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ESTADUAL DE LONDRINA

VIRGINIA MARCIA CONCATO-LOPES

**ATIVIDADE ANTITUMORAL DO 3,3',5,5'-
TETRAMETOXIBIFEENIL-4,4'-DIOL E TRILOBOLIDO-6-O-
ISOBUTIRATO EM MODELO EXPERIMENTAL EM CÉLULAS
A549 E NCI-H460:
EXPLORANDO OS EFEITOS METABÓLICOS E OS
MECANISMOS DE AÇÃO**

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EM CÉLULAS A549 E NCI-H460:
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MECANISMOS DE AÇÃO**

Tese apresentada ao Programa de Pós-Graduação em Patologia Experimental da Universidade Estadual de Londrina, como requisito à obtenção do título de Doutora em Patologia Experimental.

Orientador: Prof. Dr. Wander R. Pavanelli.

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Londrina, 20 de outubro de 2023.

Dedico este trabalho à minha mãe, meu esposo, minha filha e minha família, por todo seu amor incondicional e por sempre acreditar em mim, motivos estes que me fazem seguir em frente.

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“Não vos inquieteis com nada! Em todas as circunstâncias apresentai a Deus as vossas preocupações, mediante a oração, as súplicas e a ação de graças. E a paz de Deus, que excede toda a inteligência, haverá de guardar vossos corações e vossos pensamentos, em Jesus Cristo. ”

Filipenses 4:6,7

“Não importa o quanto a vida possa ser ruim, sempre existe algo que você pode fazer, e triunfar. Enquanto há vida, há esperança. ”

Stephen Hawking

RESUMO

CONCATO-LOPES, Virgínia Márcia. **Atividade antitumoral do 3,3',5,5'-tetrametoxibifeenil-4,4'-diol e trilobolido-6-O-isobutirato em um modelo experimental *in vitro* em células A549 e NCI-H460:** explorando os efeitos metabólicos e os mecanismos de ação. 2023. 135 p. Tese (Doutorado em Patologia Experimental) - Universidade Estadual de Londrina, Londrina, 2023.

O câncer de pulmão, uma doença crônica e heterogênea, é a principal causa de morte relacionada ao câncer mundialmente. Apesar da disponibilidade de diversas opções terapêuticas, sua eficácia continua limitada, frequentemente acompanhada de alta toxicidade, efeitos colaterais significativos, além de resistência medicamentosa que as células cancerígenas desenvolvem. O composto fenólico 3,3',5,5'-tetrametoxibifenil-4,4'-diol (TMBP), obtido por meio da biotransformação do 2,6-dimetoxifenol, mediada por lacase fungica, e a lactona sesquiterpênica extraída de *Sphagneticola trilobata*, trilobolide-6-O-isobutirato (TBB), demonstraram na literatura efeitos leishmanicida, antitumorais e imunomoduladores. No entanto, seus mecanismos de ação em câncer de pulmão de células não pequenas (CPCNP) não foram investigadas. Neste estudo, avaliamos o efeito antitumoral *in vitro* do TMBP e do TBB nas linhagens A549 e NCI-H460, elucidando os mecanismos indutores de morte celular. O tratamento com TMBP (12,5-200 μM) sobre as células NCI-H460 resultou na redução da proliferação em 24, 48 e 72 h. O IC_{50} (154 μM) foi obtido e utilizado para os demais experimentos. Após 24 h de exposição, o tratamento com IC_{50} , induziu diversas alterações morfológicas e ultraestruturais nas células NCI-H460, reduziu a migração e a marcação de N-caderina e β -catenina por imunofluorescência, aumento de espécies reativas de oxigênio (ERO) e óxido nítrico (ON), juntamente com a diminuição dos radicais ânion superóxido e atividade reduzida da glutathione redutase (GSH). Além disso, o TMBP induziu estresse metabólico, evidenciado pela redução da captação de glicose, da lactato desidrogenase intracelular e dos níveis de lactato. Outros efeitos observados incluíram a despolarização mitocondrial, aumento na formação de gotículas lipídicas e a presença de vacúolos autofágicos. Notavelmente, o TMBP induziu a parada do ciclo celular na fase G2/M e a morte celular semelhante a apoptose, conforme indicado pelo aumento da atividade da caspase-3/7 e pela redução da marcação de STAT-3 por imunofluorescência. O impacto antitumoral do TMBP foi acompanhado por uma redução nos níveis proteicos de PI3K, AKT, ARG-1 e NF- κB , juntamente com um aumento de iNOS. Em relação ao tratamento com TBB, as previsões *in silico* revelaram um potencial promissor de similaridade com medicamentos para o composto, indicando alta biodisponibilidade oral e eficiente absorção intestinal. O tratamento *in vitro*, sob células humanas A549 e NCI-H460 (12-100 μM) reduziu a viabilidade celular, obtendo valores de IC_{50} de 73 e 54 μM para A549 e NCI-H460, respectivamente, os quais foram usados nas experimentações seguintes. Após a exposição ao tratamento com IC_{50} de 73 e 54 μM , as células apresentaram alterações morfológicas e estruturais, capacidade migratória reduzida e aumento do estresse oxidativo, caracterizado pelo aumento nos níveis de ERO, radicais ânion superóxido e ON, juntamente com uma redução nos marcadores antioxidantes chave, como superóxido dismutase e GSH. Encontramos também estresse celular induzido pelo TBB após tratamento, o qual resultou em mudanças notáveis no metabolismo celular, levando ao acúmulo de gotículas lipídicas e vacúolos

autofágicos, bem como o comprometimento da integridade mitocondrial, um aspecto crucial da função celular. Adicionalmente, o TBB desencadeou uma morte celular semelhante a apoptose por meio da ativação das caspases 3/7. Essas descobertas destacam o grande potencial dos compostos TMBP e TBB como candidatos promissores para estudos futuros, consolidando sua posição como componentes adicionais no desenvolvimento de protótipos de novos medicamentos anticâncer.

Palavras-chave: Câncer de Pulmão; Compostos fenólicos; Lactona Sesquiterpênica; PI3K/AKT; Estresse oxidativo.

ABSTRACT

CONCATO-LOPES, Virgínia Márcia. **Antitumor activity of 3,3',5,5'-tetramethoxybiphenyl-4,4'-diol and trilobolido-6-O-isobutyrate in an *in vitro* experimental model in A549 and NCI-H460 cells:** exploration of metabolic effects and mechanisms of action. 2023. 135 p. Tese (Doutorado em Patologia Experimental) - Universidade Estadual de Londrina, Londrina, 2023.

Lung cancer, a chronic and heterogeneous disease, is the leading cause of cancer-related deaths worldwide. Despite the availability of several therapeutic options, their efficacy remains limited, often associated with high toxicity, significant side effects, and drug resistance developed by cancer cells. The phenolic compound 3,3',5,5'-tetramethoxybiphenyl-4,4'-diol (TMBP), obtained by the biotransformation of 2,6-dimethoxyphenol mediated by fungal laccase, and the sesquiterpene lactone extracted from *Sphagneticola trilobata*, trilobolide-6-O-isobutyrate (TBB), have been reported in the literature to have leishmanicidal, antitumor and immunomodulatory activities. However, their mechanisms of action in non-small cell lung cancer (NSCLC) have not been investigated. In this study, we evaluated the *in vitro* antitumor effects of TMBP and TBB on A549 and NCI-H460 cell lines and elucidated the mechanisms that induce cell death. Treatment of NCI-H460 cells with TMBP (12.5-200 μ M) resulted in reduced proliferation at 24-72 h. The IC₅₀ (154 μ M) was obtained and used for further experiments. After 24 h of exposure, treatment with IC₅₀ induced several morphological and ultrastructural changes in NCI-H460 cells, decreased migration and labeling of N-cadherin and β -catenin by immunofluorescence, increased reactive oxygen species (ROS) and nitric oxide (NO) along with decreased superoxide anion radicals and reduced glutathione reductase (GSH) activity. In addition, TMBP induced metabolic stress as evidenced by decreased glucose uptake, intracellular lactate dehydrogenase, and lactate levels. Other effects observed included mitochondrial depolarization, increased lipid droplet formation and the presence of autophagic vacuoles. Notably, TMBP induced cell cycle arrest in the G2/M phase and apoptosis-like cell death as indicated by increased caspase-3/7 activity and decreased immunofluorescence labeling of STAT-3. The antitumor effect of TMBP was accompanied by a decrease in protein levels of PI3K, AKT, ARG-1 and NF- κ B and an increase in iNOS. With respect to TBB treatment, *in silico* predictions revealed a promising drug-like potential for the compound, indicating high oral bioavailability and efficient intestinal absorption. *In vitro* treatment of human A549 and NCI-H460 cells (12-100 μ M) reduced cell viability, yielding IC₅₀ values of 73 and 54 μ M for A549 and NCI-H460, respectively, which were used in the following experiments. After exposure to treatment with IC₅₀ of 73 and 54 μ M, cells showed morphological and structural changes, reduced migratory capacity, and increased oxidative stress, characterized by an increase in the levels of ROS, superoxide anion radicals and NO, together with a reduction in markers of key antioxidants such as superoxide dismutase and GSH. We also found cellular stress induced by TBB after treatment, which resulted in significant changes in cellular metabolism, leading to the accumulation of lipid droplets and autophagic vacuoles, as well as the impairment of mitochondrial integrity, a crucial aspect of cellular function. In addition, TBB induced apoptosis-like cell death through the activation of caspases 3/7. These findings highlight the great potential of TMBP and TBB compounds as promising candidates for future studies, thus consolidating

their position as additional components in the development of new anticancer drug prototypes.

Key-words: Lung Cancer; Phenolic Compounds; Sesquiterpene Lactona; PI3K/AKT; Oxidative Stress.

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LISTA DE ABREVIATURAS E SIGLAS

ATP	Adenosina trifosfato
CAT	Catalase
CCG	Carcinoma de Células Grandes
CDDP	cisplatina
CTR1	Proteína transpostadora de cobre 1
DNA	Ácido desoxirribonucleico
eNOS	Óxido Nítrico Sintase endotelial
EMT	Transição epitélio-mesenquimal
FDA	<i>Food and Drug Administration</i>
GSH	Glutaciona reduzida
GSSG	Glutaciona dissulfeto
GSH-Px	Glutaciona Peroxidase
HMG	Proteína do grupo de alta mobilidade
HNO ₂	Ácido nitroso
HOCl	Hipocloreto
H ₂ O ₂	Peróxido de hidrogênio
IARC	<i>International Agency for Research on Cancer</i>
IFN	Interferon
IKK α	I κ B-Quinase -alfa
IL	Interleucina
iNOS	Óxido Nítrico Sintase induzível
LDH	Lactato Desidrogenase
LPS	Lipopolissacarídeo
LS	Lactona Sesquiterpênica
MAPK	Proteína quinase ativada por mitógeno
MMP	Metaloproteinase de matriz
MMP2/9	Metaloproteinase 2/9
MMR	Proteínas de reparo de incompatibilidade
NADPH	Fosfato de dinucleotídeo de nicotinamida e adenina
NBR	Norma Brasileira
NER	Reparo por excisão de nucleotídeos
NF- κ B	Fator Nuclear- <i>kappa</i> B

nNOS	Óxido Nítrico Sintase neuronal
NO•	Óxido Nítrico
NO ₂ -	Dióxido de nitrogênio
NOS	Óxido Nítrico Sintase
NSCLC	<i>Non-Small Cell Lung Cancer</i>
N ₂ O ₃	Trióxido de dinitrogênio
O ₂ ^{•-}	Ânion superóxido
•OH	Hidroxila
OMS	Organização Mundial da Saúde
ONOO ⁻	Peroxinitrito
OXPHOS	Fosforilação oxidativa
PI3K	Fosfoinosítideo 3-quinase
RNS	Espécies reativas de nitrogênio
ROS	Espécies reativas de oxigênio
SCLC	<i>Small Cell Lung Cancer</i>
SOD	Superóxido dismutase
TBB	Trilobolido-6-O-isobutirato
TLR	Receptor toll-like
TMBP	3,3',5,5'-tetrametoxibifenil-4,4'-diol
TNF	Fator de Necrose Tumoral
VEGF	Fator de crescimento do endotélio vascular

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1. INTRODUÇÃO

1.1 ASPECTOS GERAIS DO CÂNCER

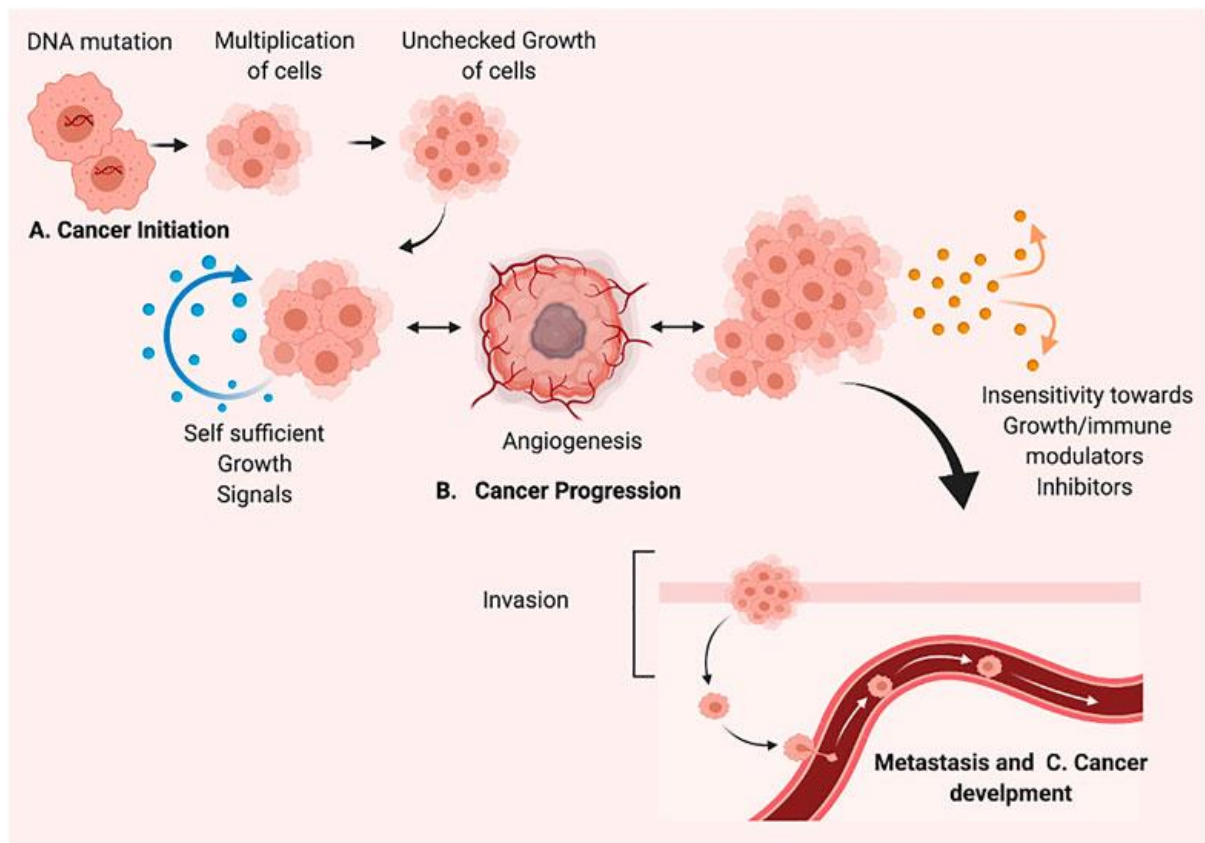
A origem das células cancerígenas envolve um conjunto complexo de eventos genéticos, moleculares e celulares. O câncer inicia quando uma única célula normal sofre uma série de mutações genéticas que a tornam anormal e desencadeiam o crescimento descontrolado (STRATTON; CAMPBELL; FUTREAL, 2009). Essas células aberrantes têm a capacidade de invadir tecidos e órgãos circundantes, podendo se espalhar para outras partes do corpo, em um processo chamado metástase. A causa fundamental do câncer reside em mutações genéticas que ocorrem no DNA. Tais mutações podem ser adquiridas ao longo da vida devido a diversos fatores, como exposição a substâncias carcinogênicas (tabaco, radiações ionizantes e produtos químicos tóxicos), fatores genéticos herdados e erros que podem ocorrer durante a replicação do DNA. Além disso, podem afetar os mecanismos de controle de crescimento celular e inibir a capacidade das células se autorregularem, o que resulta em um crescimento descontrolado. As células cancerígenas também podem ignorar os sinais do sistema imunológico que normalmente reconhecem e eliminam células anormais (HANAHAN; WEINBERG, 2011; DE VISSER; JOYCE, 2023).

A carcinogênese é um processo intrincado que pode ser subdividido em quatro estágios cruciais: iniciação, promoção, conversão maligna e progressão tumoral. No estágio inicial, agentes carcinogênicos ou moléculas iniciadoras entram em contato com as células e provocam alterações irreversíveis no material genético, resultando em mutações genéticas. Estas mutações são causadas pela ação direta dos agentes iniciadores, e elas representam o ponto de partida para o desenvolvimento do câncer. Os agentes promotores entram em ação no estágio de promoção, estimulando a proliferação das células mutadas. Esse estímulo provoca um aumento significativo na taxa de crescimento celular, resultando em um maior número de células com mutações genéticas. Isso cria um ambiente propício para a progressão do câncer, principalmente com o acúmulo de mutações nas células. A conversão maligna marca o terceiro estágio, no qual as células pré-neoplásicas que acumularam várias mutações genéticas distintas e diversificadas sofrem uma transformação em células malignas. Essa transformação envolve uma série complexa de eventos moleculares que conferem às células a capacidade de se comportarem de maneira maligna e invasiva. Finalmente, a progressão tumoral engloba a expressão

1 do fenótipo maligno adquirido, bem como a tendência dessas células malignas em
 2 adquirirem características ainda mais agressivas ao longo do tempo. Esse processo
 3 é acompanhado pelo desenvolvimento de características invasivas, o que permite que
 4 as células tumorais secretem proteases que facilitam a disseminação através da
 5 corrente sanguínea e a invasão de novos tecidos (HAHN; WEINBERG, 2002;
 6 PIRETTO et al., 2019).

7

8 **Figura 1:** Processo de desenvolvimento/carcinogênese do câncer.



9

10 **Fonte:** CHAUDHRY et al. (2022), criado com BioRender.com. (A) iniciação, (B) progressão, (C)
 11 metástase e desenvolvimento do câncer.

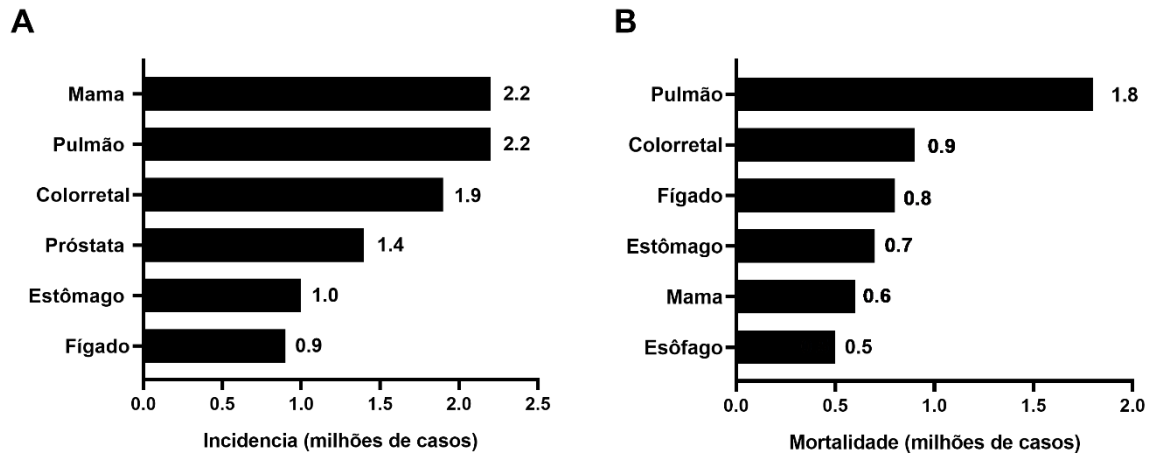
12

13 O câncer é uma das quatro doenças com a maior taxa de
 14 mortalidade, considerado um dos principais problemas de saúde pública (SUNG et al.,
 15 2021). De acordo com dados da IARC (International Agency for Research on Cancer),
 16 em 2020 os cânceres de maior incidência foram de mama (4,7 milhões) e pulmão (2,2
 17 milhões) (**Figura 2A**). Em relação às taxas de mortalidade, o câncer de pulmão levou
 18 à morte de 1,8 milhões de pessoas por ano, e o câncer de mama 1,3 milhões de
 19 mortes, ficando em primeiro e segundo lugar em mortalidade respectivamente (OMS,

1 2020) (Figura 2B).

2

3 **Figura 2** – Número estimado de novos casos e de mortalidade em todo o mundo,
4 considerando ambos os sexos e todas as faixas etárias.



5

6 **Fonte:** Adaptado de OMS (2020).

7

8 1.2 CÂNCER DE PULMÃO

9 A classificação dos tumores pulmonares segue a diretriz da
10 Organização Mundial da Saúde (OMS) de 2015. Este sistema de classificação se
11 baseia na imuno-histoquímica e na análise microscópica para orientar o tratamento e
12 estabelecer um prognóstico. Existem duas categorias principais de câncer de pulmão:
13 carcinoma de células pequenas (*Small Cell Lung Cancer* - SCLC) e carcinoma de
14 células não pequenas (*Non-Small Cell Lung Cancer* - NSCLC). O SCLC é um subtipo
15 altamente agressivo que afeta aproximadamente 13 a 15% da população com câncer
16 de pulmão. Ele tem uma tendência significativa de se espalhar para outros órgãos.
17 Por outro lado, o NSCLC é responsável por cerca de 85% dos casos de câncer de
18 pulmão e tem menor probabilidade de metástase, embora seja menos responsivo a
19 terapias citotóxicas. O NSCLC inclui subtipos como o adenocarcinoma, o carcinoma
20 de células escamosas e o carcinoma de grandes células, cada um com características
21 morfológicas distintas. Em resumo, a classificação dos tumores pulmonares de acordo
22 com a OMS considera o tipo histológico, morfologia macro e microscópica, e ajuda a
23 orientar o tratamento e a prever o prognóstico de pacientes com câncer de pulmão
24 (CLARK; ALSUBAIT, 2022; DUMA; SANTANA-DAVILA; MOLINA, 2019; GAZDAR,
25 2010; TRAVIS, 2011).

26

O adenocarcinoma, representando cerca de 40% dos casos de câncer

1 de pulmão, forma estruturas glandulares e se origina nos bronquíolos, avançando nos
2 alvéolos; por isso, geralmente estão localizados na região periférica do pulmão e
3 penetram em vasos linfáticos, por onde podem chegar à pleura e linfonodos hilares e
4 formar metástases. Sintomas locais muitas vezes são ausentes ou aparecem
5 tardiamente, e este tipo de câncer não está fortemente associado ao tabagismo
6 (ZAMBONI, 2002; BALDOTTO et al., 2016; DUMA; SANTANA-DAVILA; MOLINA,
7 2019; SIDDIQUI; VAQAR; SIDDIQUI, 2023).

8 O carcinoma de células escamosas (ou epidermóide) abrange 25 a
9 30% dos casos de câncer de pulmão. Ele geralmente se desenvolve nos brônquios
10 centrais e tem crescimento endobrônquico. A presença de pérolas de queratina é uma
11 característica histológica marcante, tornando-o facilmente identificável, especialmente
12 em tumores maiores. Os sintomas, como tosse e hemoptise, costumam aparecer
13 precocemente, permitindo um diagnóstico nas fases iniciais, mas possa ocorrer
14 disseminação para linfonodos regionais (TRAVIS, 2011; ZAPPA; MOUSA, 2016;
15 DUMA; SANTANA-DAVILA; MOLINA, 2019; SIDDIQUI; VAQAR; SIDDIQUI, 2023).

16 Por fim, o carcinoma de células grandes (CCG) é o mais raro dos tipos
17 de NSCLC, representando de 5% a 10% dos casos. Este subtipo é caracterizado pela
18 falta de características morfológicas específicas dos outros carcinomas e, muitas
19 vezes, é diagnosticado por exclusão de outras possibilidades. Os CCGs tendem a ter
20 um prognóstico desfavorável devido à sua agressividade e rápida disseminação.
21 Histologicamente, eles são compostos por células grandes com citoplasma moderado
22 e nucléolos proeminentes, podendo ocorrer em qualquer parte do pulmão
23 (ROSKOSKI, 2017; DUMA; SANTANA-DAVILA; MOLINA, 2019; SIDDIQUI; VAQAR;
24 SIDDIQUI, 2023).

25 O principal fator de risco para o desenvolvimento do câncer de pulmão
26 é o tabagismo, que é responsável por, aproximadamente, 6 milhões de mortes anuais
27 no mundo, e aproximadamente 147 mil mortes no Brasil, incluindo o câncer (INCA,
28 2018). Além disso, outros fatores estão associados com a evolução desta doença,
29 dentre eles, os fatores genéticos e ambientais, como exposição à compostos químicos
30 como o asbesto, arsênico, cádmio, hidrocarbonetos, gás mostarda, entre outros, ainda
31 mais quando associados ao hábito de fumar (SUN; SCHILLER; GAZDAR, 2007; DELA
32 CRUZ; TANOUE; MATTHAY, 2011; SIDDIQUI; VAQAR; SIDDIQUI, 2023).

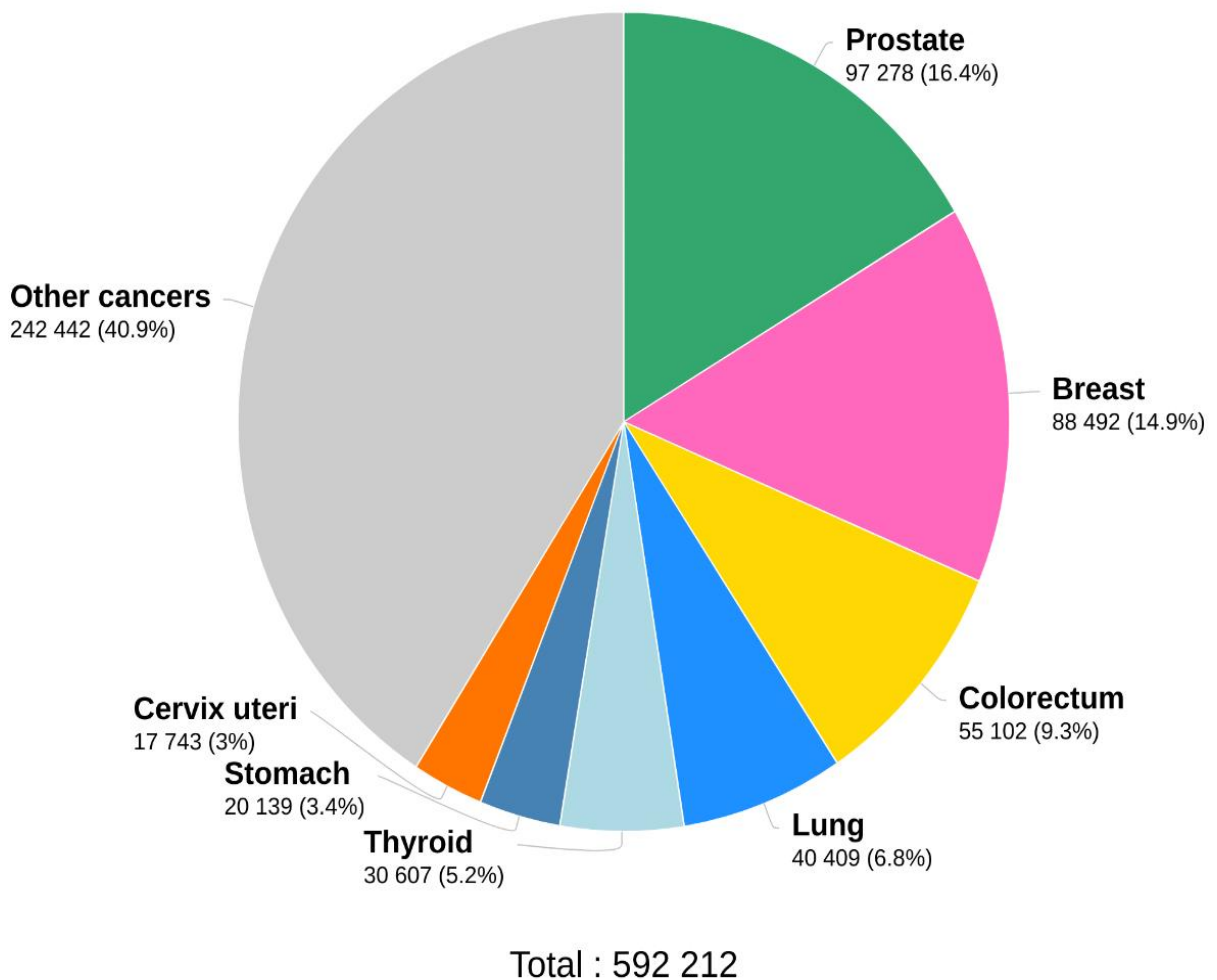
33 O câncer de pulmão é reconhecido como um dos tipos mais
34 agressivos de câncer, com uma presença significativa em todo o mundo (FERLAY et

1 al., 2013). No Brasil, no ano de 2020, os tipos de câncer mais comuns relatados
 2 incluíram próstata (com 97.278 novos casos), mama (com 88.492 novos casos),
 3 colorretal (com 55.102 novos casos), pulmão (com 40.409 novos casos), tireoide (com
 4 30.607 novos casos), estômago (com 20.139 novos casos) e colo do útero (com
 5 17.743 novos casos) (**Figura 3**) (OMS, 2020).

