UNIVERSIDADE FEDERAL DE PELOTAS

Programa de Pós-Graduação em Biotecnologia



Tese

Avaliação *in vitro* de novas moléculas (metabólitos secundários e lectinas) com potencial na terapêutica do câncer

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Pelotas, 2012

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AVALIAÇÃO IN VITRO DE NOVAS MOLÉCULAS (METABÓLITOS SECUNDÁRIOS E LECTINAS) COM POTENCIAL NA TERAPÊUTICA DO CÂNCER

Tese apresentada ao Programa de Pós-Graduação em Biotecnologia da Universidade Federal de Pelotas, como requisito parcial à obtenção do título de Doutor em Ciências (área do conhecimento: Biologia Celular, Câncer e Nanotecnologia).

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Dados de catalogação na fonte: Ubirajara Buddin Cruz – CRB 10/901 Biblioteca de Ciência & Tecnologia - UFPel

N371a Nedel, Fernanda

Avaliação in vitro de novas moléculas (metabólitos secundários e lectinas) com potencial na terapêutica do câncer / Fernanda Nedel. — 89f. — Tese (Doutorado). Programa de Pós-Graduação em Biotecnologia. Universidade Federal de Pelotas. Centro de Desenvolvimento Tecnológico, 2012. — Orientador Fabiana Kömmling Seixas ; co-orientador Tiago Veiras Collares.

1.Biotecnologia. 2.Câncer. 3.Nanotecnologia. 4.Nanotubos de carbono. 5.Lectinas. 6.Disseleneto de diarila e seus derivados substituídos. I.Seixas, Fabiana Kömmling. II.Collares, Tiago Veiras. III.Título.

CDD:616.994

Banca examinadora:

Prof. Dr. Luciano da Silva Pinto

Prof. Dr. Benildo Sousa Cavada

Prof. Dra. Sandra Beatriz Chaves Tarquinio

Prof. Dra. Fabiana Kömmling Seixas

Dedicatória

Ao meu Pai pelo seu amor incondicional com o qual continua a me iluminar.

Agradecimentos

A Universidade Federal de Pelotas que viabilizou a minha formação e despertou em mim um grande afeto por esta instituição.

A minha orientadora e o meu co-orientador, Fabiana K. Seixas e Tiago V. Collares, agradeço por terem abraçado esta oportunidade comigo de ingressar diretamente no Doutorado. E me oferecerem uma grande oportunidade de crescimento junto a disciplina de Engenharia Tecidual.

Um agradecimento especial ao Prof. Odri A. Dellagostin, que sugeriu o meu nome para esta vaga de Doutorado e por quem sempre nutri um grande carinho.

A família Demarco, grandes amigos, que continuaram me acompanhando, orientado e me servindo como modelos durante esta trajetória.

Ao grupo GPO, em especial a quatro amigos muito queridos Priscila de Leon, Helena Thurow, Vinicius Campos e Samuel Ribeiro. Eu guardo com muito carinho momentos maravilhosos com vocês.

As estagiárias e amigas Fausto Gomes, Vanessa Penna, Fernanda Rodrigues, Júlia Sallaberry, Karine Duarte e Stéphanie Caruccio, pelo apoio e dedicação.

Aos grandes amigos do Programa de Pós-Graduação em Odontologia, em especial Alessandro Menna, Guilherme Antonello, Eliana Torre, Marcus Conde, Luísa Oliveira, Fernanda Jostmeier, agradeço pelo apoio, os sorrisos e carinho de vocês.

Ao meu noivo Mateus da Costa, que esteve sempre presente, de forma ainda mais intensa nos momentos difíceis, caminhando ao meu lado e me deixando segura para que pudesse seguir o meu caminho. Serei sempre a tua admiradora.

A minha mãe e irmã, Clair Nedel e Ana Paula Nedel, que foram incansáveis e perseguiram os meus sonhos como se fossem seus. A minha eterna gratidão.

Ao meu Pai, Jorge Nedel. Pai és a minha inspiração, aquilo que me ensinate guardo dentro de mim, e se hoje finaliza-se mais uma etapa é porque tive um anjo que iluminou o meu caminho.

A todos que de alguma forma participaram deste período a minha gratidão e o meu carinho.

RESUMO

NEDEL, Fernanda. **Seleção de novas moléculas e modalidades de tratamento no combate ao câncer.** 2012. 89 f. Tese (doutorado) – Programa de Pós-Graduação em Biotecnologia. Universidade Federal de Pelotas, Pelotas.

O câncer é uma das principais causas de morte no mundo, onde os índices devem aumentar 50% até 2020. Embora a ressecção cirúrgica e terapias adicionais (como a quimioterapias e radioterapias) sejam capazes de curar tumores primários bem delimitados, o mesmo não se aplica a metástase devido ao seu envolvimento sistêmico e a resistência a terapias convencionais. Portanto, atualmente o desfio clínico é desenvolver novas drogas e modalidades de tratamentos que irão impactar significativamente as taxas de cura do câncer. Neste sentido, o presente trabalho objetivou avaliar o efeito antineoplásico e investigar a rota de apoptose induzido pelo disseleneto de diarila e seus derivados substituídos - (4-CIC₆H₄Se)₂, (3-CF₃C₆H₄Se)₂ e (4-MeOC₆H₄Se)₂ - em células de adenocarcinoma de colorretal humano (HT-29). Verificamos que os compostos (3-CF₃C₆H₄Se)₂ e (4-MeOC₆H₄Se)₂ induziram um efeito citotoxidade por meio de apoptose, onde os genes pró-apoptoticos (Bax, caspase-9, caspase-8, fator indutor de apoptose (AIF) e endonuclease G (EndoG)) foram altamente expressos e os genes anti-apoptótico (Bcl-2 e survivin) mostraram uma redução na sua expressão. Em um segundo momento avaliamos o potencial antineoplásico das lectinas Canavalia brasiliensis (ConBr), Canavalia boliviana (ConBol) e Canavalia ensiformis (ConA) em células HT-29, as quais se mostraram efetivas em reduzir a viabilidade celular. Uma vez confirmado o efeito antineoplásico, as lectinas forma marcadas com FITC e a sua interação com as células tumorais foi investigado. As lectinas FITC-ConA e FITC-ConBol demonstraram potencial de se ligar as células HT-29 ao contrário da FITC-ConBr. A fim de investigar uma nova modalidade de tratamento foi avaliada a interação entre as respectivas lectinas com as células HT-29 quando associadas à nanotubos de carbonos funcionalizados de paredes múltiplas (f-MWCNTs). Quando os f-MWCNTs foram incorporados as lectinas FITC-ConA e FITC-ConBol houve um aumentaram na intensidade de fluorescência.

Palavras-chaves: adenocarcinoma de colorretal humano; disseleneto de diarila e seus derivados substituídos; lectinas; nanotubos de carbono.

ABSTRACT

NEDEL, Fernanda. **Selection of new molecules and treatment modalities to fight cancer.** 2012. 89 p. Tese (doutorado) – Programa de Pós-Graduação em Biotecnologia. Universidade Federal de Pelotas, Pelotas.

Cancer is a leading cause of death and its rates are expected to increase 50% by 2020. Although surgical resection and additional therapies (such as chemotherapy and radiotherapy) are able to cure well-confined, primary tumors, the same does not apply during metastasis due to the systemic involvement and its resistance to conventional therapies. Therefore, the current clinical challenge is to develop new drugs and treatment modalities that will significantly impact the cure rates. In this sense, the present study aimed to evaluate the anticancer effect and study the underlying cell death mechanisms of diaryl diselenides and its substituted structures - $(4-CIC_6H_4Se)_2$, $(3-CF_3C_6H_4Se)_2$ e $(4-MeOC_6H_4Se)_2$ - on the human colon adenocarcinoma cell line (HT-29). We verified that (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ induced cytotoxicity through apoptosis mechanisms in HT-29 cells, where pro-apoptotic genes were up-regulated (Bax, caspase-9, caspase-8, apoptosis-inducing factor (AIF) and endonuclease G (EndoG), and anti-apoptotic genes were down-regulated (Bcl-2 and survivin). In a second moment we evaluated the anticancer potential of Canavalia brasiliensis (ConBr), Canavalia boliviana (ConBol) and Canavalia ensiformis (ConA) lectins in HT-29 cells, which showed an effective capacity to reduce cell viability. Once the anticancer effect was confirmed, lectins were labeled with FITC and its interaction with the tumor cells was investigated. The FITC-ConA and FITC-ConBol demonstrated the potential to bind to HT-29 cells unlike FITC-ConBr. In order to investigate a new treatment modality, the interaction between the respective lectins with HT-29 was evaluated when associated with functionalized multi-walled carbon nanotubes (f-MWCNTs). When f-MWNT was incorporated to FITC-ConBol and FITC-ConA lectins there was an increase in fluorescence intensity.

Keywords: human colon adenocarcinoma; substituted diaryl diselenides; lectins; carbon nanotubes.

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1. INTRODUÇÃO GERAL

Atualmente o câncer é uma das principais causas de morte no mundo, onde esse número deve aumentar significativamente, em parte, devido ao envelhecimento da população global (VISVADER,J.E., 2011). Segundo a Organização Mundial de Saúde (OMS), a incidência de câncer, pode aumentar em 50% nos próximos 20 anos, passando de 10 milhões de pessoas acometidas em 2000, para 15 milhões em 2020. Com base no Relatório Mundial sobre o Câncer, recentemente divulgado, a OMS solicitou que governos, autoridades e público em geral empreendam ações urgentes para evitar a ampliação do número de vítimas dessa enfermidade.

Os tratamentos mais comuns contra o câncer são restritos e envolvem a quimioterapia, radioterapia e intervenções cirúrgicas (MISRA,R. et al., 2010). Estes tratamentos são capazes de curar tumores primários bem delimitados, no entanto, o mesmo não se aplica a metástase devido ao seu envolvimento sistêmico e resistência a terapias convencionais (CHAFFER,C.L.;WEINBERG,R.A., 2011;VALASTYAN,S.;WEINBERG,R.A., 2011). Portanto, atualmente o desafio clínico é desenvolver novos medicamentos e modalidades de tratamento que irão impactar significativamente as taxas de cura, eliminando à resistência as drogas pelas células tumorais, promovendo o mínimo de toxicidade (MIURA,K. et al., 2011;SANTANDREU,F.M. et al., 2011).

O câncer se desenvolve em função de mutações em genes envolvidos no controle da proliferação e apoptose celular (morte celular programada), permitindo que as células obtenham a habilidade de invadir tecidos promovendo a formação de metástases (BARBELLIDO,S.A. et al., 2008). Considerando que um dos processos mais importantes na regulação do equilíbrio entre o crescimento e morte células é a apoptose, esta tem atraído uma crescente atenção no desenvolvimento de terapias para o tratamento do câncer (LIU,J.J. et al., 2011). De fato as terapias que envolvem a eliminação de células tumorais por meio da citotoxidade, como por exemplo, a quimioterapia, imunoterapia e radiação γ, são predominantemente mediadas pela indução de morte celular programada (FULDA,S.;DEBATIN,K.M., 2004).

Uma molécula crucial para a apoptose são as caspases, pois são tanto iniciadoras como executoras do processo. Existem três vias através das quais podem ser ativadas as caspases. As duas vias de inicialização mais comumente

descritas são a via extrínseca (receptor da morte) e a via intrínseca (mitocondrial). A via extrínseca é desencadeada pelo receptor da morte Fas, o qual é dependente da combinação com o Fas ligante (Fas-L). Quando o estímulo para a morte celular ocorre, Faz-L pode se ligar diretamente com o receptor Faz, formando um complexo, o qual promove o recrutamento dos adaptadores de proteínas intracelulares que ligam e agregam moléculas de procaspase-8, as quais clivam e ativam uma a outra. Esse complexo, por sua vez, irá ativar a procaspase-3 para induzirem o processo de apoptose (LIU,J.J. et al., 2011; WONG,R.S., 2011). A via intrínseca é uma outra estratégia que leva a apoptose, onde a mitocôndria desempenha um papel central. Quando a célula é exposta a um estímulo extracelular (por exemplo, UV, Raio-X e toxina) ou sinais intracelular (danos ao DNA, instabilidade nuclear) a membrana mitocondrial externa torna-se permeável, liberando citocromos-c. A liberação do citocromo-c promove o recrutamento de uma proteína adaptadora Apaf-1 e a procaspase-9 formando o apoptossomo e desencadeando a morte celular. Esta via é rigorosamente regulada por um grupo de proteínas pertencentes à família Bcl-2. Existem dois grupos principais de proteínas Bcl-2 uma pró-apoptotica como exemplo o Bax, e outra anti-apoptótica como o Bcl-2. Enquanto as proteínas anti-apoptóticas regulam a apoptose através do bloqueio da liberação do citocromo-c pela mitocondria, as proteínas pró-apoptóticas agem estimulando a sua liberação (LIU,J.J. et al., 2011;WONG,R.S., 2011).

Assim, durante algumas vias da apoptose a membrana externa da mitocôndria pode ser despolarizada, tornando-se parcialmente permeável a proteínas. O resultado deste processo pode acarretar na liberação de proteínas como a endonuclease G (EndoG) e fator indutor de apoptose (apoptosis-inducing factor - AIF). Acredita-se que ambas as proteínas uma vez liberadas da membrana mitocondrial translouquem para o núcleo da célula, onde elas participam da degradação da cromatina, atuando, portanto, em uma via independente de caspases (VARECHA,M. et al., 2012). Portanto, a indução da apoptose tem sido reconhecida como uma estratégia ideal para quimioterápicos, onde compostos com a habilidade de induzir apoptose em tumores possuem o potencial de serem utilizados na terapia do câncer (YAN,Q. et al., 2009).

O selênio é um elemento traço essencial, o qual sob a forma de selenocisteína compõe proteínas e enzimas antioxidantes, as quais são exigidas para as diversas funções biológicas (ZENG,H.;COMBS,G.F., 2008). Os compostos

de selênio demonstraram a capacidade de prevenir o câncer em diversos modelos animais e aumentar a eficácia quimiopreventiva em humanos com câncer de pulmão, colorretal, cabeça e pescoço e próstata, embora a mesma taxa de sucesso não tenha sido observado com o câncer de mama e de pele (ROSA,R.M. et al., 2007;SUZUKI,M. et al., 2010). O selênio parece ter uma ação antimutagênico durante os estágios iniciais do câncer, prevenindo a ativação de oncogenes e a transformação de células normais em um fenótipo maligno (ROSA,R.M. et al., 2007). Além da atividade quimiopreventiva, evidências recentes tem apontado para o potencial quimioterapêutico destes compostos (SUZUKI,M. et al., 2010). Os compostos derivados de selênio podem ser inorgânicos, tais como o selenito e o selenato, ou compostos orgânicos, como o selenometionina e disselentos de diarila (NOGUEIRA, C.W.; ROCHA, J.B., 2011). O selenito de sódio pode induzir a apoptose em linhagens de células tumorais. No entanto, a apoptose ocorre como resultado da toxicidade do selênio, causando quebras no DNA e danos cromossomais nas células de linhagens tumorais e linfócitos humanos, respectivamente (SUZUKI,M. et al., 2010). Por outro lado, os compostos orgânicos de selênio têm menos efeitos colaterais e ausência de efeitos genotóxico quando comparados com os selênios inorgânicos (RIKIISHI,H., 2007;SUZUKI,M. et al., 2010). O disseleneto de diarila, um composto orgânico de selênio, é estável e vem sendo utilizado na síntese de uma variedade de compostos organoselênio farmacologicamente ativos. Estudos experimentais tem demonstrado o potencial de proteção do disseleneto de diarila, com propriedade antioxidantes, anti-hiperglicêmicas e anti-inflamatórias (ROSA,R.M. et al., 2007;SAVEGNAGO,L. et al., 2008;NOGUEIRA,C.W.;ROCHA,J.B., 2011). Recentemente este composto tem demonstrado a capacidade de induzir efeitos citotóxicos na linhagem de neuroblastoma humano (SH-SY5Y), o qual é possivelmente mediado pela via de ERK1/2 (POSSER,T. et al., 2011). No entanto, a introdução de grupos funcionais (por exemplo, cloro, flúor e metoxila) no anel aromático do disseleneto de diarila pode alterar as propriedades da molécula proporcionando uma alternativa aos atuais agentes terapêuticos (MACHADO M.A., et al., 2009;SAVEGNAGO,L. et al., 2009). De fato a introdução do grupamento cloro no grupo aril do disseleneto de diarila confere um menor efeito citotóxico sobre as células V79 (células de fibroblastos de pulmão de hamster chinês) em comparação com o dissleneto de diarila (MACHADO M.A., et al., 2009).

