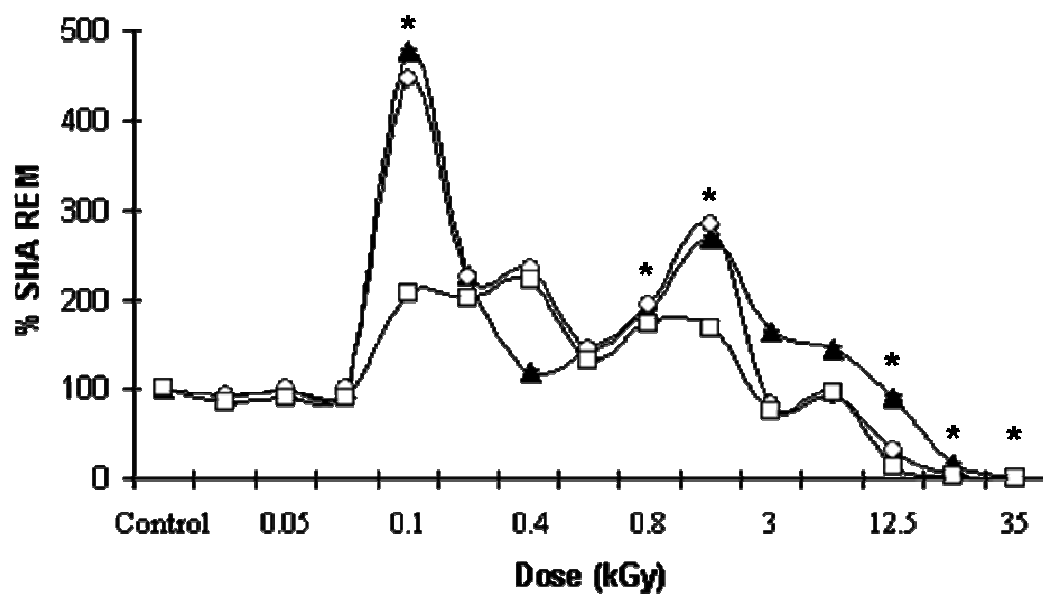


Figure 1

Figure

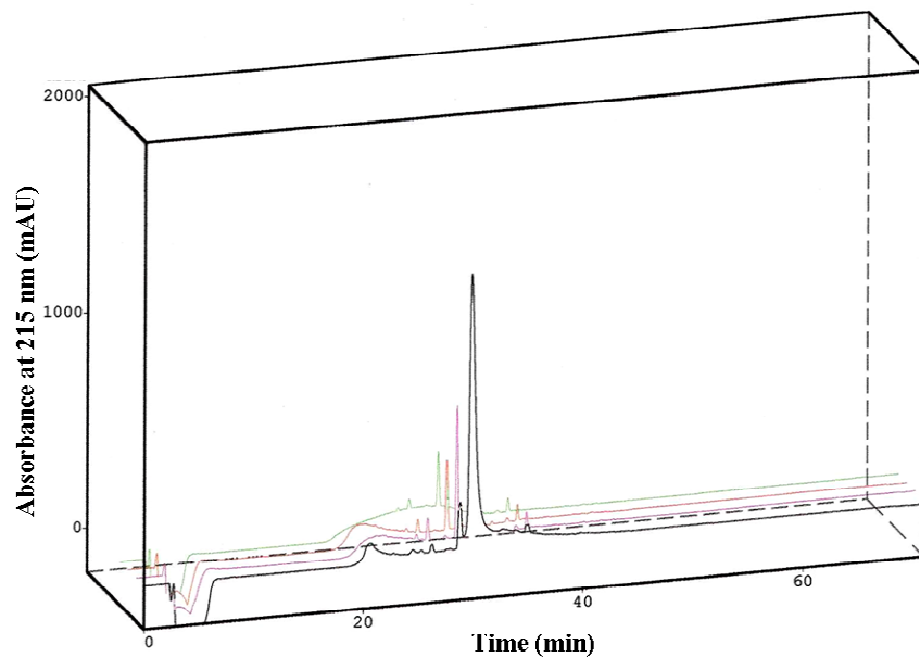


Figure 2

Figure

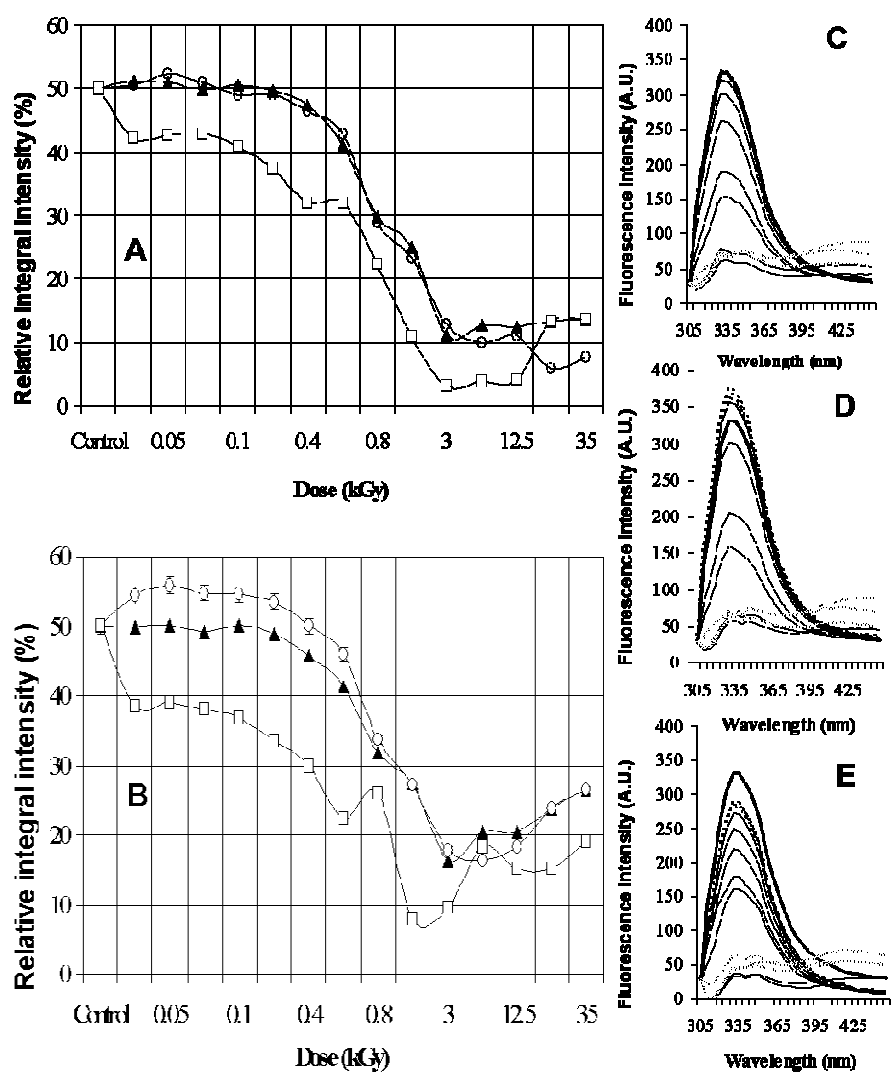


Figure 3

Figure

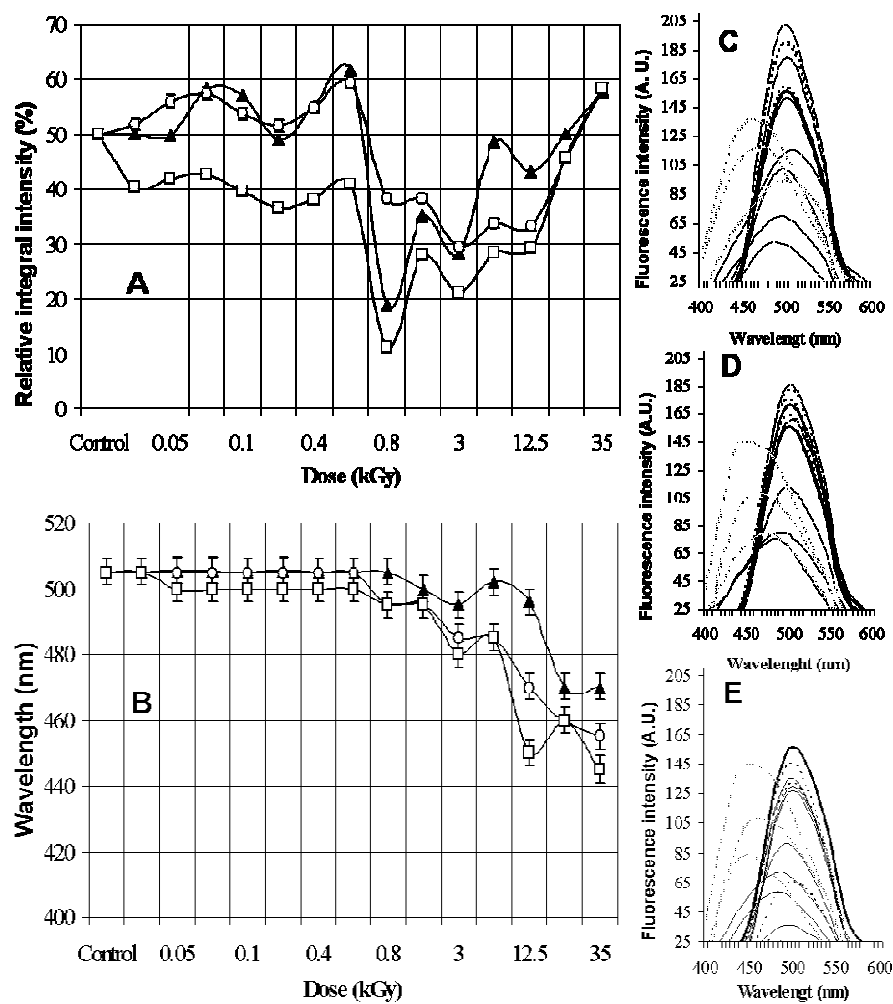


Figure 4

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Radical scavenger ability of flavonoids isolated from *Sebastiania jacobinensis* with protective action on protein oxidation after gamma irradiation

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Abbreviations used: CM-cellulose, Carboxymethyl cellulose; DEAE-Cellulose, diethylaminoethyl cellulose; DPPH, 1,1-diphenyl-2-picrylhydrazyl; SDS–PAGE, Sodium dodecyl sulphate–polyacrylamide gel electrophoresis; SejaBL, *Sebastiania jacobinensis* bark Lectin; SejaBF, *Sebastiania jacobinensis* bark Flavonoids. TLC, Thin Layer Chromatographic; %HA, percentage relative Hemagglutinating Activity.