6

7 **Figura 3:** Número estimado de novos casos de câncer em 2020 no Brasil, para ambos
 8 os sexos e todas as idades.

9



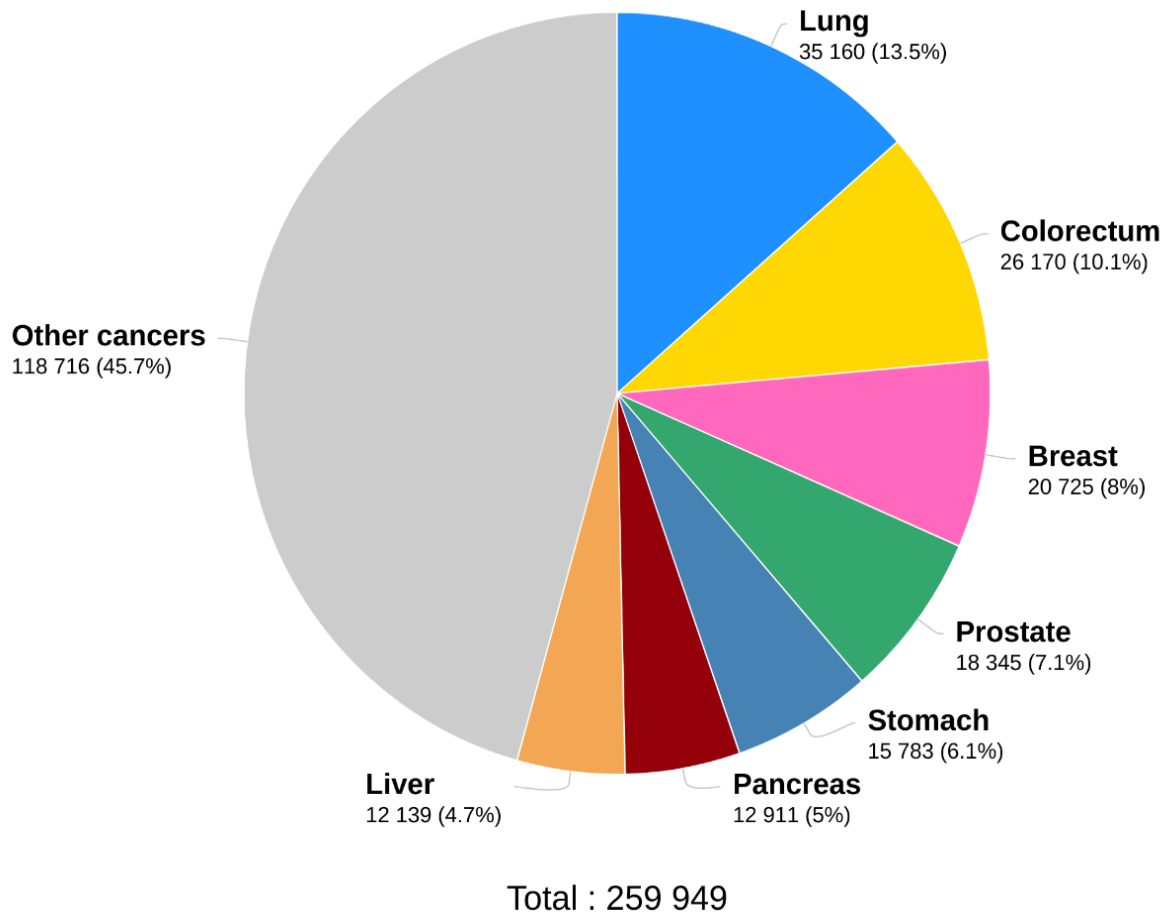
10

11 **Fonte:** Adaptado de OMS (2020).

12

13 Quanto às estimativas de mortalidade para o mesmo ano, o câncer de
 14 pulmão permaneceu no topo da lista, com 35.160 óbitos, seguido por câncer colorretal
 15 (26.170 óbitos), câncer de mama (20.725 óbitos), câncer de próstata (18.345 óbitos),
 16 câncer de estômago (15.783 óbitos), câncer de pâncreas (12.911 óbitos) e câncer de
 17 fígado (12.139 óbitos) (**Figura 4**) (OMS, 2020).

- 1 **Figura 4:** Número estimado de óbitos em 2020 no Brasil, para ambos os sexos e todas
2 as idades.



3

4 **Fonte:** Adaptado de OMS (2020).

5

6 **1.3 PATOGÊNESE DO CÂNCER DE PULMÃO NSCLC**

7 A patogênese do NSCLC envolve uma interação complexa de fatores
8 genéticos, ambientais e moleculares que levam ao desenvolvimento de células
9 cancerígenas nos pulmões. Tais fenômenos resultam na autossuficiência em sinais
10 de crescimento, insensibilidade aos sinais antiproliferativos, evasão de apoptose,
11 potencial replicativo ilimitado devido à ativação da telomerase, invasão tecidual e
12 disseminação em locais distantes (PANOV, 2005; SINGH; KATHIRESAN, 2014). A
13 interferência entre o estresse oxidativo, o estresse metabólico e as principais vias de
14 sinalização, como PI3K (fosfoinositídeo 3-quinase), Akt (proteína quinase B), mTOR
15 (proteína alvo da rifampicina em mamíferos) e NF-κB (fator nuclear-*kappa* B)
16 desempenham um papel significativo na patogênese do câncer de pulmão (WENG et

1 al., 2019; SANAEI et al., 2022). A seguir será apresentada uma visão geral de como
2 esses fatores se interconectam no contexto do desenvolvimento do câncer de pulmão.

3 4 **1.3.1 VIA DE SINALIZAÇÃO PI3K/AKT/MTOR E NF-κB**

5 A via de sinalização PI3K/Akt/mTOR desempenha papéis cruciais na
6 regulação de processos celulares, incluindo sobrevivência, proliferação, crescimento
7 e metabolismo. Essa via é frequentemente hiperativada em células cancerígenas,
8 contribuindo para a angiogênese, metastátase e resistência a terapias (WENG et al.,
9 2019) A proteína PI3K, classificada em classes I, II e III, é um dímero pertencente à
10 família de lipídios-quinases, e é responsável por fosforilar o grupo 3'-OH dos
11 fosfolipídios do inositol. PI3Ks de classe I consistem em subunidades catalíticas e
12 adaptadoras/reguladoras que formam heterodímeros. PI3Ks de classe II incluem a
13 subclasse IB, ativadas por receptores associados a proteínas G, e as isoformas
14 catalíticas que não possuem subunidades reguladoras e são ativadas por receptores
15 com atividade de tirosina quinase. Enquanto PI3Ks de classe III (PIK3C3)
16 desempenham papel na regulação da fagocitose e autofagia de macrófagos,
17 interagindo com um complexo proteico composto por uma subunidade reguladora e
18 uma subunidade catalítica. Akt é um membro da família da proteína quinase AGC
19 (PKA/PKG/PKC) e consiste em três homólogos (Akt1, Akt2 e Akt3). O mTOR é uma
20 proteína quinase pertencente à família de quinases relacionadas à PI3K, encontrado
21 em complexos como o mTORC1 (composto por mTOR, Raptor, mLST8 e PRAS40) e
22 o mTORC2 (composto por mTOR, Rictor, Sin1 e mLST8) (PORTA; PAGLINO;
23 MOSCA, 2014; ERSAHIN; TUNCBAG; CETIN-ATALAY, 2015; HE et al., 2021).

24 Os eixos PI3K/Akt e mTOR estão interconectados de tal forma que
25 podem ser considerados como uma única via. A hiperatividade dessa via está
26 associada ao aparecimento e desenvolvimento de tumores (FANG et al., 2020).
27 Alterações nos componentes desse eixo podem contribuir para a progressão do
28 câncer de pulmão, promover a metástase tumoral e desencadear uma série de
29 processos celulares, como sobrevivência, proliferação, migração, metástase,
30 angiogênese, metabolismo celular, senescência celular, integridade genômica e auto-
31 renovação de células-tronco. A Akt está envolvida em processos como apoptose,
32 proliferação, transcrição e migração. Ela desempenha um papel crucial na regulação
33 da progressão do ciclo celular, impedindo a transição da fase G1 para S, entre outros
34 processos (TAN, 2020). Além disso, a via de sinalização PI3K/Akt/mTOR pode afetar

1 negativamente proteínas epiteliais, como a E-caderina, e positivamente proteínas
2 mesenquimais, como N-caderina, vimentina e β -catenina, através da inibição do
3 homólogo de fosfatase tensina (PTEN). Essas alterações são responsáveis pela
4 indução da transição epitélio-mesenquimal (do inglês, *epithelial-mesenchymal*
5 *transition*, EMT) (SANAEL et al., 2022), um processo biológico que desempenha papel
6 crítico no desenvolvimento do câncer, pois envolve a transformação de células
7 epiteliais (forte adesão célula a célula e natureza relativamente estacionária) em
8 células semelhantes a mesenquimais (frouxamente conectadas, mais móveis e
9 invasivas), sendo um passo crucial na disseminação metastática do câncer
10 (PÉGLION; ETIENNE-MANNEVILLE, 2012; BUSH et al., 2019; JANISZEWSKA;
11 PRIMI; IZARD, 2020).

12 A interação entre a via PI3K-Akt-mTOR e o fator de transcrição NF-
13 κ B é reconhecida por contribuir para a sobrevivência e proliferação das células de
14 câncer de pulmão (CHEN et al., 2011; WENG et al., 2019). O NF- κ B é um fator de
15 transcrição crucial que desempenha um papel fundamental em respostas
16 imunológicas, inflamações, adesão celular, diferenciação, proliferação e apoptose
17 (JOMOVA et al., 2023). Ele existe em várias formas, incluindo NF- κ B/p105, NF-
18 κ B2/p100, RelA/p65, RelB e c-Rel, que podem formar heterodímeros ou homodímeros
19 e se ligar a sequências consenso de DNA (XIA; SHEN; VERMA, 2014). Sua ativação
20 descontrolada está associada a várias condições patológicas, incluindo câncer. Essa
21 ativação pode ocorrer por meio de vias canônicas e não canônicas, desencadeadas
22 por diferentes estímulos (JOMOVA et al., 2023; REYNAERT et al., 2006).

23 O NF- κ B desempenha um papel crítico na prevenção da apoptose,
24 induzindo a expressão de proteínas antiapoptóticas e suprimindo a apoptose mediada
25 pelo estresse celular através da remoção de espécies reativas de oxigênio (do inglês,
26 *reactive oxygen species* - ROS) e do aumento da expressão da superóxido dismutase
27 de manganês (MnSOD). Assim, o NF- κ B inibe a via mitocondrial (intrínseca) e do
28 receptor de morte (extrínseca). Além disso, ele pode regular negativamente a ativação
29 apoptótica de c-Jun N-Terminal quinase (JNK) e antagonizar a ação de p53. A via
30 PI3K/Akt também está envolvida na ativação do NF- κ B de maneira dependente de I κ B
31 kinase α (IKK α), contribuindo para a sobrevivência das células de câncer de pulmão
32 (WENG et al., 2019; SANAEL et al., 2022). Além disso, alguns fatores de crescimento,
33 como o fator de crescimento epidérmico (EGF), podem ativar o NF- κ B independente
34 de IKK, favorecendo o crescimento de células de adenocarcinoma pulmonar de

1 células não pequenas (CHEN et al., 2011).

3 1.3.2 ESTRESSE OXIDATIVO

4 As espécies reativas de oxigênio (ROS) e as espécies reativas de
5 nitrogênio (*reactive nitrogen species* - RNS), conhecidas como RONS, são moléculas
6 altamente reativas que contêm oxigênio e nitrogênio, respectivamente. Elas são
7 produzidas naturalmente no organismo como subprodutos do metabolismo celular e
8 têm papéis importantes em processos biológicos normais, como sinalização celular e
9 resposta imunológica (LIOU; STORZ, 2010; KALUDERCIC; GIORGIO, 2016).

10 A maioria dos RONS contém elétrons desemparelhados e são
11 denominados radicais livres. Em mamíferos, enzimas especializadas, como NADPH-
12 oxidase, mieloperoxidase e óxido nítrico sintase (do inglês, *nitric oxide sintase* - NOS),
13 desempenham um papel fundamental na geração controlada de RONS no ambiente
14 intra e extracelular, como parte do sistema imunológico inato para combater bactérias.
15 No entanto, a produção excessiva de RONS da origem ao estresse oxidativo, o que
16 pode contribuir para a indução de danos às estruturas biológicas do hospedeiro (LIOU;
17 STORZ, 2010; KALUDERCIC; GIORGIO, 2016).

18 As ROS podem ser divididas em duas categorias distintas: radicais
19 livres de oxigênio e ROS não radicais. Entre os radicais livres de oxigênio, destacam-
20 se: ânion superóxido ($O_2^{\cdot-}$), hidroxila ($\cdot OH$), peróxido de hidrogênio (H_2O_2), hipocloreto
21 (HOCl) e compostos carbonílicos altamente reativos derivados de lipídios ou
22 carboidratos. Destes, os radicais superóxido, peróxido de hidrogênio e hidroxila são
23 particularmente estudados em relação ao câncer. As RNS, por sua vez, incluem: óxido
24 nítrico (do inglês, *nitric oxide* - $NO\cdot$), dióxido de nitrogênio ($NO_2^{\cdot-}$), peroxinitrito
25 ($ONOO^-$), trióxido de dinitrogênio (N_2O_3) e ácido nitroso (HNO_2), que também
26 contribuem para o estresse oxidativo (WEIDINGER; KOZLOV, 2015; CANTON et al.,
27 2021).

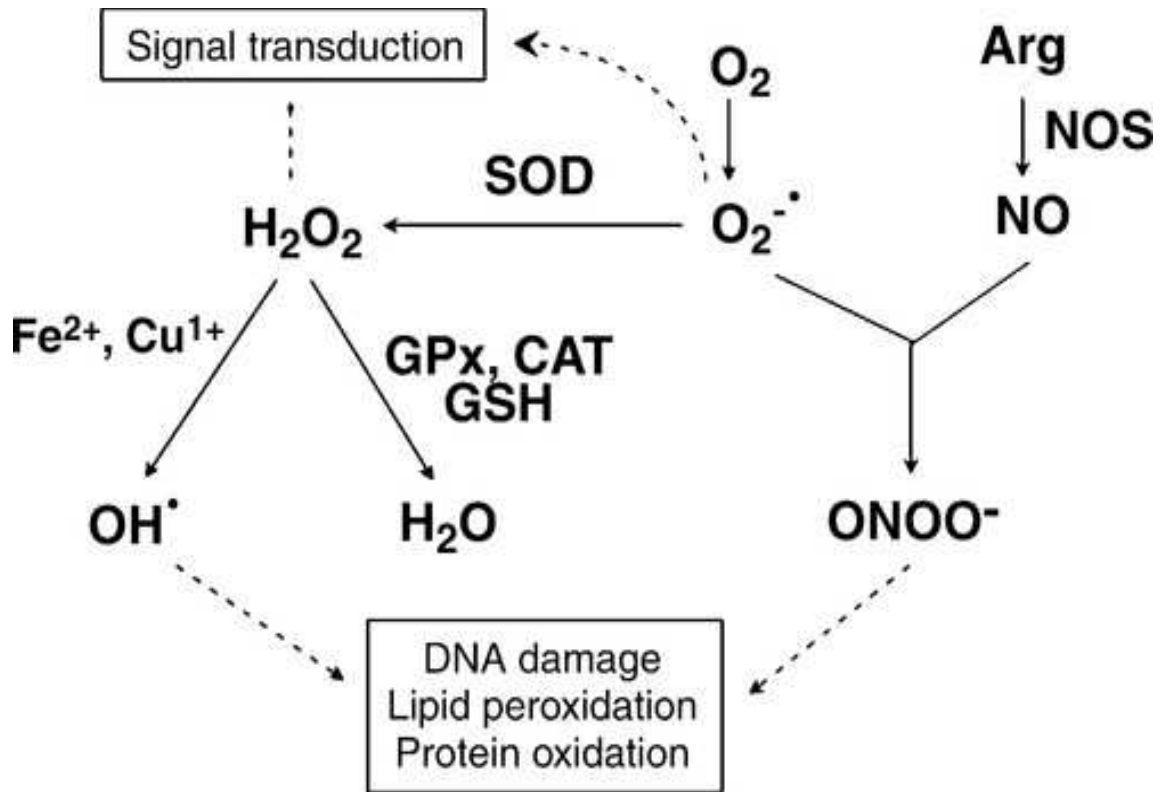
28 Várias fontes intracelulares de ROS/RNS foram identificadas e estão
29 associadas a diversas patologias. Essas fontes incluem NADPH oxidases, xantina
30 oxidase, óxido nítrico sintase desacoplada (NOS) e mitocôndrias, que em condições
31 patológicas podem contribuir para danos oxidativos em células e tecidos. Em
32 concentrações baixas, RONS desempenham um papel crucial como mediadores
33 regulatórios em processos de sinalização. No entanto, em concentrações moderadas
34 ou elevadas, podem ser prejudiciais para os organismos vivos, inativando moléculas

1 celulares importantes. Além disso, o aumento excessivo dessas espécies reativas,
2 quando induzido por quimioterápicos, pode resultar na interrupção do ciclo celular,
3 senescência ou morte de células tumorais (MIJATOVIĆ et al., 2020; CANTON et al.,
4 2021).

5 Espécies reativas de oxigênio são produzidas durante o metabolismo
6 celular aeróbio normal. Cerca de 85–90% do oxigênio (O_2) consumido é metabolizado
7 nas mitocôndrias pela cadeia de transporte de elétrons e os 10–15% restantes são
8 utilizados por diversas enzimas oxidases e oxigenases e por reações químicas de
9 oxidação direta. O oxigênio é reduzido a ânion superóxido ($O_2^{\cdot-}$) por diferentes
10 mecanismos celulares. O superóxido ($O_2^{\cdot-}$) produzido por diversas fontes atua como
11 espécie reativa primária. $O_2^{\cdot-}$ reage rapidamente com o óxido nítrico ($NO\cdot$) para
12 produzir peroxinitrito ($ONOO^-$) ou é catalisado pela superóxido dismutase (SOD) para
13 produzir peróxido de hidrogênio (H_2O_2). O H_2O_2 pode ser neutralizado pela catalase
14 ou glutathione peroxidase (GSH). No entanto, na presença de íons de metais de
15 transição, como ferro (Fe^{2+}) e cobre (Cu^+), radicais livres hidroxila ($OH\cdot$) altamente
16 tóxicos podem ser produzidos a partir de H_2O_2 (reação de Fenton). As ROS podem
17 atuar como um segundo mensageiro para ativar o fator nuclear kappa-B (NF- κ B), que
18 orchestra um espectro de expressão gênica envolvida na resposta inflamatória e na
19 transcrição de citocinas pró-inflamatórias (COSTA et al., 2018; FUJII; HOMMA;
20 OSAKI, 2022; ANDRÉS et al., 2023) (**Figura 5**).

21

22 **Figura 5:** Vias de ação do $O_2^{\cdot-}$ e $ONOO^-$.



1
2 **Fonte:** BUETLER; KRAUSKOPF; RUEGG (2004).
3

4 O óxido nítrico é uma molécula gasosa altamente reativa de vida
5 curta, desempenhando um papel fundamental na regulação de diversos processos
6 fisiológicos e fisiopatológicos. Isso inclui funções vasculares, como angiogênese,
7 controle do fluxo sanguíneo, permeabilidade vascular, interação entre leucócitos e
8 células endoteliais, agregação plaquetária e fluxo microlinfático. Além disso, o NO
9 desempenha um papel importante em funções neurológicas, como neurotransmissão
10 e desenvolvimento do sistema nervoso. Em concentrações elevadas, o NO também
11 tem funções citotóxicas, como a indução de citostase e citólise (FUKUMURA;
12 KASHIWAGI; JAIN, 2006). A produção de NO é mediada por uma família de enzimas
13 chamadas NOS, que incluem três isoformas: NOS neuronal (nNOS/NOS1), NOS
14 endotelial (eNOS/NOS3) e NOS induzível (iNOS/NOS2). A expressão da iNOS é
15 regulada por fatores pró-inflamatórios, como interferon-gama (IFN- γ), interleucina-1 β
16 (IL-1 β), fator de necrose tumoral-alfa (TNF- α), prostaglandinas, lipopolissacarídeos
17 (LPS) e/ou hipóxia. A produção de NO ocorre por meio da conversão de L-arginina
18 em citrulina, utilizando fosfato de dinucleotídeo de nicotinamida e adenina (NADPH)
19 como cofator (KASHFI; KANNIKAL; NATH, 2021; HUANG et al., 2023).

20 O NO e seus metabólitos, como nitrito, nitrato, S-nitrosotióis,

1 nitrosaminas e peroxinitrito, têm papéis importantes nos efeitos citotóxicos e/ou
2 genotóxicos. Eles induzem o estresse oxidativo e nitrosativo, resultando em danos ao
3 DNA, incluindo desaminação nitrosativa de bases de ácidos nucleicos, alquilação e
4 quebra da cadeia de DNA. Além disso, o NO pode inibir enzimas responsáveis pelo
5 reparo do DNA, como alquiltransferase e DNA ligase. Portanto, vários estudos
6 indicaram que o NO pode desencadear e/ou promover o desenvolvimento de tumores
7 (FUKUMURA; KASHIWAGI; JAIN, 2006).

8 Taxas elevadas de ROS também foram detectadas em quase todos
9 os tipos de câncer, induzindo o desenvolvimento e progressão do tumor. No entanto,
10 as células tumorais também expressam níveis aumentados de proteínas
11 antioxidantes, a fim de desintoxicar ROS, sugerindo que um delicado equilíbrio dos
12 níveis intracelulares é necessário para a função das células cancerígenas (LI et al.,
13 2023).

14 A homeostase redox celular é um processo essencial e dinâmico que
15 garante o equilíbrio entre as reações de redução e oxidação dentro das células (isto
16 é, ocorre transferência de elétrons de um doador (“agente redutor”) para um aceitador
17 (“agente oxidante”) e regula uma infinidade de respostas e eventos biológicos,
18 incluindo metabolismo, morte celular, diferenciação e desenvolvimento, respostas
19 imunes, ritmo circadiano (TRETTER et al., 2021; LE GAL; SCHMIDT; SAYIN, 2021),
20 manutenção da sobrevivência e função normal das células, sendo estabelecido um
21 equilíbrio entre produção e eliminação de RONS (LENNICKE; COCHEMÉ, 2021; LI et
22 al., 2023). As defesas antioxidantes incluem moléculas enzimáticas e não-
23 enzimáticas, que exercem efeito protetor contra os danos oxidativos, com destaque
24 para a superóxido dismutase e glutathione (BECKER; INDRA, 2023). A superóxido
25 dismutase (SOD) é uma metaloenzima responsável por catalisar a dismutação de
26 duas moléculas de ânion superóxido em peróxido de hidrogênio e oxigênio molecular.
27 Nos mamíferos, existem três formas distintas das quais requerem presença de metal
28 para sua ativação: SOD1 (Cu/ZnSOD) presente no citoplasma; SOD2 (MnSOD)
29 mitocondrial; e SOD3 (Cu/ZnSOD) extracelular. SOD1 é a principal forma intracelular,
30 representando cerca de 80% de SOD total (JOMOVA et al., 2023). A glutathione (GSH),
31 por sua vez, consiste em um antioxidante não-enzimático, composta por aminoácidos
32 (glutamina, cisteína e glicina), responsável pela desintoxicação de radicais livres e
33 xenobióticos. Para realizar essa função, as glutathione peroxidases (GSH-Px)
34 reduzem o peróxido de hidrogênio em água, convertendo GSH em glutathione

1 dissulfeto (GSSG) (AHMADI; SHADBOORESTAN, 2016).

2 Embora a perda de GSH, ou uma diminuição na relação
3 glutaciona/dissulfeto de glutaciona (GSH/GSSG), leve a um aumento da suscetibilidade
4 ao estresse oxidativo e à carcinogênese, níveis elevados de GSH aumentam a
5 capacidade antioxidante de muitas células cancerígenas, aumentando sua resistência
6 ao estresse oxidativo (TRAVERSO et al., 2013). Notavelmente, a inibição desses
7 sistemas antioxidantes induz morte das células tumorais, demonstrando a importância
8 dos sistemas antioxidantes em favorecer a progressão do tumor.

9 10 **1.3.3 METABOLISMO ENERGÉTICO DA CÉLULA TUMORAL**

11 Tanto células saudáveis como células tumorais contam com as
12 mesmas maquinarias enzimáticas para manter a homeostase celular. Na maioria das
13 células de mamíferos, a glicose é a principal fonte de combustível. A glicose passa
14 por um processo metabólico chamado glicólise, que envolve uma série de reações em
15 várias etapas, culminando na produção de piruvato. Em condições normais de
16 oxigênio, a maior parte desse piruvato é encaminhada para as mitocôndrias, onde
17 passa pelo Ciclo de Krebs, um processo de oxidação que resulta na geração de ATP
18 para suprir as necessidades energéticas da célula (HSU; SABATINI, 2008). Nas
19 células tumorais, as mutações genéticas e as alterações nas interações entre as
20 moléculas estão estreitamente ligadas a modificações complexas no metabolismo
21 energético celular. Isso resulta em uma reprogramação intrincada das vias metabólicas
22 que envolvem a glicose, lipídeos e aminoácidos. Essas transformações metabólicas
23 beneficiam o crescimento, a sobrevivência, a capacidade invasiva e o avanço do tumor
24 (PAYEN et al., 2016).

25 O fenômeno metabólico em questão, denominado "*Efeito Warburg*", é
26 uma notável característica observada em células tumorais que tem sido objeto de
27 extensa pesquisa (LIN et al., 2020). Este fenômeno se manifesta como um aumento
28 substancial na taxa de glicólise aeróbica nas células cancerosas, mesmo quando há
29 presença adequada de oxigênio, uma situação que normalmente estimula o uso de
30 processos metabólicos mais eficientes, como a fosforilação oxidativa (OXPHOS)
31 (LIBERTI; LOCASALE, 2016). A glicólise aeróbica persistente em células tumorais é
32 notável porque contrasta com o comportamento metabólico das células normais, que
33 normalmente utilizam a OXPHOS como principal via para gerar energia na presença
34 de oxigênio adequado (LIN et al., 2020; PAYEN et al., 2015).

1 Neste contexto, o lactato desempenha um papel fundamental.
2 Durante a glicólise aeróbica, que é o processo metabólico onde a glicose é convertida
3 em piruvato na presença de oxigênio, o piruvato normalmente entra na cadeia de
4 transporte de elétrons e é oxidado na mitocôndria para produzir energia na forma de
5 ATP (KENNEDY et al., 2013). No entanto, nas células tumorais que exibem o *Efeito*
6 *Warburg*, há uma mudança notável nesse processo. Nas células tumorais afetadas
7 pelo *Efeito Warburg*, o piruvato gerado a partir da glicólise aeróbica não é
8 completamente direcionado para a OXPHOS. Em vez disso, uma parte significativa
9 do piruvato é convertida em lactato por meio da enzima lactato desidrogenase (LDH),
10 mesmo na presença de oxigênio suficiente. Esse processo é conhecido como a
11 fermentação do piruvato para lactato (MARTINEZ-OUTSCHOORN et al., 2016)

12 A acumulação de lactato nas células tumorais e no ambiente
13 circundante é uma característica distintiva do *Efeito Warburg*. Isso ocorre porque a
14 glicólise aeróbica é um processo menos eficiente em termos de produção de ATP em
15 comparação com a OXPHOS, mas fornece uma vantagem metabólica para as células
16 cancerosas ao permitir uma produção mais rápida de ATP e intermediários
17 metabólicos que podem ser usados para a biossíntese de macromoléculas
18 necessárias para o rápido crescimento e proliferação das células tumorais. A
19 acumulação de lactato também tem implicações no ambiente tumoral, criando um
20 ambiente ácido que pode promover a invasão e metástase das células cancerosas,
21 além de desencadear respostas imunossupressoras (AKRAM, 2013; VAZQUEZ et al.,
22 2010).