No entanto, embora muitas pesquisas tem sido realizadas com a finalidade de identificar novas drogas para o tratamento do câncer, alguns importantes inconvenientes tem limitado o seu sucesso. Dentre estes estão: a falta de seletividade adequada contra as células tumorais e consequente toxicidade para os tecidos saudáveis; biodisponibilidade limitada para o tecido tumoral, exigindo altas dosagens de medicamentos; incapacidade de algumas drogas de atravessarem as barreiras celulares; e especialmente, o aparecimento frequente de resistência a múltiplas drogas (JI,S.R. et al., 2010;SHAPIRA,A. et al., 2011), fenômeno que contribui para a falha do tratamento em 90% dos pacientes com metástase (GAO,Z. et al., 2012). A fim de superar estas questões, não somente o desenvolvimento de novos medicamentos deve ser considerado, mas também os sistemas pelo qual as drogas podem ser entregues (drug delivering systems). A partir desta perspectiva, uma maneira plausível de entrega eficiente de drogas para o tecido tumoral seria associar drogas antineoplásica com nanopartículas (HU,C.M.;ZHANG,L., 2009;SHAPIRA,A. et al., 2011).

De fato a nonotecnologia - área que envolve o desenho e construção de estruturas com um espectro de tamanho de 1-1.000 nm - fornece uma alternativa promissora quimioterapias convencionais inovadora е а (PETROS,R.A.; DESIMONE, J.M., 2010). Esta estratégia emergente tem demonstrado uma melhor eficácia terapêutica com efeitos colaterais reduzidos, em comparação com modalidades terapêuticas clássicas, envolvendo drogas não alvo específicas (HU,C.M.;ZHANG,L., 2009;SHAPIRA,A. et al., 2011). Devido a sua capacidade de encapsular, aderir e/ou conjugar drogas ou produtos biológicos terapêuticos a sua estrutura, as nanopartículas tem sido vislumbradas como passiveis de sobrepujarem a resistência das células tumorais a múltiplas drogas. Simultaneamente nanocarreadores multifuncionais os podem aumentar penetrabilidade por barreiras fisiológicas e proteger os fármacos ou produtos biológicos terapêuticos do processo de degradação. Ainda, os nanocarreadores podem auxiliar na solubilização de drogas hidrofóbicas, diminuir a depuração de fármacos, regular a liberação de drogas assim como garantir a entrega no alvo específico de interesse (GAO,Z. et al., 2012;HU,C.J. et al., 2012).

Entre as diversas classes de nanomateriais os nanotubos de carbono (CNT) tem atraído atenção especial devido as suas propriedades espectroscópicas, térmicas e elétricas (JI,S.R. et al., 2010;SHAPIRA,A. et al., 2011) Vários estudos tem

demonstrado que CNT funcionalizados são capazes de internalizar em uma variedade de tipos celulares, através do cruzamento da membrana celular por endocitose ou por meio de outros mecanismos (JI,S.R. et al., 2010). Este processo se torna possível uma vez que as dimensões destas estruturas tubulares variam tipicamente de 0,4-2 nm de diâmetro nos nanotubos de carbonos de parede única (SWCNTs), 1-3 nm para os nanotubos de carbono de parede dupla (DWCNTs) e 2-100 nm de nanotubos de carbono de paredes múltiplas (MWCNTs). A despeito das suas nanodimensões, os CNT tem uma elevada área superficial que permite o carregamento de componentes ativos em densidades elevadas no seu interior, e/ou através de ligações funcionais do composto as paredes externas dos CTNs. No entanto, os CNTs possuem uma superfície muito hidrofóbica e, portanto, não são facilmente suspensos em solventes fisiológicos, o que pode ser uma desvantagem importante na aplicação clínica (RYBAK-SMITH,M.J.;SIM,R.B, 2011). Desta forma as paredes laterais dos CNTs têm sido alteradas de forma não covalente e covalente. Através da introdução de grupos funcionais polares as forças de van der Waals entre os CNTs individualizados ou agregados são eliminados, modificando a sua interação com os fluídos biológicos e permitindo a introdução de moléculas alvos, tais como fragmentos de anticopos (CHENG, W. et al., 2008; ZHANG, X. et al., 2010; VEDALA, H. et al., 2011;XUE,Y. et al., 2011).

Os CNTs têm sido utilizados como sistemas de entrega para uma variedade de componentes, incluindo tipicamente fármacos anticancerígenos e antifúngicos; bimoléculas, tais como proteínas, peptídeos, DNA e siRNA; ligando alvos tais como vitaminas, peptídeos e anticorpos que tem possibilitado o fornecimento de drogas a um tecido ou subpopulação de células específicas. Alguns estudos têm associado lectinas, CNTs, e células tumorais. No entanto o foco principal tem sido no desenvolvimento de tecnologias de monitoramento que sejam práticas e de alta sensibilidade e rendimento, para a análise do status de glicosilação das células tumorais e para fornecer uma ferramenta de diagnóstico que possa guiar o tratamento do câncer (CHENG,W. et al., 2008;ZHANG,X. et al., 2010;VEDALA,H. et al., 2011;XUE,Y. et al., 2011).

As lectinas são proteínas ou glicoproteínas de origem não imune que contêm pelo menos um domínio não catalítico, o que lhes permite seletivamente reconhecer e se ligar reversívelmente a açucares livres ou glicanos específicos presentes em glicoproteínas e glicolipídeos, sem alterar a estrutura do carboidrato (KOMATH,S.S.

et al., 2006;MONIRA,P. et al., 2009;FU,L.L. et al., 2011;RUSSI,M.A. et al., 2012). Esta especificidade vem sendo profundamente correlacionada com o câncer, uma vez que a transformação neoplásica das células é geralmente associada com alterações nos glicoconjugados da superfície celular, tais como alterações na ramificação de carboidratos complexos e ocasionalmente o aparecimento de estruturas incomuns (MODY,R. et al., 1995;KAUR,M. et al., 2006;SUJATHAN,K. et al., 2009). De fato, as lectinas de plantas foram utilizadas como ferramentas simples de reconhecimento tumoral para diferenciar tumores malignos de tumores benignos e avaliar o grau de glicosilação associada à metástase (MODY,R. et al., 1995;KAUR,M. et al., 2006;SUJATHAN,K. et al., 2009). Recentemente, as lectinas têm sido utilizadas em análises de *microarrays* para melhorar o reconhecimento de tumores malignos (GUPTA,G. et al., 2010;FU,L.L. et al., 2011). Além disso, as lectinas de plantas possuem uma atividade antitumoral, induzido apoptose predominantemente através de vias dependentes de caspases (LIU,B. et al., 2009).

Assim este trabalho foi delineado visando avaliar o potencial antineoplásicos de moléculas distintas: do disseleneto de diarila e seus derivados substituídos e de três tipos de lectinas - Canavalia brasiliensis (ConBr), Canavalia boliviana (ConBol) e Canavalia ensiformis (ConA) -. Mediante os resultados de citotoxidade para o disseleneto de diarila e seus derivados substituídos objetivamos avaliar a possível rota de apoptose pelo qual as moléculas poderiam estar promovendo a morte celular. Para as três lectinas uma vez comprovada a sua citotoxidade avaliamos se estas eram capazes de interagir com as células tumorais. E considerando os avanços na área de nanotecnologia no câncer investigamos se as lectinas associadas à nanotubos de carbono de paredes múltiplas funcionalizados (f-MWNT) poderia aumentar a interação célula-lectina.

Os dados gerados nesta tese estão na forma de artigos científicos, visando, assim, proporcionar uma divulgação objetiva e rápida dos resultados obtidos. Neste contexto, o artigo 1 teve por objetivo investigar os efeitos e o mecanismo de ação do disseleneto de diarila e seus derivados substituídos sobre as células de adenocarcinoma de colorretal (HT-29). Este trabalho está formatado de acordo com as normas do periódico **Life Sciences**.

O artigo 2 descreve a capacidade antineoplásica das lectinas Canavalia brasiliensis (ConBr), Canavalia boliviana (ConBol) e Canavalia ensiformis (ConA) e a

sua associação a f-MWNT. Este trabalho está formatado segundo as normas do periódico **Carbon**.

2. ARTIGO 1

Substituted diaryl diselenides: Cytotoxic and apoptotic effect in human colon adenocarcinoma cells

(Artigo formatado segundo as normas do periódico Life Sciences)

Substituted diaryl diselenides: cytotoxic and apoptotic effect in human colon

adenocarcinoma cells

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ABSTRACT

Aims: To investigate the effects and study the underlying cell death mechanisms of diaryl diselenides, including: diphenyl diselenide $(C_6H_5Se)_2$; 4-chlorodiphenyl diselenide (4- $ClC_6H_4Se)_2$; 3-(trifluoromethyl)-diphenyl diselenide (3- $CF_3C_6H_4Se)_2$ and 4-methoxydiphenyl diselenide (4- $MeOC_6H_4Se)_2$, on the human colon adenocarcinoma cell line HT-29.

Main methods: The viability of HT-29 cells after exposure to the diaryl diselenides and its substituted structures was based on the MTT assay. To verify if cell death was mediated throughout apoptosis mechanisms, flow cytometry and real-time PCR (qPCR) analyses were conducted.

Key findings: The MTT assay and flow cytometry analyses showed that (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ induced cytotoxicity through apoptosis mechanisms in HT-29 cells. qPCR revealed there was an up-regulation of pro-apoptotic (Bax, casapase-9, caspase-8, apoptosis-inducing factor (AIF) and Endonuclease G (EndoG) and cell-cycle arrest genes (p53 and p21) and down-regulation of anti-apoptotic (Bcl-2 and survivin) and Myc genes.

Significance: These results demonstrate that (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ have the potential to induce apoptosis in HT-29 cells through the activation of caspase-dependent and independent pathways and through cell-cycle arrest.

Keywords: Human colon adenocarcinoma; Apoptosis; Substituted diaryl diselenides; Selenium; Cancer

Introduction

Colorectal cancer is one of the leading causes of cancer mortality (Limami et al. 2011), corresponding to 9.4% of all cases of cancer worldwide (Cantero-Muñoz et al. 2011). Fifty percent of all recently diagnosed patients ultimately develop metastatic disease. Regardless of the advances in developing new chemotherapy agents, no drug has been able to treat colorectal cancer metastasis with a non-relapsing cure rate. Currently the clinical challenge is to develop new drugs that will have a significant impact on cure rates, by reversing drug resistance, and with minimal toxicity (Miura et al. 2011).

Selenium is an essential trace element, (Zeng and Combs 2008) that has the ability to prevent cancer in several animal models and to enhance chemopreventive efficacy in human lung, colorectal, head and neck and prostate cancer (Suzuki et al. 2010). The chemopreventive role of selenium is well supported by epidemiological, preclinical, and clinical evidence (Clark et al. 1998). Furthermore, emerging evidence has indicated the potential of selenium compounds in cancer chemotherapy (Suzuki et al. 2010).

Diphenyl diselenide (C_6H_5Se)₂, an organic selenium compound, has raised great interest due to its antioxidant, antidepressant-like, neuroprotective and antinociceptive properties (Nogueira and Rocha 2011; Savegnago et al. 2008a; Savegnago et al. 2008b; Savegnago et al. 2007). Recently, Posser et al. (2011) showed, for the first time, that (C_6H_5Se)₂ was cytotoxic to human cancer cells (SH-SY5Y) in vitro, possibly mediated by the ERK1/2 pathway (Posser et al. 2011). However, to date no study has evaluated the cytotoxic effect of (C_6H_5Se)₂ in other human cancer cell types.

In addition, studies have demonstrated that the introduction of a substitute (e.g., chloro, fluor or methoxyl) in the aromatic ring of $(C_6H_5Se)_2$ can alter its molecules properties (Machado et al. 2009; Savegnago et al. 2009; Wilhelm et al. 2009). The introduction of chloro into the aryl group of diaryl diselenide conferred a weak cytotoxic effect on V79 cells (Chinese hamster

lung fibroblast cells) compared to $(C_6H_5Se)_2$ (Machado et al. 2009; Savegnago et al. 2009; Wilhelm et al. 2009). Although this substitute could alter the biological effects of $(C_6H_5Se)_2$, their potential as cytotoxic agents for cancer chemotherapy has not yet been explored.

Therefore, our objective was to investigate the effect and the underlying cell death mechanisms of $(C_6H_5Se)_2$ and its substituted structures, 4-chlorodiphenyl diselenide (4- $ClC_6H_4Se)_2$, 3-(trifluoromethyl)-diphenyl diselenide (3- $CF_3C_6H_4Se)_2$ and 4-methoxydiphenyl diselenide (4- $MeOC_6H_4Se)_2$ on the human colon adenocarcinoma cell line (HT-29). In addition, we also verified whether the introduction of an electron donating (-methoxyl) or an electron withdrawing group (-chloro and -trifluoromethyl) into the aryl group of diaryl diselenide altered its biological effect. To the best of our knowledge this is the first study that demonstrates the effect of $(C_6H_5Se)_2$ and its substituted structures on HT-29 cells.

Materials and Methods

Chemicals

(C₆H₅Se)₂, (4-ClC₆H₄Se)₂, (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ (Fig. 1) were prepared according to methods in the literature. Analysis of ¹H and ¹³C NMR spectra showed that the analytical and spectroscopic data was in full agreement with its assigned structure. The chemical purity of these compounds was determined by gas chromatography/mass spectrometry.

Cell Culture

The HT-29 cells were obtained from the Rio de Janeiro Cell Bank (PABCAM, Federal University of Rio de Janeiro, RJ, Brazil). The cells were cultured in Dulbecco's modified Eagle's medium (DMEM), supplemented with 10% foetal bovine serum (FBS), purchased from Vitrocell Embriolife (Campinas, Brazil) and Gibco (Grand Island, NY, USA),

respectively. Cells were grown at 37 °C in an atmosphere of 95% humidified air and 5% CO₂. The experiments were performed with cells in the logarithmic phase of growth.

Determination of cytotoxicity

The viability of the HT-29 cells was determined by measuring the reduction of soluble MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] to water insoluble formazan (Ali et al. 2010; Henn et al. 2011). Briefly, cells were seeded at a density of 2 x 10⁴ cell per well in a volume of 100 µL in 96-well plates and grown at 37 °C in a humidified atmosphere of 5% CO₂/95% air for 24 h before being used in the MTT assay. Cells were incubated with different concentration of $(C_6H_5Se)_2$, $(4-ClC_6H_4Se)_2$, $(3-CF_3C_6H_4Se)_2$ or $(4-MeOC_6H_4Se)_2$ (5 - 80 μM) for 24, 48 and 72 h. These compounds were dissolved in dimethyl sulfoxide (DMSO) and added to the DMEM supplemented with 10% FBS to the desired concentrations. The final DMSO concentration in the culture medium never exceeded 0.8% and a control group exposed to an equivalent concentration of DMSO was evaluated. After incubation the media was removed and 180 µL of DMEM and 20 µL MTT (5 mg MTT/mL solution) was added to each well. The plates were incubated for an additional 3 h and the medium was discarded. 200 µL of DMSO was added to each well, and the formazan was solubilised on a shaker for 5 min at $100 \times g$. The absorbance of each well was read on a microplate reader (MR-96A, Mindray Shenzhen, China) at a wavelength of 492 nm. The percentage inhibition of cell growth was determined as follows: inhibitory rate = $(1 - Abs_{4treated cells}/Abs_{492control cells}) \times$ 100 (Zheng et al. 2011). All observations were validated by at least three independent experiments and for each experiment the analyses were performed in triplicate.

Apoptotic assay

The Guava Nexin assay (Guava Technologies) was conducted following the manufacturer's instructions. Briefly, 2.0×10^4 to 1.0×10^5 of the treated HT-29 cells (100 μ L) were added to 100 μ L of Guava Nexin reagent. Cells were incubated in the dark at room temperature for 20 min and samples (2000 cells per well) were acquired on the flow cytometry Guava EasyCyte System. In this assay, an annexin V-negative and 7-AAD-positive result indicated nuclear debris, an annexin V-positive and 7-AAD-positive result indicated late apoptotic cells, while an annexin V-negative and 7-AAD-negative result indicated live healthy cells and annexin V-positive and 7-AAD-negative result indicated live healthy cells and annexin V-positive and 7-AAD-negative result indicated the presence of early apoptotic cells.

Gene expression evaluation by real-time PCR

The HT-29 cells were seeded in 6-well flat bottom plate at a density of 2 x 10⁵ per well and grown at 37 °C in a humidified atmosphere of 5% CO₂/95% air for 24 h. Cells were then exposed to 20, 40 and 80 μM of (C₆H₅Se)₂, (3-CF₃C₆H₄Se)₂ or (4-MeOC₆H₄Se)₂ for 48 h. After this period the cells were washed with 1x phosphate-buffered saline (PBS; Gibco) and the RNA was extracted from the cells. Total RNA extraction, cDNA synthesis and real-time PCR (qPCR) were carried out as previously described (Campos et al. 2010). Briefly, RNA samples were isolated using TRIzol Reagent (Invitrogen) and samples were DNase-treated with a DNA-free kit (Ambion, USA) following the manufacturer's protocol. First-strand cDNA synthesis was performed with 2 μg of RNA using High Capacity cDNA Reverse Transcription kit (Applied Biosystems, UK) according to the manufacturer's protocol. The qPCR reactions were run on a Stratagene Mx3005P Real-Time PCR System (Agilent Technologies, Santa Clara, CA, USA) using SYBR Green PCR Master Mix (Applied Biosystems, UK) using the primers described in Table 1.