Abstract: Flavonoids are ubiquitous plant secondary products with radical scavenger ability. *Sebastiania jacobinensis* bark lectin (SejaBL), protein capable of binding to carbohydrates, has their hydrophobic surface modified by free radicals produced via water radiolysis. In the present study the protective effect of *S. jacobinensis* bark flavonoids (SejaBF) on SejaBL activity and their antioxidant stability after gamma irradiation were evaluated. SejaBF showed fast scavenger ability measured by 2,2-diphenyl-1-picrylhydrazyl radical (DPPH) after high doses of radiation and reduced by 55% the damage caused by irradiation on SejaBL activity. The results showed radiostability of antioxidant capacity of the isolated flavonoids (SejaBF) protecting the lectin from free radicals formed after the irradiation.

Keywords: Lectin; Flavonoids; DPPH radical scavenger capacity; Gamma irradiation

1. Introduction

Radiation treatments of biological materials have been applied in various fields such as sterilization of materials and food for reducing microbial contamination. Gamma irradiation can damage biomolecule by initial rupture covalent bonds as a direct result of the depositing photon energy, and indirectly, by producing free radicals and other non-radical reactive oxygen species (Kempner, 2001). Investigations showed that free radicals cause lipid peroxidation (Terao et al., 1994), structural damage proteins with biological activity abolished (Cho and Song, 2000), DNA damage (Wallace, 1998), and oxidative stress (Aruoma, 1998) making the recovery of biologically active materials unlikely. However, chemical changes caused by irradiation can be prevented or reversed by use effective and radiostable antioxidant agents.

Flavonoids are among the most ubiquitous of plant secondary metabolites groups acting on reproduction and protection of plants (Winkel-Shirley, 2002). Their low toxicity (Zhang et al., 2005), high antioxidant activity (Garcia-Alonso et al., 2005), ultraviolet protection (Bieza and Lois, 2001), anti-inflammatory (Lin et al., 2008), antiviral (Yamaguchi et al., 2001) and antitumour activities (Hou, 2003) have been documented. Flavonoids are

subdivided into flavones, flavonol, flavanone, and flavanol types depending on the pattern of substitution: hydroxylation or methoxylation.

Irradiated flavonoid solutions produce chemical reactions between the solute and reactive species formed from the radiolyzed solvent operating in the fast repair of chemical modifications on biomolecules (Hanasaki et al., 1994; Kozłowski et al., 2007). One of the most important probes, to evaluate the fast repair antioxidant capacity is the 2,2-diphenyl-1-picrylhydrazyl radical (DPPH), a stable nitrogen synthetic radical (Brand-Williams et al., 1995).

Lectins constitute a class of proteins that possess a non-catalytic domain capable of specific recognition and binding with carbohydrates (Sharon and Lis, 2004). The number of roles has been proposed, including stimulation of cell proliferation (Peumans and Van Damme, 1998), carbohydrate transport (Hsu and Liu, 2004), plant-microbe interactions (Limpens and Bisseling, 2003), and in plant defense (Peumans and Van Damme, 1995). So far, lectins have been isolated and characterized on the basis of hemagglutinating activity.

S. jacobinensis bark lectin (SejaBL) is a protein which loses their activate front free radicals produced by water radiolysis due to modification in hydrophobic surfaces. *Sebastiania jacobinensis* Müll. Arg. (Euphorbiaceae family) is a tree which is resistant to the arid climate and common in tropical regions of Brazil. Therefore, the aim of our study was to evaluate the protective effect on SejaBL, free radical scavenging ability and radiostability of flavonoids isolated from *S. jacobinensis* bark after ample dose of γ -radiation.

2. Experimental

2.1. Materials

S. jacobinensis bark was collected from trees in the semi-arid region, state of Pernambuco, Brazil. Reference samples of (+)-catechin, 1,1-Diphenyl-2-picrylhydrazyl radical (DPPH), DEAE-Cellulose and ascorbate were purchased from Sigma Chemical Co., (St. Louis, MO). All the solvents and other chemicals used were of analytical grade from Merck (Darmstadt, Germany). All solutions were made with water purified by the Milli-Q system.