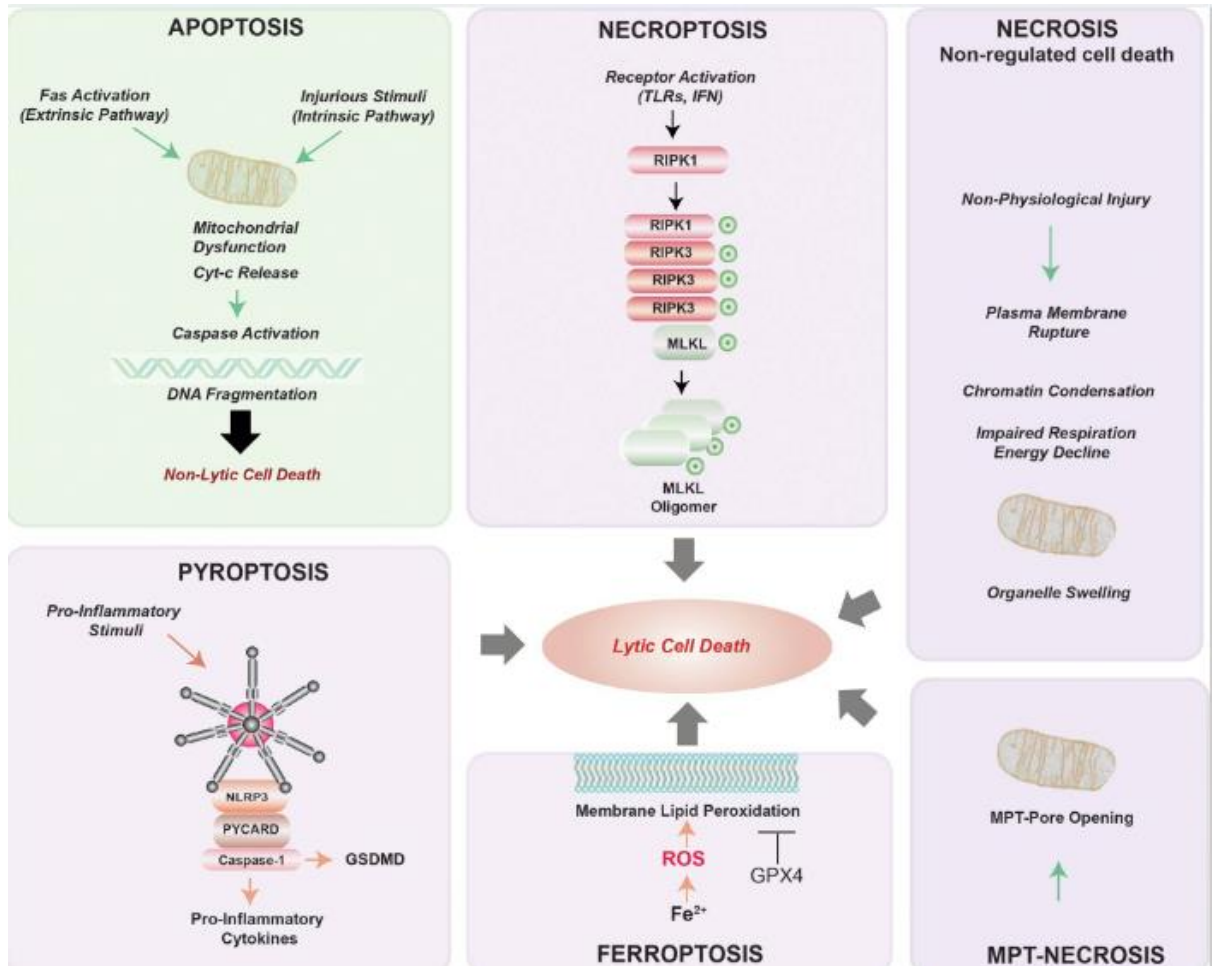
23 A compreensão detalhada do *Efeito Warburg* e suas implicações nas
24 células tumorais é de grande importância na pesquisa sobre câncer, uma vez que
25 essa adaptação metabólica singular é vista como uma estratégia central para atender
26 às demandas bioenergéticas e biosintéticas das células de câncer, contribuindo para
27 a sua proliferação descontrolada. A investigação contínua desse fenômeno
28 metabólico tem o potencial de levar a abordagens terapêuticas inovadoras no
29 tratamento do câncer.

30 31 **1.3.4 PRINCIPAIS TIPOS DE MORTE CELULAR**

32 A morte celular é um processo biológico fundamental que ocorre tanto
33 em situações normais quanto patológicas. No contexto do câncer, as células
34 conseguem evitar os mecanismos indutores de morte celular e promover o

1 crescimento descontrolado, bem como, manter a sobrevivência dessas células
 2 aberrantes. Existem vários tipos de morte celular, cada um com características e
 3 mecanismos distintos (**Figura 6**).

4
 5 **Figura 6:** Tipos de morte celular e suas características gerais.
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7
 8 **Fonte:** CHOI et al. (2019).
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10 1.3.4.1 AUTOFAGIA

11 A autofagia é um mecanismo intracelular de degradação no qual
 12 proteínas e organelas são encapsuladas em vesículas especializadas e
 13 posteriormente decompostas por enzimas lisossomais. Esse processo tem como
 14 objetivo a reciclagem de componentes celulares e a provisão de nutrientes essenciais
 15 para a célula. Em situações normais, a autofagia ocorre em níveis basais,
 16 desempenhando um papel fundamental na manutenção da homeostase celular. No
 17 entanto, em resposta ao estresse celular, como a falta de nutrientes, hipóxia, estresse

1 no retículo endoplasmático, estresse oxidativo ou exposição a substâncias tóxicas, a
2 autofagia é ativada como uma via pró-sobrevivência, impedindo a morte celular (YUN;
3 LEE, 2018; MULCAHY LEVY; THORBURN, 2020; LIM; MOHAMAD HANIF; CHIN,
4 2021).

5 Existem três tipos principais de autofagia que resultam na degradação
6 lisossomal: Macroautofagia (ou simplesmente autofagia): caracterizada pela formação
7 de autofagossomos, que são vacúolos de membrana dupla capazes de capturar
8 proteínas citoplasmáticas, mitocôndrias, retículo endoplasmático e ribossomos. Esses
9 autofagossomos, uma vez fundidos com lisossomos, formam os autofagolisossomos,
10 onde ocorre a degradação das substâncias encapsuladas. Os produtos de
11 degradação resultantes, como aminoácidos, nucleotídeos, ácidos graxos e açúcares,
12 são reciclados para sustentar o metabolismo celular. Microautofagia: nesse tipo, o
13 material citoplasmático é diretamente incorporado pelo lisossomo, sem a formação de
14 vesículas de membrana dupla. Autofagia mediada por chaperonas: neste caso,
15 proteínas específicas são translocadas através da membrana lisossomal e
16 subsequentemente desdobradas e degradadas no interior do lisossomo (YUN; LEE,
17 2018; LIM; MOHAMAD HANIF; CHIN, 2021; MULCAHY LEVY; THORBURN, 2020).

18 A influência da autofagia no desenvolvimento de tumores ainda não
19 está completamente esclarecida, pois pode desempenhar papéis contraditórios. Por
20 um lado, a autofagia pode promover a sobrevivência das células tumorais, fornecendo
21 nutrientes essenciais durante o estresse e permitindo a resistência à quimioterapia.
22 Por outro lado, a autofagia pode suprimir o desenvolvimento de tumores, eliminando
23 células de câncer que não podem passar pelo processo apoptótico. Além disso, em
24 alguns casos, bloquear a autofagia pode ser necessário para manter o fenótipo
25 maligno de certas células de câncer. Em resumo, a autofagia desempenha um papel
26 complexo no contexto do câncer, fornecendo uma vantagem de sobrevivência às
27 células tumorais em algumas situações, enquanto suprime o desenvolvimento tumoral
28 em outras (YUN; LEE, 2018; LIM; MOHAMAD HANIF; CHIN, 2021; MULCAHY LEVY;
29 THORBURN, 2020).

30

31

1.3.4.2 MORTE CELULAR PROGRAMADA - APOPTOSE

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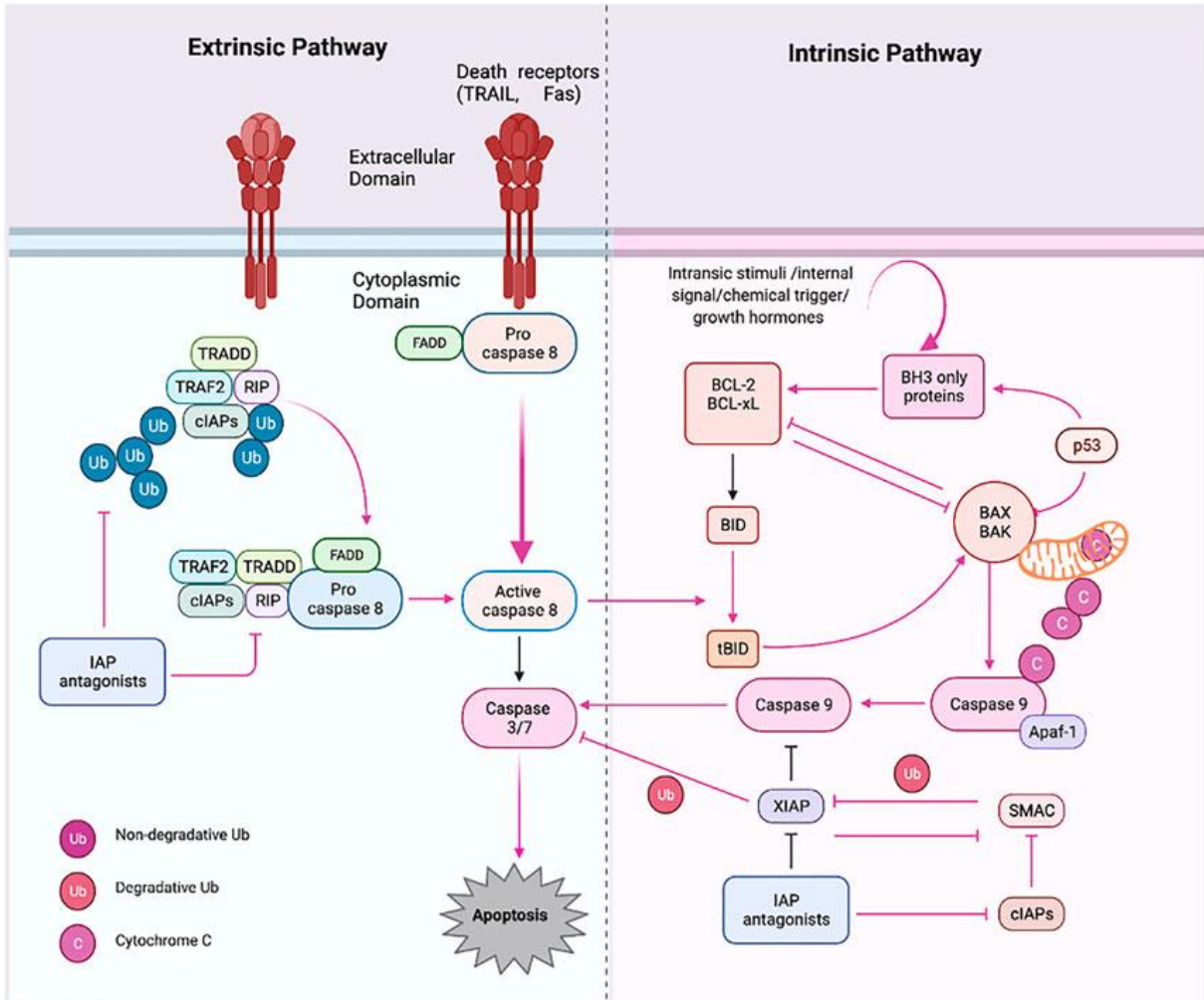
33

A morte celular pode ocorrer através de um processo ativo e regulado,
conhecido como morte celular programada, ou de um curso passivo e descontrolado.

1 Devido às dramáticas diferenças morfológicas desses dois tipos de morte celular, eles
2 foram originalmente denominados em apoptose e necrose, respectivamente.

3 A apoptose ou morte celular programada é um processo fisiológico
4 que promove de maneira eficaz a remoção de células em excesso ou danificadas,
5 sem a geração de resposta inflamatória, garantindo assim a homeostase do tecido. A
6 apoptose é o principal tipo de morte celular que ocorre quando o dano ao DNA é
7 irreparável e pode ser ativada de duas maneiras, pela via intrínseca ou mitocondrial e
8 pela via extrínseca ou do receptor de morte. A via intrínseca é desencadeada pela
9 perda do potencial de membrana mitocondrial, pelo seu rompimento decorrentes de
10 danos no DNA causados por agentes citotóxicos, privação de fator de crescimento e
11 privação de citocinas o que resulta na liberação do citocromo C no citoplasma. A via
12 extrínseca é mediada por receptores transmembranares, chamados de receptores de
13 morte, que são membros da superfamília do gene do receptor do fator de necrose
14 tumoral (TNF) produzidos por células T citotóxicas do sistema imunológico em
15 resposta a células danificadas ou infectadas. Os caminhos convergem nas caspases
16 executoras (PFEFFER; SINGH, 2018; CARNEIRO; EL-DEIRY, 2020; CHAUDHRY et
17 al., 2022) (**Figura 7**).

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19 **Figura 7:** Vias de apoptose.



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Fonte: CHAUDHRY et al. (2022), criado com BioRender.com. A apoptose tem duas vias principais conhecidas como i) via extrínseca e ii) intrínseca. Primeiro, estímulos externos ou moléculas ligantes ativam a transdução, incluindo receptores de morte, levando à ativação da caspase 3/7 ou 8. Via intrínseca; através da inserção de moléculas pró-apoptóticas BAX (proteína) na membrana mitocondrial resulta na geração de citocromo c, formando um apoptossomo, que desencadeia ainda mais as cascatas apoptóticas começando com a ativação pró-apoptótica da caspase 9 e ou então da caspase 3.

Assim que a apoptose é sinalizada, mudanças começam a ocorrer dentro da célula. Estas alterações incluem a ativação de caspases que clivam componentes celulares necessários para a função celular normal, tais como proteínas do citoesqueleto e nucleares. Como resultado da atividade das caspases, as células apoptóticas começam a encolher e sofrem alterações na membrana plasmática que sinalizam a resposta dos macrófagos. A apoptose é realizada por caspases (proteases específicas da cisteína aspartil), que são uma classe de proteínas cisteínas que clivam proteínas alvo. A atividade da caspase protease é essencial para o sucesso da

1 apoptose, pois clivam centenas de várias proteínas. Existem quatro caspases
2 iniciadoras (caspase-2, -8, -9, 10) e três caspases executoras (caspase-3, -6, -7). As
3 caspases executoras clivam as proteínas alvo, o que eventualmente leva à morte da
4 célula. As vias são altamente reguladas, de modo que a apoptose só ocorrerá se for
5 sinalizada. A via intrínseca, em particular, é regulada pela família de proteínas do
6 linfoma-2 de células B (BCL-2), que inclui proteínas efetoras pró-apoptóticas,
7 proteínas pró-apoptóticas BH3 e proteínas BCL-2 anti-apoptóticas. As proteínas
8 antiapoptóticas BCL-2 inibem a apoptose através da inibição das proteínas BCL-2 pró-
9 apoptóticas, proteína X associada a BCL-2 (BAX) e antagonista homólogo BCL-2
10 (BAK). As proteínas BH3 inibem as proteínas antiapoptóticas BCL-2 (PFEFFER;
11 SINGH, 2018; CARNEIRO; EL-DEIRY, 2020; CHAUDHRY et al., 2022).

12 O termo apoptose descreve um processo de morte celular, cujas
13 alterações incluem arredondamento da célula, redução do volume celular (picnose),
14 formação de bolhas na membrana plasmática (*blebbing*), perda de adesão celular,
15 condensação nuclear e, por fim, fagocitose por macrófagos residentes sem
16 desencadear inflamação. As alterações moleculares observadas neste processo
17 correspondem à exposição de fosftidilserina, fragmentação oligonucleossomal de
18 DNA, ativação de caspases, perda do potencial de membrana mitocondrial com
19 manutenção da integridade da membrana plasmática (PFEFFER; SINGH, 2018;
20 CARNEIRO; EL-DEIRY, 2020; CHAUDHRY et al., 2022).

21 A desregulação da apoptose é um fenômeno encontrado em uma
22 ampla variedade de doenças. Evidências acumuladas mostraram que a expressão
23 anormal de alguns fatores regulatórios da apoptose pode levar ao câncer, indicando
24 as intrincadas relações entre apoptose e câncer, como por exemplo, a molécula Bcl-
25 2, a qual é super expressa em muitos tipos de células cancerígenas e o seu aumento
26 ainda conduz a resistência a fármacos quimioterapêuticos e terapia de radiação
27 (PFEFFER; SINGH, 2018; CARNEIRO; EL-DEIRY, 2020; CHAUDHRY et al., 2022).
28 Portanto, a regulação da apoptose padrão poderia beneficiar o tratamento e a
29 prevenção do câncer.

30

31 **1.3.4.3 MORTE CELULAR PROGRAMADA – PIROPTOSE**

32 A piroptose, uma forma de morte celular programada, é uma via
33 recentemente identificada de morte celular que é estimulada por uma série de
34 infecções microbianas (por exemplo, *Salmonella*, *Francisella* e *Legionella*) e estímulos

1 não infecciosos, incluindo fatores do hospedeiro produzidos durante o enfarte do
2 miocárdio. É um processo inflamatório altamente regulado que desempenha um papel
3 crucial na defesa do organismo contra infecções. Quando as células imunológicas
4 detectam sinais de infecção, elas podem desencadear piroptose nas células
5 infectadas para eliminar o patógeno (BERGSBAKEN; FINK; COOKSON, 2009; TAN
6 et al., 2021; ZHENG et al., 2021; YU et al., 2021).

7 A piroptose é morfológica e mecanicamente distinta de outras formas
8 de morte celular. O início da piroptose começa com o reconhecimento de uma
9 ameaça, que pode ser um patógeno intracelular (por exemplo, bactérias ou vírus) ou
10 sinais de estresse celular. Vários receptores de reconhecimento de padrões (RRPs)
11 na superfície ou dentro da célula, incluindo receptores do tipo *Toll* (TLRs) e receptores
12 do tipo Nod (NLRs), podem detectar a presença da ameaça. Em resposta à ameaça,
13 RRP's específicos podem ativar um complexo proteico denominado inflamassoma.
14 Uma vez ativado o inflamassoma, ele leva à ativação da caspase-1, uma enzima
15 protease mediadora chave na via da piroptose. A caspase-1, cliva as proteínas
16 gasdermina, particularmente gasdermina-D (GSDMD). As gasderminas ativadas
17 sofrem uma mudança conformacional, expondo seu domínio N-terminal, formando
18 poros na membrana celular. Esses poros permitem que íons e fluidos entrem na célula
19 e perturbem o equilíbrio osmótico, levando ao inchaço celular e, por fim, à ruptura da
20 membrana. Esta ruptura leva à liberação de conteúdos celulares, incluindo citocinas
21 pró-inflamatórias, como interleucina-1 beta (IL-1 β) e interleucina-18 (IL-18), no
22 ambiente extracelular. Essas citocinas ativam respostas imunológicas e atraem
23 células do sistema imune para o local da infecção ou dano celular. Este processo
24 ajuda a conter a infecção (BERGSBAKEN; FINK; COOKSON, 2009; TAN et al., 2021;
25 ZHENG et al., 2021; YU et al., 2021).

26 Este processo contrasta com a apoptose, onde as células encolhem
27 e se fragmentam em corpos apoptóticos que são silenciosamente engolidos por
28 células vizinhas ou fagócitos.

30 **1.3.4.4 MORTE CELULAR PROGRAMADA - NECROPTOSE**

31 A necroptose é uma forma de morte celular programada que
32 compartilha características tanto com a apoptose quanto com a necrose (morte celular
33 não programada). É um processo altamente regulado que pode ser iniciado sob

1 condições específicas e desempenha um papel em vários contextos fisiológicos e
2 patológicos (DHURIYA; SHARMA, 2018; CHOI et al., 2019).

3 A necroptose é iniciada quando as células recebem sinais que
4 normalmente desencadeariam a apoptose, mas a apoptose é inibida ou bloqueada.
5 Isto pode ocorrer em resposta a certos receptores de morte, infecções virais ou
6 estresse celular. O processo começa com a ativação de receptores de superfície
7 celular, como os receptores do fator de necrose tumoral (TNF). Esses receptores
8 podem ativar uma cascata de sinalização que inclui proteínas como a proteína quinase
9 1 de interação com o receptor (RIPK1) e a proteína quinase 3 de interação com o
10 receptor (RIPK3). O complexo de sinalização do receptor ativado forma uma estrutura
11 conhecida como necrossoma, que inclui RIPK1 e RIPK3. O necrossoma é um dos
12 principais impulsionadores da necroptose (DHURIYA; SHARMA, 2018; CHOI et al.,
13 2019).

14 Dentro do necrossoma, o RIPK3 fosforila a pseudoquinase
15 semelhante ao domínio da quinase de linhagem mista (MLKL), uma molécula efetora
16 chave na necroptose. O MLKL ativado e fosforilado transloca-se para a membrana
17 celular e forma poros, perturbando a integridade da membrana plasmática. Isso leva
18 à perda da integridade da membrana, fazendo com que a célula inche e se rompa. À
19 medida que ocorre o rompimento, o conteúdo celular, incluindo moléculas pró-
20 inflamatórias e padrões moleculares associados a danos (DAMPs), é liberado no
21 espaço extracelular, o que pode resultar na ativação de respostas imunológicas e
22 contribuir para a inflamação no tecido circundante (DHURIYA; SHARMA, 2018; CHOI
23 et al., 2019).

24 **1.3.4.5 MORTE CELULAR NÃO-PROGRAMADA - NECROSE**

25 O processo de morte celular por necrose ocorre em resposta a muitos
26 tipos de danos, como trauma, infarto, toxinas, e, portanto, é normalmente o resultado
27 de um processo patológico. Por este motivo, a necrose foi considerada morte celular
28 quase "acidental", um processo aleatório e descontrolado (ELMORE, 2007; JABOIN;
29 HWANG; LU, 2009; LIU et al., 2017).

30 O processo necrótico por muitos anos era anteriormente considerado
31 uma morte celular catastrófica e desordenada, caracterizado por integridade da
32 membrana plasmática comprometida, inchaço das organelas celulares, degradação
33 aleatória do DNA e liberação descontrolada de moléculas inflamatórias. Estudos
34 realizados na última década descobriram que a necrose pode ocorrer de forma

1 regulada e inclui vários mecanismos de morte celular conhecidos como necroptose,
2 ocitose, ferroptose, NETose, pironecrose e piroptose. Esses mecanismos de morte
3 celular têm sido empregados para explicar uma variedade de necroses observadas *in*
4 *vivo* em diversas situações patológicas. Ainda assim, se a necrose tumoral é regulada
5 através destes mecanismos e como a necrose se desenvolve no desenvolvimento do
6 tumor permanecem em grande parte obscuros (LIU; JIAO, 2020; YEE; LI, 2021).

7 As principais alterações morfológicas que ocorrem no processo de
8 necrose incluem inchaço celular, formação de vacúolos citoplasmáticos, retículo
9 endoplasmático distendido, formação de bolhas citoplasmáticas, mitocôndrias
10 condensadas, inchadas ou rompidas, desagregação e desprendimento dos
11 ribossomos, membranas de organelas rompidas, lisossomas inchados e rompidos, e,
12 finalmente, a ruptura da membrana celular. Essa perda de integridade da membrana
13 celular resulta na liberação do conteúdo citoplasmático no tecido circundante,
14 enviando sinais quimiotáticos com eventual recrutamento de células inflamatórias.
15 Além disso, a inflamação local subsequente induzida pela necrose pode promover o
16 crescimento do tumor e correlaciona-se com mau prognóstico em tumores sólidos
17 (KARSCH-BLUMAN et al., 2019).

18 Em resumo, o estresse oxidativo, o estresse metabólico e a
19 desregulação das vias de sinalização como PI3K/AKT e NF-κB são processos
20 interligados que contribuem para a patogênese do câncer de pulmão. O estresse
21 oxidativo pode levar a danos no DNA e à disfunção mitocondrial, enquanto o
22 metabolismo alterado, particularmente o *efeito Warburg*, fornece às células
23 cancerígenas a energia e os suprimentos necessários para o crescimento
24 descontrolado. A via PI3K/AKT é frequentemente hiperativada no câncer de pulmão e
25 desempenha um papel fundamental na regulação do metabolismo, sobrevivência e
26 proliferação celular. O NF-κB, associado à inflamação e à sobrevivência celular,
27 também interage com essas vias, promovendo ainda mais o desenvolvimento e
28 progressão do câncer de pulmão. A evasão ou escape da morte celular proporciona
29 as células de câncer capacidade de resistir ou evitar os mecanismos normais de morte
30 celular que normalmente eliminariam células danificadas ou prejudiciais do corpo,
31 contribuindo para o crescimento descontrolado e a sobrevivência das células
32 cancerígenas. A compreensão dessas interações complexas pode fornecer insights
33 sobre potenciais alvos terapêuticos para o tratamento do câncer de pulmão.

1.4 TRATAMENTO

O câncer de pulmão de células não pequenas é considerado uma doença heterogênea. Em vista disso, seu tratamento deve ser realizado de acordo com a condição clínica do paciente (*performance status*), estágio da doença, tipo histológico e perfil molecular. De maneira geral, o NSCLC pode ser classificado em ressecável, localmente avançado e avançado, desta forma, o tratamento pode envolver cirurgia, radiação e quimioterapia, separados ou em combinação, dos quais visam a remoção ou destruição das células tumorais, sem afetar células normais. Em tumores ressecáveis, a ressecção cirúrgica é considerada a opção mais eficaz, no entanto, em tumores de estágio avançado e irressecáveis, é necessária combinação de radiação e quimioterapia (CHAFT et al., 2021). Recentemente, a imunoterapia tem sido amplamente avaliada, envolvendo inibidores de checkpoints imunológicos anti-antígeno 4 de linfócito T citotóxico (anti-CTLA-4) e anti-proteína de morte celular programada 1 (anti-PD-1) (STEVEN; FISHER; ROBINSON, 2016; ONOI et al., 2020).

O quimioterápico de primeira escolha para NSCLC consiste na cisplatina (*cis*-diaminodicloroplatina-II) (CDDP), composto organometálico a base de platina. A CDDP liga-se ao DNA genômico (gDNA) ou mitocondrial (mtDNA) através da formação de adutos, e conseqüentemente, induz distorção da dupla hélice e bloqueio da replicação e transcrição, interferindo nos mecanismos de reparo e levando a morte celular (RANASINGHE; MATHAI; ZULLI, 2022). Inicialmente, o influxo de CDDP nas células ocorre através de difusão passiva ou por meio da proteína transportadora de cobre (CTR1). Dentro das células, ocorre processo de adequação da cisplatina, no qual grupos de cloreto são substituídos por moléculas de água, sendo capaz de reagir com componentes de membrana, citoplasmáticos e DNA.

Compostos à base de platina hidratados, como a cisplatina, são conhecidos por sua capacidade de reagir com o átomo de nitrogênio na posição 7 (N7) das purinas, que são uma das duas principais classes de bases nitrogenadas encontradas no DNA (a outra classe é a das pirimidinas). As purinas incluem adenina e guanina, que são componentes essenciais das moléculas de DNA. A cisplatina, portanto, quando presente no ambiente celular, pode formar ligações covalentes com o átomo N7 das purinas no DNA e induzir ligações cruzadas intracadeias, entre duas purinas adjacentes (1,2 ou 1,3), ou formar ligações intercadeias, entre duas bases de fitas opostas de DNA. A ligação CDDP-DNA pode desencadear mecanismos de reparo por excisão de nucleotídeos (NER, do inglês, *nucleotide excision repair*)

1 permitindo a sobrevivência da célula, no entanto, proteínas HMG (do inglês, *high*
2 *mobility group*) são capazes de proteger do processo de reparo, induzindo parada do
3 ciclo celular. Além do dano ao DNA, a citotoxicidade da cisplatina pode ser induzida
4 através de estresse oxidativo, regulação positiva de p53, ativação de proteína quinase
5 ativada por mitógeno (MAPK), e consequente morte por apoptose (DASARI;
6 BERNARD TCHOUNWOU, 2014; GHOSH, 2019).