Data analysis

Data sets from the MTT assay and qPCR were analysed using a two-way ANOVA followed by a Tukey test for multiple comparisons. Two factors were considered: the compound used (four levels) and the concentration of the compound (three levels). Significance was considered at p < 0.05 in all analyses. The data are expressed as the means \pm SEM.

Results

Determination of cytotoxicity

Both the $(C_6H_5Se)_2$ and $(4\text{-ClC}_6H_4Se)_2$ compounds had a significant cytotoxic effect on the HT-29 cells at 80 μ M and this effect improved significantly with exposure time (Fig. 2). Both the $(3\text{-CF}_3C_6H_4Se)_2$ and $(4\text{-MeOC}_6H_4Se)_2$ compounds achieved significant cytotoxicity at a concentration of 20 μ M. After 48 h exposure to 20 μ M $(3\text{-CF}_3C_6H_4Se)_2$, cytotoxicity was 24% (p < 0.05) and this increased significantly to 96% at 80 μ M (Fig. 2). The cytotoxicity of the $(4\text{-MeOC}_6H_4Se)_2$ compound at 20 μ M, after 24 h exposure, was 44% and further increases in the concentration of the compound resulted in significant reduction in the viability of the HT-29 cells (62 and 75% cytotoxicity, Fig. 2). The exposure time had no significant effect on the cytotoxicity of the $(3\text{-CF}_3C_6H_4Se)_2$ compound. Only the $(4\text{-MeOC}_6H_4Se)_2$ compound showed a significant improvement with exposure time, for example, at 20 μ M and after 24 and 48 h exposure, cytotoxicity increased from 44 to 65%, respectively, although there was no further improvement at 72 h (Fig. 2). The presence of 0.8% DMSO in the culture medium had no effect on cell viability, as compared to the control cells without DMSO.

Apoptosis analysis

The Annexin-PE staining assay was performed to further characterize the observation that the (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ compounds could induce apoptosis in HT-29 cells after exposure for 48h. Annexin V binds to those cells that express phosphatidylserine on the outer

layer of the cell membrane, a characteristic feature of cells entering apoptosis. The results indicated that $(C_6H_5Se)_2$ induced apoptosis at a concentration of 80 μ M (22.5%, Fig. 3B). The lower concentrations (20 and 40 μ M) of $(C_6H_5Se)_2$ where not effective in causing cell death through apoptosis, inducing similar levels of apoptosis (5.2 and 6.1%, respectively) seen in the control groups (3.0 and 6.1%, respectively). The (3-CF₃C₆H₄Se)₂ compound induced a higher percentage of apoptosis at the 40 and 80 μ M concentrations (22.3 and 84.7%, respectively) compared to the controls and the $(C_6H_5Se)_2$ compound. At the 20 μ M concentration the percentage of apoptotic cells was 7.8%, similar to that observed in the control groups. The (4-MeOC₆H₄Se)₂ compound was able to induce significant apoptosis in the HT-29 cells at 20 μ M (38.6%), this increased to 58.9% upon exposure to a concentration of 40 μ M, although a further increase in concentration to 80 μ M did not increase apoptosis (54.7%). Apoptosis induction from exposure of the HT-29 cells to 0.8% DMSO had no effect.

Gene expression

In order to evaluate the likely apoptosis pathways activated by $(3\text{-}CF_3C_6H_4Se)_2$ and $(4\text{-}MeOC_6H_4Se)_2$ in HT-29 cells (48 h exposure), anti-apoptotic and pro-apoptotic gene expression were investigated. Bax mRNA levels were significantly higher (p<0.05) in cells exposed to $(3\text{-}CF_3C_6H_4Se)_2$ (80 μ M) and $(4\text{-}MeOC_6H_4Se)_2$ (20, 40 and 80 μ M) when compared to the control groups (Fig. 4A). However, $(C_6H_5Se)_2$ had no effect on Bax mRNA levels when compared to the control groups (p>0.05). Bcl-2 mRNA levels decreased significantly (p<0.05) in cells exposed to $(3\text{-}CF_3C_6H_4Se)_2$ (80 μ M) and $(4\text{-}MeOC_6H_4Se)_2$ (40 and 80 μ M) when compared to control groups. HT-29 cells exposed to $(3\text{-}CF_3C_6H_4Se)_2$ (40 and 80 μ M) decreased Bcl-2 mRNA levels when compared to control groups (p<0.05) (Fig. 4B). Caspase 9 was up-regulated (p<0.05) in cells treated with $(3\text{-}CF_3C_6H_4Se)_2$ (80 μ M), $(4\text{-}MeOC_6H_4Se)_2$ (40 and 80 μ M)

(Fig 4C). Exposure to $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})_2$ (20 and 40 μM), $(4\text{-MeOC}_6\text{H}_4\text{Se})_2$ (20 μM) and $(C_6\text{H}_5\text{Se})_2$ (20, 40 and 80 μM) had no effect on caspase 9 gene expression (p>0.05). However, caspase 8 mRNA levels were significantly higher (p<0.05) in cells exposed to (4-MeOC $_6\text{H}_4\text{Se})_2$ (40 and 80 μM) when compared to the control groups. $(C_6\text{H}_5\text{Se})_2$, (3-CF $_3\text{C}_6\text{H}_4\text{Se})_2$ and $(4\text{-MeOC}_6\text{H}_4\text{Se})_2$ (20 μM) did not affect caspase 8 gene expression (p>0.05) (Fig. 4D). Survivin expression was significantly down-regulated (p<0.05) in HT-29 cells treated with $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})_2$ (40 and 80 μM), $(4\text{-MeOC}_6\text{H}_4\text{Se})_2$ (20, 40 and 80 μM) and $(C_6\text{H}_5\text{Se})_2$ (80 μM) when compared to the control group (Fig. 4E). The $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})_2$ (20 μM) and $(C_6\text{H}_5\text{Se})_2$ (20 and 40 μM) compounds had no effect on survivin expression (p>0.05).

The mRNA levels for AIF and EndoG were also evaluated. AIF expression was significantly up-regulated (p<0.05) upon exposure to (3-CF₃C₆H₄Se)₂ (80 μ M) and (4-MeOC₆H₄Se)₂ (20, 40 and 80 μ M) when compared to the control group (Fig. 4F). However, (C₆H₅Se)₂ and (3-CF₃C₆H₄Se)₂ (20 and 40 μ M) had no effect on AIF mRNA levels when compared to control groups (p>0.05). EndoG mRNA expression was up-regulated (p<0.05) when the HT-29 cells were treated with (C₆H₅Se)₂ (20, 40 and 80 μ M), (3-CF₃C₆H₄Se)₂ (20, 40 and 80 μ M) and (4-MeOC₆H₄Se)₂ (20, 40 and 80 μ M) compared to the control group (Fig. 4G). HT-29 cells treated with (3-CF₃C₆H₄Se)₂ (80 μ M) and (4-MeOC₆H₄Se)₂ (40 and 80 μ M) had altered levels of cell cycle-related gene expression, p53 expression was significantly up-regulated (p<0.05), in comparison to the control groups. (C₆H₅Se)₂, at all concentrations tested, had no effect on p53 mRNA levels (Fig. 5A). p21 gene expression showed the same expression pattern as p53, where (3-CF₃C₆H₄Se)₂ (80 μ M) and (4-MeOC₆H₄Se)₂ (40 and 80 μ M) caused significant up-regulation (p<0.05) and (C₆H₅Se)₂ had no effect (Fig. 5B). MYC gene expression was significantly reduced (p<0.05) in cells treated with (3-CF₃C₆H₄Se)₂ (80 μ M) and (4-MeOC₆H₄Se)₂ (40 and 80 μ M). (C₆H₅Se)₂ had no effect on MYC gene expression (Fig. 5C).

Gene expression upon exposure to 0.8% DMSO was similar to the control group in all experiments.

Discussion

Previous studies have confirmed that organoselenium compounds, such as $(C_6H_5Se)_2$ and its substituted structures, exhibit a remarkable spectrum of pharmacological properties (Machado et al. 2009; Savegnago et al. 2009; Wilhelm et al. 2009). Indeed, $(C_6H_5Se)_2$ has exhibited antioxidant, antidepressant-like, neuroprotective and antinociceptive properties and recently it was demonstrated that $(C_6H_5Se)_2$ had a cytotoxic effect, mediated by the ERK1/2 pathway, on SH-SY5Y cancer cells (Posser et al. 2011). Posser et al. (2011) reported that 30 μ M $(C_6H_5Se)_2$ significantly decreased cell viability in 50% of cells and, at a concentration of 10 μ M, induced changes in cell morphology (Posser et al. 2011). To the best of our knowledge no study has evaluated the effect of $(C_6H_5Se)_2$ and the substituted diaryl diselenides (4-ClC₆H₄Se)₂, (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ as cytotoxic and apoptotic agents against cancer cells in vitro or in vivo.

In the present study, $(C_6H_5Se)_2$ and one of its substituted structures, $(4\text{-CIC}_6H_4Se)_2$, only presented significant cytotoxic effects against the HT-209 cells at a concentration of 80 μ M. A similar study that used a neuroblastoma cell line reported cytotoxic effects at lower concentrations $(10\text{-}30~\mu\text{M}~(C_6H_5Se)_2)$. However, this discrepancy may be related to differences between the SH-SY5Y and HT-29 tumour cell lines, as they exhibit different gene profiles when exposed to potent toxic substances (Thirunavukkarasusx et al. 2011). These results suggest that $(C_6H_5Se)_2$ has a selective action and therefore offers an opportunity to investigate its use as a therapeutic agent. This selectivity has been observed with other selenium compounds, where cancer cells, including lung (A549) and head and neck (HSC-3), were substantially more sensitive to selenite and prone to induction of apoptosis than the

breast cancer cell line MCF-7 (Suzuki et al. 2010). The $(3-CF_3C_6H_4Se)_2$ and (4-MeOC₆H₄Se)₂ compounds induced cytotoxicity and alterations in cell morphology in HT-29 cells in a dose-dependent manner: 20 µM (24.4 vs. 65.2%), 40 µM (81.8 vs. 81.7%) and 80 μM (91.2 vs. 96.1%), respectively. A recent study evaluated the ability of different selenium compounds (selenate, selenite, MeSeA, MeSeCys and SeMet) to induce cell death in HT-29 cells (Lunge et al. 2011). The most effective compound was selenite, an inorganic selenium, the percentage of cell death was 21 (10 µM) and 39% (100 µM), followed by two organic selenium compounds, MeSeA (methylseleninic acid) 2 (10 µM) and 14% (100 µM), and MeSeCys (Se-methylselenocysteine) 3% (100 μM). This suggests that the (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ compounds evaluated in the current study are potentially cytotoxic against human colon adenocarcinoma cells, albeit in vitro. The substitution of a hydrogen atom on the aryl group of diaryl diselenide by an electron withdrawing group (trifluoromethyl) or an electron donating group (-methoxyl) altered the cytotoxicity when compared to diphenyl diselenide. However, these effects were independent of the nature of the aromatic ring in the diaryl diselenide. Both molecules demonstrated greater cytotoxicity compared to $(C_6H_5Se)_2$ and $(4-ClC_6H_4Se)_2$. It has been reported that selenium can inhibit cell proliferation, inducing injury via generation of reactive oxygen species (ROS) (Rudolf et al. 2008). ROS levels can activate the JNK pathway and caspases-3 and 9 via cytochrome c, with down-regulation of Bcl-2 and up-regulation of Bax (Chen et al. 2011). Also, it has been demonstrated that (C₆H₅Se)₂ and (4-ClC₆H₄Se)₂ present higher thiol peroxidase activity and an improved antioxidant potential than (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ in vivo (Meotti et al. 2004). Since, selenium-induced apoptosis in cancer cells can be suppressed by antioxidants (Wu et al. 2010), it is possible that the higher antioxidant potential of (C₆H₅Se)₂ and (4-ClC₆H₄Se)₂ could trigger a less effective cytotoxic effect on HT-29 cells than (3- $CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$.

Since apoptosis is thought to be the mediator of selenium anticancer activity, we verified, by an Annexin-PE staining assay, that the cytotoxicity effect caused by the $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})_2$ and $(4\text{-MeOC}_6\text{H}_4\text{Se})_2$ compounds was mediated by apoptosis. Caspases are central to the mechanism of apoptosis as they are both the initiators and executioners. One pathway by which caspases can be activated involves the extrinsic death receptor pathway, where death ligands bind to death receptors, activating caspase 8 and subsequently initiating apoptosis by cleaving other downstream or executioner caspases (Wong 2011). When $(C_6H_5Se)_2$ and its substituted structures were tested for their ability to stimulate expression of caspase-8, $(4\text{-MeOC}_6H_4\text{Se})_2$ (40 and 80 μ M) was the only compound that induced high levels of caspase-8 mRNA. Since the upstream caspase for the extrinsic death receptor pathway is caspase-8, this suggests that $(4\text{-MeOC}_6H_4\text{Se})_2$ could be activating a death receptor and therefore contributing to apoptosis in the HT-29 cells. In addition, $(4\text{-MeOC}_6H_4\text{Se})_2$ could present a different biological effect from the other substituted structures due to its electron donating group (-methoxyl).

A second pathway involved in caspase activation is the mitochondrial release of cytochrome c (Wong 2011). Cytoplasmatic release of cytochrome c activates capase-3 via formation of a complex (apoptosome) which is made of cytochrome c, APAF-1 and caspase-9 (Jackson and Combs 2008). Bcl-2 (anti-apoptotic) and Bax (pro-apoptotic) are closely involved in this process, an increase in Bcl-2 expression prevents cytochrome c release from the mitochondria, inhibiting the activation of caspase-9 and caspase-3, and preventing apoptosis (Santandreu et al. 2011). In the present study, Bcl-2 expression was down-regulated by (3-CF₃C₆H₄Se)₂ (80 μ M) and (4-MeOC₆H₄Se)₂ (40 and 80 μ M), whereas Bax expression was up-regulated. These findings suggest that Bax and Bcl-2 were involved in mediating the apoptotic effects associated with the cytotoxicity of (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ in HT-29 cells. In addition, caspase-9 mRNA levels were significantly increased by treatment

with (3-CF₃C₆H₄Se)₂ (80 μM) and (4-MeOC₆H₄Se)₂ (40 and 80 μM) showing that caspase-9 was involved in mediating the apoptotic effects associated with these compounds. Apoptosis induced by selenium has been reported to involve activation of the caspases. It was shown that MeSeA induced apoptosis in human prostate cancer (Yamaguchi et al. 2005) and leukaemia cells (Kim et al. 2001) by the activation of multiple caspases (caspase-3, -7, -8 and -9), mitochondrial release of cytochrome c and DNA fragmentation. Other organic and inorganic selenium compounds have been shown to induce caspase-mediated apoptosis, including MeSeCys, selenite (Suzuki et al., 2010), sodium selenite (Chen et al. 2011), selenium dioxide (SeO₂) (Rikiishi 2007).

Additional apoptotic factors that can be released from the mitochondrial intermembrane space into the cytosol are AIF and EndoG, which translocate to the nucleus, triggering chromatin condensation and DNA degradation in a caspase-independent manner (Vařecha et al. 2012; Wong 2011). In the current study AIF gene expression was up-regulated by (3-CF₃C₆H₄Se)₂ (80 μM) and (4-MeOC₆H₄Se)₂ (20, 40 and 80 μM) and EndoG was up-regulated by exposure to the two substituted diaryl diselenides as well as to (C₆H₅Se)₂. These results suggest that diaryl diselenide and its substituted structures could induce apoptosis not only through the activation of multiple caspases but also through a caspase-independent pathway.