2.2. Flavonoids isolation

S. jacobinensis bark was mixed in a beaker with 250 mL of ethanol/acetic acid/water (70:4:26, v/v/v) for 2 h at 4 °C (this was carried out three times to extract all phenolic compounds; a fourth extraction did not extract any phenolic compounds detectable in the amount). The three extracts were combined (750 mL) vacuum-filtered through filter paper (Whatman N°. 1), concentrated to dryness under reduced pressure (40 °C) and re-dissolved in 10 mL ethanol/water (70:30, v/v). The extract containing flavonoids was chromatographed on an anion-exchange DEAE-Cellulose column (40 x 2.5 cm) equilibrated with ethanol-water (70:30, v/v). Elution was performed with ethanol-acetic acid-water (70:4:26, v/v/v) at a flow rate of 2 mL min⁻¹. Each fraction collected was checked by TLC for their compositions using ethyl acetate/formic acid/acetic acid/water (100:11:11:26, v/v/v/v) for development and vanillin dipping in chloridic acid was used for detection under UV 254 nm. TLC conditions were optimized using (+)-catechin standard. The obtained flavonoids were denominated SejaBF.

2.3. LC-ESI-MS

The analysis of SejaBF was revealed by LC analyses (Waters) with a diode-array detector (LC-UV), in Deltapak C18 column (2.0 mm x 150 mm; 3.5 µm size; 60 Å; Walters) at a flow rate of 0.4 mL min⁻¹ keeping the column at a constant temperature (30 °C) and using 0.1% TFA/H₂O (solvent A) and 90% ACN/0.1% TFA/H₂O (solvent B) as solvents. A linear gradient elution was used with 5-95% B in A at 30 min. The molecular mass was determined by electrospray ionization ion-trap mass spectrometry on a Waters LC/ESI+/MS scanning of 200-2500 m/z.

2.4. Purified lectin from *Sebastiania jacobinensis* and Analyses

S. jacobinensis bark was homogenized overnight at 4 °C in 10 mM Tris-HCl buffer (pH 8.5) with 0.2% (v/v) Triton X-100 detergent. The homogenate was centrifuged 5,000g for

20 min (crude extract) followed by lyophilization. In brief, two volumes of cold acetone were added, mixed well and kept on ice for 10 min. The solution was centrifuged at 10,000g for 20 min at 4 °C. The acetone precipitate was separate. The supernatant was concentrated under vacuum until total evaporation of the acetone. The supernatant obtained was precipitated with 60% (NH₄)₂SO₄. The precipitate was dialyzed against Tris–HCl buffer. This fraction was applied to a CM-Cellulose column (2 x 20 cm) previously equilibrated with Tris–HCl buffer into 150 mM of NaCl. Following removal of the unadsorbed proteins, the column was eluted with 1 M acetic acid. The largest fractions of absorbance eluted with acetic acid were pooled, dialyzed and concentrated by ultrafiltration (Amicon ultra-15, Mr 10,000 cut-off) and chromatographed on a Sephadex G-100 column (2 x 70 cm) equilibrated with 150 mM NaCl. The column was eluted with the same solution at a flow rate of 0.5 mL min⁻¹. The primary unadsorbed fraction (SejaBL) was dialyzed by ultrafiltration against 10 mM phosphate buffer and stored. Hemagglutinating activity (HA) was evaluated as described by Correia and Coelho (1995), the lowest sample dilution showing hemagglutination. The percentage of remaining HA (%HA_{REM}) was calculated according to equation:

$$\%HA_{REM} = [HA]_G / [HA]_{G_0} \times 100$$

where G is the lectin HA of each irradiated dose (0.02-35 kGy) and G₀ is the non-irradiated lectin HA (control).

The molecular mass profile of irradiated lectin was determined using SDS-PAGE (Laemmli, 1970). Equal amounts of protein samples (200 µg) were loaded, resolved on a 10% separation gel and stained with silver kit (Bio-Rad). Protein Markers: Rabbit muscle myosin (205 kDa), E. coli β-galactosidase (116 kDa), rabbit muscle phosphorylase b (97.4 kDa), rabbit muscle fructose-6-phosphate kinase (84 kDa), bovine serum albumin (66 kDa), bovine liver glutamic dehydrogenase (55 kDa), egg albumin (45 kDa), rabbit muscle glyceraldehyde-3-phosphate dehydrogenase (36 kDa), bovine erythrocyte carbonic anhydrase (29 kDa), bovine pancreas trypsinogen (24 kDa), soybean trypsin inhibitor (20 kDa), bovine milk α-lactalbumin (14.2 kDa) and bovine lung aprotinin (6.5 kDa).

2.5. Gamma irradiation

The lectin 0.5 mg mL⁻¹ in 10 mM phosphate buffer (pH 7.0) was added in borosilicate glass vials (16-125 mm) with and without SejaBF (1 mg mL⁻¹). The mixtures were frozen and irradiated into atmospheric O₂, using a ⁶⁰Co gamma ray irradiator Gammacell 220 Excel (Ontario, Canada). The doses were 0.020; 0.050; 0.075; 0.1; 0.2; 0.4; 0.6; 0.8; 1.0; 3.0; 6.0; 12.5; 25; 35 kGy. The dose rate was 8.8 kGy/h. Each sample was analyzed 1 h, 24 h and 7 days after exposition to HA.