7 Os danos induzidos pela cisplatina nas células cancerígenas podem,
8 no entanto, desencadear mecanismos de quimiorresistência, como é o caso de células
9 não pequenas de câncer pulmão, das quais ativam processos que removem ou
10 reparam as lesões do DNA. Assim, a ligação cisplatina-DNA ativa vias de reparo NER,
11 levando ao reconhecimento, desenrolamento e remoção de adutos, ou ainda pode
12 ocorrer correção de bases mal pareadas através da ativação de proteínas de reparo
13 de incompatibilidade (MMR, do inglês, *mismatch repair*) (ROCHA et al., 2018). Além
14 disso, uma vez que o influxo de cisplatina depende de transportadores, a redução da
15 expressão de CTR1 ou rápida degradação pode resultar em sua menor
16 disponibilidade intracelular. A superexpressão de transportadores da família ABC
17 (MRP2) também pode favorecer a quimiorresistência, aumentando a liberação de
18 cisplatina das células (AMABLE, 2016). A CDDP em sua forma adequada é capaz de
19 reagir com diversas proteínas, incluindo a enzima GSH, que pode servir como sistema
20 de desintoxicação, removendo a cisplatina intracelular. Por fim, sabe-se que a
21 exposição contínua a cisplatina também é capaz de promover anormalidades nas
22 diferentes vias ativadas, induzindo capacidade antiapoptótica e quimiorresistência,
23 assim, a inativação da via MAPK, por exemplo, pode favorecer a sobrevivência celular
24 (CHEN; CHANG, 2019).

25 A quimioterapia a base de CDDP também é reconhecida por induzir
26 uma série de efeitos adversos. Dentre os principais efeitos consiste a nefrotoxicidade,
27 desencadeando lesão renal aguda e perda de função, devido ao seu acúmulo em
28 células tubulares, inflamação e lesão vascular. A ototoxicidade é outro efeito bem
29 conhecido induzido por cisplatina, no qual leva a perda auditiva neurossensorial
30 bilateral, progressiva e irreversível, muitas vezes associada a vertigem e zumbido.
31 Isso ocorre devido ao estresse oxidativo, inflamação e morte celular das células
32 presentes no ouvido interno (GHOSH, 2019).

1 **1.5 PRODUTOS NATURAIS COMO ALTERNATIVA TERAPÊUTICA PARA O CÂNCER DE** 2 **PULMÃO**

3 Ao longo da história, os produtos naturais desempenharam um papel
4 dominante no tratamento de doenças humanas. Por exemplo, a lendária descoberta
5 da penicilina por Alexander Fleming, em 1928, transformou a existência global, pois
6 antes da sua introdução não existia tratamento eficaz para diversas infecções. Outro
7 medicamento, o ácido salicílico, que foi derivado de uma planta medicinal, deu origem
8 ao ácido acetilsalicílico (AAS; conhecido como aspirina) em 1897, através do trabalho
9 de Felix Hoffman. O AAS foi considerado uma "droga milagrosa" por suas notáveis
10 propriedades medicinais (SNEADER, 2000; MINER; HOFFHINES, 2007; AMERICAN
11 CHEMICAL SOCIETY, 2023).

12 Atualmente, os produtos naturais constituem uma grande parte dos
13 agentes farmacêuticos atuais e são amplamente reconhecidos como fontes
14 abundantes de compostos bioativos com potencial terapêutico. Nos últimos anos, tem
15 havido esforços significativos para isolar novos produtos naturais de microrganismos
16 e plantas terrestres e aquáticas a fim de avaliar suas propriedades biológicas em
17 várias doenças, incluindo o câncer (ATANASOV et al., 2021; HASHEM et al., 2022;
18 HUANG; LU; DING, 2021). Pesquisas recentes revelaram que 32% de todos os
19 medicamentos de moléculas pequenas aprovados entre janeiro de 1981 e setembro
20 de 2019 eram produtos naturais e seus derivados (NEWMAN; CRAGG, 2020).

21 Uma das fontes mais importantes de compostos biologicamente ativos
22 é o reino vegetal. Atualmente, existem mais de 350.000 espécies de plantas
23 vasculares registradas no mundo, e novas espécies são adicionadas a cada ano.

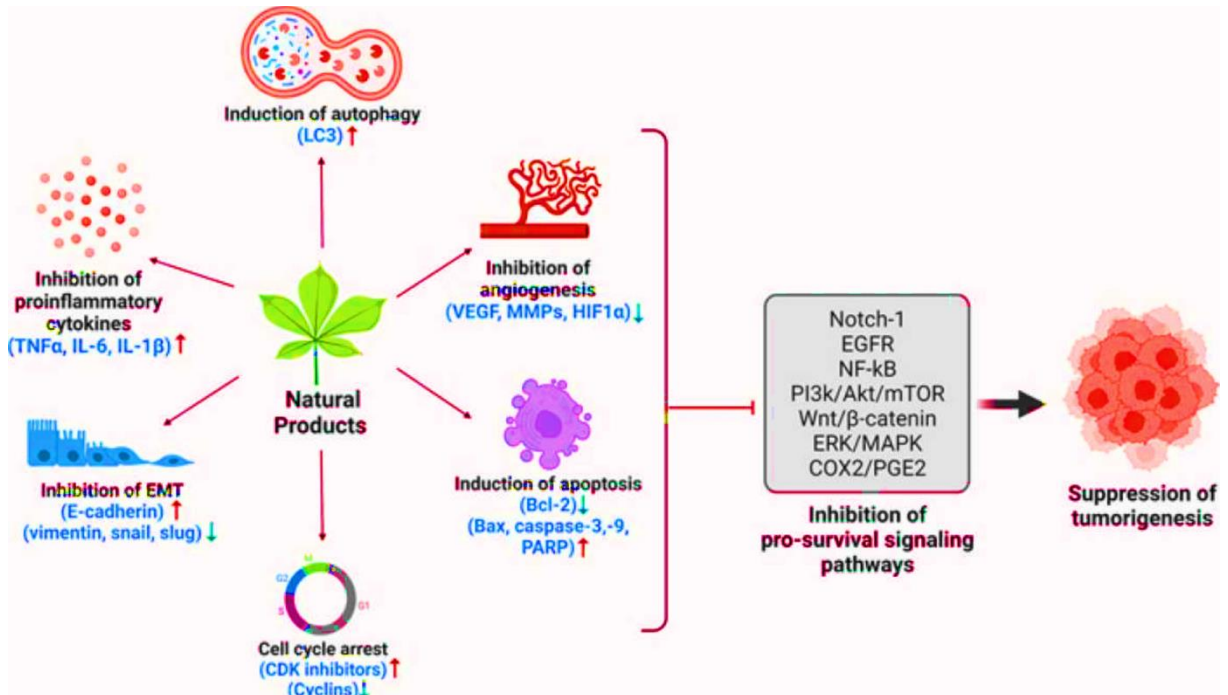
24 As plantas podem ser usadas terapêuticamente em diversas formas,
25 como chá, onde várias partes das plantas, incluindo folhas, flores, caules e raízes, são
26 mergulhadas em água quente para criar infusões que podem ser consumidas como
27 bebidas para aliviar doenças comuns, como problemas digestivos, estresse, insônia e
28 até mesmo por fornecer efeitos antioxidantes e anti-inflamatórios; corantes, os quais
29 podem ser frequentemente derivados de raízes, folhas ou frutos de certas plantas e
30 têm sido usados para diversas aplicações terapêuticas e cosméticas, como no caso
31 da cúrcuma, um corante amarelo-laranja, usada na medicina por suas propriedades
32 anti-inflamatórias e antioxidantes e henna, usada na coloração capilar, esmaltes e
33 maquiagens; e extratos, onde as plantas são processadas para extrair compostos
34 bioativos específicos (BRUDZYŃSKA; SIONKOWSKA; GRISEL, 2021; SINGH, Tripti

1 et al., 2023). Além disso, seus compostos ativos podem ser isolados e utilizados como
2 medicamentos ou como precursores de drogas sintéticas e semissintéticas. Existe
3 uma grande lista de fitoquímicos (ou seja, compostos químicos produzidos por
4 plantas) com atividade terapêutica, incluindo alcaloides, óleos essenciais, flavonoides,
5 carotenoides, terpenos e compostos fenólicos. Esses compostos têm sido objeto de
6 investigação extensa devido à sua segurança, à acessibilidade, à aplicabilidade e à
7 redução da citotoxicidade (HASHM et al., 2022).

8 De maneira geral, esses compostos podem ser usados para
9 quimioprevenção e quimioterapia, pois atingem várias vias de sinalização, impactando
10 assim a atividade molecular das células, interferindo na iniciação, desenvolvimento e
11 progressão do câncer e modulando diversos mecanismos. Entre os mecanismos
12 principais encontrados, tem-se a supressão da proliferação celular, regulação do ciclo
13 e interferência em diversas vias de sinalização tumorigênica, como fosfoinositídeo 3-
14 quinase (PI3K), metaloproteinase de matriz (MMP), MAPK/ERK conhecida como a via
15 Ras-Raf-MEK-ERK, via do receptor toll-like (TLR) e via AKT. Além disso, os produtos
16 naturais podem estimular mecanismos de reparo do DNA através da ação dos
17 produtos dos genes p21, p27, p51 e/ou p53, como as proteínas Bax, Bak e Bid, que
18 causam a síntese de enzimas protetoras, como as caspases 3, 7, 8, 9, 10, 12 e modula
19 enzimas antioxidantes como GST, GSH e GPx, bem como apoptose, angiogênese e
20 metástase (NAEEM et al., 2022). Consequentemente, eles podem ser usados como
21 potenciais adjuvantes na terapia do câncer (**Figure 8**).

22
23 **Figura 8:** Importância dos produtos naturais na inibição das várias vias de sinalização
24 que servem como impulsionadores da carcinogênese.

25



1
2 **Fonte:** HASHEM et al. (2022).

3
4 De acordo com a Food and Drug Administration (FDA), foram
5 aprovados cerca de 300 medicamentos quimioterápicos para o tratamento do câncer,
6 incluindo taxol, vinca e seus derivados, análogos de platina, 5-fluorouracil,
7 bevacizumabe, erlotinibe, nivolumabe, ipilimumabe, sunitinibe e olaparibe, com outras
8 combinações diferentes de medicamentos quimioterápicos (PATRIDGE et al., 2016).
9 No entanto, a maioria deles é altamente tóxica e induz resistência, o que acaba
10 resultando em recorrência tumoral e metástase, limitando suas atividades
11 anticancerígenas devido ao desenvolvimento de resistência adaptativa por células
12 tumorais direcionadas e toxicidades não específicas para células normais (NAEEM et
13 al., 2022). Portanto, são necessárias alternativas melhores, mais seguras e mais
14 eficazes para ter uma atividade anticancerígena mais proeminente e superar a
15 quimiorresistência encontrada no câncer de pulmão.

16 17 **1.5.1 3,3',5,5'-TETRAMETOXIBIFENIL-4,4'-DIOL (TMBP)**

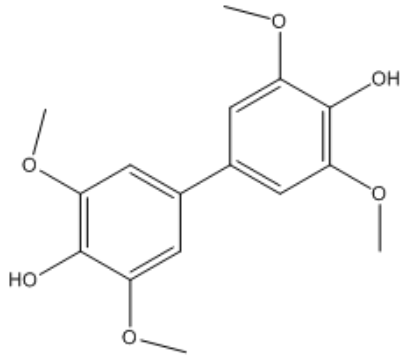
18 Compostos fenólicos são encontrados naturalmente em plantas, mas
19 a obtenção de compostos dessa classe também pode ser alcançada por diferentes
20 metodologias de sínteses. A biocatálise é uma metodologia baseada no uso de
21 enzimas para a transformação estrutural de moléculas. Neste processo ocorre a
22 síntese de moléculas estruturalmente diversas através da utilização de lacases.

1 As lacases são enzimas polifenol oxidases, que podem ser obtidas a
2 partir de fungos, plantas, insetos e bactérias (CANNATELLI; RAGAUSKAS, 2017;
3 WAN; DU; MIYAKOSHI, 2008). Estas enzimas oxidam uma extensa variedade de
4 compostos fenólicos, promovendo a formação de derivados que podem apresentar
5 propriedades bioativas potencializadas em relação aos substratos de origem
6 (CANNATELLI; RAGAUSKAS, 2017; MOGHARABI; FARAMARZI, 2014). A oxidação
7 do ácido ferúlico catalisada por lacase levou a formação de derivados fenólicos com
8 maior atividade antioxidante (ADELAKUN et al., 2012). Na oxidação do 2,6-
9 dimetoxifenol catalisada por lacases, o composto bifenólico formado, TMBP,
10 apresentou potencial antioxidante superior ao substrato (ADELAKUN et al., 2012;
11 SCHIRMANN et al., 2019).

12 Os compostos fenólicos têm sido extensivamente estudados quanto
13 às suas atividades biológicas, inclusive como agentes anticâncer (ABOTALEB et al.,
14 2020). Nesse sentido, os compostos bifenólicos, constituídos por dois anéis de
15 benzeno unidos ligados através de suas posições 1,1', são considerados um
16 importante intermediário na química orgânica que constitui a porção estrutural de
17 muitos compostos apresentando diversas atividades farmacológicas, como
18 antiinflamatória, diurética, anti-diabética e antimicrobiana (JAIN; GIDE; KANKATE,
19 2017).

20 O bifenil, 3,3',5,5'-tetrametoxibifenil-4,4'-diol (TMBP), é um composto
21 fenólico, que foi obtido através da oxidação do 2,6-dimetoxifenol (do inglês, 2,6-DMP)
22 utilizando a enzima lacase produzida pelo fungo ascomicetos *Botryosphaeria rhodina*
23 MAMB-05 (SCHIRMANN et al., 2018, 2019). Do ponto de vista estrutural, o TMBP é
24 composto por dois anéis aromáticos, duas hidroxilas e quatro grupos metoxilas
25 (RIZWANUL FATTAH et al., 2014), conferindo à molécula propriedades antioxidantes
26 (**Figura 9**) (FOTI et al., 2005; RICE-EVANS; MILLER; PAGANGA, 1996; SCHIRMANN
27 et al., 2018, 2019; SOARES, 2002). O TMBP foi descrito como um antioxidante com
28 potencial para inibir processos oxidativos e influência estabilizadora no biodiesel de
29 soja (FOTI et al., 2005; SCHIRMANN et al., 2018, 2019).

30 **Figura 9:** Estrutura química bidimensional do 3,3',5,5'-tetrametoxibifenil-4,4'-diol
31 (TMBP).



1

2 **Fonte:** O próprio autor.

3 No estudo realizado por nosso grupo de pesquisa foi relatado que o
4 tratamento com TMBP sobre a linhagem celular A549 de câncer de pulmão foi capaz
5 de reduzir a viabilidade celular, por ação citotóxica e promover a redução do tamanho
6 celular. Posteriormente, descobrimos que o tratamento com TMBP aumentou ROS,
7 causando despolarização mitocondrial e aumento de gotículas lipídicas. O tratamento
8 também foi capaz de promover a parada do ciclo celular na fase G2/M e induzir a
9 morte por apoptose direta. Além disso, o tratamento com TMBP sobre as células
10 primárias, como macrófagos peritoneais e eritrócitos de ovelhas não causou
11 toxicidade. Assim, este estudo revelou que o TMBP poderia ser um candidato
12 promissor para o desenvolvimento de medicamentos antitumorais direcionados ao
13 câncer de pulmão (Concato et al. 2020).

14 No trabalho de Schirmann et al. (2023), os autores mostraram pela
15 primeira vez a atividade biológica do TMBP contra as formas promastigotas e
16 amastigotas de *L. amazonensis* utilizando diferentes abordagens experimentais. O
17 tratamento com TMBP inibiu a proliferação e induziu a morte de formas promastigotas
18 gerando ERO e disfunção mitocondrial. Nas amastigotas intracelulares, o TMBP
19 reduziu a porcentagem de macrófagos infectados, sendo 62,7 vezes mais seletivo ao
20 parasita. O TMBP não hemolisou eritrócitos de ovelhas; indicativo de baixa
21 citotoxicidade. Além disso, análise de docking molecular em dois alvos enzimáticos
22 de *L. amazonensis*: tripanotiona redutase (TR) e leishmanolisina (Gp63), sugeriram
23 que o grupo hidroxila poderia ser um grupo farmacofórico devido à sua afinidade de
24 ligação por ligações de hidrogênio com resíduos no sítio ativo de ambas as enzimas.
25 O TMBP foi mais seletivo ao alvo Gp63 do que o TR. Dessa forma, torna-se promissor
26 para atuar como agente anti-*Leishmania*.

1 Os efeitos antiproliferativos e indutores de apoptose de outro
2 composto bifenílico, viz., 2,4,3',4'-tetrametoxibifenil, também relataram apoptose
3 agindo através de uma via mitocondrial/caspase em células MGC-803 de câncer
4 gástrico humano (SUI et al., 2014).

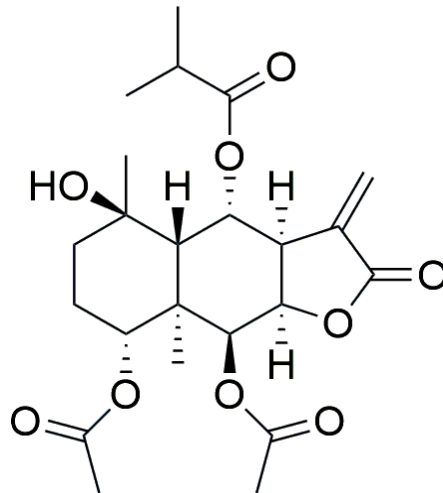
6 1.5.2 TRILOBOLIDO-6-O-ISOBUTIRATO (TBB)

7 As lactonas sesquiterpênicas (LS) são um grupo grande e
8 diversificado de produtos naturais presentes em mais de 100 famílias de plantas. Entre
9 eles o maior número de SLs é derivado da família Compositae (Asteraceae), com mais
10 de 3.000 estruturas relatadas. Uma das plantas medicinais importantes desta família,
11 é a *Sphagneticola trilobata* (L.) J.F. Pruski (HUANG; JIANG; OOI, 2003; ZHANG et
12 al., 2016).

13 *S. trilobata*, também conhecida popularmente como “pseudo-arnica”,
14 “margaridão”, “pingo-de-ouro”, “picão-da-praia” ou apenas “Wedelia”, pertence à
15 família de plantas Asteraceae e tem sido um recurso medicinal tradicional na Índia,
16 China, Caribe e Américas Central e do Sul (MARDINA et al., 2020). Apesar da sua
17 classificação como erva daninha invasora, esta planta medicinal destaca-se pela sua
18 elevada diversidade molecular e pela sua diversificada atividade biológica que
19 engloba antibacteriana (SHANKAR; THOMAS, 2014), antifúngica (HUSAIN;
20 SRIVASTAVA, 2018), antiplasmodial (SRAVYA et al., 2011), atividades antidiabética,
21 hepatoprotetora, antipirético-analgésica e anticancerígena (MARDINA et al., 2020).

22 Os principais constituintes químicos de *S. trilobata* são os diterpenos
23 ent-kaurano, lactonas sesquiterpênicas eudesmane e triterpenos (HUI et al., 2019).
24 Investigações recentes demonstraram o potencial terapêutico das lactonas
25 sesquiterpênicas (SLs), especialmente no domínio da pesquisa do câncer (DHYANI
26 et al., 2022; DONG; QIAN; LI, 2022; SHOAIB et al., 2017). Entre estes está o
27 trilobolido-6-O-isobutirato (TBB), fórmula molecular $C_{23}H_{32}O_9$ (**Figura 10**), uma
28 notável lactona sesquiterpênica natural extraída de *S. trilobata* (Bohlmann et al.,
29 1981). O TBB é construído a partir da fusão de dois anéis ciclohexano e um anel de
30 lactona. Este último anel contém oxigênio com função carbonila, e é o grupo mais
31 responsável por efeitos biológicos, como tratamento de tumor, e redução da pressão
32 arterial em humanos (FERREIRA et al., 1994; HUANG; JIANG; OOI, 2003).

33 **Figura 10.** Representação bidimensional da estrutura química do trilobolido-6-O-
34 isobutirato (TBB).



1

2 **Fonte:** O próprio autor.

3

4 A lipofilicidade também foi considerada um aspecto importante na
 5 atividade das LS, potencialmente devido à maior permeabilidade da membrana,
 6 permitindo que os sesquiterpenos atinjam células e núcleos onde exercem seus
 7 principais efeitos. Outras funções biológicas de grande relevância, como
 8 antimicrobiana, anti-herbicidas, antioxidante, antimalárica, imunomodulador e até
 9 mesmo anti-metástase são também descritas (CHADWICK et al., 2013; GHANTOUS
 10 et al., 2010; MERFORT, 2011; TOPPO et al., 2013)

11 No trabalho de Guzman et al., (2005) observaram que as LS
 12 partenólídeo induziu apoptose em células estaminais e progenitoras de leucemia
 13 mielóide aguda humana. Uma das principais razões para isto é que LS tem preferência
 14 por Fe (II), o qual é sequestrado em altas concentrações em células cancerígenas e
 15 o partenólídeo é clivado a oxidantes citotóxicos em contato com Fe (II) que danificam
 16 as células tumorais próximas. Além disso, foi constatado que o partenólídeo
 17 sensibiliza as células tumorais a medicamentos contra o câncer, como tamoxifeno
 18 paclitaxel, além do TNF e, conseqüentemente, promove a morte celular quando
 19 aplicado em níveis baixos o suficiente para não ser tóxico (CHADWICK et al., 2013;
 20 REN; YU; KINGHORN, 2016.)

21 LS ainda possuem a capacidade de inibir a expressão de NF-κB,
 22 proteína que medeia a resposta imune em humanos, controlando a resposta de outros
 23 efetores, como citocinas, moléculas inflamatórias e moléculas de adesão celular, além
 24 de regular mais de 150 genes inflamatórios, sendo alguns responsáveis pela
 25 proliferação celular (CHADWICK et al., 2013; GHANTOUS et al., 2010; MERFORT,

1 2011)

2 Em relação ao TBB, uma recente investigação *in vitro* utilizando TBB,
3 isolado de *S. trilobata*, contra as linhagens de células tumorais de hepatocarcinoma
4 humano: HuH7, HepG2, SK-Hep1 e Bel7402, descobriu-se que o tratamento inibiu o
5 crescimento e a formação de colônias em um tempo- e forma dependente da
6 concentração. O tratamento com TBB exibiu efeitos inibitórios sobre a migração,
7 invasão e glicólise, culminando em última análise na paragem do ciclo celular na fase
8 G2 / M, disfunção mitocondrial e morte celular induzida por apoptose. Além disso, o
9 TBB também foi capaz de regular a expressão de genes essenciais associados à
10 proliferação, migração e metástase como PCNA (antígeno nuclear de células em
11 proliferação), Ki67, Ciclina B1, Ciclina E, Bax, Bcl2, MMP2/9 (metaloproteinase 2/9) e
12 PGK1 (fosfoglicerato quinase 1), principalmente através da inibição dos transdutores
13 de interleucina-6/sinal e ativadores da via de sinalização da transcrição-3 (IL-
14 6/STAT3). A eficácia *in vivo* dos atributos anticancerígenos do TBB foi validada
15 através de experimentação com um modelo de tumor xenoenxertado em
16 camundongos nus BALB/c machos, onde o TBB foi eficaz na eliminação do
17 crescimento tumoral (peso e volume) sem causar toxicidade aos animais
18 experimentais (ZHOU et al., 2021).

19

20 **1.6 MODELO EXPERIMENTAL *IN VITRO***

21 Um modelo experimental *in vitro* refere-se a um ambiente controlado
22 e artificial onde experimentos são conduzidos usando células, tecidos ou
23 componentes biológicos isolados fora de seu contexto natural, normalmente em um
24 ambiente de laboratório. O termo "*in vitro*" significa "*em vidro*" em latim, refletindo o
25 uso histórico de recipientes de vidro para tais experimentos (KATT et al., 2016;
26 COSTA et al., 2020; MARTINEZ-PACHECO; O'DRISCOLL, 2021).

27

28 Os modelos *in vitro* são representações simplificadas de processos
29 biológicos em comparação com modelos *in vivo* (em um organismo vivo). Os
30 pesquisadores podem isolar elementos específicos e focar em aspectos específicos
31 de um sistema biológico, como sinalização celular, expressão genética ou testes de
32 drogas. Neste modelo experimental, as variáveis externas são minimizadas ou
33 controladas, podem ser facilmente reproduzidos, aumentando a confiabilidade e
34 validade dos resultados. Além disso, são comumente usados em pesquisas
farmacêuticas para testar a segurança e eficácia de novos medicamentos ou terapias,

1 o que permite a triagem inicial antes de avançar para estudos *in vivo* de maior
2 complexidade, além de ser crucial na investigação do câncer para decifrar
3 mecanismos moleculares de eventos-chave, como crescimento tumoral, metástase,
4 resistência a medicamentos e aspectos de evasão imunológica (KATT et al., 2016;
5 COSTA et al., 2020; MARTINEZ-PACHECO; O'DRISCOLL, 2021).

6 No entanto, este modelo experimental possui algumas desvantagens.
7 Apresentam limitações por não replicarem totalmente a complexidade dos sistemas
8 biológicos nos organismos vivos e os resultados precisam ser validados em modelos
9 *in vivo* para garantir a sua relevância para situações da vida real (KATT et al., 2016;
10 COSTA et al., 2020; MARTINEZ-PACHECO; O'DRISCOLL, 2021).

12 **2 JUSTIFICATIVA**

13 A atual quimioterapia para o câncer não é satisfatória devido à alta
14 toxicidade dos fármacos utilizados, tais como toxicidade gastrointestinal,
15 nefrotoxicidade, neurotoxicidade, ototoxicidade, hipersensibilidade e
16 mielossupressão. Além disso, os pacientes continuam a apresentar recorrência da
17 doença e resistência aos medicamentos durante o curso da terapia à base de platina,
18 o que reduz a eficácia do tratamento do NSCLC. Sabendo da importância desta
19 doença e de suas complicações anatomofisiológicas, a busca por novas alternativas
20 para o tratamento do câncer é necessária.

21 Uma série de estudos tem demonstrado o efeito anticâncer de
22 compostos isolados naturais e sintéticos e seus múltiplos mecanismos de ação em
23 experimentos *in vitro* e *in vivo*. Esses compostos são capazes de atuar nas células
24 tumorais através de vias importantes relacionadas a proliferação, invasão, metástase
25 e de morte celular.

26 Esses achados nos encorajaram a investigar o potencial anticâncer
27 do bifenil TMBP e da lactona sesquiterpênica TBB, por diferentes mecanismos de
28 ação, sobre linhagens tumorais de câncer de pulmão (A549 e NCI-H460) que ainda
29 não foram relatados na literatura.

31 **3 OBJETIVOS**

32 **3.1 Objetivo Geral**

33 Avaliar *in vitro* a atividade antiproliferativa dos compostos 3,3',5,5'-
34 tetrametoxibifenil-4,4'-diol (TMBP) e trilobolido-6-O-isobutirato (TBB) sobre as

1 linhagens tumorais de câncer de pulmão, A549 (adenocarcinoma pulmonar) e NCI-
2 H460 (carcinoma de células grandes), bem como elucidar os principais mecanismos
3 de ação e propriedades metabólicas envolvidas.