Survivin has been implicated in the inhibition of apoptosis, cell proliferation, angiogenesis, and cellular stress response. In HT-29 cells, (3-CF₃C₆H₄Se)₂ (40 and 80 μM), (4-MeOC₆H₄Se)₂ (20, 40 and 80 μM) and (C₆H₅Se)₂ (80 μM) down-regulated the gene expression of survivin. Survivin expression was down-regulated in cell lines derived from prostate cancer cells, such as LNCaP, C4-2 (Chun et al. 2007), DU145 and PC-3 (Hu et al. 2008) treated with selenium. However, when the same selenium compound was tested with a metastatic cell line derived from PC-3 (PC-3M) and two other prostate cancer cell lines (C4-2B and 22Rv1), it had no effect on survivin expression, indicating that the apoptosis induced

by selenium was not mediated by decreasing survivin expression (Liu et al. 2010). These results indicated that selenium could trigger different responses depending on the type of cell. Furthermore, p53 and p21 mRNA expression levels were increased while MYC gene expression was down-regulated upon exposure to (3-CF₃C₆H₄Se)₂ (80 μM), (4-MeOC₆H₄Se)₂ (40 and 80 μM). The expression of p53, p21 and MYC induced by (C₆H₅Se)₂ did not differ from that of the control groups. Investigators have shown that cells deficient in p21 escaped G2/M phase cell cycle arrest when exposed to DNA damaging agents (Rosa et al. 2007b), and that p53 arrested the cell cycle by lowering Cyclin B1 levels (Rosa et al. 2007a). In addition, reduction of MYC expression was associated with cell cycle arrest in SH-SY5Y cells (Posser et al. 2011). Our results suggest that (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ influenced the expression of p53, p21 and MYC and that they could be effective as anti-proliferative agents by inducing G2/M cell cycle arrest. Selenite was shown to elevate the levels of phosphorylated p53 protein at Ser-15 and concomitantly increase the expression of p21. In addition, the pro-apoptotic Bax levels were elevated and when a p53-specific inhibitor was used Bax expression was reduced by 50%, suggesting that selenium compounds could mediate tumour cell death by the p53 pathway. However, other mechanisms may also contribute to the expression of Bax. In addition, it was observed that cytochrome c, capspases-9 and -8 did not participate in the execution of apoptosis in selenite-exposed cells (Rudolf et al. 2008). In the present study, the $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$ compounds appeared to mediate apoptosis in a caspase-dependent manner, since the expression of caspase-9 was significantly higher in treated HT-29 cells. However, p53 phosphorylation could also contribute to elevated Bax expression leading to apoptosis.

It is important to clarify that the benefit of selenium compounds is related to its bioavailability in the intestine and its ability to enter the bloodstream where it can be distributed to the various organs and tissues. Of note, the bioavailability of selenium is closely related to its chemical form (Thiry et al. 2012). In this study the most cytotoxic compound, (4-MeOC₆H₄Se)₂, exhibited a significant inhibitory effect (> 40%) on HT-29 cells at a concentration of 20 μ M that increased to >75% at a concentration of 80 μ M following exposure for 24 h. Furthermore, these concentrations are similar to those used in other studies that reported induction of apoptosis in cancer cells with similar doses (10-100 μ M) of selenium compounds (Lunøe et al. 2011; Posser et al. 2011). Further work will need to be carried out to verify the cytotoxic effects of the compounds in animal models and to confirm their bioavailability at these concentrations.

Conclusion

In summary, for the first time the cytotoxic potential of (3-CF₃C₆H₄Se)₂ and (4-MeOC₆H₄Se)₂ was demonstrated in human colon adenocarcinoma cells and the cytotoxic effect was likely mediated through the induction of apoptosis. In addition, several molecular targets of these compounds were investigated and the evidence suggests that apoptosis was stimulated by a caspase-dependant pathway as well as by a caspase-independent pathway and that cell-cycle arrest was mediated by the p53, p21 and MYC genes.

Acknowledgments

This work was supported by CNPq (Grant 472644/2010-6), CAPES and FAPERGS (PRONEX 10/0027-4, PqG 1012043). L.S, D.A., A.J.A.M and O.A.D. are recipients of CNPq fellowships and F.N. has a fellowship from CAPES.

Conflict of interest

The authors declare that there is no conflict of interest.

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Figure legends

$$\begin{array}{c} \text{Se.}_{Se} \\ \text{diphenyl diselenide} \\ \text{(C_6H_5Se)}_2 \\ \text{Se.}_{Se} \\ \text{CI} \\ \text{Se.}_{Se} \\ \text{CI} \\ \text{4-chlorodiphenyl diselenide} \\ \text{($4-MeOC_6H_4Se$)}_2 \\ \text{3-(trifluoromethyl)-diphenyl diselenide} \\ \text{($3-CF_3C_6H_4Se$)}_2 \\ \end{array}$$

Fig. 1. Chemical structure of diaryl diselenides

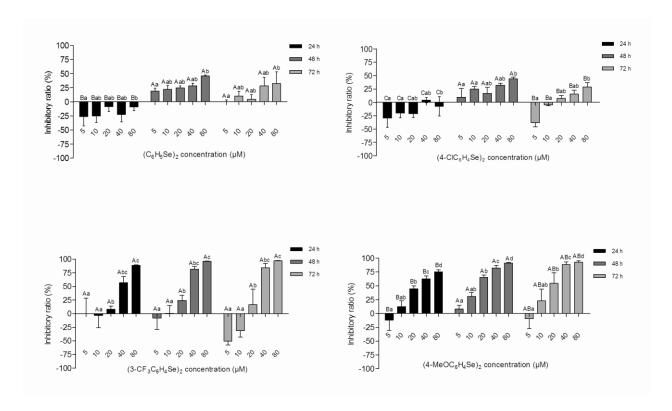


Fig. 2. Effect of the different concentration of substituted diaryl diselenides, $(C_6H_5Se)_2$ (4- $ClC_6H_4Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$ following exposure for 24, 48 and 72 h on the inhibition of HT-29 cells. Data are expressed as the means \pm SEM. Uppercase letters indicate significant differences between treatment times and lowercase letters indicate significant differences in the concentrations used. A p-value < 0.05 was considered significant (Tukey test).

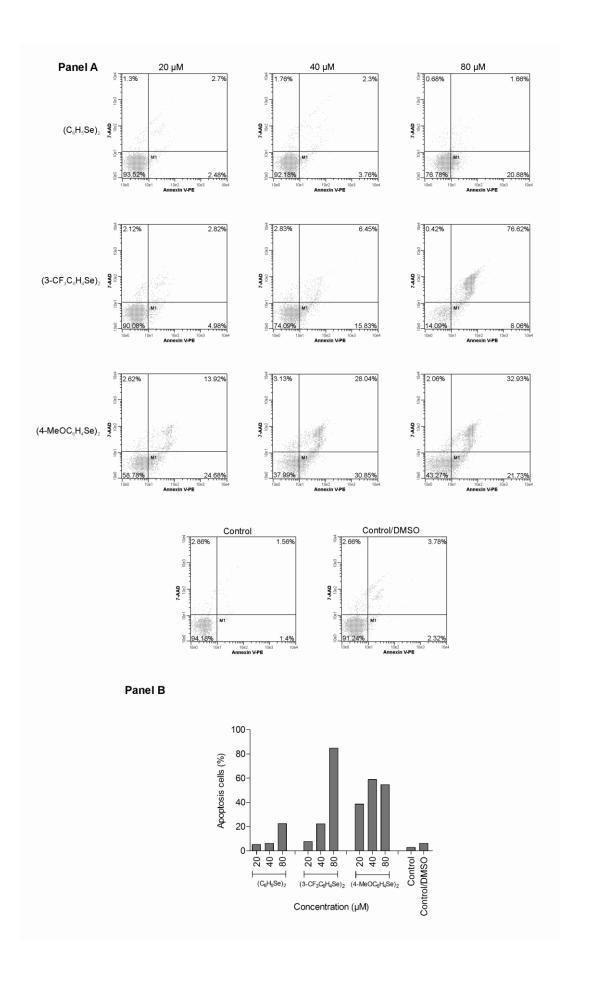


Fig. 3. Annexin V-PE analysis of HT-29 cells treated with 20, 40 and 80 μ M of $(C_6H_5Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$, and control groups after exposure for 48 h. Panel A. Flow cytometry graphs. Panel B. Percentage of apoptotic cells.

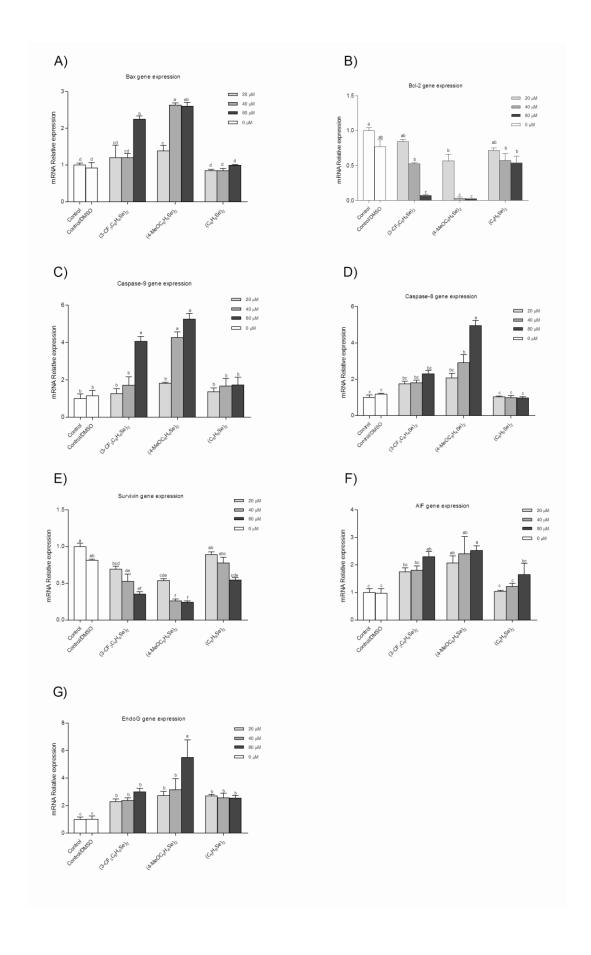


Fig. 4. Effect of $(C_6H_5Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$, in apoptotic-related gene expression. A - Bax, B - Bcl-2, C - Caspase 9, D - Caspase 8, E - Survivin, F - AIF and G - EndoG. The data shown are expressed as the means \pm SEM of a representative experiment performed in triplicate (n = 3). Letters above the bars indicate significant differences. A p-value < 0.05 was considered significant (Tukey test).

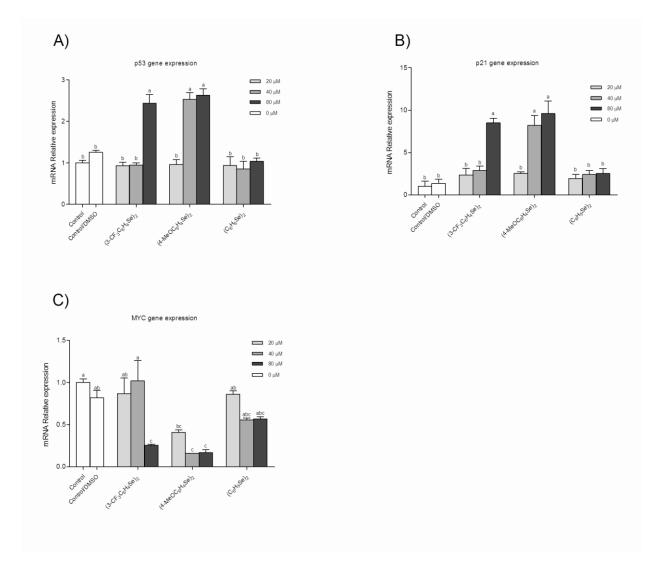


Fig. 5. Effect of $(C_6H_5Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$, in cell-cycle arrest-related gene expression. A – p53, B – p21 and C - Myc. The data shown are expressed as the means \pm SEM of a representative experiment performed in triplicate (n = 3). Letters above the bars indicate significant differences. A p-value < 0.05 was considered significant (Tukey test).

Table 1. Primers sequences used in this study.

Primers	Sequence 5'→ 3'	Reference	
p53 For	AGCGAGCACTGCCCAACA	(Gochhait et al. 2009)	
p53 Rev	CACGCCCACGGATCTGAA		
Bcl-2 For	GTGTGGAGAGCGTCAACC	(Chen et al. 2010)	
Bcl-2 Rev	CTTCAGAGACAGCCAGGAG		
Bax For	ATGCGTCCACCAAGAAGC	(Chen et al. 2010)	
Bax Rev	ACGGCGGCAATCATCCTC		
Casp9 For	CCAGAGATTCGCAAACCAGAGG	(Huang et al. 2007)	
Casp9 Rev	GAGCACCGACATCACCAAATCC		
Survivin For	CTGTGGGCCCCTTAGCAAT	(Wang et al. 2008)	
Survivin Rev	TAAGCCCGGGAATCAAAACA		
p21 For	CCTAATCCGCCCACAGGAA	(War a at al. 2000)	
p21 Rev	ACCTCCGGGAGAGAGGAAAA	(Wang et al. 2008)	
MYC For	TCAGCAACAACCGAAAATGC	(Wang et al. 2008)	
MYC Rev	TTCCGTAGCTGTTCAAGTTTGTG		
GAPDH For	GGATTTGGTCGTATTGGG	(Hu et al. 2010)	
GAPDH Rev	TCGCTCCTGGAAGATGG		
Casp8 For	GGATGGCCACTGTGAATAACTG	(Lin et al. 2011)	
Casp8 Rev	TCGAGGACATCGCTCTCTCA		
AIF For	GGGAGGACTACGGCAAAGGT	(Lu et al. 2010)	
AIF Rev	CTTCCTTGCTATTGGCATTCG		
EndG For	GTACCAGGTCATCGGCAAGAA	(Lin et al. 2008)	
EndG Rev	CGTAGGTGCGGAGCTCAATT		

3. ARTIGO 2

CNTLectins: new insights in cancer therapy.

(Artigo formatado segundo as normas do periódico Carbon)

CNTLectins: new insights in cancer therapy

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Abstract

Considering the independent potential of plant lectins and carbon nanotubes (CNTs) in cancer therapy, we proposed a combination of the two for a cancer therapeutic approach: CNTLectin. Additionally, we investigated the antineoplastic effects of three plant lectins isolated from the seeds of *Canavalia brasiliensis* (ConBr), *Canavalia boliviana* (ConBol) and *Canavalia ensiformis* (ConA) in human colon adenocarcinoma (HT-29) cells, as well as the lectin-cell interaction. The viability of the HT-29 cells towards the three lectins was determined using the MTT assay. To verify the lectin-cell interaction, the lectins were labeled with FITC and associated with f-MWCNTs. The results indicated that all lectins were cytotoxic to HT-29 cells. The FITC-ConBol conjugates demonstrated an intense fluorescence signal associated with the extracellular membrane. Controversially, the FITC-ConBr conjugate did not bind to the HT-29 cells. The interaction of ConBol and ConA with cells improved when the fluorescent lectins were associated with the f-MWCNTs. These results demonstrate that ConA, ConBr and ConBol are potential anti-neoplastic agents for HT-29 cells. Additionally, our study demonstrates, for the first time, that ConBol and ConA enhance the lectin-cell interaction when associated with f-MWCNTs.

1. Introduction

Cancer is a leading cause of death and is a major health problem in both industrialized and developing nations [1, 2]; its rates are expected to increase 50% by 2020 [3, 4]. Although surgical resection and additional therapies are able to cure well-confined, primary tumors, the same does not apply during metastasis due to the systemic involvement and the resistance of metastatic tumors to conventional therapies [5, 6]. Therefore, the current clinical challenge is to develop new drugs and treatment modalities that will significantly impact the cure rates by reversing drug resistance with minimal toxicity [7, 8]

Neoplastic cell transformations are usually associated with alterations in cell surface glycoconjugates, such as increases in sialylation, alterations in the branching of complex carbohydrates and occasionally the emergence of unusual structures. Therefore, researchers have focused on using such variations in future therapies and as diagnostic or prognostic targets [9-11]. Lectins are proteins or glycoproteins of non-immune origin that contain at least one non-catalytic domain, which enables them to selectively recognize and reversibly bind specific free sugars or glycans present on glycoproteins and glycolipids without altering the structure of the carbohydrate [12-15]. Previously, plant lectins have been used as simple tumor recognition tools to differentiate malignant tumors from benign tumors and to evaluate the degree of glycosylation associated with metastasis [9, 15]. Recently, plant lectins have been used in microarray analyses to enhance the recognition of malignant tumors [15, 16]. Additionally, plant lectins possess antitumor activities, inducing apoptosis, predominantly through caspase-dependent pathways [17]. This programmed cell death is a highly regulated mechanism that allows a cell to self-degrade [18]; the failure of this process is responsible for tumor promotion and progression, as well as treatment resistance [7]. Thus, apoptosis signaling systems serve as promising targets for the development of novel anticancer agents [18].

Although extensive and profound research has been conducted to identify new drugs for cancer treatment, some major drawbacks have limited the success of chemotherapy. Some treatments lack sufficient selectivity towards cancer cells and therefore are toxic to healthy tissues. Some drugs have limited bioaccessibility to the tumor tissues, requiring high drug doses, while other drugs are unable to cross cellular barriers. Another factor limiting successful cancer treatment is the frequent emergence of drug resistance [19, 20]. To overcome these issues, not only the development of novel pharmaceuticals but also the system of drug delivery require consideration. From this perspective, a plausible way to efficiently deliver cancer drugs is to associate anticancer drugs with nanoparticles. The emerging use of nanoparticle-based strategies has demonstrated enhanced therapeutic efficacy with reduced side effects compared with the classic, non-targeted therapeutic drug combination modalities currently used [20, 21].