2.6. Determination of antioxidant activity

Spectrophotometric analysis was used to determine the inhibition concentration (IC₅₀) and the inhibition percentage (IP). IC₅₀ is the amount of antioxidant necessary to decrease the initial concentration of DPPH radical (75 µM) by 50%. IP is the percentage of total DPPH radical (75 µM) which reacted with antioxidant at the steady state (20 min). The experiments were performed using the SmartSpec 3000 spectrophotometer (Bio-Rad) according to the methods of Soler-Rivas et al. (2000) and Moure et al. (2001). An aliquot (20 µL) of SejaBF irradiated in doses of 0.02; 0.8; 3; 35 kGy and not irradiated (control) at 0.5 mg mL⁻¹ were mixed separately with 75 µM methanolic DPPH radical solution to a final volume of 1 mL. Pure methanol was used as negative control. (+)-Catechin and ascorbate in methanol (1 mg mL⁻¹) were used as positive control. The disappearance of the DPPH radical was monitored by the decrease in absorbance at 515 nm, which was recorded after 0; 1; 2; 3; 4 and 5 min, and subsequently every 5 min up to 30 min, during which time the radical was stable. The DPPH radical concentration in the reaction mixture was calculated by the calibration curve according to the following nonlinear regression equation (R= 0.997): $A_{515\text{ nm}} = 0.01029 [\text{DPPH}] - 0.0177$, where [DPPH] is expressed in mg mL⁻¹. The percentage of remaining DPPH (%DPPH_{REM}) was calculated according to Brand-Williams et al. (1995), as follows:

$$\% \text{DPPH}_{\text{REM}} = [\text{DPPH}]_{\text{T}} / [\text{DPPH}]_{\text{T}_0} \times 100$$

where T is the time when absorbance was determined (1–30 min) and T_0 is the time zero.

IC_{50} was calculated by plotting the %DPPH_{REM} at the steady state (20 min) against various concentrations of each sample (1, 0.5, 0.25, 0.125, 0.0625 mg mL⁻¹). The results were expressed by mg antioxidant/g DPPH \pm standard deviation. For determination of IP, an aliquot (50 μ L) of SejaBF was added to 2 mL of 75 μ M methanolic solution of the DPPH radical and the absorbance was determined at 515 nm at the steady state (20 min). IP was calculated according to the expression:

$$IP = [(A_{T_0} - A_{T_S})/A_{T_0} \times 100$$

where A_{T_0} is the absorbance at time zero and A_{T_S} is the absorbance at the steady state (defined as the state when the absorbance of the sample remained stationary). The results were expressed by % of inhibition.

2.7. Data Analysis

Data are presented as mean \pm S.E.M. The curve and its 95% confidence intervals (CI 95%) were obtained by nonlinear regression using the GraphPrism® (GraphPad Software Inc., San Diego, CA, USA). Student's t-test for paired data was developed and significant difference between groups was analyzed at $p < 0.05$.

3. Results and Discussion

Two flavonoids were isolated from *S. jacobinensis* bark extracts by combination of anion-exchange DEAE-Cellulose, HPLC and TLC chromatographys (Fig. 1), denominated *S. jacobinensis* bark flavonoids (SejaBF). Figure 2 shows a positive molecular ion [(m/z) = $M+H^+$] to 613 m.u. corresponding to epigallocatechin dimmer and a non-identified peak [(m/z) = $M+H^+$ 702 m.u.] that can be heterogeneous dimer of monogalloylated procyanidin. The identification of heterogeneous dimers by antioxidant capacity and HPLC/MS protocol was demonstrated for analysis of the procyanidins (Ferreira and Slade, 2002).

SejaBF exhibited high antioxidant capacity by DPPH method when compared to ascorbate, antioxidant widely used as standard, however, its antioxidant activity decreased after irradiation, in relation to the IC_{50} and IP, available in 20 min (Table I). IP reduction (19.2% to high dose) was significant although the IC_{50} reduction was not.

DPPH radical scavenger capacity (%DPPH_{REM}) was significantly reduced only in the initial 5 min after high dose of radiation (Fig. 3). The DPPH radical is scavenged by antioxidants through donation of hydrogen to form stable reduced DPPH molecule. The radical scavenger is reduced after high doses of gamma radiation due to displacement from the B-ring to the 2,3-double bond, indicating a relatively small reactivity (because of the absence of high density spin on a given site) for these intermediate species (Zhao et al., 2002; Marfak et al., 2003). Our result is in good agreement with the presence of stabilized quinones, which has been observed after H-abstraction from the B-ring of flavonoids in radiolytic solutions (Pannala et al., 2001).

Gamma irradiation of SejaBL aqueous solution showed significant loss ($p < 0.05$) of HA with high dose of gamma rays compared to the control (non-irradiated), but the damage caused by irradiation on the activity of lectin was minimized by 55% in the presence of SejaBF in high dose (Fig. 4). The lectin activate protecting can be explained by the high radical scavenger capacity of flavonoids always after irradiation. This antiradical mechanism in phenol-protein aggregates might be due to the ability of phenolic compounds to transfer oxidative damage from one phenolic site to another, protecting proteins and amino acids from oxidation (Franzini et al., 1993; Xu et al., 2003; Štajner et al., 2008). In a study with 42 lectins, Barre et al. (2002) evidenced common hydrophobic domains, suggesting strong structural conservation, with secondary metabolites binding nature of these domains. Thus, flavonoids might have a protective effect on hydrophobic domains easily attacked by active species of water radiolysis (Steinhart, 1991).