4 5 **3.1.1 Objetivos específicos**

- 6
- 7 ● Efetuar o estudo *in silico* da estrutura de TBB quanto às regras de Lipinski
- 8 e Veber e das propriedades de ADME;
- 9 ● Avaliar *in vitro* a viabilidade do TBB sobre as linhagens epiteliais, monocitos
- 10 e macrófagos alveolares;
- 11 ● Avaliar *in vitro* a viabilidade hemolítica do TMBP em eritrócitos extraídos de
- 12 carneiro;
- 13 ● Determinar as concentrações de TBB e TMBP capazes de inibir o
- 14 crescimento das linhagens tumorais A549 e NCI-H460;
- 15 ● Determinar a concentração inibitória (IC₅₀) das linhagens normais e
- 16 tumorais de TBB e TMBP e índice de seletividade tumoral para TBB;
- 17 ● Analisar os aspectos morfológicos das células tumorais após o tratamento
- 18 com TBB e TMBP;
- 19 ● Avaliar parâmetros relacionados ao perfil redox e metabólico das células
- 20 tumorais após o tratamento com TBB e TMBP;
- 21 ● Avaliar se o tratamento com TBB e TMBP alteram a capacidade migratória
- 22 das células tumorais;
- 23 ● Analisar se o TMBP é capaz de interferir na regulação do ciclo celular e
- 24 caracterizar os mecanismos de morte envolvidos durante a ação dos
- 25 tratamentos com TBB e TMBP sob as células tumorais;
- 26 ● Analisar as proteínas PI3K, AKT, Pakt, ARG-1, iNOS, NF-Kb, pNF-Kb e
- 27 Nrf2 após o tratamento com TMBP;
- 28

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4 PRODUÇÃO CIENTÍFICA

4.1 Manuscrito: 3,3',5,5'-Tetramethoxybiphenyl-4,4'diol triggers oxidative stress, metabolic changes and apoptosis-like process by reducing the PI3K/AKT/NF- κ B pathway in the NCI-H460 lung cancer cell line

Aceito na revista: *Biomedicine & pharmacological*

4.2 Manuscrito: Trilobolide-6-O-isobutyrate from *Sphagneticola trilobata* acts by inducing oxidative stress, metabolic changes and apoptosis-like processes by caspase 3/7 activation of human lung cancer cell lines

Aceito na revista: *Phytomedicine*

33 **3,3',5,5'-Tetramethoxybiphenyl-4,4'diol triggers oxidative stress, metabolic**
34 **changes, and apoptosis-like process by reducing the PI3K/AKT/NF- κ B**
35 **pathway in the NCI-H460 lung cancer cell line**

36

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61 **Highlights**

- 62 • TMBP treatment reduces cell viability and alters morphology in NCI-H460.
- 63 • The treatment reduces migratory capacity and N-cadherin and β -catenin proteins.
- 64 • TMBP treatment promotes oxidative and metabolic stress on lung cancer cell-line.
- 65 • TMBP induces the G2/M arrest phase, cell death, and reduces STAT-3 in NCI-H460.
- 66 • The treatment act in NCI-H460 decreases PI3K/AKT/ARG-1/NF- κ B and increases
- 67 iNOS.

68

69 **Abstract**

70 Lung cancer is one of the leading causes of cancer-related deaths in men and women worldwide.
71 Current treatments have limited efficacy, cause significant side effects, and cells can develop
72 drug resistance. New therapeutic strategies are needed to discover alternative anticancer agents
73 with high efficacy and low-toxicity. TMBP, a biphenyl obtained by laccase-biotransformation
74 of 2,6-dimethoxyphenol, possesses antitumor activity against A549 adenocarcinoma cells.
75 Without causing damage to sheep erythrocytes and mouse peritoneal macrophages of BALB/c
76 mice. In addition to being classified as a good oral drug according to *in-silico* studies. This
77 study evaluated the *in-vitro* cytotoxic effect of TMBP on lung-cancer cell-line NCI-H460 and
78 reports mechanisms on immunomodulation and cell death. TMBP treatment (12.5-200 μ M)
79 inhibited cell proliferation at 24, 48, and 72 h. After 24-h treatment, TMBP at IC₅₀ (154 μ M)
80 induced various morphological and ultrastructural changes in NCI-H460, reduced migration
81 and immunofluorescence staining of N-cadherin and β -catenin, induced increased reactive
82 oxygen species and nitric oxide with reduced superoxide radical-anion, increased superoxide
83 dismutase activity and reduced glutathione reductase. Treatment also caused metabolic stress,
84 reduced glucose-uptake, intracellular lactate dehydrogenase and lactate levels, mitochondrial
85 depolarization, increased lipid droplets, and autophagic vacuoles. TMBP induced cell-cycle
86 arrest in the G2/M phase, death by apoptosis, increased caspase-3/7, and reduced STAT-3
87 immunofluorescence staining. The anticancer effect was accompanied by decreasing PI3K,
88 AKT, ARG-1, and NF- κ B levels, and increasing iNOS. These results suggest its potential as a
89 candidate for use in future lung anticancer drug design studies.

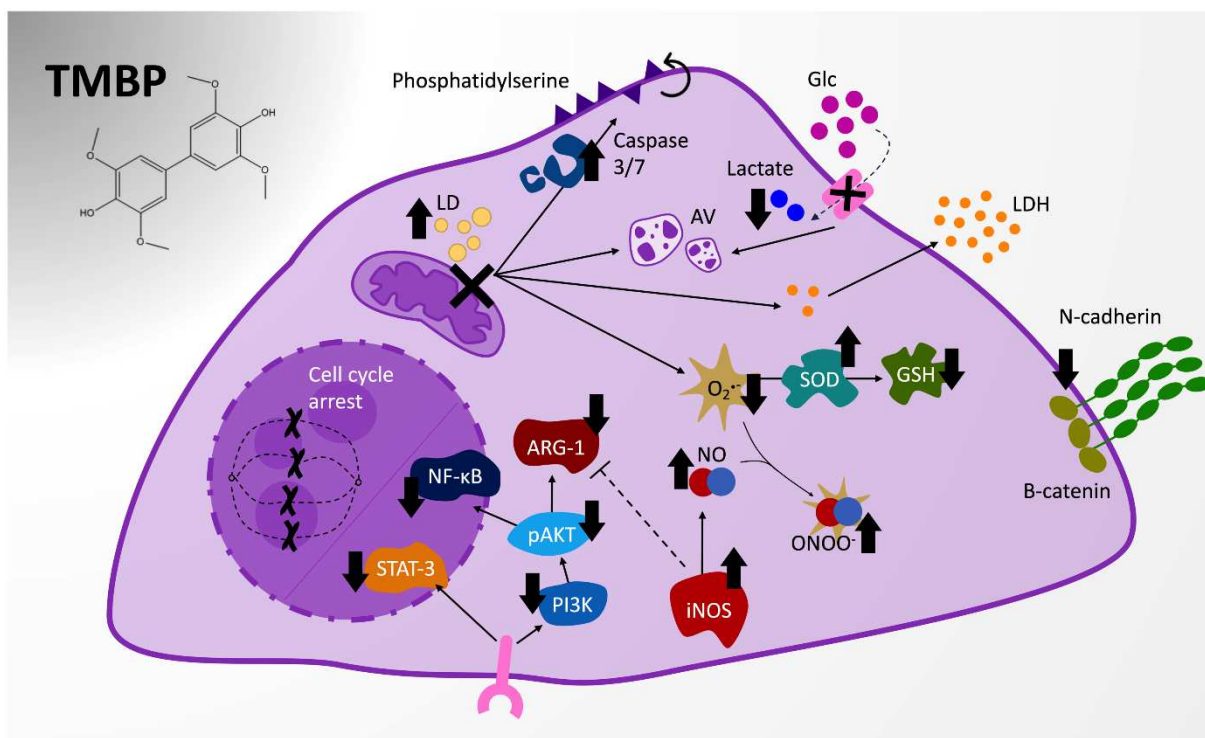
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92

93 **Keywords:** Treatment; Cancer; Cell Death; Redox Disruption;

94 Graphical Abstract



95 **Schematic model of the lung anticancer effect of TMBP on the NCI-H460 large cell**
 96 **carcinoma cell line.** Caption: TMBP, 3,3',5,5'-Tetramethoxybiphenyl-4,4'-Diol; ROS,
 97 Reactive oxygen species; NO, Nitric oxide; O₂^{•-}, Superoxide anion radical; SOD, Superoxide
 98 dismutase; GSH, Reduced glutathione; Glc, Glucose; LDH, Lactate dehydrogenase; LD, Lipid
 99 droplets; LD, Lipid droplets; AV, Autophagic vacuoles; iNOS, Induzivel nitric oxide synthase;
 100 ARG-1, Arginase-1; PI3K, phosphatidylinositol-3-kinase; AKT, Protein kinase B, pAKT,
 101 Phosphoprotein kinase B; NF-κB, Nuclear factor *kappa* B; STAT-3, signal transducer and
 102 activator of transcription 3; ONOO⁻, Peroxynitrite anion;

103

104 **Abbreviations:** LC, Lung Cancer; NSCLC- Non-small cell lung cancer; SCLC- Small cell lung
 105 cancer; TMBP, 3,3',5,5'-Tetramethoxybiphenyl-4,4'-Diol; 2,6-DMP, 2,6-Dimethoxyphenol;
 106 ROS, Reactive Oxygen Species; DMSO, Dimethyl sulfoxide; SEM, Scanning Electron
 107 Microscopy; TEM, Transmission Electron Microscopy; H₂DCFDA, 2',7'-dichlorofluorescein
 108 diacetate; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; PBS,
 109 Phosphate-buffered saline; SBF, fetal bovine serum; Glc, Glucose; TMRE,
 110 tetramethylrhodamine-ethyl-ester; H₂O₂, Hydrogen peroxide; CCCP, Carbonylcyanide *m*-
 111 chlorophenylhydrazone; NAC, N-Acetylcysteine; NO, Nitric Oxide; L-NAME, NG-nitro-L-
 112 arginine methyl ester; O₂^{•-}, Superoxide anion radical; SOD, Superoxide dismutase; GSH,
 113 Reduced Glutathione; LDH, Lactate dehydrogenase; Δψ_m, Mitochondrial membrane potential;

114 LD, Lipid droplets; NR, Nile red; MDC, monodansylcadaverine; CPT, Camptothecin; ELISA,
115 Enzyme-linked immunosorbent assay; iNOS, Induzivel nitric oxide sintase; ARG-1, Arginase-
116 1; PI3K, phosphatidylinositol-3-kinase; AKT, Protein kinase B, pAKT, Phospho Protein kinase
117 B; NF- κ B, Nuclear factor *kappa* B; p NF- κ B, Phospho nuclear factor *kappa* B; NRF2,
118 Erythroid-derived nuclear factor 2; KOH, Potassium hydroxide; Z-DEVD-AMC substrate,
119 Fluorogenic substrate for caspase 3 e 7; Ac-DEVD-CHO inhibitor, Caspase-3 e 7 Inhibitor I,
120 N-Ac-Asp-Glu-Val-Asp-CHO; RIPA, Radioimmunoprecipitation Assay; EDTA,
121 Ethylenediaminetetraacetic acid; TME, tumor microenvironment; STAT-3, signal transducer
122 and activator of transcription 3; EMT, epithelial-mesenchymal transition; OH \cdot , Hydroxyl
123 radical; OXPHOS, Oxidative phosphorylation; CAT, catalase; PRDXs, Peroxiredoxins; GPXs,
124 Glutathione peroxidases; ONOO $^-$, Peroxynitrite anion; RNS, Reactive Nitrogen Species;

125 **1. Background**

126 Lung cancer (LC), a heterogeneous disease, remains one of the leading causes of cancer-
127 related mortalities worldwide, with an estimated 2.9 million cases by 2040 [1]. In addition, most
128 patients have metastatic disease at the time of diagnosis [1]. Non-small cell lung cancer
129 (NSCLC) accounts for more than 85% of all LCs and is common in both men and women [2].
130 Available chemotherapy for NSCLC has advanced significantly over the past decade, and
131 platinum-based doublet therapy (e.g., cisplatin in combination with carboplatin) has been
132 successfully implemented [3]. Several side effects, however, remain, including severe kidney
133 problems, allergic reactions, decreased immunity to infection, gastrointestinal upset, bleeding,
134 hearing loss, and also drug resistance, especially if the tumor recurs [4]. Because of the
135 challenges in treating this disease, new therapeutic strategies are needed and should be
136 developed.

137 Phenolic compounds are naturally present in plants and can be synthesized through
138 various methods, including biocatalysis, which involves the use of enzymes for the
139 transformation of molecular structures [5,6]. One of the advantages of synthesizing compounds
140 through biocatalysis is that it can be part of a green process, as it employs fewer toxic
141 methodologies. This has been highlighted in research conducted by [7–9]. 3,3',5,5'-
142 Tetramethoxybiphenyl-4,4'-diol (TMBP), a biphenyl (**Figure suppl. 1**), was obtained by
143 oxidation of 2,6-dimethoxyphenol (2,6-DMP) using laccase produced by the ascomyceteous
144 fungus *Botryosphaeria rhodina* (MAMB-05) [7,8]. Biphenolic compounds have interesting
145 chemical structures and many of these compounds play diverse biological roles [5,6], including
146 fungicidal [10,11], bactericidal [12,13], anti-inflammatory [14,15], antioxidant [16], cell anti-

147 proliferation [17,18], and anti-tumor activity. In a recent study using TMBP on the A549 (lung
148 adenocarcinoma) tumor cell line, our group showed that treatment with TMBP reduced
149 viability, increased reactive oxygen species (ROS) production, induced mitochondrial
150 depolarization, was involved in cell cycle arrest in the G2/M phase, and death by direct
151 apoptosis, without causing toxicity in BALB/c mouse peritoneal macrophages and sheep
152 erythrocytes [19].

153 In this study, using different endpoints, TMBP was shown to reduce cell viability and
154 migratory potential, promote morphological and ultrastructural changes, followed by apoptosis
155 and autophagy. These phenomena were mediated by an oxidative burst and metabolic damage
156 through decreased levels of PI3K/pAKT/ARG-1/pNF- κ B and an increase of the iNOS pathway
157 in NCI-H460 lung cancer cells.

158

159 **2. Material And Methods**

160

161 **2.1 TMBP**

162 The biphenyl 3,3',5,5'-tetramethoxybiphenyl-4,4'-diol (TMBP; C₁₆H₁₈O₆, MW 306) was
163 synthesized from 2,6-DMP using laccase from *Botryosphaeria rhodina* MAMB-05 under
164 conditions previously described [20]. The resulting yellow solid was identified by ¹H NMR
165 (CDCl₃, 400 MHz) at δ 3.98 (s, 12H), 6.73 (s, 4H), 5.56 (s, 2H), purity 99%, and melting point
166 185-188 °C. A stock solution of TMBP was prepared in 1% dimethyl sulfoxide (DMSO)
167 (Sigma-Aldrich, St. Louis, MO, USA); the concentration of DMSO was kept at a maximum of
168 0.06% in all experiments.

169

170 **2.2 Cell culture**

171 The established NSCLC line, NCI-H460 (large cell carcinoma), from Frederick Ma
172 (National Cancer Institute, Bethesda, MD, USA) was kindly provided by the Toxicological
173 Genetics Laboratory-UEL. The cell was cultured in Roswell Park Memorial Institute (RPMI)
174 1640 culture medium (Life Technologies, Carlsbad, CA, USA) supplemented with 10% fetal
175 bovine serum (FBS) (GIBCO, Invitrogen, New York, USA), 100 U/mL penicillin and 100
176 μ g/mL streptomycin (Santa Cruz Biotechnologies, Dallas, TX, USA), and incubated at 37 °C
177 under 5% CO₂.

178 **2.3 Cell viability assay**

179 NCI-H460 cells (10^4 cells/well) were treated with TMBP (range 12.5-100 μ M) or
180 cisplatin (Cis) (50 μ M), and cytotoxicity was assessed at 24, 48, and 72 h. MTT assays were
181 performed as previously described [19]. Briefly, the cells were washed, and MTT (0.33 mg/ml)
182 (Sigma-Aldrich) was added to the cells and then incubated for 3 h at 37 °C. The MTT product
183 formed (formazan crystals) was diluted with 100 μ l of DMSO and measured
184 spectrophotometrically at 550 nm. The 50% inhibitory concentration (IC_{50}) curve was
185 calculated by logarithmic regression using GraphPad Prism 8 software for the above times. The
186 IC_{50} at 24 h was used for the remaining experiments. Micrographs of TMBP-treated IC_{50} cells
187 were taken using an EVOS Microscope FL Auto Cell imaging system (Thermo-Fisher
188 Scientific, Multiskan GO, Waltham, MA, USA) at 100 \times magnification.

189

190 **2.4 Morphological and ultrastructural analysis of NCI-H460 cells by scanning electron 191 microscopy (SEM) and transmission electron microscopy (TEM)**

192 Scanning electron microscopy (SEM) of the NCI-H460 cell line was performed to
193 analyze morphological changes in the cell surface topography [21]. Briefly, cells (10^6
194 cells/well) were treated with IC_{50} -TMBP for 24 h at 37 °C. They were then fixed with 2.5%
195 glutaraldehyde in 0.1 M sodium cacodylate buffer (pH 7.4) and allowed to adhere to poly-L-
196 lysine-coated coverslips for 60 min. The cells were then dehydrated through increasing
197 concentrations of ethanol (30-100%), subjected to critical point drying (Baltec SCD-030,
198 Balzers, Liechtenstein), and metallized with gold for visualization using a high-resolution FEI
199 SCIOS (Oregon, USA) double beam electron microscope (Thermo Fisher Scientific,
200 Massachusetts, United States).

201 To assess ultrastructural changes in the TMBP-treated cells by TEM (JEOL USA JEM
202 1400, Tokyo, Japan, transmission electron microscope), cells were treated and fixed similarly
203 as described above. The samples were then postfixed with 1% OsO_4 , 0.8% potassium
204 ferrocyanide, and 10.0 mM $CaCl_2$ in 0.1 M sodium cacodylate buffer (pH 7.4) for 1 h at room
205 temperature and protected from light. The samples were then washed in 0.1 M sodium
206 cacodylate buffer and dehydrated in increasing concentrations of acetone (30-100%). The total
207 acetone content was gradually replaced by EPONTM epoxy resin (Electron Microscopy
208 Sciences, Hatfield, United States) by cell diffusion and polymerized at 60 °C for 72 h.
209 Nanoscale sections (60-70 nm) were sliced with an ultramicrotome (PowerTomer BMC -
210 Germany) and contrasted with 5% uranyl acetate and 2% lead citrate. Finally, the samples were

211 analyzed by TEM. The NCI-H460 cell line maintained in RPMI culture medium without TMBP
212 treatment was used as a negative control.

213

214 **2.5 Cell scratch assay**

215 NCI-H460 cells (10^6 cells/well) were seeded into 6-well plates and incubated (37 °C,
216 5% CO₂) to confluence. We then created a 'scratch' in the monolayer by gently passing a 200
217 µL pipette tip over the bottom of each well. This passage created a discontinuity in the
218 monolayer whose cells at the edge tended to migrate into the empty spaces. The cells were next
219 treated with IC₅₀-TMBP, and photomicrographs (40× objective) were taken at different times
220 (0, 12, and 24 h) using an EVOS FL Auto 2 microscope (Thermo Scientific, USA). Cell
221 migration was assessed as the free area (region without cells) measured using Image-Pro plus
222 program software (ImageJ), while the percentage decrease in the area was characterized by the
223 cell migration index. Assays were performed in triplicate from each group tested.

224

225 **2.6 Immunofluorescence (IF) staining**

226 NCI-H460 cells (10^5 cells/well) were seeded in 24-well plates containing coverslips and
227 treated with the 24 h IC₅₀ value of TMBP at 37 °C. The assay was performed as described by
228 Duplancic and Kero (2021) [22]. Briefly, background blocking was performed using Abcam
229 Protein Block (ab64226; Abcam, UK) for 25 min after the antigen retrieval. The following
230 primary and secondary antibodies used for IF staining (24 h at 4 °C) were from Santa Cruz
231 Biotechnology (USA): N-cadherin (1:1000), β-catenin (1:500), and STAT-3 (1:1500).
232 Secondary antibodies were used at dilution 1:2000: anti-mouse Texas red (RED) and anti-rabbit
233 Alexa Fluor 488 (GREEN). Cell nuclei were stained with diamidino-2-phenylindole (DAPI,
234 Sigma-Aldrich, USA). The slides were mounted with glycerol and kept protected from light
235 under a fluorescence microscope for later observation. The intensity and localization of the
236 immunoreactivity were analyzed with the Motic BA 410E fluorescence microscope and
237 MOTICAM ProS5 Plus (40× objective lens). The merge of DAPI and Texas Red images was
238 performed using the Image J software (NIH, USA).

239

240 **2.6 Analysis of Reactive Oxygen Species (ROS)**

241 ROS generation was assessed by the conversion of non-fluorescent H₂DCFDA to highly
242 fluorescent 2',7'-dichlorofluorescein (DCF) by intracellular free radicals. Briefly, ROS
243 scavenger (N-acetylcysteine – NAC, 5.0 M) was added, or not added, to the NCI-H460 cells 2
244 h before they received treatment with IC₅₀ of TMBP, and were incubated for 24 h at 37 °C.

245 After this period, the cells were washed, and H₂DCFDA solution (10.0 M) was added and left
246 in the dark for 45 min. Fluorescence intensity was quantified using a Glomax®
247 spectrophotometer (Promega, USA) (excitation: 488 nm; and emission: 530 nm). Untreated
248 cells were used as a negative control, and H₂O₂ (0.04%) was used as a positive control. Data
249 were normalized and expressed in arbitrary units.

250

251 **2.7 Determination of nitrite; an estimate of nitric oxide (NO) levels**

252 NO levels were determined based on the Griess method. NCI-H460 cells (10⁵ cells/well)
253 were treated, or not, with the non-specific iNOS inhibitor (L-NAME, NG-nitro-L-arginine
254 methyl ester, Sigma-Aldrich, USA) (25μM) for 1 h before treatment with IC₅₀ of TMBP and
255 incubated for 24 h. The supernatant of the samples was collected and analyzed according to
256 Cataneo et al., (2019) [23]. Briefly, 60 μL of supernatant was added to 60 μL of Griess reagent
257 (1% sulfanilamide and 0.1% N-(1-Naphthyl) ethylenediamine in 5% orthophosphoric acid
258 (H₃PO₄)), incubated for 10 min, protected from light, at room temperature, and read on a plate
259 reader (Thermo Scientific, Multiskan GO, USA) at 550 nm. For quantification, serial dilutions
260 of NaNO₂ were used. Untreated cells were used as a negative control.

261

262 **2.8 Resazurin assay**

263 To assess whether cells treated with IC₅₀ of TMBP would alter viability in the presence
264 or absence of the inhibitors NAC and L-NAME, the resazurin assay was performed as follows.
265 NCI-H460 cells (10⁵ cells/well) were seeded in 24-well plates and incubated (24 h at 37 °C).
266 After this period, NAC (5M) or L-NAME (50μM) was added 2 h and 1 h before, respectively,
267 with or without IC₅₀-TMBP treatment and incubated (24 h at 37 °C). Resazurin (60 μM) was
268 then added and left for 2 h, and the fluorescence intensity was quantified in a Glomax®
269 spectrophotometer (520 nm excitation and 580nm emission). Data were normalized and
270 expressed in arbitrary units.

271

272 **2.9 Superoxide anion production**

273 The production of superoxide anion radical (O₂^{•-}) was determined by the reduction of
274 the Redox dye nitro blue tetrazolium (1 mg/mL; NBT, Sigma-Aldrich). NCI-H460 cells (10⁴
275 cells/well) were treated with IC₅₀-TMBP, incubated for 24 h, washed with PBS, and
276 subsequently, 100 μL of NBT solution was added and incubated at 37 °C for 1 h. The
277 supernatant was removed, and the formazan precipitate was then solubilized by adding 60 μL
278 of 2.0 M KOH and 60 μL of DMSO. The optical density was measured spectrophotometrically

279 at 600 nm in a microplate reader (Multiskan GO Microplate Spectrophotometer, Thermo
280 Scientific, and Vantaa, Finland). Untreated cells were used as a negative control, and phorbol
281 myristate acetate (PMA) treatment was used as a positive control.

282

283 **2.10 Superoxide dismutase (SOD) activity**

284 SOD activity was determined based on the enzyme capacity to inhibit auto-oxidation of
285 pyrogallol, as described by Marklund and Marklund [24]. For the determination of SOD
286 activity, NCI-H460 cells (10^5 cells/well) were seeded, treated with the IC_{50} of TMBP, and
287 incubated for 24 h at 37 °C. Subsequently, the cells submitted a cycle of freezing at – 80 °C
288 and thawing at 37 °C for 30 min each. Total protein was determined on the cell lysate by the
289 Lowry method [25]. The data were normalized as mg protein/mL. Untreated cells were used as
290 a negative control.

291

292 **2.11 Reduced glutathione (GSH)**

293 Reduced L-glutathione (GSH) was measured to determine the intensity of oxidative
294 stress. NCI-H460 cells (10^5 cells/well) were seeded and treated with TMBP as described above.
295 After 24 h, GSH quantification was performed according to the method described by Rahman
296 et al., 2007 [26]. Concentrations were determined based on a standard curve constructed with
297 serial dilutions of GSH. Untreated cells were used as a negative control.

298

299 **2.12 Glucose assay**

300 To assess glucose uptake, NCI-H460 cells (10^5 cells/well) were seeded in 24-well plates,
301 treated with IC_{50} TMBP, and incubated for 24 h at 37 °C. The enzymatic-colorimetric glucose
302 assay was performed according to the manufacturer's kit instructions (Gold Analisa
303 Diagnostics, Minas Gerais, Brazil). Briefly, 1.0 μ l aliquots of the supernatant from the treated
304 or standard samples were transferred to a 96-well plate, followed by the addition of 100 μ l of
305 the glucose kit reagent, and the mixture incubated for 10 min at 37 °C. The reading was
306 performed in a preheated spectrophotometer (37 °C) at 505 nm. Untreated cells were used as a
307 negative control.

308

309 **2.13 Lactate dehydrogenase (LDH) Activity**

310 To evaluate the LDH activity, NCI-H460 cells (10^6 cells/well) were seeded in 6-well
311 microplates and treated with IC_{50} -TMBP for 24 h at 37 °C. After this period, the cell supernatant

312 was collected to measure extracellular LDH. For the determination of intracellular LDH
313 activity, the cells that had adhered to the multiplate wells were first washed twice with PBS,
314 and water was added to perform 3 repetitions of freezing (-80 °C) and thawing in a heated
315 waterbath, in order to lyze the cells. The intra- and extracellular LDH assays were performed
316 according to the kit instructions (Gold Analisa Diagnóstico, Minas Gerais, Brazil). Untreated
317 cells were used as a negative control.

318

319 **2.14 Lactate analysis**

320 For intracellular lactate measurement, cells were lysed (as described above) and the
321 procedures were followed according to the kit datasheet (Katal Biotecnológica, Belo Horizonte,
322 MG, Brazil). Spectrophotometric reads (540 nm) were performed using a microplate reader
323 (Thermo Scientific Multiskan). Untreated cells were used as a negative control.

324

325 **2.15 Determination of mitochondrial membrane potential ($\Delta\Psi_m$)**

326 We performed the analysis of mitochondrial membrane potential through
327 tetramethylrhodamine-ethyl ester (TMRE; Sigma-Aldrich, USA). NCI-H460 cells (10^4
328 cells/well) were treated with IC₅₀-TMBP for 24 h at 37 °C. The assay was performed as
329 described by Concato et al. (2020) [19]. Untreated cells were used as a negative control and
330 carbonylcyanide *m*-chlorophenylhydrazone (CCCP) (100 μ M) was used as a positive control.
331 Data were normalized and expressed in arbitrary units.

332

333 **2.16 Detection of lipid droplets (LD)**

334 To determine LD accumulation, NCI-H460 cells (10^4 cells/well) were treated with IC₅₀-
335 TMBP for 24 h at 37 °C and stained with 10 μ g/mL Nile red (NR; Sigma-Aldrich, St. Louis,
336 MO, USA) for 30 min. The assay was performed according to Concato et al. (2020) [19].
337 Untreated cells were used as a negative control, and PBS treatment was used as a positive
338 control. Data were normalized and expressed in arbitrary units.

339

340 **2.17 Determination of the formation of autophagic vacuoles**

341 The formation of autophagic vacuoles was evaluated by staining with
342 monodansylcadaverine (MDC; Sigma-Aldrich, St. Louis, MO, USA), according to Concato et
343 al. (2020) [19]. Briefly, NCI-H460 cells (10^4 cells/well) were treated with IC₅₀-TMBP for 24 h
344 at 37 °C and MDC was then added and left for 1 h. The fluorescence intensity was quantified

345 by spectrophotometry (Glomax, Promega, USA) (excitation 380 nm and emission 525 nm).
346 Untreated cells were used as a negative control, and PBS treatment was used as a positive
347 control. Data were normalized and expressed in arbitrary units.