Diverse classes of nano-material carbone nanotubes (CNTs) have attracted particular attention due to their unique properties, such as their spectroscopic, thermal and electrical properties [19, 22]. Several studies have demonstrated that the functionalized CNTs internalize a wide variety of cell types by crossing cell membranes through endocytosis or using other mechanisms [19]. The dimensions of these tubular structures typically range from 0.4 to 2 nm in diameter for single-walled CNTs (SWCNTs), 1 to 3 nm for double-walled CNTs (DWCNTs) and 2 to 100 nm for multi-walled CNTs (MWCNTs) [22]. Aside from their nanodimension, CNTs have a large surface area that enables to load their interior with a high density of active components, or/and use their exterior walls to make functional attachments with essential compounds. CNTs have been used as delivery systems for a variety of components: typical anticancer and antifungal drugs; biomacromolecules such as proteins, peptides, DNA, siRNA and antisense oligomers; and targeting ligands such as vitamins, peptides and antibodies have all been used in complex CNT-based nanodrugs to enable

delivery into a specific tissue or cell subpopulation. In addition, some studies have associated lectins, especially ConA, with CNTs and cancer cells. However, the focus of CNT-cancer research is to develop sensitive, practical and high-throughput monitoring technologies that analyze the glycosylation status of cancer cells and provide the diagnostic tools that guide cancer treatment [23-26].

Thus, considering the independent potential of plant lectins and CNTs, we initially aimed to investigate the antineoplastic effects of three plant lectins in human colon adenocarcinoma (HT-29) cells; the plant lectins were isolated from seeds of *Canavalia brasiliensis* (ConBr), *Canavalia boliviana* (ConBol) and *Canavalia ensiformis* (ConA). In addition, we investigated the interaction between these lectins and HT-29 cells when associated with functionalized multi-walled carbone nanotubes (f-MWCNTs). To the best of our knowledge, this is the first study that demonstrates the antineoplastic potential of ConBol and ConBr and their association with CNTs. Finally, we use the association between lectins and CNTs in a cancer therapeutic approach, which can serve as a new drug and treatment modality for cancer therapy: CNTLectins.

2. Experiments

2.1 Lectins

The lectins used in this study were purified from leguminous plant seeds growing in Ceara state, Brazil. The steps for purification have been described previously [27, 28]. Briefly, the defatted (with n-hexane) seed flour from each species (C. brasiliensis, C. boliviana and C. ensiformis) was extracted with 0.15 M NaCl (1:10, m/v) for 3 h under continuous stirring at room temperature and then centrifuged at 16,000 g for 20 min at 4 °C. The clear supernatant was then applied to a Sephadex G-50 column (40×2.5 cm) that was equilibrated and eluted with 0.15 M NaCl containing 5 mM CaCl₂ and 5 mM MnCl₂. After elution of the unbound

protein, the lectin was desorbed from the column using a 0.1 M solution of the specific inhibitor sugar, d-glucose, added to the equilibrium solution or using a 0.05 M glycine–HCl buffer pH 2.6 containing 0.15 M NaCl. The fractions containing the lectin were pooled, dialyzed against 1 M acetic acid for 1 h, dialyzed exhaustively against distilled water, lyophilized and used for the preparation of the solution to be assayed. Prior to its use in the bioassays, the purity of each lectin was evaluated using denaturing electrophoresis (SDS–PAGE).

2.2 FITC-Lectins

FITC-labeled lectins were prepared in the inhibition buffer (0.1 mol L^{-1} D-mannose in 0.1 mol L^{-1} carbonate-bicarbonate buffer, pH 9.0), the conjugation buffer (0.1 mol L^{-1} carbonate-bicarbonate buffer, pH 9.0) and the washing buffer (phosphate-buffered saline (PBS): 0.01 mol L^{-1} sodium phosphate buffer, 0.027 mol L^{-1} KCl and 0.15 mol L^{-1} NaCl, pH 7.4). Initially, the lectins were dissolved in the inhibition buffer and incubated at 37 °C for 1 h. Then, 250 μ L of fluorescein isothiocyanate (FITC) (500 μ g/mL in conjugation buffer) was added dropwise. The solution was incubated for 2 h at room temperature with gentle stirring. Subsequently, unconjugated FITC was separated from FITC-lectin by size exclusion chromatography using a Sephadex G-25 column previously equilibrated and eluted with washing buffer. The absorbance for all of the fractions was determined at 280 nm (protein) and 495 nm (FITC) to verify the chromatographic efficiency. The FITC-labeled lectins were dialyzed against 1 mol L^{-1} acetic acid for 1 h to remove the inhibitor carbohydrate and then dialyzed against distilled water and lyophilized.

2.3 MWCNT-oxidation

For the MWCNT functionalization, we employed the methodology described by [29]. Briefly, the MWCNTs (Sigma-Aldrich, St. Louis, MO, USA) were refluxed in a 9.0 mol/L HNO₃ solution (200 mL) for 24 h at 150 °C. Then, the system was cooled to room temperature, filtered through a 0.2 µm PTFE membrane and washed with deionized water until a neutral pH was obtained. The HNO₃-treated MWCNTs were dried using a vacuum for 24 h and washed as described in the related protocol.

2.4 Cell Culture

The human colon adenocarcinoma cell line (HT-29) was obtained from the Rio de Janeiro Cell Bank (PABCAM, Federal University of Rio de Janeiro, RJ, Brazil). They were cultured in Dulbecco's Modified Eagle's Medium (DMEM), supplemented with 10% fetal bovine serum (FBS), which were purchased from Vitrocell Embriolife (Campinas, Brazil) and Gibco (Grand Island, NY, USA), respectively. The cells were grown at 37 °C in an atmosphere of 95% humidified air and 5% CO₂. The experiments were performed with cells that were in the logarithmic growth phase.

2.5 Determining the cytotoxicity of Lectins

The viability of HT-29 cells was determined by measuring the reduction of soluble MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] to water-insoluble formazan [30-32]. Briefly, cells were seeded at a density of 2 x 10^4 cells per well in a volume of 100 μ L in 96-well plates and grown at 37 °C in a humidified atmosphere of 5% CO₂/95% air for 24 h before being used in the cell viability assay. The HT-29 cells were then incubated with different concentrations of ConA, ConBol and ConBr (5 – 100 μ g.mL⁻¹) for 24, 48 and 72 h. The lectins were dissolved in DMEM/10% FBS to the desired concentrations. The f-MWCNTs were dissolved in PBS.

After the incubation periods, the medium was removed, and 180 μ L of medium and 20 μ L of MTT (5 mg MTT^{mL-1} solution) were added to each well. The plates were incubated for an additional 3 h, and the medium was discarded. DMSO (200 μ L) was added to each well, and the formazan was solubilized on a shaker for 5 min at 100 g. The absorbance of each well was read on a microplate reader (MR-96A, Mindray Shenzhen, China) at a test wavelength of 492 nm. The cell inhibitory growth rate was determined as follows: inhibitory rate = (1- Abs₄₉₂ treated cells/Abs₄₉₂ control cells) x 100% [33]. All observations were validated by at least two independent experiments and in triplicate for each experiment.

2.6 Lectin-cell association

The HT-29 cells were cultured in 96-well culture plates at a density of 2 x 10⁴ cells per well and grown at 37 °C in a humidified atmosphere of 5% CO₂/95% air for 24 h. Ten microliters (mg.mL⁻¹) of FITC-ConA, FITC-ConBr and FITC-ConBol was dissolved in DMEM/10% FBS, added to the cells containing DMEM/10% FBS (200 μL) and incubated for 1, 3 and 6 h. At the end of the incubation period, the cells were washed twice with PBS and then viewed and photographed using an inverted fluorescence Olympus IX71 microscope (Olympus Optical Co., Ltd. Tokyo, Japan).

2.7 CNTLectin

The HT-29 cells were cultured in 96 well culture plates at a density of 2 x 10⁴ cells per well and grown at 37 °C in a humidified atmosphere of 5% CO₂/95% air for 24 h. Functionalized-MWCNTs were added to PBS (50 μg.mL⁻¹) and subsequently added to the existent 100 μL of DMEM/10% SFB in each well to obtain a final concentration of 5 μg.mL⁻¹. The FITC-ConA, FITC-ConBr or FITC-ConBol was dissolved in DMEM/10% FBS (1 mg.mL⁻¹), added to f-MWCNT/DMEM/10% FBS in a final concentration of 100 μg.mL⁻¹ and incubated for 1, 3

and 6 h. At the end of the incubation period, the cells were washed twice with PBS and then were viewed and photographed with an inverted fluorescence Olympus IX71 microscope (Olympus Optical Co., Ltd. Tokyo, Japan).

2.8 Data analysis

The data sets from the MTT analysis were analyzed using a two-way ANOVA followed by a Tukey test for multiple comparisons. Two factors were considered: amount of used compound (four levels) and compound concentration (three levels). In all analyses, p <0.05 was considered statistically significant. The data are expressed as the mean \pm SEM.

3. Results

3.1 Determination of cytotoxicity

All of the lectins tested demonstrated a significant *in vitro* cytotoxic activity against the HT-29 cells; this cytotoxic effect increased in a time-dependent manner. The most effective period was 72 h for ConA and 48 h for ConBr and ConBol (p<0.05) (Fig. 1).

Table 1 shows a summary of the cytotoxic effects that each lectin had on the HT-29 cells, including the concentration, time and inhibition rate. Lower concentrations (5 - 25 μ g mL⁻¹) of all lectins were less effective inhibitors of HT-29 cell proliferation, as the cell inhibitory rates were lower than 50% (Fig. 1). The f-MWCNTs did not present a significant cytotoxic effect against Chinese hamster ovary cells (CHO-K1) (data not shown).

Table 1. The inhibitory ratio (%) effect of lectins on HT-29 cells after 48 and 72 h of treatment.

Lectin	% Inhibitory ratio (48 h)		% Inhibitory ratio (72 h)	
	50 μg.mL ⁻¹	100 μg.mL ⁻¹	50 μg.mL ⁻¹	100 μg.mL ⁻¹
ConBr	53.1	81	66.8	94.9
ConBol	47.1	75.7	68.2	97.1
ConA	57.3	74	85.6	97.2

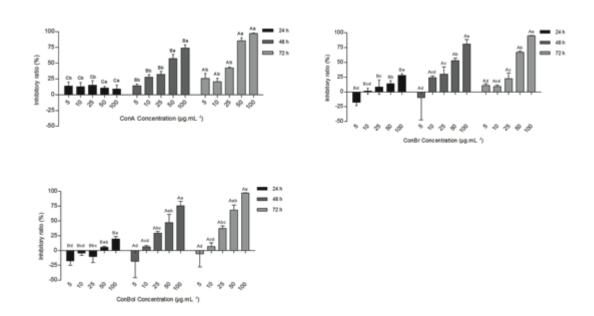


Figure 1. The effect of different concentrations of ConA, ConBol and ConBr on the inhibition of HT-29 cells after 24, 48 and 72 h. The data are expressed as the mean \pm SEM. Uppercase letters indicate differences among treatment times, and lowercase letters indicate differences among concentrations. A p <0.05 was considered statistically significant (Tukey test).

3.2 Lectin-cell interaction

ConA, ConBr and ConBol showed effective cytotoxicity against the HT-29 cells; therefore, the interaction between these cells and lectins was investigated using FITC-lectins. The results demonstrated that ConA and ConBol were able to bind the HT-29 cells within 6 h. The most intense interaction was observed with ConBol (Fig. 4), followed by a slight fluorescence with ConA after 3-6 h (Fig. 2). After 1 h and 3 h of exposure, ConBol exhibited fluorescence, mostly in the extracellular membrane, conferring a honeycombed shape to the cell cluster; however, a slight fluorescence was observed in the intracellular domains as well (Fig. 4 – A5 and B5). Between 3 and 6 h, ConBol was seen in the extracellular domain (Fig. 4 – B1-2 and C1-2). On the contrary, ConBr did not bind to the HT-29 cells during the 6 h period (Fig. 3)

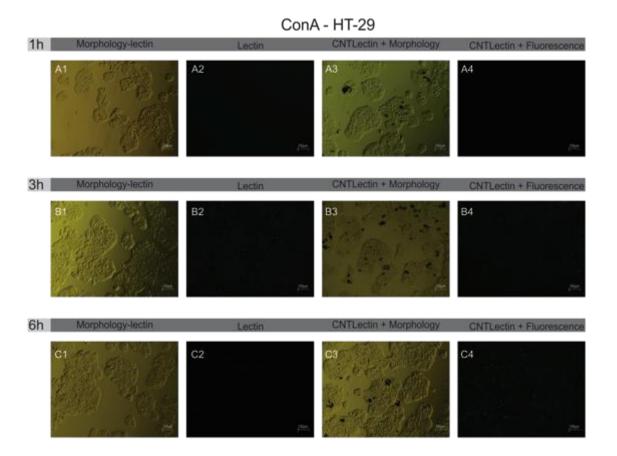


Figure 2. Fluorescent microscopy of the ConA/cell-lectin interaction. Fig. 2 represents the interaction between the HT-29 cells, the fluorescent ConA lectin, and the f-MWCNT-ConA after 1 (A1 and A2/ A3 and A4), 3 (B1 and B2/ B3 and B4) and 6 h (C1 and C2/ C3 and C4).

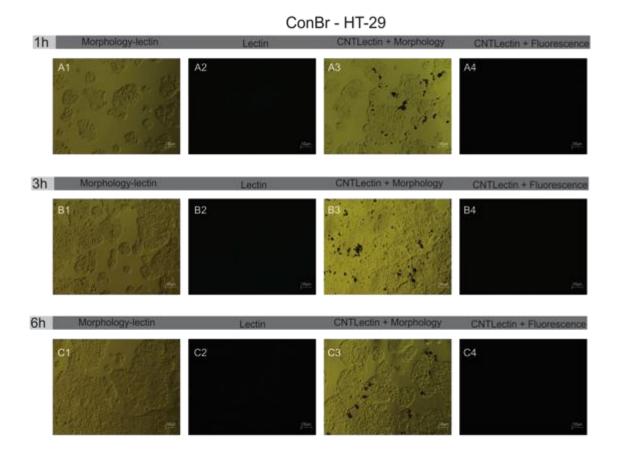


Figure 3. Fluorescent microscopy of the ConBr/cell-lectin interaction. Fig. 3 represents the interaction between the HT-29 cells, the fluorescent ConBr lectin and the f-MWCNT-ConA after 1 (A1 and A2/ A3 and A4), 3 (B1 and B2/ B3 and B4) and 6 h (C1 and C2/ C3 and C4).

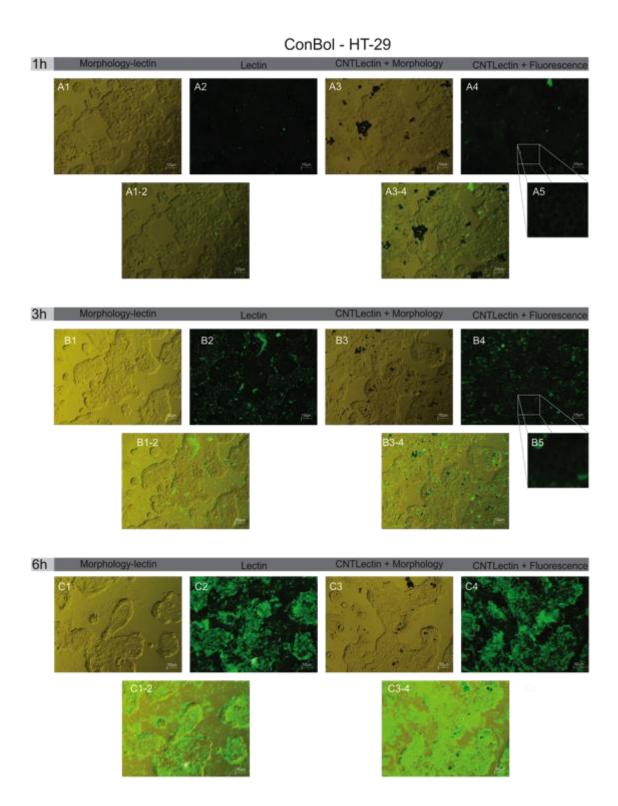


Figure 4. Fluorescent microscopy of the ConBol/cell-lectin interaction. Fig. 4 represents the interaction between the HT-29 cells, the fluorescent ConBol lectin and the f-MWCNT-ConA after 1 (A1 and A2/ A3 and A4), 3 (B1 and B2/ B3 and B4) and 6 h (C1 and C2/ C3 and C4). Fig 4.A5 and B5 show a close-up of the ConBol interaction with the HT-29 cells,

demonstrating the honeycomb shape of cell clusters, as well as the higher level of fluorescence in the extracellular membrane.