The change in terms of molecular weight pattern in irradiated lectin was significant, with the formation of degraded small molecular weight to high dose of gamma radiation (Fig. 5). The degradation of lectin (MW 52,000 Da) was lower in the samples with the antioxidant SejaBF. Usually, breakage of covalent bonds into irradiated proteins shows new bands below the major band (Le Maire et al., 1990; Xu and Chance, 2005).

4. Conclusions

SejaBF prevented the lectin oxidation by free radicals, due to possible binding to the hydrophobic domain and fast repair of groups sensitive to ROS. In summary, this study demonstrated the special protective effect on protein and the high antioxidant capacity of the obtained flavonoids after irradiation.

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Table 1

Table 1 Radical scavenging performance.

Samples	IP (%)	IC ₅₀ (mg/ g DPPH)
SejaBF 20 Gy	89.1313 ^{a#}	0.115 ± 0.002 ^a
SejaBF 800 Gy	88.3073 ^{a#}	0.112 ± 0.002 ^a
SejaBF 3 kGy	79.6359 ^{b#}	0.180 ± 0.002 ^a
SejaBF 35 kGy	70.886 ^{c#}	0.300 ± 0.004 ^a
SejaBF	90.1122 ^a	0.126 ± 0.004 ^a
(+)-catechin	90.8577 ^a	0.206 ± 0.000 ^a
Ascorbate	39.2215 ^d	1.317 ± 0.002 ^b

Values with the same superscript letter are not significantly different at $p < 0.05$ (Student's T-test). IC₅₀ and IP (0.5 mg mL⁻¹) of SejaBF were calculated at the steady state (20 min). [#] Values of IP (%) are significantly different at $p < 0.05$ (Student's T-test) in relation to non-irradiated SejaBF.

Figure

Figure captions

Fig. 1. Isolation of flavonoids from *S. jacobinensis* (SejaBF). **(a)** Analysis by thin-layer chromatography: (1) Extract ethanolic, (2) SejaBF isolated from DEAE-Cellulose chromatography and (3) (+)-catechin standard. **(b)** SejaBF reverse phase chromatography in C-18 column by HPLC system (Waters).

Fig. 2. *S. jacobinensis* flavonoids (SejaBF) ESI-MS spectra. Purified flavonoids in LC/MS with retention time of 10.04 min (a) and 9.59 min (b).

Fig. 3. Kinetic behaviour of irradiated antioxidants. The error in the determination of %DPPH_{REM} to relative values of different doses is approximately $\pm 1\%$, smaller than the size of the symbols. * Values significantly different at $p < 0.05$ (Student's T-test) in relation to (○) non-irradiated *S. jacobinensis* flavonoids (SejaBF). SejaBF after irradiation by: (x) 35 kGy; (▲) 3 kGy; (■) 800 Gy; () 20 Gy; (●) Ascorbate and (+)-catechin.

Fig. 4. Effect of γ -radiation on *S. jacobinensis* lectin (SejaBL) activity. %HA (▲) 01 h; (O) 24 h and () 7 days after irradiation. SejaBL with absence **(a)** and presence **(b)** *S. jacobinensis* flavonoids (SejaBF). * Significant difference ($p < 0.05$) in relation to the non-irradiated (i.e., the control). # Significant difference ($p < 0.05$) in relation to the groups without **(a)** and with **(b)** antioxidant (SejaBF).

Fig. 5. SDS-PAGE profile of irradiated *S. jacobinensis* lectin (SejaBL). (1) Protein markers. Irradiated SejaBL without SejaBF (even numbers): (2) 35 kGy; (4) 3 kGy; (6) 800 Gy and (8) 20 Gy. Irradiated SejaBL in SejaBF presence (odd numbers): (3) 35 kGy; (5) 3 kGy; (7) 800 Gy and (9) 20 Gy. (10) Non-irradiated SejaBL. Arrows indicate the change on irradiated SejaBL molecular weight with formation of degraded small peptide, after high dose of gamma radiation.