348

349 **2.18 Cell cycle analysis**

350 NCI-H460 cells (10^6 cells/well) were treated with IC₅₀-TMBP for 24 h at 37 °C. The
351 assay was performed as described by Concato et al. (2020) [19]. Briefly, the cell suspension
352 was subjected to centrifugation at 1500 rpm for 5 min and the cell pellet resuspended in 300 µL
353 of PBS. Subsequently, a solution containing 0.05% ribonuclease A (RNase A) (Sigma, St.
354 Louis, MO, USA) was added and incubated for 30 min at 37 °C. In the end, a solution of 0.1%
355 sodium citrate and 1% Triton-X100 was added with 50 µg propidium iodide (PI) (Sigma, St.
356 Louis, MO, USA) for 30 min. The fluorescence was estimated using a Muse Cell Analyzer
357 (Merck Millipore), with 5000 events. The DNA content was analyzed, and the percentage of
358 cells in different phases of the cycle (G1, S, and G2/M) were estimated according to the
359 fluorescence intensity.

360

361 **2.19 Cell death assay**

362 NCI-H460 cells (10^6 cells/well) were treated with IC₅₀-TMBP for 24 h, and
363 phosphatidylserine exposure was detected using Annexin-V FITC (Invitrogen, Eugene, USA),
364 and cellular membrane integrity was detected with PI (0.50 µg/ ml) (Sigma, St. Louis, MO,
365 USA). The assay was performed and described by Concato et al. (2020) [19].

366

367 **2.20 Caspase-3/7 Assay**

368 Caspase-3/7 protease activity was measured using the EnzChek™ Caspase-3/7 Assay
369 Kit (Invitrogen™). Briefly, NCI-H460 cells (10^6 cells/well) were treated with IC₅₀-TMBP for
370 24 h, washed with PBS, and incubated at -80 °C for lysis. Lysis buffer (Triton X-100 in 1%
371 PBS) was subsequently added and the solution was left for 30 min. This was followed by the
372 addition of kit reagents (i.e., reaction buffer, dithiothreitol, and 5.0 mM Z-DEVD-AMC
373 substrate) and incubated for 30 min. On completion of the reaction, the increase in fluorescence
374 intensity, which is indicative of cleavage of the Z-DEVD-AMC substrate, was fluorometrically
375 read (GloMax, Promega, USA) at excitation and emission wavelengths of 342 and 441 nm,
376 respectively. The reactions were performed with and without the Ac-DEVD-CHO inhibitor (1
377 mM). Data were normalized and expressed in arbitrary units.

378

379 2.21 Protein Extraction and Enzyme-linked Immunosorbent Assay

380 NCI-H460 cells (10^6 cells/well) were plated and treated with IC₅₀-TMBP for 24 h,
381 followed by washing with sterile PBS to remove non-adherent cells in the plate wells. The
382 adhered cells were incubated with RIPA lysis buffer (150 mM sodium chloride, 1% triton X-
383 100, 0.5% sodium deoxycholate, 0.1% sodium dodecyl sulfate, 5 mM EDTA, 50 mM Tris, pH
384 8.0) overnight at -80 °C. The cell lysate was collected and centrifuged at $13,000 \times g$ for 20 min
385 at 4 °C, the supernatant recovered and transferred to a new tube and the total protein
386 concentration was quantified using NanoVue Plus reagent (Biochrom, USA). The protein
387 concentration of all samples was normalized to 20 µg/mL, and a 100-µL aliquot was added in
388 a 96-well ELISA plate for adsorption of proteins overnight at 4 °C. This was followed by adding
389 a blocking buffer (ELISA/ELISPOT, eBioscience™, USA) and the plate was incubated for 1
390 h. The wells were then washed 3x with wash buffer (PBS in 0.5% Tween-20) and incubated
391 with primary antibodies: anti-human PI3K, AKT, pAKT, arginase-1, NF-κB, pNF-κB, iNOS,
392 and NRF2 (Santa Cruz Biotech, USA) for 2 h at room temperature. The wells were then washed
393 (wash buffer) to remove unbound antibodies, followed by the addition of universal biotinylated
394 secondary antibody (LSAB2 System-HRP, Dako, USA) and incubated for 1 h. The wells were
395 then washed (wash buffer), followed by the addition of streptavidin-HRP (LSAB2 System-
396 HRP, Dako, USA) and incubated for 1 h. After the incubation time, the wells were washed 5
397 times (wash buffer) and 100 µL of TMB substrate solution (eBioscience™, USA) was added,
398 followed by incubation for 30 min, and then 100 µL of stop solution (0.5 M sulfuric acid) was
399 added. The plate was read in a microplate reader at 450 nm (GloMax, Promega, USA).

400

401 2.22 Statistical analysis

402 Data are expressed as the mean \pm standard error of the mean. Three independent
403 experiments were performed, each with duplicate data sets, and analyzed with GraphPad Prism
404 8 statistical software (GraphPad Software, Inc., USA, 500.288). Significant differences
405 between groups were determined through t-tests or one-way ANOVA followed by Tukey's test
406 for multiple comparisons. A result was considered statistically significant if $p < 0.05$. Values
407 were also categorized fallows: * ($p < 0.05$); ** ($p \leq 0.01$); *** ($p \leq 0.001$); **** ($p \leq 0.0001$).

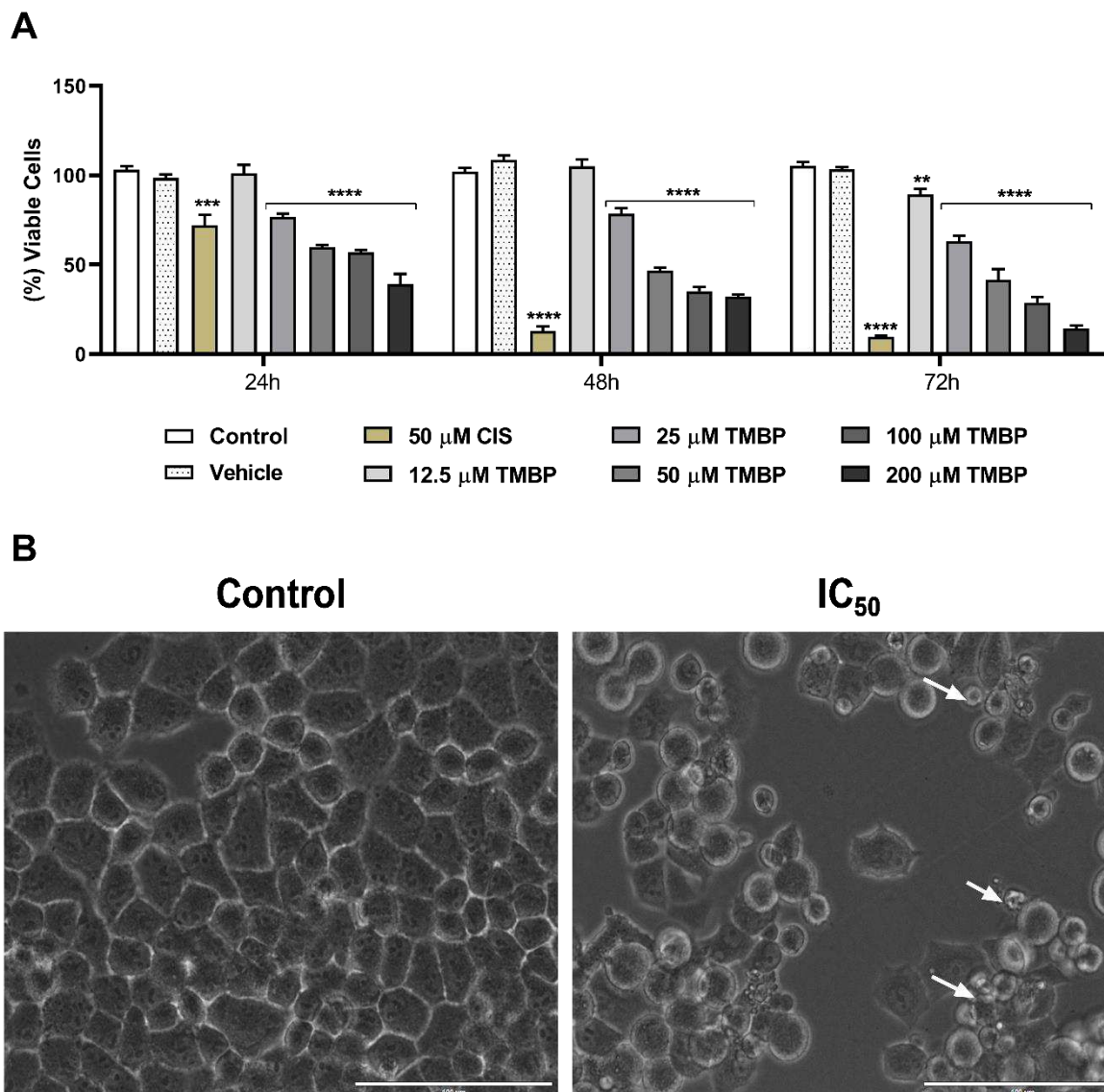
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409 3. Results

410 3.1 TMBP treatment reduces proliferative activity in NCI-H460 cell line

411 We first evaluated the cell viability after treatment with TMBP (12.5-200 µM) or
412 cisplatin (CIS) (50 µM) for 24-72 h. The vehicle (DMSO) showed no toxic effect at any of the

413 times tested. At 24, 48, and 72 h, the concentrations of TMBP within the range 25-200 μM
 414 reduced the proliferation of the NCI-H460 cells ($p \leq 0.0001$), but the effects were more
 415 pronounced at 72 h. The concentration of 12.5 μM TMBP also affected cell viability which was
 416 reduced by 20% during this period (**Fig. 1A**). The IC_{50} was calculated for all the times tested
 417 (24, 48, and 72 h), resulting in the respective values of 152 μM (± 0.07), 73 μM (± 0.03), and
 418 42 μM (± 0.03) (**Table 1**). The IC_{50} at 24 h was used for the following experiments, as
 419 previously described [19] and representative microscopic images showed morphological
 420 changes, such as rounding and apoptosis-like signs (white arrow) (**Fig. 1B**).



421

422 **Figure 1: TMBP reduced the viability of the human cancer cell line NCI-H460 at different**
 423 **concentrations.** NSCLC cells were treated with TMBP for 24, 48, and 72 h. The relative
 424 survival rate (**A**) was determined by the MTT test. After determining the half-maximal
 425 inhibitory concentration (IC_{50}) of TMBP, representative images were taken by light microscopy

426 (1000×) (B). Blebbing around the cells (white arrow). Values are expressed as mean \pm SEM.
 427 Significant difference with $**p \leq 0.01$, $***p \leq 0.001$ and $****p \leq 0.0001$ vs. control; vehicle
 428 (DMSO 0.06%); CIS, cisplatin. Scale bar = 100 μ m.

429

430 **Table 1:** The half-maximal inhibitory concentration (IC_{50}) in lung cancer NCI-H460 cell line
 431 after TMBP treatment for 24-72 h.

Cell Type	IC_{50} (\pm SEM)		
	24 h	48 h	72 h
NCI-H460	152 μ M (\pm 0.07)	73 μ M (\pm 0.03)	42 μ M (\pm 0.03)

432

433

IC_{50} – Half-maximal inhibitory concentration; SEM – standard error of the mean.

434

3.2 TMBP induces morphological changes in NCI-H460 cell line

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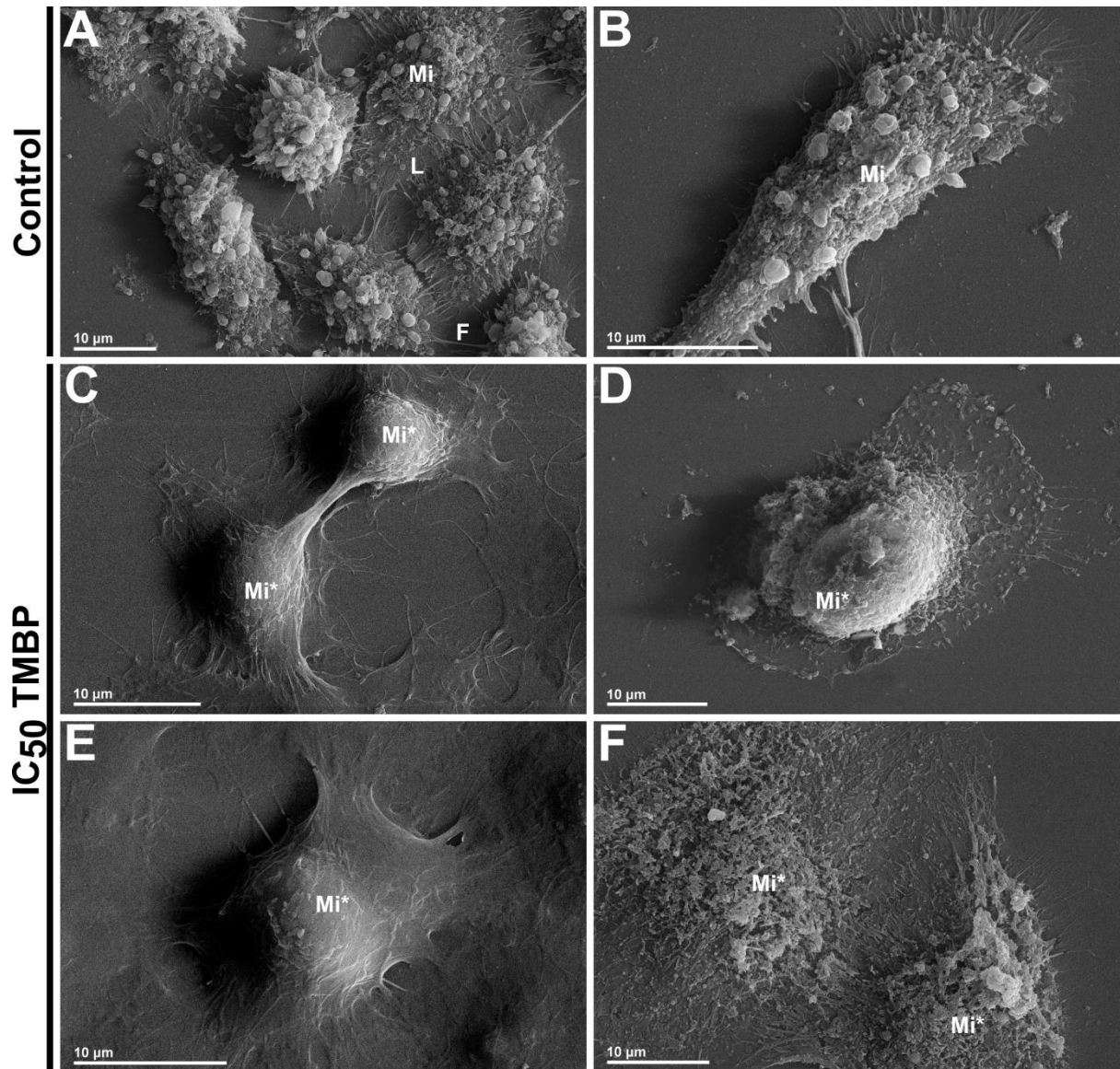
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SEM was used to determine the morphological changes induced by TMBP treatment in the NCI-H460 cell line. Untreated cells (Fig. 2 A, B) showed normal characteristics such as firm adherence to the substrate, intact cell membrane, presence of lamellipodia, filopodia, microvilli throughout the cell surface, and well-preserved structures. Cells treated with IC_{50} -TMBP, however, showed rounded shapes, reduction of lamellipodia, irregular cell surface, and absence and/or reduction of microvilli. TMBP treatment induced a reduction in cell surface blebbing, indicating the release of large oncosomes; a type of extracellular vesicle characteristic of cells with a more aggressive profile and metastatic potential (Fig. 2 C-F). Our data demonstrated that in addition to reducing proliferation, TMBP treatment was also able to alter the morphology of the cells, contributing to their inability to move and reducing malignancy.



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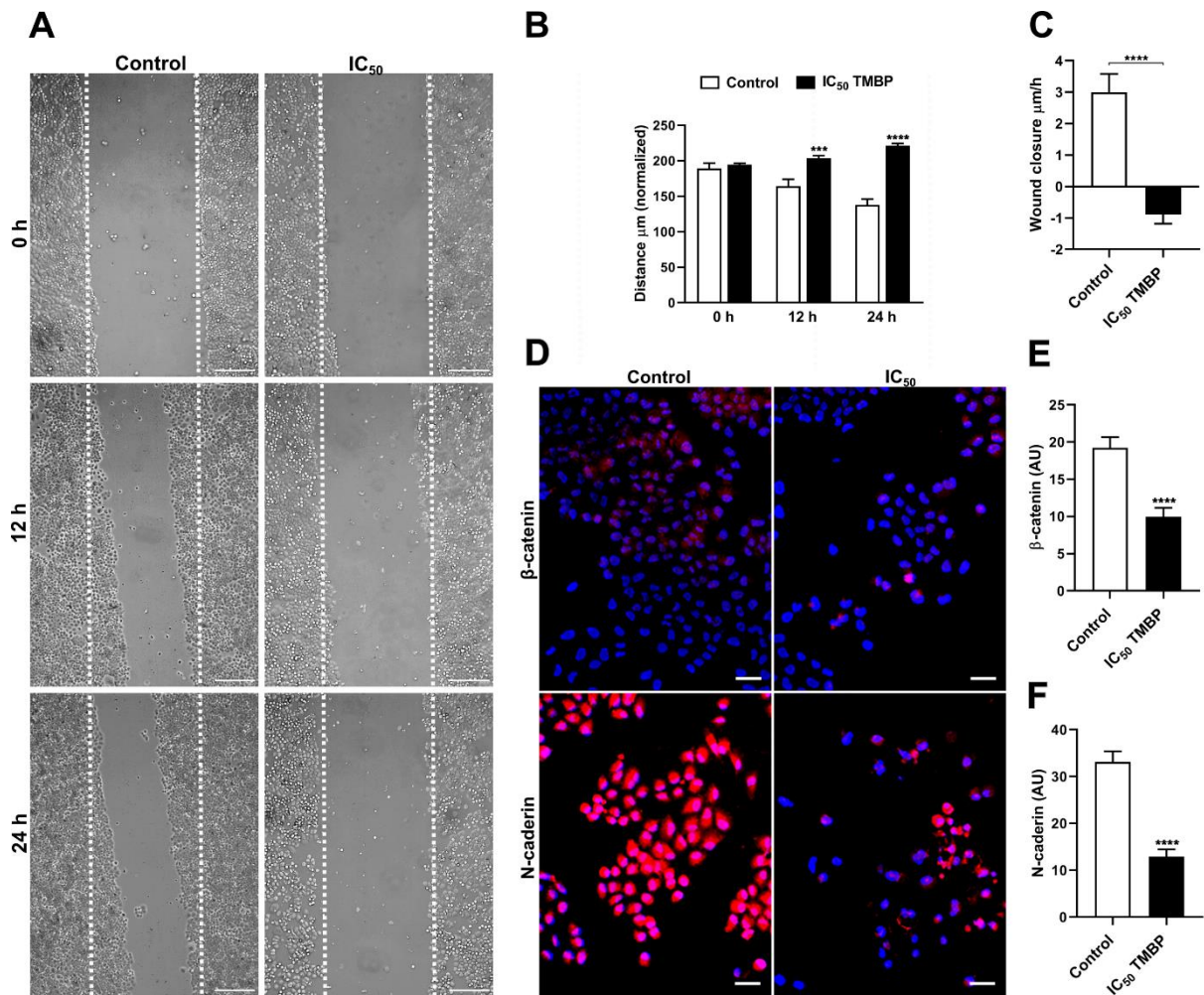
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Figure 2: Effect of TMBP treatment on the morphology of NCI-H460 cells. Morphological changes were observed by scanning electron microscopy after treatment with TMBP for 24 h. (A-B) Control, untreated cells; (C-F) cells treated with IC₅₀ of TMBP (152 μM). (Mi) microvilli, (Mi*) microvilli alterations, (L) lamellipodia and (F) filopodia. Scale bars = 10 μm (A-F).

3.3 TMBP treatment reduces migratory capacity and N-cadherin/β-catenin markers in the NCI-H460 cell line

The effect of IC₅₀-TMBP treatment on the migratory potential of NCI-H460 cells was assessed using the *in vitro* wound healing assay (**Fig. 3**). The filling of the scratched area by migrating cells was monitored at regular time intervals (0, 12, and 24 h) and illustrated by representative images (**Fig. 3A**). Quantitative analysis concerning the time traveled revealed

458 that in the control group, cells migrated to the wound site, however, those that received TMBP
 459 treatment showed an increase in the scratched area in comparison to time 0 h ($p < 0.0001 \pm$
 460 0.29) (**Fig. 3 B-C**). Corroborating these data, after 24 h of treatment with IC_{50} of TMBP, we
 461 observed reduced levels of fluorescence intensity of β -catenin ($p < 0.0001 \pm 1.22$) and N-
 462 cadherin ($p < 0.0001 \pm 1.56$) compared to the untreated group (**Fig. 3 D-F**).
 463

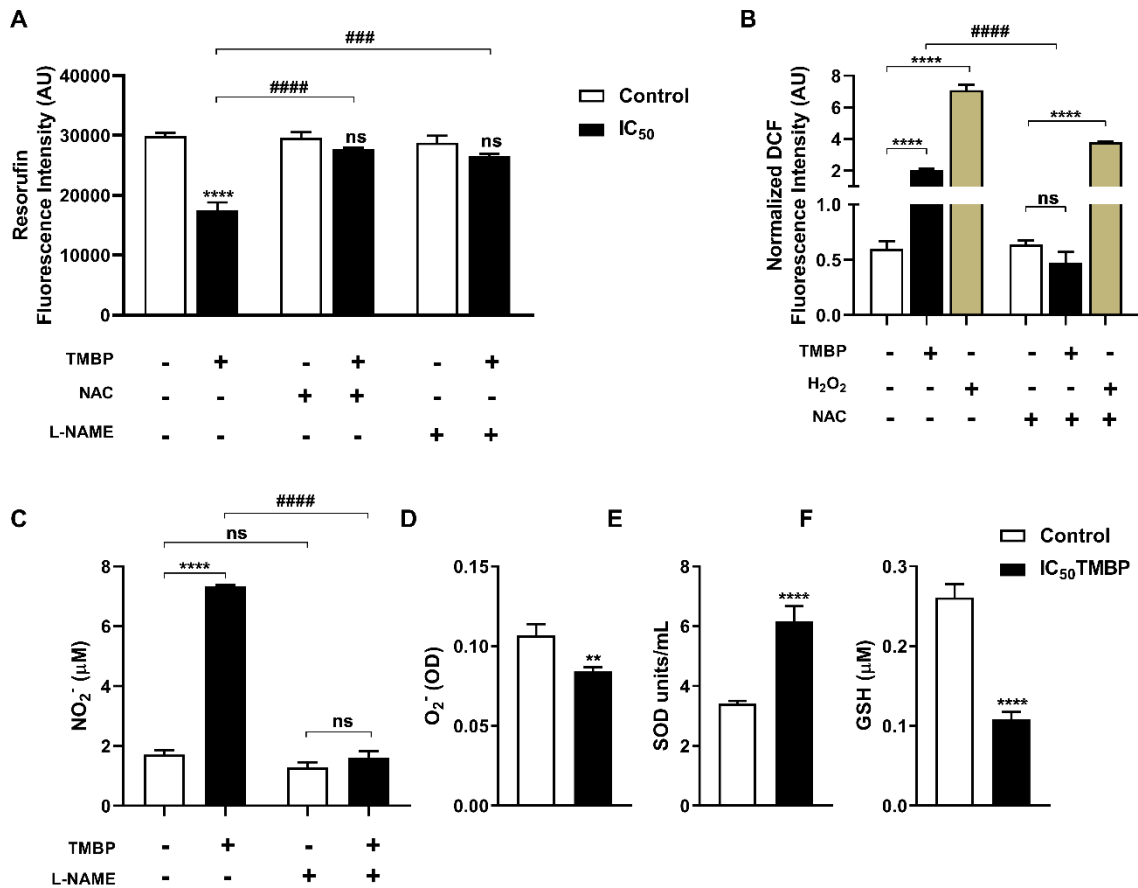


464
 465 **Figure 3: TMBP treatment inhibited NCI-H460 cell migration, with cell retraction and**
 466 **reduced N-cadherin and β -catenin staining.** A scratch assay was performed to observe the
 467 effect of treatment on cell migration over time (0, 12 and 24 h) (A). The distance of the free
 468 surface of these treated cells (B) was quantified. The time taken by the cells to reach the wound
 469 site was also quantified (C). Immunofluorescence staining of TMBP-treated and control cells
 470 to show reduced β -catenin (red) and N-cadherin (red) presence after 24 h (D) (scale bar 2 μm).
 471 (E-F) quantification of fluorescence intensity using Image J software. Values represent the
 472 mean \pm SEM of three different experiments performed in triplicate. Significant difference
 473 versus control *** $p \leq 0.001$ and **** $p \leq 0.0001$. Scale bars = 200 μm (A) and 20 μm (D).

474 **3.4 TMBP induces oxidative stress in the NCI-H460 cell line by a ROS/NO-dependent**
475 **process**

476 ROS production in TMBP-treated cells was analyzed by fluorescence. **Figure 4B**
477 demonstrated that exposure of cells to TMBP resulted in a 2-fold increase in fluorescence
478 intensity compared to the control ($p < 0.0001 \pm 0.09$). The antioxidant N-acetyl cysteine (NAC)
479 was used to confirm the elevation of ROS. The addition of NAC effectively blocked TMBP-
480 induced ROS in NCI-H460 cells and restored viability (**Fig. 4 A, B**). In the same way, when
481 the levels of NO post-treatment with TMBP were evaluated, an increase of 80% ($p < 0.0001 \pm$
482 0.041) levels of this mediator was observed. The addition of L-NAME, a nitric oxide synthase
483 (NOS) inhibitor, restored these levels and cell viability (**Fig. 4 A, C**).

484 Superoxide anion radical ($O_2^{\bullet-}$) is one of the most frequently generated ROS in the cell
485 and can react with other molecules to form potent reactive species, such as peroxynitrite, or be
486 dismutated by SOD to produce O_2 and H_2O_2 [27,28]. After treatment of the cells with TMBP,
487 we observed reduced $O_2^{\bullet-}$ levels (**Fig. 4D**) and increased SOD activity (**Fig. 4E**). Furthermore,
488 a reduction of the antioxidant GSH (**Fig. 4F**) (which is essential for ROS detoxification) was
489 also observed. These results suggest that TMBP reduces NCI-H460 cell viability mediated by
490 increasing levels of ROS/NO molecules.



491

492 **Figure 4: TMBP treatment induced oxidative stress in NCI-H460 cells.** The viability of
 493 cells treated with IC₅₀ of TMBP was assessed by resazurin in the presence or absence of NAC
 494 or L-NAME (A). ROS levels were assessed with H₂DCFDA in the presence or absence of NAC
 495 (B). Nitric oxide was determined after TMBP treatment in the presence or absence of L-NAME
 496 (C). Superoxide radical anion was determined using NBT (D). SOD and GSH activity were
 497 also determined after TMBP treatment (E-F). Data are expressed as mean ± SEM of 3
 498 independent experiments performed in triplicate. ns not significant; ** ($p < 0.01$); **** ($p \leq$
 499 0.0001) vs. control. ### ($p \leq 0.001$) and #### ($p \leq 0.0001$) treatment differences.