3.3 CNTLectin

Since the f-MWCNTs were not cytotoxic, and ConA, ConBr and ConBol inhibited the rate of HT-29 cell growth, we combined these components aiming to increase the lectin-tumor cell interaction. In the f-MWCNT-ConA group, fluorescence was increased in the HT-29 cells after 6 h of exposure (Fig. 2), and these f-MWCNT-ConA showed an affinity for the extracellular membrane (Fig. 2). The f-MWCNT-ConBol group exhibited increased fluorescence in the first hour of cell contact (Fig. 4). After 3 h, a higher amount of fluorescence of ConBol was observed in the extracellular domain; however, when the f-MWCNTs were associated with ConBol, the fluorescence was greater in the cell clusters (Fig. 4 – B1-2/B3). Meanwhile, ConBr showed no fluorescence when associated with the carbon nanotubes (Fig. 3).

4. Discussion

Plant lectins are a group of proteins/glycoproteins with important biological activities and have been used as probes and diagnostic and activator tools. Notably, their anticancer properties have been demonstrated *in vitro*, *in vivo* and in case studies, suggesting their role as therapeutic agents [34, 35]. Concanavalin A (Con A), the first reported legume lectin, was isolated from the jack bean (*Canavalia ensiformis*) and is a Ca²⁺ /Mn²⁺ - dependent, mannose/glucose-binding lectin [36]. ConA has been gaining attention due to its antitumor and antiproliferative activity; ability to trigger apoptosis, autophagy and anti-angiogenesis; and immunomodulatory effects [36]. Studies have demonstrated that ConA can trigger the apoptotic events mediated by the mitochondrial pathway in diverse cell types including

human melanoma (A375) [17] and human hepatocellular liver carcinoma (HepG2) [37]. ConBr, which is a lectin isolated from the seeds of *Canavalia brasiliensis* [38], has been shown to stimulate cultivated human lymphocytes and interferon-γ production [39], cause mast cells to release histamine *in vitro*[40] and increase the production of cytokines (IL-2, IL-6 and IFN-γ) and decrease the production of IL-10 in cultivated splenocytes [41]. ConBr stimulates the production of high levels of IFN-γ and TNFα in human peripheral blood mononuclear cells [38], and it stimulates the production of nitric oxide by murine macrophages *in vitro* and *in vivo* [42]. Finally, it releases and induces apoptosis in lymphocytes *in vivo* [43]. Meanwhile, ConBol is a lectin purified from *Canavalia boliviana* seeds, and its biological effects have not been well characterized, in contrast to ConBr and ConA [28]. ConBol exhibits antinociceptive effects of both central and peripheral origins, involving the opioid system [44], and demonstrates an inhibitory activity on the growth of *S. mutans* and biofilms [45].

Although several studies have focused on the anticancer potential of ConA, the same aspects have not been evaluated for ConBr and ConBol. In the current study, ConBr and ConBol had similar cytotoxic effects on HT-29 cells. ConA has been suggested as a potential antineoplastic agent in pre-clinical and clinical trials for cancer therapy; therefore, the similar cytotoxic results obtained with ConBr and ConBol could indicate a new potential use for both lectins.

To verify the interaction between the HT-29 cells and either ConBr or ConBol, lectins were labeled with FITC. The data showed that both ConBol and ConA bind to the HT-29 cells in a similar way. However, with ConBol, a higher fluorescence can be observed in the extracellular membrane that confers a honeycombed shape to the cell cluster. On the other hand, ConA has a high level of fluorescence that is concentrated in the intracellular space. In agreement with our observation, it has been shown that ConA binds to the cisternal space of

the nuclear envelop, the rough endoplasmic reticulum (RER) and the cisternae along the proximal face of the Golgi stack [46, 47], and it has been used to identify the ER in biological assays [48]. Our findings suggest that ConBol binds the nuclear envelope, the RER and the Golgi complex, which is similar to ConA. However, ConBol also binds to the extracellular membrane with high affinity, indicating that different receptors may be involved in the interactions between both lectins and HT-29 cells. Despite the high level of similarity shared between the Diocleinae lectins, ConBr did not bind to the HT-29 cells. These results suggest that ConBr induces its cytotoxic effects through another mechanism, as in hemagglutination, and not by directly binding HT-29 cells. Liu et al. [17] demonstrated a link between ConA's hemagglutinating activity, mannose-binding activity and antiproliferative activity. Interestingly, ConBr shares a 99% amino acid sequence identity with ConA [49] and ConBol [28]. Despite this similarity, they possess different biological activities [38].

Once we verified the cytotoxic potential of ConBol and ConBr in HT-29 cells and the lectincell interaction, we investigated whether the incorporation of a functionalized multi-walled
carbone nanotube (f-MWCNT) could increase the binding of lectin to HT-29 cells. Thus, it
was demonstrated that the incorporation of f-MWCNTs increased the interaction between HT29 cells and either ConA (6 h) or ConBol (1 – 6 h). Proteins can adsorb spontaneously onto
the sidewalls of CNTs, forming protein-CNT conjugates [50, 51]. The formation of these
conjugate seems to be very specific and depends on the protein structure [52]. SalvadorMorales et al. [51] showed that out of 35 proteins in the human complement activation
system, only fibrinogen and apolipoprotein bind in great quantity to CNTs. Recently, Ge et al.
[52] investigated the interaction of SWCNTs with human serum proteins and observed that
Tyr and Phe amino acids were present in the adsorption region and directly contacted the
surface of SWCNTs, thereby playing a critical role in determining the adsorption capacity of
these proteins. Additionally, as previously described, ConA, ConBr and ConBol display a

high degree of similarity in their primary structures and in the amino acid residues that determine their carbohydrate binding site (Tyr12, Asn14, Leu99, Tyr100, Asp208 and Arg228), their metal binding site (Glu8, Asp10, Tyr12, Asn14, Asp19, His24, Val32, Ser34, Asp208 and Arg228) and their hydrophobic cavity (Tyr54, Leu81, Leu85, Val89, Val91, Phe111, Ser113, Val179, Ile181, Phe191, Phe212 and Ile214), which are conserved in their primary structure [38, 53]. Because the hydrophobic cavity of these lectins contain Tyr54, Phe191 and Phe212, which are the preferential sites for SWCNT-binding, it is possible that ConA, ConBr and ConBol interact with the f-MWCNTs through Tyr and Phe residues, leaving the carbohydrate-binding sites of ConBol and ConA free to bind HT-29 cells. In fact, it was recently suggested that ConA can adsorb to SWCNTs in a nonspecific manner [23]. In addition, when ConBol was associated with the f-MWCNTs (3 h), the fluorescence in the extracellular domain was significantly decreased and the fluorescence in the cell clusters increased (Fig. 4 - B1-2/B3). This result indicates that f-MWCNTs provide HT-29 cells with more lectins to bind and internalize, resulting in a faster incorporation of lectin in the tumor cell. Likewise, single MWCNTs can enter the cell through direct penetration, while MWCNT bundles undergo endocytosis [54]. In addition, MWCNT-conjugated proteins form endosomes after endocytosis and undergo retrograde-transport to the ER, where they are translocated to the cytosol [55]. Therefore, the endocytosis promoted by MWCNTs may induce faster incorporation of lectin in the HT-29 cells. In fact, Weng et al. [56] demonstrated that when recombinant ricin A chain protein (RTA), a lectin from the castor bean plant Ricinus communis, was incubated with L-929 cells, the endocytosis was a slow process, and even after 15 h, there was no significant cell death. However, when RTAs were conjugated with the MWCNTs for a period of 20 h, cell death was induced in approximately 40% of the cells. This study also demonstrated that in the HeLa cell lines, some MWCNT-RTAs are translocated to the cytoplasm, while others are localized near the endoplasmic reticulum, ribosomes and the Golgi apparatus, where the localization of these conjugates was correlated to an increase in cell death. Finally, when tumor cell lines (MCF-7 e HeLa) were treated with MWCNT-RTAs, cell death was significantly increased.

ConBr showed no fluorescence when associated with the f-MWCNTs. This result demonstrates that, although CNTs can enhance lectin binding to HT-29 cells when present, their activity is linked to the existing lectin-binding properties.

5. Conclusion

In summary, we demonstrated that both ConBr and ConBol have the potential to act as antineoplastic agents in HT-29 cells. However, ConBol and ConBr seem to differ in their respective mechanisms for cytotoxic induction. ConBol exhibited a large number of cell-lectin interactions, indicating a possible mechanism for cell death via a cell receptor. However, ConBr did not bind to the HT-29 cells. Additionally, our study demonstrated that the association of ConBol or ConA with f-MWCNTs increases the cell-lectin interactions, thereby increasing the amount of lectin available for tumor cells and potentially contributing to their cytotoxicity. This interaction enables a new possible cancer treatment modality: CNTLectin.

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Figure captions

Figure 1. The effect of different concentrations of ConA, ConBol and ConBr on the inhibition of HT-29 cells after 24, 48 and 72 h. The data are expressed as the mean \pm SEM. Uppercase letters indicate differences among treatment times, and lowercase letters indicate differences among concentrations. A p <0.05 was considered statistically significant (Tukey test).

Figure 2. Fluorescent microscopy of the ConA/cell-lectin interaction. Fig. 2 represents the interaction between the HT-29 cells, the fluorescent ConA lectin, and the f-MWCNT-ConA after 1 (A1 and A2/A3 and A4), 3 (B1 and B2/B3 and B4) and 6 h (C1 and C2/C3 and C4). **Figure 3.** Fluorescent microscopy of the ConBr/cell-lectin interaction. Fig. 3 represents the interaction between the HT-29 cells, the fluorescent ConBr lectin and the f-MWCNT-ConA after 1 (A1 and A2/A3 and A4), 3 (B1 and B2/B3 and B4) and 6 h (C1 and C2/C3 and C4). **Figure 4.** Fluorescent microscopy of the ConBol/cell-lectin interaction. Fig. 4 represents the interaction between the HT-29 cells, the fluorescent ConBol lectin and the f-MWCNT-ConA after 1 (A1 and A2/A3 and A4), 3 (B1 and B2/B3 and B4) and 6 h (C1 and C2/C3 and C4).

Fig 4.A5 and B5 show a close-up of the ConBol interaction with the HT-29 cells, demonstrating the honeycomb shape of cell clusters, as well as the higher level of

Table caption

fluorescence in the extracellular membrane.

Table 1. The inhibitory ratio (%) effect of lectins on the HT-29 cells after 48 and 72 h of treatment

4. Conclusões

- **A)** Os derivados substituídos de disseleneto de diarila (3-CF₃C₆H₄Se)₂ e (4-MeOC₆H₄Se)₂, mostraram ser capazes de induzir a citotoxidade em células HT-29.
- **B)** Os compostos (3-CF₃C₆H₄Se)₂ e (4-MeOC₆H₄Se)₂ mostraram promover esta citotoxidade por meio da apoptose, por ativação de vias independentes e dependentes de caspapses.
- **C)** As lectinas ConA, ConBr e ConBol demonstraram o potencial de induzir um efeito antineoplásico sobre as células HT-29.
- **D)** As lecitnas ConA e ConBol mostraram-se capazes de se ligar as células HT-29, onde a ConBol mostrou uma maior intensidade de florescência na membrana plasmática da célula quando comparada com a ConA.
- **E)** Quando as três lectinas foram associadas aos f-MWCNTs, a ConA e ConBol demonstraram um aumento na intensidade da fluorescência.

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6. Anexos

Artigo 1: Publicado no periódico Life Sciences

LFS-13391; No of Pages 8

Life Sciences xxx (2012) xxx-xxx



Contents lists available at SciVerse ScienceDirect

Life Sciences

journal homepage: www.elsevier.com/locate/lifescie



Substituted diaryl diselenides: Cytotoxic and apoptotic effect in human colon adenocarcinoma cells

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ARTICLE INFO

Article history

Received 11 January 2012

12 Accepted 18 July 2012 13 Available online xxxx 16

17 Keywords

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41 40

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48 49

Human colon adenocarcinoma

19 Apoptosis

20 Substituted diaryl diselenides

21 Selenium

ABSTRACT

Aims: To investigate the effects and study the underlying cell death mechanisms of diaryl diselenides, including: 23 diphenyl diselenide ($C_6H_5Se_1$)₂; 4-chlorodiphenyl diselenide (4-Cl $_6H_4Se_2$); 3-(trifluoromethyl)-diphenyl diselenide 24 (3-CF $_5C_6H_4Se_2$)₂ and 4-methoxydiphenyl diselenide (4-MeO $_6H_4Se_2$), on the human colon adenocarcinoma cell line 25 HT-29.

Main methods: The viability of HT-29 cells after exposure to the diaryl diselenides and its substituted structures was 27 based on the MTT assay. To verify if cell death was mediated throughout apoptosis mechanisms, flow cytometry 28 and real-time PCR (qPCR) analyses were conducted. 29

Key findings: The MTT assay and flow cytometry analyses showed that $(3-\text{CF}_3\text{C}_6\text{H}_4\text{Se})_2$ and $(4-\text{MeOC}_6\text{H}_4\text{Se})_2$ in- 30 duced cytotoxicity through apoptosis mechanisms in HT-29 cells. qPCR revealed there was an up-regulation of 31 pro-apoptotic (Bax, casapase-9, caspase-8, apoptosis-inducing factor (AIF) and Endonuclease G (EndoG)) and 32 cell-cycle arrest genes (p53 and p21) and down-regulation of anti-apoptotic (Bcl-2 and survivin) and Myc genes. 33 Significance: These results demonstrate that $(3-\text{CF}_3\text{C}_6\text{H}_4\text{Se})$ and $(4-\text{MeOC}_6\text{H}_4\text{Se})_2$ have the potential to induce ap- 34 optosis in HT-29 cells through the activation of caspase-dependent and independent pathways and through 35 cell-cycle arrest.

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Introduction

Colorectal cancer is one of the leading causes of cancer mortality (Limami et al., 2011), corresponding to 9.4% of all cases of cancer worldwide (Cantero-Muñoz et al., 2011). Fifty percent of all recently diagnosed patients ultimately develop metastatic disease. Regardless of the advances in developing new chemotherapy agents, no drug has been able to treat colorectal cancer metastasis with a non-relapsing cure rate. Currently the clinical challenge is to develop new drugs that will have a significant impact on cure rates, by reversing drug resistance, and with minimal toxicity (Miura et al., 2011).

Selenium is an essential trace element (Zeng and Combs, 2008) that has the ability to prevent cancer in several animal models and to enhance chemopreventive efficacy in human lung, colorectal, head and neck and prostate cancer (Suzuki et al., 2010). The chemopreventive role of selenium is well supported by epidemiological, preclinical, and clinical evidence (Clark et al., 1998). Furthermore, emerging evidence has indicated the

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0024-3205/\$ – see front matter © 2012 Published by Elsevier Inc. http://dx.doi.org/10.1016/j.lfs.2012.07.023

potential of selenium compounds in cancer chemotherapy (Suzuki et al., $\, 58 \,$ 2010). $\, 59 \,$

Diphenyl diselenide $(C_6H_5Se)_2$, an organic selenium compound, 60 has raised great interest due to its antioxidant, antidepressant-like, 61 neuroprotective and antinociceptive properties (Nogueira and 62 Rocha, 2011; Savegnago et al., 2007, 2008a, 2008b). Recently, 63 Posser et al. (2011) showed, for the first time, that $(C_6H_5Se)_2$ was 64 cytotoxic to human cancer cells (SH-SY5Y) in vitro, possibly medi-65 ated by the ERK1/2 pathway (Posser et al., 2011). However, to date 60 no study has evaluated the cytotoxic effect of $(C_6H_5Se)_2$ in other 67 human cancer cell types.

In addition, studies have demonstrated that the introduction of a substitute (e.g., chloro, fluor or methoxyl) in the aromatic ring of (C_6H_5Se) 70 can alter its molecular properties (Machado et al., 2009; Savegnago et 71 al., 2009; Wilhelm et al., 2009). The introduction of chloro into the aryl 72 group of diaryl diselenide conferred a weak cytotoxic effect on V79 73 cells (Chinese hamster lung fibroblast cells) compared to (C_6H_5Se) 74 (Machado et al., 2009; Savegnago et al., 2009; Wilhelm et al., 2009). Al-75 though this substitute could alter the biological effects of (C_6H_5Se) 76 their potential as cytotoxic agents for cancer chemotherapy has not yet 77 been explored.