Figure

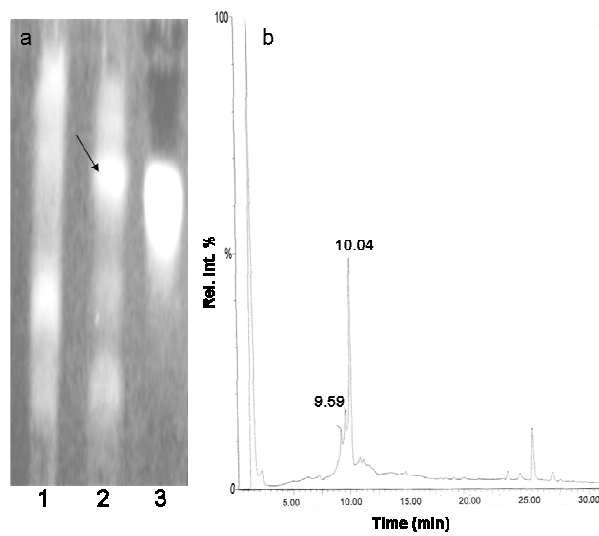


Figure 1

Figure

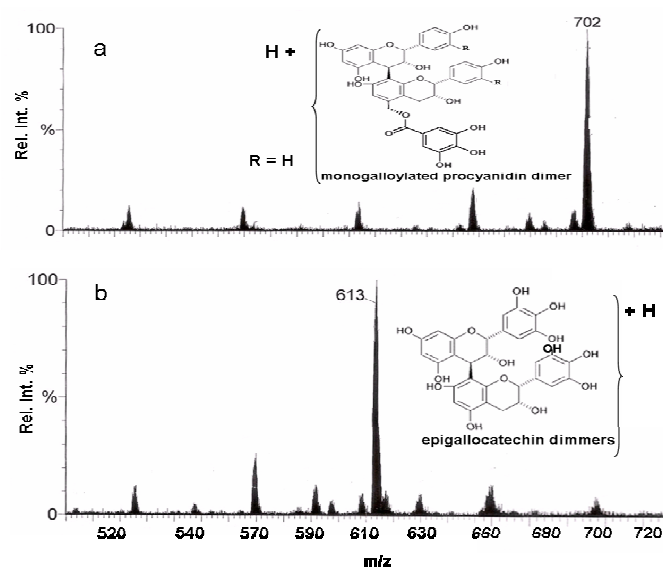


Figure 2

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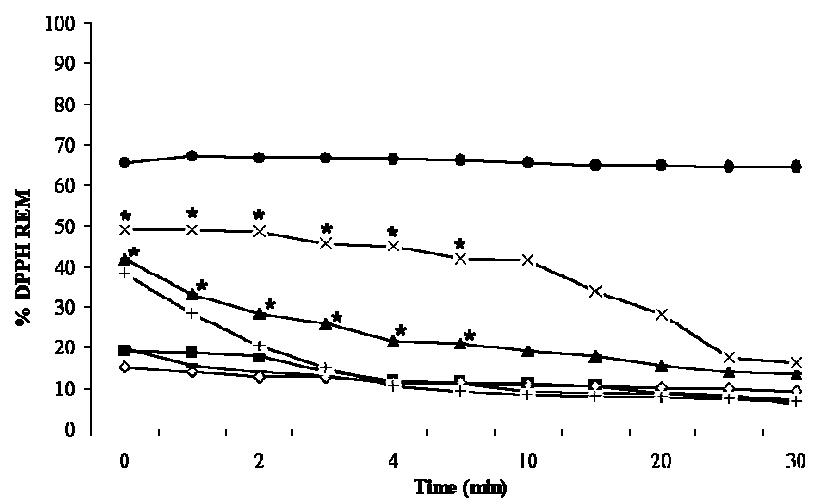


Figure 3

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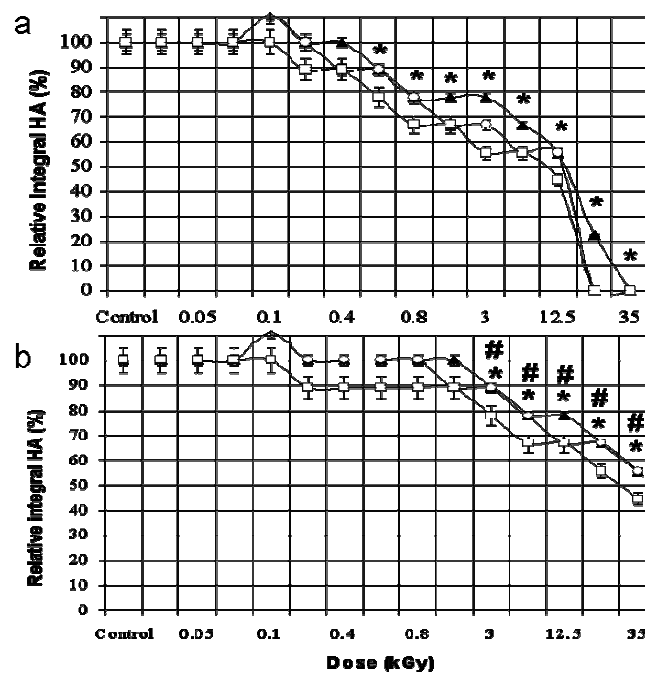


Figure 4

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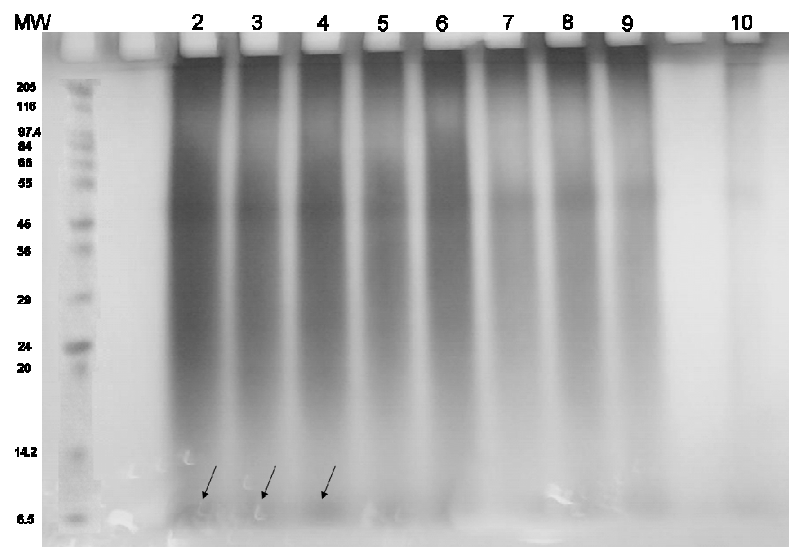