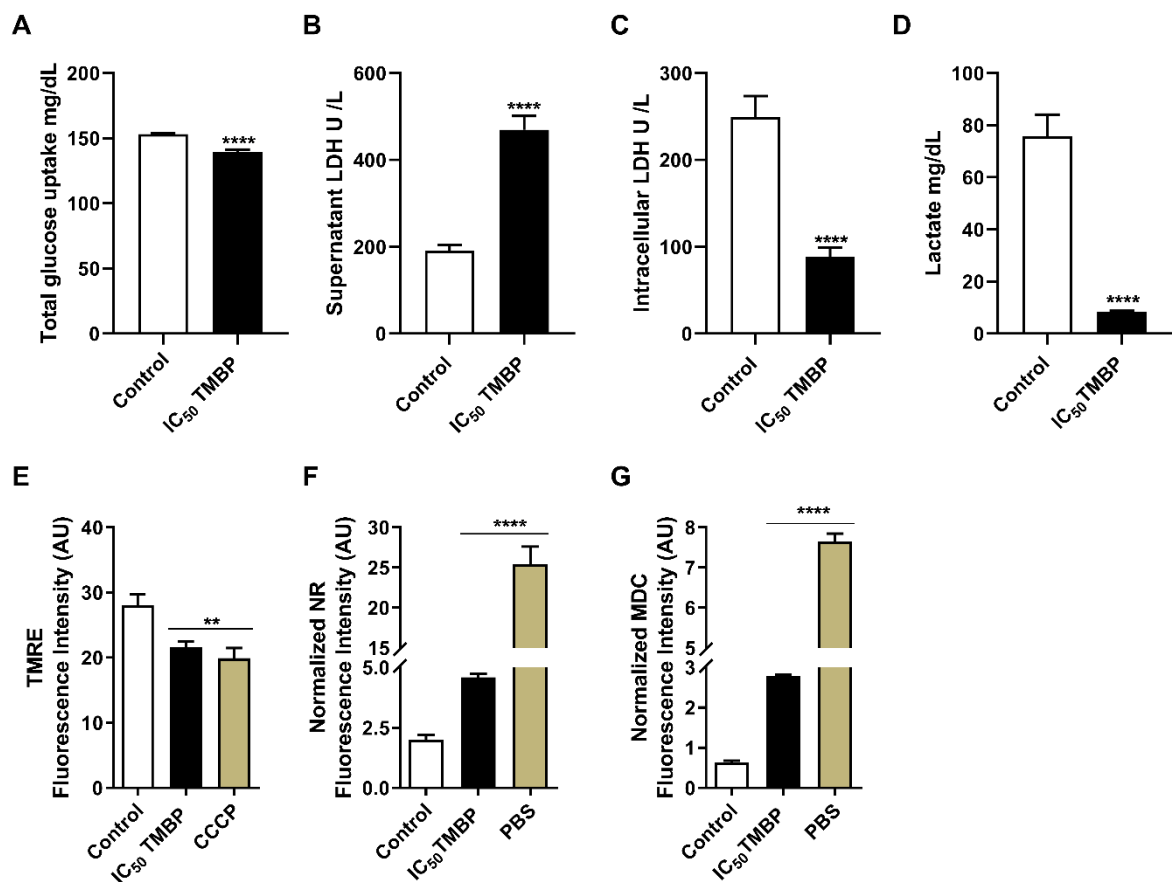
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501 3.5 TMBP promotes metabolic changes in NCI-460

502 To assess cellular metabolism, we first examined glucose levels after TMBP treatment
 503 of NCI-H460 cells. We observed a reduction in glucose uptake compared to untreated cells (p
 504 ≤ 0.0001) (Fig. 5A). When LDH (which is crucial for converting pyruvate to lactate and vice
 505 versa) enzyme activity was assessed, there was a significant 70% increase in extracellular levels
 506 and a 65% decrease in intracellular levels ($p > 0.0001$; both activities), indicating cellular
 507 damage (Fig. 5 B-C). The levels of lactate (a glucose metabolite of aerobic glycolysis) also

508 decreased compared to the control, with concentrations of 80 mg/dL in the control group and
 509 10 mg/dL in the TMBP-treated groups ($p > 0.0001$) (**Fig. 5D**).

510 The mitochondrial membrane was examined in NCI-H460 cells after treatment with IC_{50}
 511 of TMBP. A decrease in TMRE fluorescence intensity was observed, indicating a loss of
 512 mitochondrial activity ($p \leq 0.01$) (**Fig. 5E**). Excess intracellular lipids can be stored in lipid
 513 droplets (LDs) to prevent lipotoxicity. During nutrient deprivation, fatty acids are released from
 514 LDs for energy production through mitochondrial β -oxidation and the Krebs cycle apoptosis
 515 (Cruz et al., 2020). In our results, TMBP treatment induced and increased the level of LD
 516 accumulation compared to the control ($p \leq 0.0001$) (**Fig. 5F**). In addition, we investigated
 517 whether treating cells with IC_{50} of TMBP would affect autophagy. We found that the treated
 518 group increased the fluorescence intensity, indicating the formation of autophagic vacuoles in
 519 NCI-H460 cells ($p \leq 0.0001$) (**Fig. 5G**).



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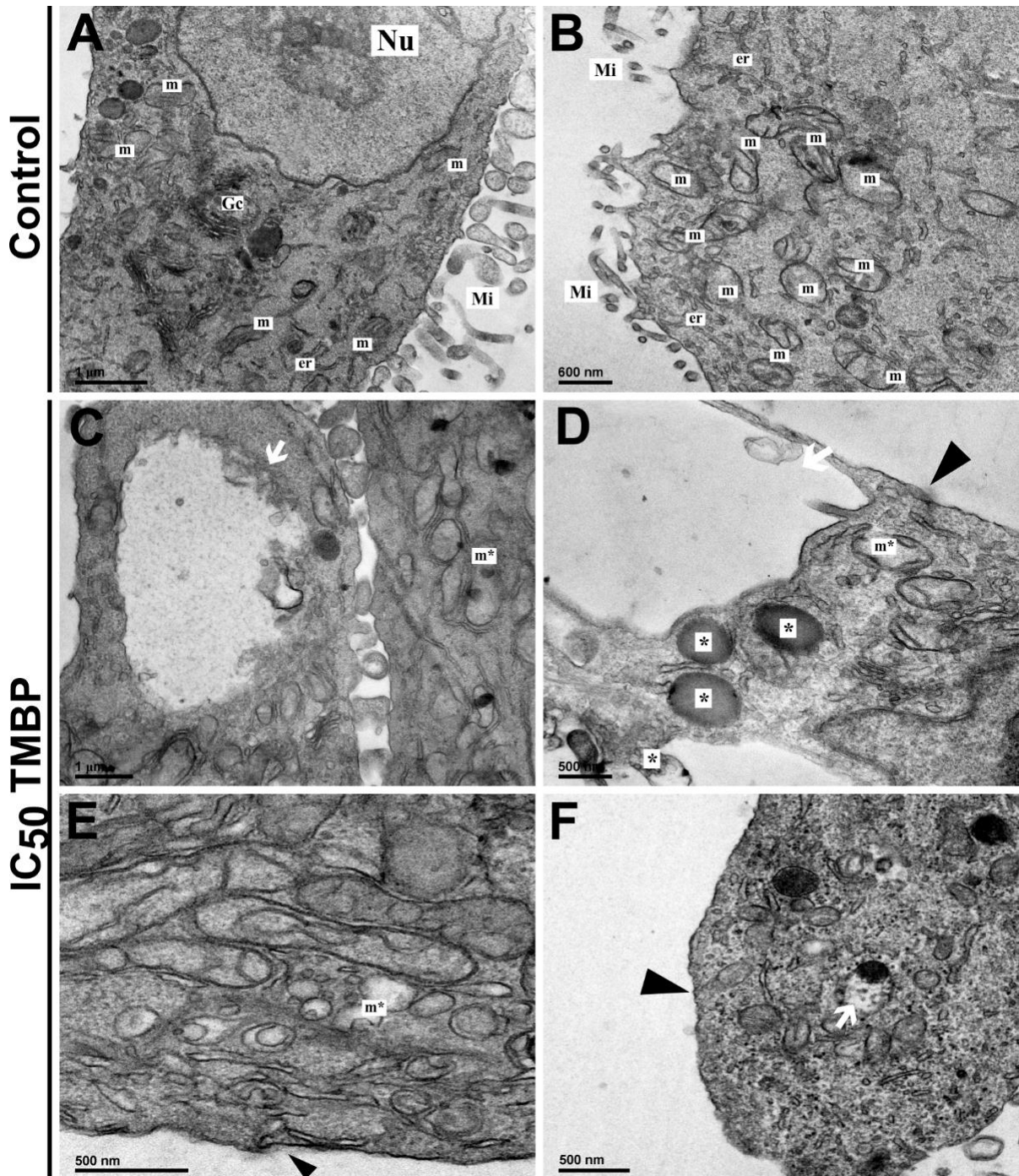
521 **Figure 5: TMBP treatment induced metabolic stress in NCI-H460 cells.** Cells were treated
 522 with TMBP for 24 h. (A) extracellular glucose uptake; (B and C) extracellular and intracellular
 523 LDH activity, respectively; and (D) intracellular lactate level. $\Delta\Psi_m$ (E), lipid droplets (F), and
 524 autophagic vacuoles (G) were quantified by fluorescence. Data are expressed as mean \pm SEM

525 of 3 independent experiments, each performed in triplicate. ** ($p < 0.01$); **** ($p \leq 0.0001$)
526 vs. control. CCCP, carbonylcyanide *m*-chlorophenylhydrazone; PBS, phosphate-buffered
527 saline

528

529 **3.6 TMBP induces structural changes in the NCI-H460 cell line**

530 Ultrastructural evaluation by TEM showed that the control group displayed a
531 homogeneous cytoplasm with regular organelles (mitochondria and endoplasmic reticulum)
532 and without the presence of vacuole-like structures, indicating cellular integrity (**Fig. 6 A-B**).
533 After TMBP treatment, changes in mitochondrial structures were observed, such as swelling
534 and enlargement of these organelles (m*) (**Fig. 6 C-E**). These changes were consistent with the
535 data presented above and indicated that mitochondria were damaged. In addition, an intense
536 accumulation of lipid storage droplets (*) and autophagic vacuoles (white arrow) were observed
537 (**Fig. 6 C, D, F**). Furthermore, the absence of microvilli, filopodia, and lamellopodia (black
538 arrowhead) were also observed (**Fig. 6 D-F**).



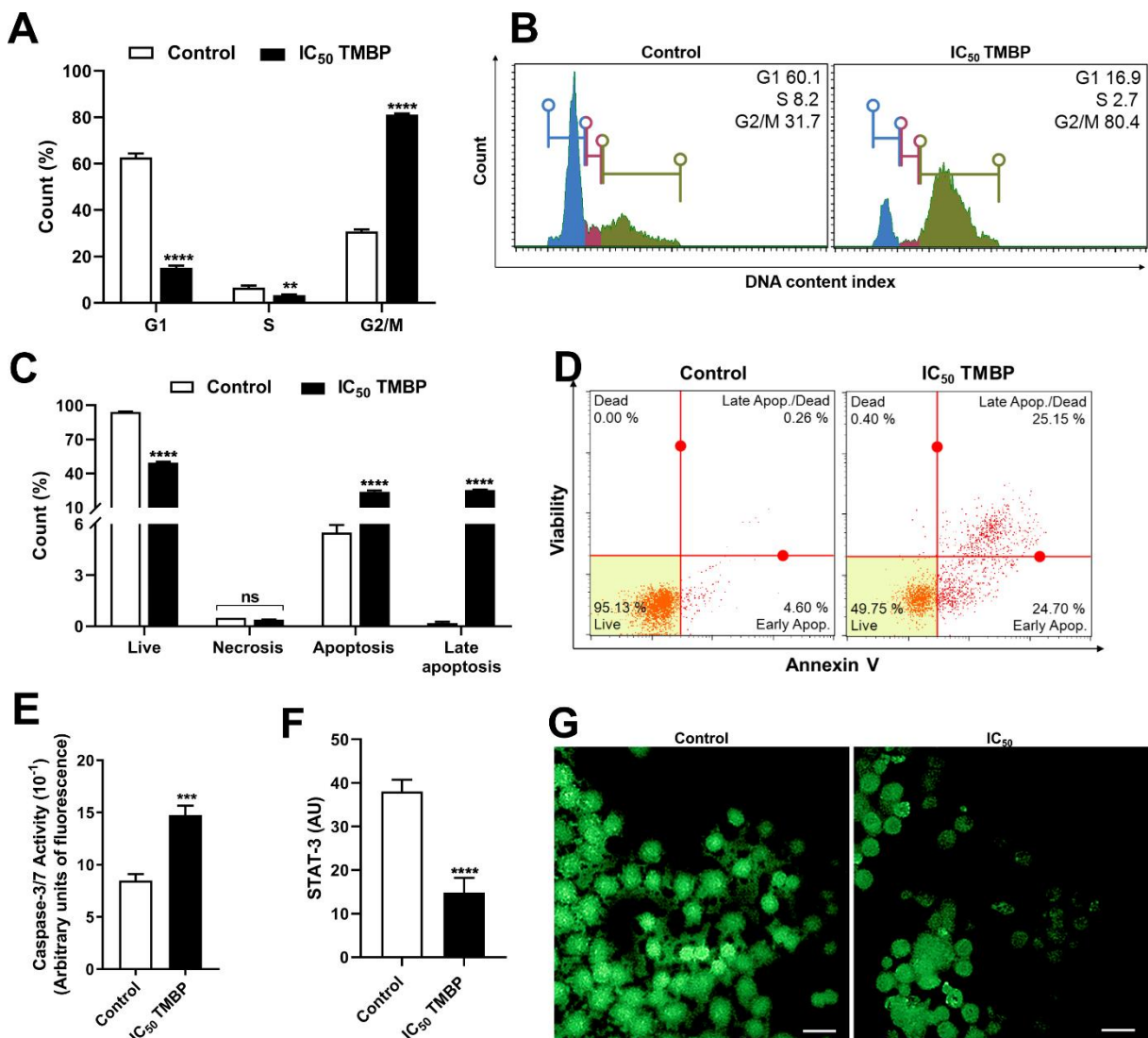
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540 **Figure 6: Ultrastructural changes in NCI-H460 cells treated with TMBP for 24 h and**
 541 **analyzed by transmission electron microscopy. (A-B) untreated cells; (C-F) cells treated**
 542 **with IC₅₀ TMBP (152 μM); (m) mitochondria; (Gc) Golgi complex; (er) endoplasmic**
 543 **reticulum; (m*) mitochondrial swelling; (Nu) nucleus; (white arrow) cytoplasmic autophagic**
 544 **vacuole (black arrowhead) absence of microvilli; (*) lipid droplets storage areas. Scale bars =**
 545 **1 μm (A, C), 600 nm (B) and 500 nm (D-F).**

546

547 **3.7 TMBP promotes cell cycle arrest in the G2/M phase and apoptosis via caspase-**
 548 **3/STAT-3**

549 Analysis of cell cycle distribution also revealed a reduction in the G1 and S phases
 550 ($51.1\% \pm 0.86$ and $9.5\% \pm 1.40$, respectively) and an accumulation of cells in the G2/M phase
 551 ($52.7\% \pm 1.31$) (**Fig. 7 A-B**). **Figure 7 C-D** shows that TMBP-treated cells promoted reduced
 552 cell viability by 49.75% (± 0.72), increased annexin (24.7% ± 0.85), and co-marking
 553 (annexin/PI) ($25.15\% \pm 0.22$) compared to the control, indicating cell death by an apoptosis-
 554 like process. This was confirmed by the increase in caspase-3/7 activity observed in these cells
 555 (**Fig. 7E**). Also, when evaluated by immunofluorescence the signal transducer and activator of
 556 transcription 3 (STAT-3), a reduction of this protein was observed in cells treated with TMBP
 557 ($p \leq 0.0001$) (**Fig. 7F-G**). In contrast, higher staining was observed in the untreated group.
 558 These data show that TMBP promoted metabolic damage that disrupted mitosis and progression
 559 leading to cell death.



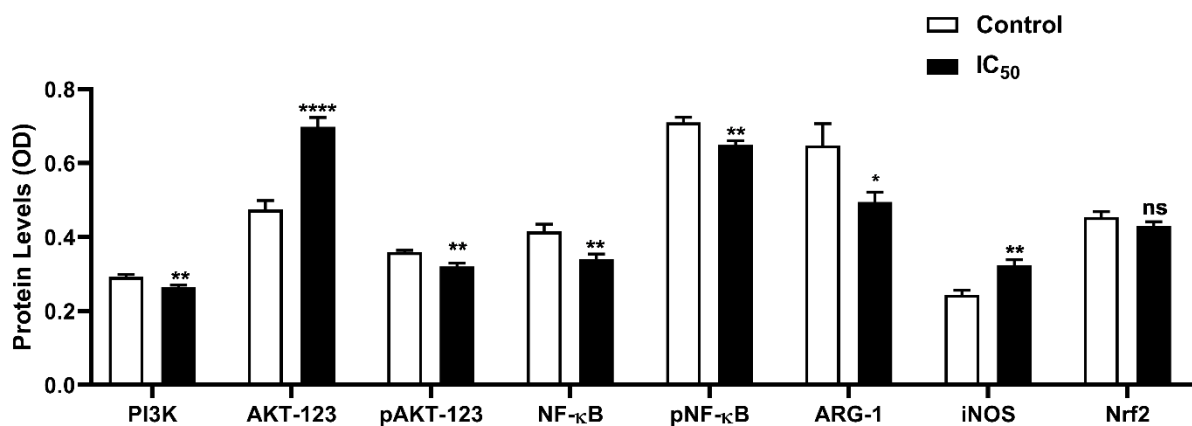
561 **Figure 7: TMBP treatment induced cell cycle arrest in the G2/M phase, apoptosis, and**
 562 **reduced STAT-3 immunofluorescence staining in NCI-H460 cells.** Cells were treated with
 563 IC₅₀ of TMBP for 24 h and cell cycle distribution and histogram were quantified (A) and
 564 analyzed (B), respectively by flow cytometry. Quantitative analysis of viable, apoptotic, and
 565 necrotic NCI-H460 cells was evaluated in 24 h (C). For analysis in dot plots of cytometry, the
 566 induction of viable cells, apoptosis, and necrosis were evaluated for 24 h (D). Caspase 3/7 was
 567 quantified by fluorescence (E). Immunofluorescence staining of TMBP-treated and control
 568 cells to show reduced STAT-3 presence (green) after 24 h (F-G). Values are expressed as mean
 569 \pm SEM. Cytometry data are expressed as percentage vs. control; and mean \pm standard deviation.
 570 $p \leq 0.01$; *** $p \leq 0.001$; **** $p \leq 0.0001$ vs. control. Scale bars = 20 μ m (G).

571

572 3.8 TMBP acts on the PI3K/AKT/NF- κ B pathway and activates iNOS mediated pro- 573 oxidant profile in NCI-H460 tumor cell line

574 The PI3K/AKT signaling pathway is frequently activated in cancer as it is involved in
 575 the regulation of proliferation, survival, inhibition of apoptosis, and cell invasion [34]. These
 576 proteins were therefore evaluated by ELISA after treatment with TMBP (Fig. 8). A decrease in
 577 PI3K ($p \leq 0.01$) and an increase in total AKT ($p \leq 0.0001$) were observed. The pAKT ($p \leq 0.01$)
 578 phosphorylated activated form, however, decreased.

579 After treatment with TMBP, we observed reduced levels of arginase-1 (ARG-1) ($p <$
 580 0.05) and increased levels of inducible nitric oxide synthase (iNOS) ($p \leq 0.01$). Next, nuclear
 581 factor *kappa* B (NF- κ B) was evaluated after treatment with TMBP. We observed a decrease in
 582 total NF- κ B protein levels ($p \leq 0.01$) and its activated phosphorylated form (pNF- κ B) ($p \leq$
 583 0.04). Furthermore, when the nuclear factor erythroid 2 related factor 2 (Nrf2) was analyzed,
 584 we found that its levels did not significantly differ from the control group ($p > 0.05$).



585

586 **Figure 8: Evaluation of intracellular signaling mediators involved in the pathogenesis of**
587 **lung cancer cells.** ELISA measurement of PI3K, AKT, pAKT, ARG-1, iNOS, NF- κ B, pNF-
588 κ B and Nrf2 levels, expressed as the natural logarithm of absorbance. White bars represent
589 untreated NCI-H460 cells, while black bars represent 24-h TMBP-treated cells. Values
590 represent the mean \pm SEM. * ($p < 0.05$); ** ($p \leq 0.01$); **** ($p \leq 0.0001$); ns (not significant).
591

592 4. Discussion

593 Chemoresistance is a common feature and a major cause of treatment failure in NSCLC,
594 leading to tumor recurrence and disease progression. This ability is associated with most
595 available drugs, including cisplatin. The main factors associated with cisplatin resistance are
596 reduced accumulation and inactivation of the drug by binding to various proteins, increased
597 DNA repair, and alteration of various proteins signaling apoptosis [35–37]. In this context, there
598 is a growing interest in microbial biotransformation of drugs and natural products with
599 antitumor activity. This process allows the modification of the relevant chemical molecules,
600 generating new compounds that may be more effective and less toxic. In addition, it is possible
601 to take advantage of the mild conditions under which the reactions take place, with most
602 transformations occurring at temperatures between 20°C and 40°C and with aqueous solvents.
603 Although yields are sometimes limited, several options for biological reagents can be
604 manipulated through mutants, aeration, temperature, and culture medium to improve these
605 results [38, 39]. Thus, the development of therapeutic strategies that are effective in eliminating
606 aberrant cells and preventing malignant growth is of great interest as it may can expand the
607 possibilities of new approaches for therapeutic applications.

608 In a previous study, we demonstrated that TMBP exhibited direct cytotoxic effects on
609 the A549 cell line. In addition, *in silico* predictions have shown promising drug-likeness
610 potential for TMBP, indicating high oral bioavailability and intestinal absorption [19]. For the
611 first time, the present study provides new insights into the type of cell death induced by TMBP
612 against the NCI-H460 cell line, demonstrating anti-proliferative activity, metabolic and
613 structural changes (reduced lamellipodia, filopodia, microvilli, and release of large oncosomes).
614 A series of biochemical processes showed that TMBP acts by producing ROS and NO that
615 potentially lead to cell death. Furthermore, TMBP treatment of the cell line reduced protein
616 levels of PI3K/pAKT/ARG-1/pNF- κ B and increased iNOS, which may favor increased
617 oxidative stress and promote death in these tumor cells. The significance of our *in vitro* findings
618 is reinforced by the fact that key components of the TMBP-mediated pathway described have

619 identified cellular self-destruction in lung cancer, and may lead to improved oncotherapy.

620 Cell proliferation and migration are considered an essential and regulated behavior in
621 several normal physiological processes, such as morphogenesis, embryogenesis, immune cell
622 responses, angiogenesis, tissue repair, cell differentiation, and tissue regeneration [40]. In
623 cancer, cell migration, invasion, and adhesion are phenomena that determine cell plasticity and
624 epithelial-mesenchymal transition (EMT). During the process of EMT, which involves the
625 transformation of cancer cells from a non-motile epithelial phenotype to a migratory
626 mesenchymal-like phenotype, several changes occur in the markers expression. Epithelial
627 markers such as E-cadherin, α -catenin, and β -catenin are downregulated, while mesenchymal
628 markers like N-cadherin and vimentin are upregulated. This transition closely correlates with
629 increased cell migration and invasion, contributing to cancer progression [41–43].

630 It is worth noting that in the context of cancer, the stabilization of β -catenin and its
631 subsequent translocation into the nucleus can promote cancer progression, cell proliferation,
632 and metastasis [44–47]. By reducing β -catenin levels, the compound TMBP may actually
633 inhibit the pro-metastatic effects associated with β -catenin. Similarly, the decrease in N-
634 cadherin is in line with our hypothesis that TMBP counteracts the EMT process. N-cadherin is
635 typically upregulated during EMT and contributes to cell migration and invasion [44,48–50].
636 Therefore, the reduction in N-cadherin levels supports the potential anti-metastatic activity of
637 TMBP [51–53]. In addition to its effects on metastasis-related markers, our experimental data
638 demonstrates that TMBP not only reduces cell proliferation but also diminishes the cells'
639 migratory capacity. Moreover, we observed that TMBP decreases cell protrusion extensions,
640 including lamellipodia and filopodia, as well as microvilli. These observations suggest that
641 TMBP interferes with cell locomotion by altering their invasive morphology, leading to reduced
642 actin filament content, which subsequently impacts their motility and cell-cell adhesion.

643 Oxidative stress is defined as an excess of oxidants over antioxidant activity and leads
644 to many diseases. Understanding the role of oxidative stress in cancer pathophysiology is
645 necessary since these biomolecules impact the whole organism. Regulation of the redox state
646 in a normal environment is critical for cellular maintenance, including viability, proliferation,
647 and metabolism. Cells generate ROS and reactive nitrogen species (RNS), as an unavoidable
648 consequence of physiological processes and for intracellular signaling. The major components
649 comprising ROS are superoxide radical anion ($O_2^{\bullet-}$), hydroxyl radical (OH^{\bullet}), and hydrogen
650 peroxide (H_2O_2). The main site of $O_2^{\bullet-}$ production is the mitochondria, which are used for
651 energy production and result from the process of OXPHOS. Superoxide anion radical can be
652 dismutated spontaneously to H_2O_2 or by the action of SODs, mainly SOD2. H_2O_2 can be

653 catalytically converted to H₂O and O₂ by catalase, peroxiredoxins (PRDXs), and glutathione
654 peroxidases (GPXs). GPXs require glutathione (GSH); reducing the levels of H₂O₂. Superoxide
655 radical anion can also react with H₂O₂ to form OH•, a highly reactive radical that can damage
656 proteins, lipids, carbohydrates, and DNA [54–56]. It should also be noted that when SOD2
657 activity is increased, proliferating cells go into quiescence, *i.e.* they do not multiply, but remain
658 metabolically active, resulting in increased H₂O₂ activity [57].

659 Nitric oxide (NO) and peroxynitrite anion (ONOO⁻) are the major RNS. GSH is highly
660 abundant in all cellular compartments and is the main soluble antioxidant and determinant of
661 oxidative stress. GSH also contributes to the scavenging of NO, the reduction of ONOO⁻ and
662 the denitrosylation of proteins. Thus, O₂•⁻ and ONOO⁻ are highly reactive species that can
663 damage intracellular macromolecules, including polyunsaturated fatty acids and nucleic acids
664 [58, 59]. Therefore, in our study, TMBP was able to reduce cancer cell viability via a ROS- and
665 NO-dependent mechanism. Furthermore, TMBP treatment caused a reduction in O₂•⁻ levels,
666 increased SOD2 activity, and decreased GSH, suggesting that the highly unstable superoxide
667 radical anions may be converted to peroxynitrite and/or hydroxyl radical.

668 Cancer cells exhibit aberrant metabolism, which is used to maintain their vital functions.
669 This leads to the activation of pro-tumorigenic signaling, increased cell proliferation, migration,
670 invasion, inhibition of apoptosis, therapeutic resistance, progression, and other aggressive
671 features. Therefore, when cancer cells are presented with a high demand for oxidative stress,
672 they initiate the process of adaptation in order to mitigate the damage and escape death [60, 61].
673 Another prominent feature is the Warburg effect, which refers to the metabolic shift in cancer
674 cells from OXPHOS to aerobic glycolysis. This glycolytic shift leads to a dramatic increase in
675 glucose uptake and consequent lactate production, which are essential in providing substrates
676 for cancer cells to sustain a high rate of proliferation [62–64]. Furthermore, glucose metabolism
677 is a hallmark of NSCLC [65, 66].

678 In addition to glycolysis, mitochondrial function, lipid biogenesis, and autophagy,
679 although essential in physiological processes, also play a critical role in cancer development by
680 providing cells with considerable flexibility, such as changes in utilization and energy supply,
681 and oxidative stress management. It has also been reported that when these processes become
682 irreparably damaged, the cell tends to undergo apoptosis [67–74]; [29–33]. In our study, we
683 observed that TMBP treatment of NCI-H460 cells was capable of reducing glucose and lactate
684 consumption, indicating that the cancer cells were under metabolic stress. We also found that
685 TMBP altered mitochondrial function, as demonstrated by [19], suggesting that the cells were
686 damaged, as confirmed by the intra- and extra-cellular LDH levels. Additionally, TMBP

687 treatment increased lipid droplets and autophagic vacuoles; indicative of a metabolic stress
688 profile, as confirmed by the TEM images. Taken together, these data suggest that TMBP-treated
689 NCI-H460 cells were damaged and initiated a process of metabolic stress in an attempt for the
690 cells to regain viability.

691 Controlling or terminating the uncontrolled growth of cancer cells using their own death
692 mechanism is a highly effective method of treating cancer. Apoptosis is an essential event in
693 the regulation of a wide range of physiological cellular processes. However, dysfunction of the
694 apoptotic machinery is implicated in many diseases, particularly cancer. Failure to activate the
695 apoptotic pathway can lead to chemotherapeutic resistance, tumor recurrence, and cell
696 migration [75, 76]. DNA damage, cellular and metabolic stress are known triggers of cell death
697 by apoptosis or senescence that can eliminate dysfunctional and dangerous conditions in cells.
698 According to our data, TMBP was able to induce cell cycle arrest, culminating in apoptosis
699 with activation of caspase-3 and -7. The functions of these proteases in apoptosis overlap,
700 although they have different functions [77], with caspase-3 controlling DNA fragmentation and
701 morphological changes, while caspase-7 is associated with loss of cell viability [78,79]. TMBP
702 treatment was also able to reduce STAT-3 protein immunolabeling, which favors reduced
703 migration, invasion, and metastasis.