Therefore, our objective was to investigate the effect and the underly- 79 ing cell death mechanisms of $(C_6H_5Se)_2$ and its substituted structures, 80

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4-chlorodiphenyl diselenide (4-ClC₆H₄Se)₂, 3-(trifluoromethyl)-diphenyl diselenide (3-CF₃C₆H₄Se)₂ and 4-methoxydiphenyl diselenide (4-MeOC₆H₄Se)₂ on the human colon adenocarcinoma cell line (HT-29). In addition, we also verified whether the introduction of an electron donating (-methoxyl) or an electron withdrawing group (-chloro and -trifluoromethyl) into the aryl group of diaryl diselenide altered its biological effect. To the best of our knowledge this is the first study that demonstrates the effect of (C_6H_5Se)₂ and its substituted structures on HT-29 cells.

Materials and methods

91 Chemicals

 $(C_6H_5Se)_2$, $(4\text{-ClC}_6H_4Se)_2$, $(3\text{-CF}_3C_6H_4Se)_2$ and $(4\text{-MeOC}_6H_4Se)_2$ (Fig. 1) were prepared according to methods in the literature. Analysis of ^{1}H and ^{13}C NMR spectra showed that the analytical and spectroscopic data was in full agreement with its assigned structure. The chemical purity of these compounds was determined by gas chromatography/mass spectrometry.

98 Cell culture

The HT-29 cells were obtained from the Rio de Janeiro Cell Bank (PABCAM, Federal University of Rio de Janeiro, RJ, Brazil). The cells were cultured in Dulbecco's modified Eagle's medium (DMEM), supplemented with 10% foetal bovine serum (FBS), purchased from Vitrocell Embriolife (Campinas, Brazil) and Gibco (Grand Island, NY, USA), respectively. Cells were grown at 37 °C in an atmosphere of 95% humidified air and 5% CO₂. The experiments were performed with cells in the logarithmic phase of growth.

Determination of cytotoxicity

The viability of the HT-29 cells was determined by measuring the reduction of soluble MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] to water insoluble formazan (Ali et al., 2010; Henn et al., 2011). Briefly, cells were seeded at a density of 2×10^4 cells per well in a volume of $100~\mu L$ in 96-well plates and grown at 37 °C in a humidified atmosphere of 5% CO₂/95% air for 24 h before being used in the MTT assay. Cells were incubated with different concentrations of $(C_6H_5Se)_2$, $(4-ClC_6H_4Se)_2$, $(3-CF_3C_6H_4Se)_2$ or $(4-MeOC_6H_4Se)_2$ (5-80 μ M) for 24, 48 and 72 h. These compounds were dissolved in dimethyl sulfoxide (DMSO) and added to the

DMEM supplemented with 10% FBS to the desired concentrations. The 118 final DMSO concentration in the culture medium never exceeded 119 0.8% and a control group exposed to an equivalent concentration 120 of DMSO was evaluated. After incubation the media were re- 121 moved and 180 μ L of DMEM and 20 μ L MTT (5 mg MTT/mL solution) were added to each well. The plates were incubated for an 123 additional 3 h and the medium was discarded. 200 μ L of DMSO was 124 added to each well, and the formazan was solubilized on a shaker for 125 min at 100 × g. The absorbance of each well was read on a microplate 126 reader (MR-96A, Mindray Shenzhen, China) at a wavelength of 492 nm. 127 The percentage inhibition of cell growth was determined as follows: 128 inhibitory rate = (1 – Abs_4treated_cells/Abs_492control_cells) × 100 (Zheng et 129 al., 2011). All observations were validated by at least three independent 130 experiments and for each experiment the analyses were performed in 131 triplicate.

Apoptotic assay 133

The Guava Nexin assay (Guava Technologies) was conducted fol- 134 lowing the manufacturer's instructions. Briefly, 2.0×10^4 to 1.0×10^5 135 of the treated HT-29 cells ($100 \, \mu$ L) were added to $100 \, \mu$ L of Guava 136 Nexin reagent. Cells were incubated in the dark at room temperature 137 for 20 min and samples ($2000 \,$ cells per well) were acquired on the 138 flow cytometry Guava EasyCyte System. In this assay, an annexin 139 V-negative and 7-AAD-positive result indicated late apoptotic 141 cells, while an annexin V-negative and 7-AAD-negative result indicated late apoptotic tells while an annexin V-negative and 7-AAD-negative result indicated live healthy cells and annexin V-positive and 7-AAD-negative result indicated the presence of early apoptotic cells.

Gene expression evaluation by real-time PCR

The HT-29 cells were seeded in a 6-well flat bottom plate at a density $146 \text{ fo} \times 10^5 \text{ per}$ well and grown at 37 °C in a humidified atmosphere of 5% $147 \text{ CO}_2/95\%$ air for 24 h. Cells were then exposed to 20, 40 and 80 µM of $148 \text{ (}C_6H_5Se)_2$, $(3\text{-}CF_3C_6H_4Se)_2$ or $(4\text{-}MeOC_6H_4Se)_2$ for 48 h. After this period the cells were washed with phosphate-buffered saline (PBS; Gibco) 150 and the RNA was extracted from the cells. Total RNA extraction, cDNA 151 synthesis and real-time PCR (qPCR) were carried out as previously described (Campos et al., 2010). Briefly, RNA samples were isolated 153 using TRIzol Reagent (Invitrogen) and samples were DNase-treated with a DNA-free kit (Ambion, USA) following the manufacturer's protocol. First-strand cDNA synthesis was performed with 2 µg of RNA using 156 High Capacity cDNA Reverse Transcription kit (Applied Biosystems, UK) 157 cm

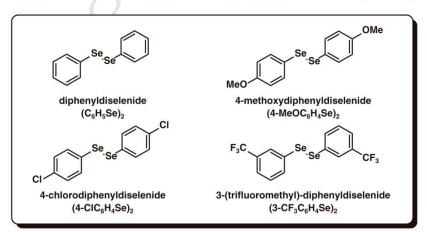


Fig. 1. Chemical structure of diaryl diselenides.

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according to the manufacturer's protocol. The qPCR reactions were run
 on a Stratagene Mx3005P real-time PCR system (Agilent Technologies,
 Santa Clara, CA, USA) using SYBR Green PCR Master Mix (Applied
 Biosystems, UK) using the primers described in Table 1.

162 Data analysis

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Data sets from the MTT assay and qPCR were analysed using a two-way ANOVA followed by a Tukey test for multiple comparisons. Two factors were considered: the compound used (four levels) and the concentration of the compound (three levels). Significance was considered at p<0.05 in all analyses. The data are expressed as the means \pm SEM.

169 Results

170 Determination of cytotoxicity

Both the (C₆H₅Se)₂ and (4-ClC₆H₄Se)₂ compounds had a significant cytotoxic effect on the HT-29 cells at 80 µM and this effect improved significantly with exposure time (Fig. 2). Both the $(3-CF_3C_6H_4Se)$ and (4-MeOC₆H₄Se) compounds achieved significant cytotoxicity at a concentration of 20 μ M. After 48 h exposure to 20 μ M (3-CF₃C₆H₄Se), cytotoxicity was 24% (p<0.05) and this increased significantly to 96% at 80 μM (Fig. 2). The cytotoxicity of the (4-MeOC₆H₄Se) compound at $20\,\mu M\text{,}$ after $24\,h$ exposure, was 44% and further increases in the concentration of the compound resulted in a significant reduction in the viability of the HT-29 cells (62 and 75% cytotoxicity, Fig. 2). The exposure time had no significant effect on the cytotoxicity of the (3-CF₃C₆H₄Se) compound. Only the (4-MeOC₆H₄Se) compound showed a significant improvement with exposure time, for example, at 20 µM and after 24 and 48 h exposure, cytotoxicity increased from 44 to 65%, respectively, although there was no further improvement at 72 h (Fig. 2). The presence of 0.8% DMSO in the culture medium had no effect on cell viability, as compared to the control cells without DMSO.

189 Apoptosis analysis

The annexin-PE staining assay was performed to further characterize the observation that the $(3-CF_3C_6H_4Se)$ and $(4-MeOC_6H_4Se)_2$

Table 1
Primer sequences used in this study.

3	Primers	Sequence 5'→3'	Reference	
.4 p53 for		AGCGAGCACTGCCCAACA	Gochhait et al. (2009)	
,	p53 rev	CACGCCCACGGATCTGAA		
5	Bcl-2 for	GTGTGGAGAGCGTCAACC	Chen et al. (2010)	
	Bcl-2 rev	CTTCAGAGACAGCCAGGAG		
	Bax for	ATGCGTCCACCAAGAAGC	Chen et al. (2010)	
	Bax rev	ACGGCGGCAATCATCCTC		
	Casp9 for	CCAGAGATTCGCAAACCAGAGG	Huang et al. (2007)	
	Casp9 rev	GAGCACCGACATCACCAAATCC		
	Survivin for	CTGTGGGCCCCTTAGCAAT	Wang et al. (2008)	
	Survivin rev	TAAGCCCGGGAATCAAAACA		
	p21 for	CCTAATCCGCCCACAGGAA	Wang et al. (2008)	
	p21 rev	ACCTCCGGGAGAGAGGAAAA		
	MYC for	TCAGCAACAACCGAAAATGC	Wang et al. (2008)	
	MYC rev	TTCCGTAGCTGTTCAAGTTTGTG		
	GAPDH for	GGATTTGGTCGTATTGGG	Hu et al. (2010)	
	GAPDH rev	TCGCTCCTGGAAGATGG		
	Casp8 for	GGATGGCCACTGTGAATAACTG	Lin et al. (2011)	
	Casp8 rev	TCGAGGACATCGCTCTCTCA		
	AIF for	GGGAGGACTACGGCAAAGGT	Lu et al. (2010)	
	AIF rev	CTTCCTTGCTATTGGCATTCG		
	EndG for	GTACCAGGTCATCGGCAAGAA	Lin et al. (2008)	
	EndG rev	CGTAGGTGCGGAGCTCAATT		

compounds could induce apoptosis in HT-29 cells after exposure for 192 48 h. Annexin V binds to those cells that express phosphatidylserine 193 on the outer layer of the cell membrane, a characteristic feature of 194 cells entering apoptosis. The results indicated that (C₆H₅Se) induced 195 apoptosis at a concentration of 80 µM (22.5%, Fig. 3B). The lower concentrations (20 and 40 μM) of (C₆H₅Se) were not effective in causing 197 cell death through apoptosis, inducing similar levels of apoptosis (5.2 198 and 6.1%, respectively) seen in the control groups (3.0 and 6.1%, re- 199 spectively). The (3-CF₃C₆H₄Se) compound induced a higher percent- 200 age of apoptosis at the 40 and 80 µM concentrations (22.3 and 84.7%, 201 respectively) compared to the controls and the (C₆H₅Se) compound. 202 At the 20 μM concentration the percentage of apoptotic cells was 7.8%, 203 similar to that observed in the control groups. The (4-MeOC₆H₄Se) 204 compound was able to induce significant apoptosis in the HT-29 cells 205 at 20 µM (38.6%), this increased to 58.9% upon exposure to a concentra- 206 tion of 40 μ M, although a further increase in concentration to 80 μ M did 207 not increase apoptosis (54.7%). Apoptosis induction from exposure of 208 the HT-29 cells to 0.8% DMSO had no effect.

Gene expression

In order to evaluate the likely apoptosis pathways activated by 211 (3-CF₃C₆H₄Se) and (4-MeOC₆H₄Se) in HT-29 cells (48 h expo- 212 sure), anti-apoptotic and pro-apoptotic gene expressions were in- 213 vestigated. Bax mRNA levels were significantly higher (p<0.05) in 214 cells exposed to (3-CF₃C₆H₄Se) (80 µM) and (4-MeOC₆H₄Se) (20, 215 40 and 80 µM) when compared to the control groups (Fig. 4A), 216 However, (C₆H₅Se) had no effect on Bax mRNA levels when com- 217 pared to the control groups (p>0.05). Bcl-2 mRNA levels decreased 218 significantly (p<0.05) in cells exposed to (3-CF₃C₆H₄Se) (80 μ M) and 219 (4-MeOC₆H₄Se) (40 and 80 μM) when compared to control groups. 220 HT-29 cells exposed to $(3-CF_3C_6H_4Se)$ $(40 \mu M)$, $(4-MeOC_6H_4Se)$ 221 (20 μ M) and (C₆H₅Se) (40 and 80 μ M) decreased Bcl-2 mRNA levels 222 when compared to control groups (p<0.05) (Fig. 4B). Caspase 9 was 223 up-regulated (p<0.05) in cells treated with (3-CF₃C₆H₄Se) (80 μM), 224 (4-MeOC₆H₄Se) (40 and 80 μ M) (Fig. 4C). Exposure to (3-CF₃C₆H₄Se) (20 and 40 µM), (4-MeOC₆H₄Se) (20 µM) and (C₆H₅Se) (20, 40 and 226 80 μ M) had no effect on caspase 9 gene expression (p>0.05). However, 227 caspase 8 mRNA levels were significantly higher (p<0.05) in cells ex- 228 posed to (4-MeOC₆H₄Se) (40 and 80 μ M) when compared to the control 229 groups. (C_6H_5Se), (3- $CF_3C_6H_4Se$) and (4- $MeOC_6H_4Se$) (20 μM) did not 230 affect caspase 8 gene expression (p>0.05) (Fig. 4D). Survivin expression 231 was significantly down-regulated (p<0.05) in HT-29 cells treated with 232 $(3-CF_3C_6H_4Se)$ (40 and 80 μ M), (4-MeOC₆H₄Se) (20, 40 and 80 μ M) 233 and (C₆H₅Se) (80 µM) when compared to the control group (Fig. 4E). 234 The $(3-CF_3C_6H_4Se)$ (20 μ M) and (C_6H_5Se) (20 and 40 μ M) compounds 235 had no effect on survivin expression (p>0.05).

The mRNA levels for AIF and EndoG were also evaluated. AIF expression 237 was significantly up-regulated (p<0.05) upon exposure to (3-CF₃C₆H₄Se) 238 (80 μ M) and (4-MeOC₆H₄Se) (20, 40 and 80 μ M) when compared to the 239 control group (Fig. 4F), However, (C₆H₅Se) and 3-CF₂C₆H₄Se) (20 and 240 40 μM) had no effect on AIF mRNA levels when compared to control groups 241 (p>0.05). EndoG mRNA expression was up-regulated (p<0.05) when the 242 HT-29 cells were treated with $(C_6H_5Se)_2$ (20, 40 and 80 μ M), (3-CF₃C₆H₄Se) 243 $(20, 40 \text{ and } 80 \mu\text{M}) \text{ and } (4\text{-MeOC}_6\text{H}_4\text{Se}) (20, 40 \text{ and } 80 \mu\text{M}) \text{ compared to the } 244$ control group (Fig. 4G). HT-29 cells treated with (3-CF₃C₆H₄Se) (80 µM) and 245 $(4-MeOC_6H_4Se)$ (40 and 80 μ M) had altered levels of cell cycle-related gene 246 expression, p53 expression was significantly up-regulated (p<0.05), in com- 247 parison to the control groups. (C₆H₅Se), at all concentrations tested, had no 248 effect on p53 mRNA levels (Fig. 5A). p21gene expression showed the same ex- 249 pression pattern as p53, where (3-CF $_3$ C $_6$ H $_4$ Se) (80 μ M) and (4-MeOC $_6$ H $_4$ Se) $_{250}$ (40 and 80 μ M) caused significant up-regulation (p<0.05) and (C₆H₅Se) had 251 no effect (Fig. 5B). MYC gene expression was significantly reduced (p<0.05) 252 in cells treated with $(3-CF_3C_6H_4Se)$ (80 μ M) and $(4-MeOC_6H_4Se)$ (40 and 253 80 μ M). (C₆H₅Se) had no effect on MYC gene expression (Fig. 5C). Gene 254

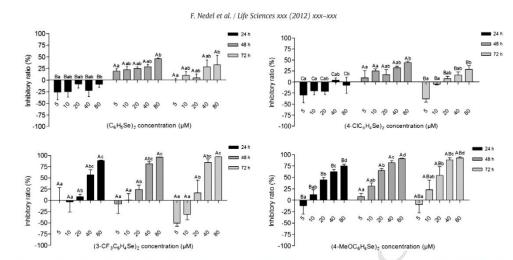


Fig. 2. Effect of the different concentration of substituted diaryl diselenides, $(C_6H_5Se)_2$ (4-Cl $_6H_4Se)_2$, (3-CF $_3C_6H_4Se)_2$ and (4-MeOC $_6H_4Se)_2$ following exposure for 24, 48 and 72 h on the inhibition of HT-29 cells. Data are expressed as the means \pm SEM. Uppercase letters indicate significant differences between treatment times and lowercase letters indicate significant differences in the concentrations used. A p-value <0.05 was considered significant (Tukey test).

expression upon exposure to 0.8% DMSO was similar to the control group in all experiments.