Figure 5

6. CONCLUSÃO

- ❖ Num protocolo eficiente, cromatografias em CM-Celulose e Sephadex G-100 isolam uma lectina da entrecasca de *Sebastiania jacobinensis* (SejaBL);
- ❖ SejaBL é termoresistente, não dependente de íons, estável em pH ácido, glicosilada, com resistência a proteinases e inibe fortemente tripsina;
- ❖ SejaBL possui massa molecular (52 kDa) composta por duas subunidades de 24 kDa não homologas, superfície hidrofóbica e perfil homogêneo;
- ❖ A lectina em baixa concentração promove ação antifúngica, atuando como um inibidor do crescimento de espécies de *Fusarium*; SejaBL também promove ação antifúngica seletiva, pois organismos não alvos não foram afetados podendo ser uma ferramenta útil em estudos de biotecnologia vegetal sem agredir a diversidade ambiental;
- ❖ SejaBL perdeu totalmente AHE após exposição a altas doses de radiação gama;
- ❖ SejaBL exibiu a partir da dose de 6 kGy desestruturação molecular e perda de AHE;
- ❖ A fluorescência intrínseca mostrou uma eliminação da emissão de fluorescência quando excitada a 280 e 295 nm, indicando alterações químicas em aminoácidos aromáticos;
- ❖ A fluorescência do bis-ANS mostrou uma emissão blue shifts a 490 nm em altas doses de radiação evidenciando a ação de radicais livres sobre superfície hidrofóbica protéica;
- ❖ Dois flavonóides foram isolados exibindo alta capacidade para capturar radicais;
- ❖ Os flavonóides exibiram boa estabilidade frente irradiação gama;
- ❖ SejaBL exibiu um menor queda da AH na presença dos flavonóides após irradiação;
- ❖ A lectina mostrou diferentes perfis de oxidação protéica na presença do antioxidante pela possível conservação de domínios hidrofóbicos.

ANEXOS

Anexo 1: Normas do periódico especializado, ao qual o trabalho da dissertação será submetido (BIOCHEMICAL AND BIOPHYSICAL RESEARCH COMMUNICATIONS).

Anexo 2: Normas do periódico especializado, ao qual o trabalho da dissertação será submetido (ARCHIVES OF BIOCHEMISTRY AND BIOPHYSICS).

Anexo 3: Normas do periódico especializado, ao qual o trabalho da dissertação será submetido (RADIATION PHYSICS AND CHEMISTRY).

7.1. Anexo 1: BIOCHEMICAL AND BIOPHYSICAL RESEARCH COMMUNICATIONS

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The abstract (p. 2) must be a single paragraph that summarizes the main findings of the paper in less than 150 words. After the abstract a list of up to 10 keywords that will be useful for indexing or searching should be included.

The introduction should be as concise as possible, without subheadings.

Materials and methods should be sufficiently detailed to enable the experiments to be reproduced.

Results and Discussion may be combined and may be organized into subheadings.

Acknowledgments should be brief and should precede the references.

References to the literature should be cited by numbers in square brackets in the text and listed in numerical order at the end. Use the most recent edition of the Chemical Abstracts Service Source Index for abbreviations of journal titles. Only articles that have been published or are in press should be included in the references. Unpublished results or personal communications should be cited as such in the text. Please note the following examples.

[1] W.D. Strayhorn, B.E. Wadzinski, Arch. Biochem. Biophys. 400 (2002) 76-84.

[2] R. Hesketh, The Oncogene FactsBook, Academic Press, San Diego, 1995.

[3] O.R. Mettam, L.B. Adams, in: E.S. Jones, R.Z. Smith (Eds.), Introduction to the Electronic Age, E-Publishing Inc., New York, 1999, pp. 218-304.

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March 2008 version

7.3. Anexo 3: Radiation Physics and Chemistry

The Journal for Radiation Physics, Radiation Chemistry and Radiation Processing

Guide for Authors

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Subdivision

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Title, Authors, Affiliations, Abstract, Keywords.

Introduction, Experimental, Results, Discussion, Conclusion.

Acknowledgements, Appendix, References, Figure legends, Tables. The introduction should briefly describe the background and reason for doing the work described. The experimental should provide enough information for readers to repeat the experiment. The discussion should evaluate the results and their relationship to comparable works. The conclusion should briefly summarize the obtained results.

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If there is more than one appendix, they should be identified as A, B, etc. Formulae and equations in appendices should be given separate numbering: Eq. (A.1), Eq. (A.2), etc.; in a subsequent appendix, Eq. (B.1) and so on.

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Acknowledgements

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