Discussion

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Previous studies have confirmed that organoselenium compounds, such as (C_6H_5Se) and its substituted structures, exhibit a remarkable spectrum of pharmacological properties (Machado et al., 2009; Savegnago et al., 2009; Wilhelm et al., 2009). Indeed, (C_6H_5Se) has exhibited antioxidant, antidepressant-like, neuroprotective and antinociceptive properties and recently it was demonstrated that (C_6H_5Se) had a cytotoxic effect, mediated by the ERK1/2 pathway, on SH-SY5Y cancer cells (Posser et al., 2011). Posser et al. (2011) reported that 30 μ M (C_6H_5Se) significantly decreased cell viability in 50% of cells and, at a concentration of 10 μ M, induced changes in cell morphology (Posser et al., 2011). To the best of our knowledge no study has evaluated the effect of (C_6H_5Se)₂ and the substituted diaryl diselenides (4-CIC₆-H₄Se)₂, (3-CF₃C₆H₄Se) and (4-MeOC₆H₄Se) as cytotoxic and apoptotic agents against cancer cells in vitro or in vivo.

In the present study, (C_6H_5Se) and one of its substituted structures, (4-ClC₆H₄Se)₂, only presented significant cytotoxic effects against the HT-209 cells at a concentration of 80 µM. A similar study that used a neuroblastoma cell line reported cytotoxic effects at lower concentrations (10–30 μ M (C₆H₅Se)). However, this discrepancy may be related to differences between the SH-SY5Y and HT-29 tumor cell lines, as they exhibit different gene profiles when exposed to potent toxic substances (Thirunavukkarasusx et al., 2011). These results suggest that (C₆H₅Se) has a selective action and therefore offers an opportunity to investigate its use as a therapeutic agent. This selectivity has been observed with other selenium compounds, where cancer cells, including lung (A549) and head and neck (HSC-3), were substantially more sensitive to selenite and prone to induction of apoptosis than the breast cancer cell line MCF-7 (Suzuki et al., 2010). The $(3-CF_3C_6H_4Se)$ and $(4-MeOC_6H_4-Re)$ Se) compounds induced cytotoxicity and alterations in cell morphology in HT-29 cells in a dose-dependent manner: 20 µM (24.4 vs. 65.2%), 40 μM (81.8 vs. 81.7%) and 80 μM (91.2 vs. 96.1%), respectively. A recent study evaluated the ability of different selenium compounds (selenate, selenite, MeSeA, MeSeCys and SeMet) to induce cell death in HT-29 cells (Lunøe et al., 2011). The most effective compound was selenite, an inorganic selenium, the percentage of cell death was 21 (10 µM) and 39% (100 µM), followed by two organic selenium compounds, MeSeA (methylseleninic acid) 2 (10 μM) and 14% (100 μM),

and MeSeCys (Se-methylselenocysteine) 3% (100 µM). This suggests 295 that the (3-CF₃C₆H₄Se) and (4-MeOC₆H₄Se) compounds evaluated in 296 the current study are potentially cytotoxic against human colon adeno- 297 carcinoma cells, albeit in vitro. The substitution of a hydrogen atom on 298 the aryl group of diaryl diselenide by an electron withdrawing group 299 (-trifluoromethyl) or an electron donating group (-methoxyl) altered 300 the cytotoxicity when compared to diphenyl diselenide. However, 301 these effects were independent of the nature of the aromatic ring in 302 the diaryl diselenide. Both molecules demonstrated greater cytotoxicity 303 compared to (C₆H₅Se) and (4-ClC₆H₄Se)₂. It has been reported that se- 304 lenium can inhibit cell proliferation, inducing injury via generation of 305 reactive oxygen species (ROS) (Rudolf et al., 2008). ROS levels can acti- 306 vate the INK pathway and caspases-3 and 9 via cytochrome c, with 307 down-regulation of Bcl-2 and up-regulation of Bax (Chen et al., 2011). 308 Also, it has been demonstrated that (C₆H₅Se) and (4-ClC₆H₄Se)₂ pres- 309 ent higher thiol peroxidase activity and an improved antioxidant poten- 310 tial than (3-CF₃C₆H₄Se) and (4-MeOC₆H₄Se) in vivo (Meotti et al., 311 2004). Since, selenium-induced apoptosis in cancer cells can be 312 suppressed by antioxidants (Wu et al., 2010), it is possible that the 313 higher antioxidant potential of (C₆H₅Se) and (4-ClC₆H₄Se)₂ could trig- 314 ger a less effective cytotoxic effect on HT-29 cells than (3-CF₃C₆H₄Se) 315 and (4-MeOC₆H₄Se). 316

Since apoptosis is thought to be the mediator of selenium anticancer 317 activity, we verified, by an Annexin-PE staining assay, that the cytotoxicity 318 effect caused by the (3-CF₃C₆H₄Se) and (4-MeOC₆H₄Se) compounds was 319 mediated by apoptosis. Caspases are central to the mechanism of apoptosis as they are both the initiators and executioners. One pathway by 321 which caspases can be activated involves the extrinsic death receptor 322 pathway, where death ligands bind to death receptors, activating caspase 323 8 and subsequently initiating apoptosis by cleaving other downstream or $\,324$ executioner caspases (Wong, 2011). When (C₆H₅Se)₂ and its substituted 325 structures were tested for their ability to stimulate expression of caspase- 326 8, (4-MeOC₆H₄Se) (40 and 80 μ M) was the only compound that induced 327 high levels of caspase-8 mRNA. Since the upstream caspase for the extrinsic death receptor pathway is caspase-8, this suggests that (4-MeOC₆H₄₋₃₂₉ Se) could be activating a death receptor and therefore contributing to 330 apoptosis in the HT-29 cells. In addition, (4-MeOC₆H₄Se) could present 331 a different biological effect from the other substituted structures due to 332 its electron donating group (-methoxyl).

A second pathway involved in caspase activation is the mitochondrial 334 release of cytochrome c (Wong, 2011). The cytoplasmatic release of cytochrome c activates capase-3 via the formation of a complex (apoptosome) 336

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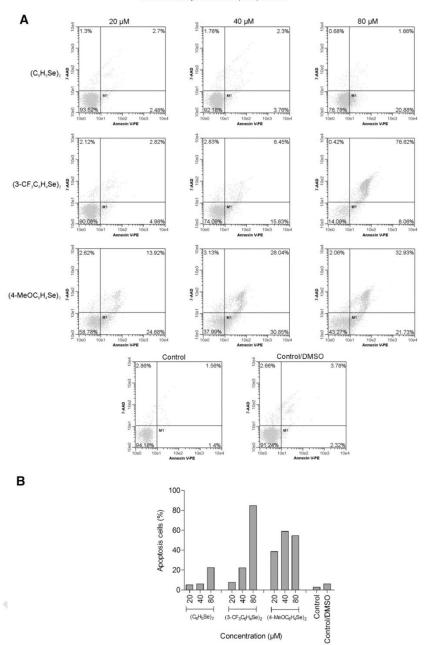


Fig. 3. Annexin V-PE analysis of HT-29 cells treated with 20, 40 and 80 μ M of $(C_6H_5Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$, and control groups after exposure for 48 h. Panel A. Flow cytometry graphs. Panel B. Percentage of apoptotic cells.

which is made of cytochrome c, APAF-1 and caspase-9 (Jackson and Combs, 2008). Bcl-2 (anti-apoptotic) and Bax (pro-apoptotic) are closely involved in this process, an increase in Bcl-2 expression prevents cytochrome c release from the mitochondria, inhibiting the activation of caspase-9 and caspase-3, and preventing apoptosis (Santandreu et al., 2011). In the present study, Bcl-2 expression was down-regulated by

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 $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})$ (80 µM) and (4-MeOC₆H₄Se) (40 and 80 µM), whereas 343 Bax expression was up-regulated. These findings suggest that Bax and 344 Bcl-2 were involved in mediating the apoptotic effects associated with 345 the cytotoxicity of $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})$ and $(4\text{-MeOC}_6\text{H}_4\text{Se})$ in HT-29 cells. 346 In addition, caspase-9 mRNA levels were significantly increased by treatment with $(3\text{-CF}_3\text{C}_6\text{H}_4\text{Se})$ (80 µM) and $(4\text{-MeOC}_6\text{H}_4\text{Se})$ (40 and 80 µM) 348

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showing that caspase-9 was involved in mediating the apoptotic effects
 associated with these compounds. Apoptosis induced by selenium has
 been reported to involve the activation of caspases. It was shown that

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MeSeA induced apoptosis in human prostate cancer (Yamaguchi et al., 352 2005) and leukemia cells (Kim et al., 2001) by the activation of multiple 353 caspases (caspases-3, -7, -8 and -9), mitochondrial release of cytochrome 354

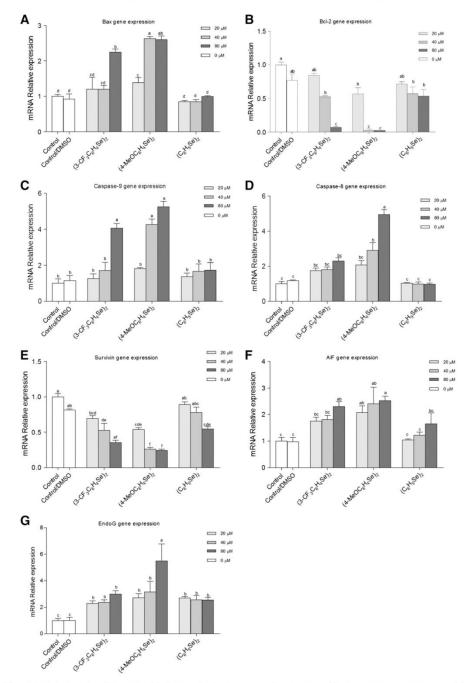
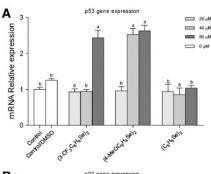
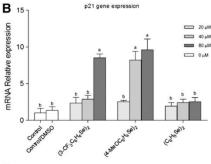


Fig. 4. Effect of $(C_6H_5Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$, in apoptotic-related gene expression. A-Bax, B-Bcl-2, C-caspase 9, D-caspase 8, E-survivin, F-AIF and G-EndoG. The data shown are expressed as the means \pm SEM of a representative experiment performed in triplicate (n=3). Letters above the bars indicate significant differences. A p-value < 0.05 was considered significant (Tukey test).





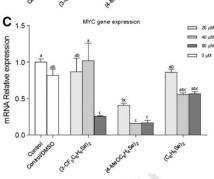


Fig. 5. Effect of $(C_6H_5Se)_2$, $(3-CF_3C_6H_4Se)_2$ and $(4-MeOC_6H_4Se)_2$, in cell-cycle arrest-related gene expression. A-p53, B-p21 and C-Myc. The data shown are expressed as the means \pm SEM of a representative experiment performed in triplicate (n=3). Letters above the bars indicate significant differences. A p-value <0.05 was considered significant (Tukey test).

c and DNA fragmentation. Other organic and inorganic selenium compounds have been shown to induce caspase-mediated apoptosis, including MeSeCys, selenite (Suzuki et al., 2010), sodium selenite (Chen et al., 2011), and selenium dioxide (SeO $_2$) (Rikiishi, 2007).

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Additional apoptotic factors that can be released from the mitochondrial intermembrane space into the cytosol are AIF and EndoG, which translocate to the nucleus, triggering chromatin condensation and DNA degradation in a caspase-independent manner (Vařecha et al., 2012; Wong, 2011). In the current study AIF gene expression was up-regulated by (3-CF₃C₆H₄Se) (80 μ M) and (4-MeOC₆H₄Se) (20, 40 and 80 μ M) and EndoG was up-regulated by exposure to the two substituted diaryl diselenides as well as to (C₆H₅Se)₂. These results suggest time that diaryl diselenide and its substituted structures could induce apoptosis not only through the activation of multiple caspases but also through a caspase-independent pathway.

Survivin has been implicated in the inhibition of apoptosis, cell proliferation, angiogenesis, and cellular stress response. In HT-29 cells,

 $(3-CF_3C_6H_4Se)$ (40 and 80 μ M), (4-MeOC₆H₄Se) (20, 40 and 80 μ M) 372 and (C₆H₅Se) (80 μM) down-regulated the gene expression of survivin. 373 Survivin expression was down-regulated in cell lines derived from pros- 374 tate cancer cells, such as LNCaP, C4-2 (Chun et al., 2007), DU145 and 375 PC-3 (Hu et al., 2008) treated with selenium. However, when the same se- 376 lenium compound was tested with a metastatic cell line derived from 377 PC-3 (PC-3M) and two other prostate cancer cell lines (C4-2B and 378 22Rv1), it had no effect on survivin expression, indicating that the apoptosis induced by selenium was not mediated by decreasing survivin expression (Liu et al., 2010). These results indicated that selenium could trigger 381 different responses depending on the type of cell. Furthermore, p53 and 382 p21 mRNA expression levels were increased while MYC gene expression 383 was down-regulated upon exposure to (3-CF₃C₆H₄Se) (80 µM) and 384 (4-MeOC₆H₄Se) (40 and 80 μ M). The expression of p53, p21 and MYC in- $_{385}$ duced by (C₆H₅Se) did not differ from that of the control groups. Investigators have shown that cells deficient in p21 escaped G2/M phase cell 387 cycle arrest when exposed to DNA damaging agents (Rosa et al., 2007b), 388 and that p53 arrested the cell cycle by lowering cyclin B1 levels (Rosa et 389 al., 2007a). In addition, reduction of MYC expression was associated 390 with cell cycle arrest in SH-SY5Y cells (Posser et al., 2011). Our results sug- 391 gest that (3-CF₃C₆H₄Se) and (4-MeOC₆H₄Se) influenced the expression 392 of p53, p21 and MYC and that they could be effective as anti- 393 proliferative agents by inducing G2/M cell cycle arrest. Selenite was 394 shown to elevate the levels of phosphorylated p53 protein at Ser-15 395 and concomitantly increase the expression of p21. In addition, the 396 pro-apoptotic Bax levels were elevated and when a p53-specific inhibitor 397 was used Bax expression was reduced by 50%, suggesting that selenium 398 compounds could mediate tumor cell death by the p53 pathway. Howev- 399 er, other mechanisms may also contribute to the expression of Bax. In ad-400 dition, it was observed that cytochrome c, capspases-9 and -8 did not 401 participate in the execution of apoptosis in selenite-exposed cells 402 (Rudolf et al., 2008). In the present study, the (3-CF₃C₆H₄Se) and (4-403 MeOC₆H₄Se) compounds appeared to mediate apoptosis in a caspase- 404 dependent manner, since the expression of caspase-9 was significantly 405 higher in treated HT-29 cells. However, p53 phosphorylation could also 406 contribute to elevated Bax expression leading to apoptosis. 407

Of note, the role of apoptosis in the current study was determined 408 using real-time PCR and this is a potential limitation as it is known 409 that mRNA does not necessarily reflect protein concentration, this 410 will be part of future work on these compounds. Furthermore, it is 411 important to clarify that the benefit of selenium compounds is related 412 to its bioavailability in the intestine and its ability to enter the blood- 413 stream where it can be distributed to various organs and tissues. Of 414 note, the bioavailability of selenium is closely related to its chemical 415 form (Thiry et al., 2012). In this study the most cytotoxic compound, 416 (4-MeOC₆H₄Se), exhibited a significant inhibitory effect (> 40%) on 417 HT-29 cells at a concentration of 20 µM that increased to >75% at a 418 concentration of 80 µM following exposure for 24 h. Furthermore, 419 these concentrations are similar to those used in other studies that 420 reported induction of apoptosis in cancer cells with similar doses 421 (10-100 µM) of selenium compounds (Lunøe et al., 2011; Posser et 422 al., 2011). Further work will need to be carried out to verify the cyto- 423 toxic effects of the compounds in animal models and to confirm their 424 bioavailability at these concentrations. 425

Conclusion 426

In summary, for the first time the cytotoxic potential of $(3\text{-CF}_3C_6H_4\text{Se})$ 427 and $(4\text{-MeOC}_6H_4\text{Se})$ was demonstrated in human colon adenocarcinoma 428 cells and the cytotoxic effect was likely mediated through the induction of 429 apoptosis. In addition, several molecular targets of these compounds were 430 investigated and the evidence suggests that apoptosis was stimulated by 431 caspase-dependant pathway as well as by a caspase-independent pathway and that cell-cycle arrest was mediated by the p53, p21 and MYC 433 genes. However, mRNA levels do not necessarily reflect protein concentration and further work will be required to confirm these findings. 435

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		Conflict	of	interest	statement
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The authors declare that there is no conflict of interest

438 Acknowledgments

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This work was supported by CNPq (Grant 472644/2010-6), CAPES and FAPERGS (PRONEX 10/0027-4 and PqG 1012043). L.S, D.A., A.J.A.M and 440 O.A.D. are recipients of CNPq fellowships and F.N. has a fellowship from 442 CAPES

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