704 Another important transcription factor involved with the evolution of neoplastic cells is
705 NF- κ B, which manifests a response against invading pathogens, and also promotes a metabolic
706 switch to glycolysis and controls the tumor microenvironment in several types of cancer [80].
707 Furthermore, NF- κ B communicates with PI3K/AKT signaling to promote the survival and
708 proliferation of lung cancer cells [81]. The PI3K/AKT/mTOR pathway represents an attractive
709 target for novel anticancer therapies in NSCLC, as it is also involved in the regulation of cell
710 growth, metabolism, and resistance to therapeutic agents [34]. Judging from our data, TMBP
711 was able to act on this pathway by reducing the levels of PI3K/AKT/NF- κ B.

712 Another molecule of interest in cancer is the enzyme ARG-1, which has been implicated
713 in cancer progression through several signaling pathways. ARG-1, when positively regulated,
714 increases the levels of phosphorylated AKT and ERK, leading to cancer cell viability and
715 proliferation. In addition, phosphorylated STAT-3 binds directly to the promoter region of
716 ARG-1, promoting a more aggressive cancer condition. Moreover, there is evidence that
717 patients with NSCLC have high plasma levels of ARG-1, which may be a promising target for
718 clinical use. ARG-1 may also regulate cellular NO production by competing with iNOS for the
719 substrate L-arginine. When produced by M2-type macrophages, ARG-1 can modulate NO
720 production and promote tissue repair and remodeling, production of collagen and other

721 components of the extracellular matrix. However, when iNOS metabolizes arginine to L-
722 citrulline and NO, the latter at high concentrations (μM range) induce cellular cytotoxicity,
723 DNA damage, nitrosation/oxidative stress, and apoptosis [82]. Thus, TMBP treatment was able
724 to decrease ARG-1 levels and increase iNOS in NCI-H460 cells.

725 The effective response of TMBP against the NCI-H460 tumor cell line was due to the
726 reduction in cell proliferation, invasive and metastatic activity, and induced morphological and
727 ultrastructural changes in the lung cancer cells by increasing toxic mediators as oxidative
728 species and metabolic stress, leading to a resolution of the tumor environment.

729

730 **5. Conclusion**

731 The development of therapies is a recent research trend as these are capable of reducing
732 side effects and can improve efficacy. The use of the biotransformed compound TMBP in lung
733 cancer has not been reported, and our study is the first to evaluate the effect of TMBP treatment
734 on the viability of lung cancer cells and its biological mechanism and focused on the
735 development of cell growth and apoptosis. TMBP was capable of reducing proliferation,
736 migration, and invasion, as well as morphological and ultrastructural changes in NCI-H460
737 cells. TMBP treatment increased oxidative stress parameters and induced metabolic changes
738 culminating in apoptotic death. TMBP negatively regulated proliferation-related proteins such
739 as PI3K/AKT/NF- κ B/ARG-1 and positively regulated the pro-oxidant profile mediated by the
740 iNOS pathway.

741 Considering that non-small cell lung cancer (NSCLC) encompasses diverse histological
742 subtypes, including large cell carcinoma, squamous cell carcinoma, adenocarcinoma, and
743 adenosquamous carcinoma, our study focuses primarily on the NCI-H460 large cell carcinoma
744 cell line and the A549 adenocarcinoma cell line, as indicated by previous research (Concato et
745 al., 2020). These results suggest the potential broad-spectrum efficacy of TMBP against various
746 lung cancer cells. However, it is essential to emphasize that additional investigations in different
747 NSCLC cell lines are imperative to confirm these findings.

748 In light of these promising outcomes, compounds with the ability to induce cancer cell
749 death are potential candidates for oncotherapy. Future studies should deeper into exploring the
750 potential tumor-suppressing effects of TMBP, thereby advancing our understanding and
751 potentially contributing to the development of novel therapeutic approaches for lung cancer.

752

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784

785 **Competing interests**

786 The authors declare that they have no competing interests.

787

788 **References**

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1046 **Trilobolide-6-*O*-isobutyrate from *Sphagneticola trilobata* acts by inducing**
1047 **oxidative stress, metabolic changes and apoptosis-like processes by caspase 3/7**
1048 **activation of human lung cancer cell lines**

1049

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1074 **Highlights**

- 1075 • TBB exhibits drug-likeness parameters and pharmacological characteristics.
- 1076 • TBB treatment reduces cell viability and alters morphology in A549 and NCI-H460.
- 1077 • The treatment reduces migratory capacity in tumor cell lines.
- 1078 • TBB treatment promotes oxidative and metabolic stress in A549 and NCI-H460.
- 1079 • TBB induces cell death through caspase activation in A549 and NCI-H460.

1080

1081 **Abstract**

1082 *Background:* Lung cancer, a chronic and heterogeneous disease, is the leading cause of cancer-
1083 related death on a global scale. Presently, despite a variety of available treatments, their
1084 effectiveness is limited, often resulting in considerable toxicity and adverse effects. Additionally,
1085 the development of chemoresistance in cancer cells poses a challenge. Trilobolide-6-*O*-
1086 isobutyrate (TBB), a natural sesquiterpene lactone extracted from *Sphagneticola trilobata*, has
1087 exhibited antitumor effects. Its pharmacological properties in NSCLC lung cancer, however,
1088 have not been explored.

1089 *Purpose:* This study evaluated the impact of TBB on the A549 and NCI-H460 tumor cell lines
1090 in vitro, examining its antiproliferative properties and initial mechanisms of cell death.

1091 *Methods:* TBB, obtained at 98% purity from *S. trilobata* leaves, was characterized using
1092 chromatographic techniques. Subsequently, its impact on inhibiting tumor cell proliferation in
1093 vitro, TBB-induced cytotoxicity in LLC-MK2, THP-1, AMJ2-C11 cells, as well as its effects on
1094 sheep erythrocytes, and the underlying mechanisms of cell death, were assessed.

1095 *Results:* *In silico* predictions have shown promising *drug-likeness* potential for TBB, indicating
1096 high oral bioavailability and intestinal absorption. Treatment of A549 and NCI-H460 human
1097 tumor cells with TBB demonstrated a direct impact, inducing significant morphological and
1098 structural alterations. TBB also reduced migratory capacity without causing toxicity at lower
1099 concentrations to LLC-MK2, THP-1 and AMJ2-C11 cell lines. This antiproliferative effect
1100 correlated with elevated oxidative stress, characterized by increased levels of ROS, superoxide
1101 anion radicals and NO, accompanied by a decrease in antioxidant markers: SOD and GSH. TBB-
1102 stress-induced led to changes in cell metabolism, fostering the accumulation of lipid droplets and
1103 autophagic vacuoles. Stress also resulted in compromised mitochondrial integrity, a crucial
1104 aspect of cellular function. Additionally, TBB prompted apoptosis-like cell death through
1105 activation of caspase 3/7 stressors.

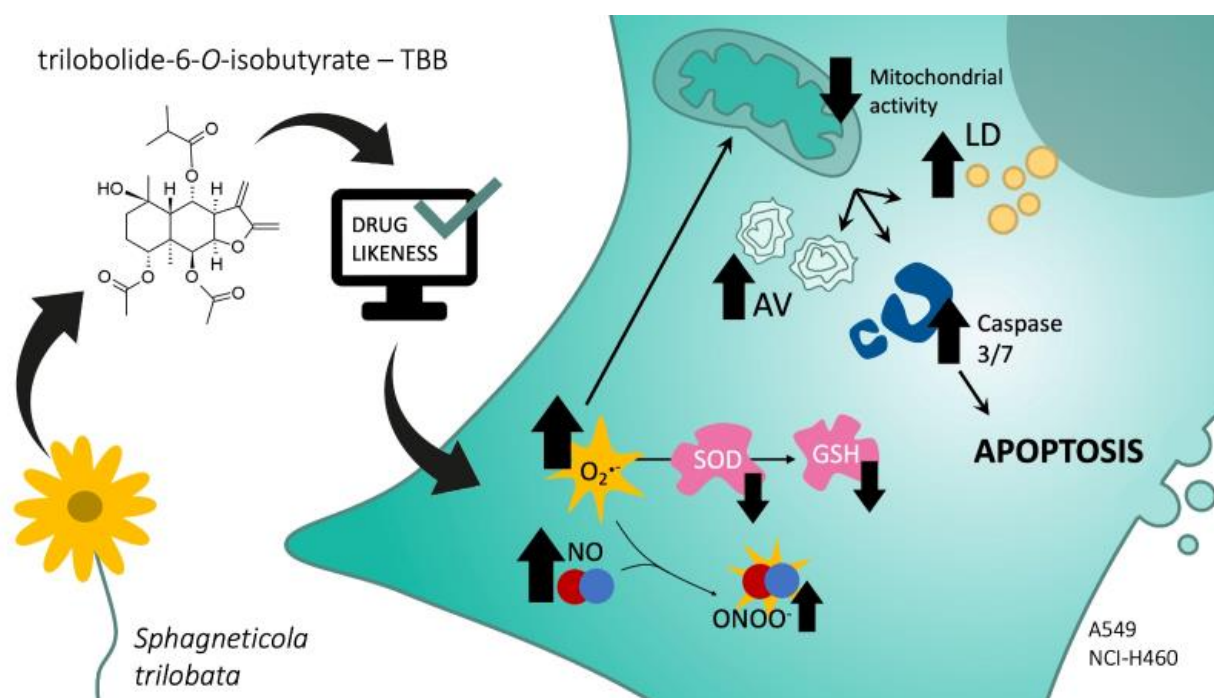
1106 *Conclusion:* These findings underscore the potential of TBB as a promising candidate for future
1107 studies and suggest its viability as an additional component in the development of novel
1108 anticancer drugs prototypes.

1109
1110

1111 **Key words:** NSCLC - A549, NCI-H460 cell lines; Natural Product; Oral bioavailability;
1112 Cytotoxicity; Cell Death.

1113

1114 Graphical abstract



1115

1116

1117 **Abbreviations:** ADMET, Absorption, distribution, metabolism, excretion and toxicity; CCCP,
1118 Carbonyl cyanide *m*-chlorophenylhydrazone; DMEM, Dulbecco's Modified Eagle Medium;
1119 DMSO, Dimethyl sulfoxide; DNA, Deoxyribonucleic acid; FBS, Fetal bovine serum; FDA, Food
1120 and Drug Administration; GSH, Reduced L-glutathione; H₂DCFDA, 2',7'-dichlorofluorescein
1121 diacetate; H₂O₂, Hydrogen peroxide; H-Acc, Number of Hydrogen bond acceptor; H-Don,
1122 Number of Hydrogen bond donor; IC₅₀, Inhibitory concentration at 50%; IL-6, Interleucin-6; LD,
1123 lipid droplets; MDC, Monodansylcadaverine; miLog P, Log of partition-coefficient; m.p,
1124 Melting point; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; MW,
1125 Molecular weight; NBT, Nitro blue tetrazolium; NH, Non-hemolytic; NMR, Nuclear magnetic
1126 resonance; NO, Nitric oxide; NR, Nile red; NSCLC, Non-small cell lung cancer; O₂^{•-},
1127 Superoxide anion radical; PBS, Phosphate-buffered saline; NMR, Magnetic resonance; PCNA,
1128 Proliferating cell nuclear antigen; P-gp, Permeability glycoprotein; PI, Propidium iodide; PMA,

1129 Phorbol myristate acetate; RB, Number of rotatable bonds; ROS, Reactive Oxygen Species;
1130 SCLC, Small cell lung cancer; SEM, Scanning electron microscopy; SL, Sesquiterpene lactone;
1131 SOD, Superoxide dismutase; TBB, Trilobolide-6-*O*-isobutyrate; TEM, Transmission electron
1132 microscopy; TMRE, Tetramethyl rhodamine-ethyl-ester; TPSA, Molecular Polar Surface Area;
1133 TSI, Tumor selectivity index; VLC, Vacuum liquid chromatography; CIS, cisplatin.

1134

1135 **1. Background**

1136 Lung-related cancers have presented more than 2.2 million novel cases worldwide, and
1137 in 2020 resulted in an estimated 1.8 million deaths (Sung et al., 2021, ranking this as the primary
1138 cause of cancer-related mortality globally. Lung cancer is a heterogeneous disease that
1139 encompasses various biologically distinct tumors. Broadly categorized into non-small cell lung
1140 cancer (NSCLC, 85% of total diagnoses) and small cell lung cancer (SCLC, 15% of total
1141 diagnoses). NSCLC is further segregated into adenocarcinoma - the prevailing subtype and is
1142 succeeded by squamous-cell carcinoma and large-cell carcinoma (Thai et al., 2021; Travis et al.,
1143 2015).

1144 Although there is a plethora of therapeutic options (chemotherapy, target therapy and
1145 immunotherapy) for systemic treatment, chemotherapy remains a critical component of lung
1146 cancer treatment, especially for patients with advanced stages of the disease, pre-existing
1147 autoimmune disease, and toxicity associated with combination immunotherapy. Consequently,
1148 platinum-based duplex regimens (e.g., cisplatin in combination with carboplatin or paclitaxel, or
1149 another cytotoxic drug) are commonly employed in this setting. The combination of cisplatin is
1150 a preferred treatment regimen, especially in combination with new-generation drugs, but it still
1151 results in adverse side-effects for patients, such as gastrointestinal toxicity, nephrotoxicity,
1152 neurotoxicity, ototoxicity, hypersensitivity, and myelosuppression. In addition, patients continue
1153 to experience disease recurrence and drug resistance during platinum-based therapy, which
1154 reduces the efficacy of NSCLC treatment (Guo et al., 2022; Lee, 2019; Mithoowani and
1155 Febbraro, 2022; Ryan et al., 2021).

1156 Plant-derived compounds possess a broad spectrum of biomedical applications, primarily
1157 attributed to their diminished adverse effects, cost-effectiveness relative to pharmaceuticals, and
1158 their endorsement by the Food and Drug Administration (FDA) (Atanasov et al., 2021; Dutta et
1159 al., 2019). *Sphagneticola trilobata* (L.) J.F. Pruski, also popularly known as "*margaridão*" and
1160 "*Wedelia*", belongs to the *Asteraceae* plant family and has been a traditional medicinal resource
1161 in India, China, the Caribbean, and Central and South America (Mardina et al., 2020) . Despite

1162 its classification as an invasive weed, this medicinal plant is noted for its high molecular diversity
1163 and its diverse biological activity that encompasses antibacterial, antidiabetic, hepatoprotective,
1164 antipyretic-analgesic, and anticancer activities (Shankar and Thomas, 2014; Mardina et al., 2020;
1165 Mardina et al., 2021; Zhang et al., 2021).

1166 The principal chemical constituents from *S. trilobata* are the ent-kaurane diterpenes,
1167 eudesmane sesquiterpene lactones, and triterpenes (Hui et al., 2019) . Recent investigations have
1168 demonstrated the therapeutic potential of sesquiterpene lactones (SLs), especially within the
1169 domain of cancer research (Dhyani et al., 2022; Dong et al., 2022; Shoaib et al., 2017) . Among
1170 these is trilobolide-6-*O*-isobutyrate (TBB), molecular formula $C_{23}H_{32}O_9$, a notable natural
1171 sesquiterpene lactone extracted from *S. trilobata* (Bohlmann et al., 1981).

1172 In a recent *in vitro* investigation utilizing TBB against the human hepatocellular carcinoma
1173 cell lines: HuH7, HepG2, SK-Hep1 and Bel7402, TBB from *S. trilobata* was found to inhibit
1174 growth and colony formation in a time- and concentration- dependent manner. TBB treatment
1175 exhibited inhibitory effects on migration, invasion, and glycolysis, ultimately culminating in cell
1176 cycle arrest within the G2/M phase, mitochondrial dysfunction, and apoptosis-driven cell death.
1177 TBB was furthermore, also able to regulate the expression of pivotal genes associated with
1178 proliferation, migration, and metastasis as PCNA (proliferating cell nuclear antigen), Ki67,
1179 Cyclin B1, Cyclin E, Bax, Bcl2, MMP2/9 (metalloproteinase 2/9) and PGK1 (phosphoglycerate
1180 kinase 1), primarily via the inhibition of the interleukin-6/signal transducers and activators of the
1181 transcription-3 (IL-6/STAT3) signaling pathway. The *in vivo* efficacy of TBB's anticancer
1182 attributes was validated through experimentation with a xenograft tumor model in male BALB/c
1183 nude mice, where TBB was effective in eliminating tumor growth (weight and volume) without
1184 causing toxicity to the experimental animals (Zhou et al., 2021).

1185 Here, we report for the first time, the biological activity of TBB isolated from *S. trilobata*
1186 in NSCLC cell lines, A549 (lung adenocarcinoma) and NCI-H460 (large cell carcinoma),
1187 employing diverse experimental methodologies. This study encompasses an evaluation of TBB's
1188 theoretical drug-likeness predictions and an investigation into its *in vitro* antiproliferative effects
1189 on NSCLC cell lines. Furthermore, we explore the potential mechanistic actions of TBB in
1190 inducing cell death and assess its cytotoxicity across a spectrum of cell types, including epithelial
1191 cells, monocytes, and erythrocytes.

1192

1193 **2. Material and Methods**

1194 **2.1 Animals and ethics committee**

1195 This study was approved by the Animal Ethics Committee of the State University of
1196 Londrina (approval N° 8286.2016.60 on 1/6/2016) and all methods performed were conducted
1197 following the Arouca Guideline and all ethical international regulations.

1198

1199 **2.1.1 Hemolytic Assay**

1200 Sheep red blood cells were used to evaluate direct membrane toxicity through the
1201 hemolytic assay. In this assay, erythrocytes were treated with TBB (3, 6, 12, 25, 50, 100, 200
1202 µM) or cisplatin (CIS) (12, 25, 50, 100, 200 µM) following the protocol established by Concato
1203 et al., 2020).

1204

1205 **2.2 Cell lines and reagents**

1206 We conducted a comprehensive characterization of a panel of cell lines, including Rhesus
1207 monkey renal epithelial cells (LLC-MK2), human leukemia monocytic cells (THP-1), murine
1208 alveolar macrophages (AMJ2-C11), and the established NSCLC lines, A549 and NCI-H460,
1209 sourced from the ATCC. All cells were cultured in Roswell Park Memorial Institute (RPMI)
1210 1640 medium (Life Technologies, CA, USA) supplemented with 10% fetal bovine serum (FBS)
1211 (GIBCO, Invitrogen, New York, USA), 100 U/mL Penicillin and 100 µg/mL Streptomycin
1212 (Santa Cruz Biotechnologies, Dallas, TX, USA) and maintained at 37 °C in 5% CO₂.

1213

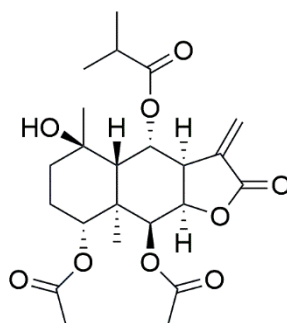
1214 **2.3 Plant material and isolation of trilobolide-6-O-isobutyrate (TBB)**

1215 The leaves of *S. trilobata* were collected in February 2013 at Horto de Plantas Medicinai
1216 do Centro de Ciências Agrárias da Universidade Estadual de Londrina (23°32' 64" N, 51°20' 63"
1217 W). A voucher specimen (N°. 49306, collected by one of the authors N.S.A.) was deposited at
1218 the Herbarium of the State University of Londrina, Brazil.

1219 After drying the leaves of *S. trilobata* to a constant mass, a crude extract was obtained
1220 through maceration with dichloromethane until the total exhaustion of the plant material.
1221 Subsequently, the extract was dried under reduced pressure using a rotary evaporator. The
1222 resulting crude extract was re-dissolved in a solvent mixture of 10% (v/v) methanol (MeOH) in
1223 water and subjected to sequential partitioning with hexane (hex) and dichloromethane. The
1224 solvent fractions were subsequently dried under reduced pressure. The dichloromethane fraction
1225 was further subjected to vacuum liquid chromatography (VLC) using an increasing polarity
1226 gradient (Hex: ethyl acetate (EtOAc): MeOH) and monitored by Comparative Thin Layer
1227 Chromatography (TLC). The fraction obtained with a ratio of 7:3 (EtOAc: Hex) was subjected

1228 to a subsequent round of VLC. Fraction F54, obtained during this process, yielded crystals (495
1229 mg) which were subjected to a low temperature acetone wash.

1230 The isolated compound (75 mg, 97% purity) (**Fig. 1**) by Ian Lucas Alves Cardoso was
1231 identified through ^1H and ^{13}C nuclear magnetic resonance (NMR) spectroscopy (Bruker Avance
1232 Model III, 400 MHz for ^1H and 100 MHz for ^{13}C), in CDCl_3 . The compound's identity was
1233 confirmed by comparison with spectral data obtained through literature values (Ferreira et al.,
1234 1994). The melting point (m.p) was determined using a hotplate device (Microquímica MQAPF
1235 302, Campinas, SP, Brazil). A stock solution of TBB was prepared in 1% dimethyl sulfoxide
1236 (DMSO) (GIBCO). The concentration of DMSO in the experiments never exceeded 0.06%.



1237

1238 **Figure 1:** Chemical structure of trilobolide-6-*O*-isobutyrate (TBB).

1239

1240 2.4 *In silico* studies for drug-likeness analysis

1241 An in-silico study was conducted to analyze the structure of TBB and assess various
1242 theoretical parameters. Initially, the Molinspiration Property Calculator program (accessible at:
1243 www.molinspiration.com) was used to determine parameters associated with oral bioavailability,
1244 in alignment with Lipinski's Rule of Five (Ro5) (Lipinski, 2004). Subsequently, Veber's rule
1245 (Veber et al., 2002) employed for further analysis. Following this, pharmacokinetic predictions
1246 were made regarding the similarity of drugs in terms of oral bioavailability, absorption,
1247 distribution, metabolism, excretion, and toxicity properties. These predictions were obtained
1248 from the SwissADME online database (<http://www.swissadme.ch/index.php>). The properties
1249 assessed included blood-brain barrier penetration, human intestinal absorption, analysis of the
1250 five most critical cytochrome P450 isoforms, AMES toxicity, carcinogens and carcinogenicity,
1251 and the human ether-à-go-go related gene (hERG)—a potassium channel crucial for cardiac
1252 repolarization. Additionally, an evaluation was conducted to determine whether TBB acts as a
1253 substrate for permeability glycoprotein (P-gp), which is responsible for assessing active efflux
1254 across biological membranes (Cheng et al., 2012; Yang et al., 2019).

1255

1256 **2.5 Viability assay**

1257 To assess the impact of TBB on the viability of LLC-MK2, THP-1 and AMJ2-C11 cell
1258 lines, we employed the MTT assay, following the protocol outlined by Mosmann (1983).
1259 Specifically, cells were seeded (1×10^4 cells/well) in 96-well plates and exposed to varying
1260 concentrations of TBB (12.5, 25, 50, 100, 200, and 400 μM) for a duration of 24 h at 37 °C in a
1261 5% CO₂ atmosphere. For A549 and NCI-H460 cells (1×10^4 cells/well), TBB was administered
1262 at concentrations of 3, 6, 12, 25, 50, and 100 μM , and cytotoxicity was evaluated at 24, 48, and
1263 72 h. The MTT assay was performed at the end of each respective incubation period, wherein
1264 cells were washed and MTT solution (0.05 mg/ml) was added, followed by a 3-h incubation.
1265 Formazan crystals formed were then diluted with 100 μL of DMSO and their optical density
1266 (OD) was measured at 540 nm using a spectrophotometer (Thermo Scientific, Multiskan GO).
1267 Experimental groups in this study were categorized as follows: a control group, a vehicle group
1268 (0.06% DMSO, representing the highest vehicle concentration used in treated groups), groups
1269 exposed to the specified TBB concentrations, and a CIS group (50 μM) serving as a positive
1270 control. The results were expressed as percentage of viability compared to the control group
1271 calculated with the following formula: % (viable cells) = (OD of TBB-treated samples/OD
1272 control-sample) \times 100.

1273

1274 **2.6.1 Half-maximal concentration (IC₅₀) calculation and tumor selectivity index (TSI)**

1275 The IC₅₀ curves for all cell lines were calculated using logarithmic regression. The values
1276 obtained for A549 and NCI-H460 were utilized to conduct further experiments. The TSI was
1277 determined through the following calculation: IC₅₀ cell lines / IC₅₀ of tumor cells at 24 h.

1278

1279 **2.6.2 Morphological and ultrastructural analysis of lung cancer cell lines**

1280 Scanning electron microscopy (SEM) was performed on A549 and NCI-H460 cells to
1281 analyze morphology cell surface following the procedure outlined by Tomiotto-Pellissier et al.,
1282 (2018). Briefly, cells (1×10^6 cells/well) were exposed to IC₅₀-TBB [A549 (73 $\mu\text{M} \pm 0.03$); NCI-
1283 H460 (54 $\mu\text{M} \pm 0.06$)] for 24 h at 37 °C. Subsequently, the cells were fixed with 2.5%
1284 glutaraldehyde in 0.1 M sodium cacodylate buffer (pH 7.4) and gradually dehydrated using
1285 increasing concentrations of ethanol (30-100%). Following dehydration, the samples underwent
1286 critical point drying (Baltec SCD-030, Balzer's, Liechtenstein), were gold-coated, and visualized
1287 using a high-resolution FEI SCIOS double-beam electron microscope (Thermo Fisher Scientific,
1288 Massachusetts, USA) for imaging purposes.

1289 To assess ultrastructure in the TBB-treated cells by transmission electron microscopy
1290 (TEM) (JEOL USA JEM 1400, Tokyo, Japan), cells were treated and fixed similarly as described
1291 above. The samples were then post-fixed with 1% OsO₄, 0.8% potassium ferrocyanide and 10.0
1292 mM CaCl₂ in 0.1 M sodium cacodylate buffer (pH 7.4) for 1 h at room temperature and protected
1293 from light. The samples were subsequently rinsed in 0.1 M sodium cacodylate buffer and
1294 dehydrated using increasing concentrations of acetone (30-100%). The acetone was gradually
1295 substituted with EPON™ epoxy resin (Electron Microscopy Sciences, Hatfield, United States)
1296 through cell diffusion, followed by allowing polymerization at 60 °C for 72 h. Subsequently, the
1297 resin-embedded cells were sliced (nanoscale sections, 60-70 nm) using an ultramicrotome
1298 (Power Tomer BMC - Germany), and then the sectioned slices were contrasted using 5% uranyl
1299 acetate and 2% lead citrate. In addition, the samples were subjected to analysis using TEM. The
1300 A549 and NCI-H460 cell lines, maintained in RPMI culture medium without TBB treatment,
1301 served as the negative control.

1302

1303 **2.6.3 Wound healing assay**

1304 A549 and NCI-H460 cells (1×10^6 cells/well) were plated in 6-well plates and incubated
1305 at 37 °C with 5% CO₂ to confluence. The cell monolayer was then gently "scratched" with a 200
1306 µL pipette tip, creating a gap in the monolayer where cells at the edges began migrating into the
1307 open space. Cells were then treated with IC₅₀-TBB [A549 (73 µM ± 0.03); NCI-H460 (54 µM ±
1308 0.06)] and photomicrographs (40× objective) were taken at different time points (0, 12 and 24 h)
1309 using an EVOS FL Auto2 microscope (Thermo Scientific). The extent of cell migration was
1310 evaluated by measuring the free area (the area without cells) using Image-Pro Plus software
1311 (ImageJ). The percentage decrease in area was quantified as the cell migration index. All assays
1312 were performed in triplicate for each tested group.

1313

1314 **2.6.4 Production of Reactive Oxygen Species (ROS)**

1315 ROS generation was assessed by the conversion of non-fluorescent 2',7'-
1316 dichlorofluorescein diacetate (H₂DCFDA) to highly fluorescent 2',7'-dichlorofluorescein (DCF)
1317 by intracellular free radicals. A549 and NCI-H460 cells (1×10^4 cells/well) were treated with IC₅₀-
1318 TBB [A549 (73 µM ± 0.03); NCI-H460 (54 µM ± 0.06)] for 24 h, and the assay performed
1319 according to Concato et al., (2020). Untreated cells were used as a negative control, and H₂O₂
1320 (0.04%) was used as a positive control. Data were normalized and expressed in arbitrary units.

1321

1322 **2.6.5 Determination of nitrite, an estimate of nitric oxide (NO) levels**

1323 NO levels were determined based upon the Griess method. A549 and NCI-H460 cells
1324 (1×10^5 cells/well) were treated with IC₅₀-TBB [A549 ($73 \mu\text{M} \pm 0.03$); NCI-H460 ($54 \mu\text{M} \pm$
1325 0.06)] and incubated for 24 h. The supernatant of the samples was collected and analyzed
1326 according to Cataneo et al., (2019).

1327

1328 **2.6.6 Superoxide anion radical production**

1329 The generation of O₂^{•-} was determined through the reduction of the redox dye nitro blue
1330 tetrazolium (1 mg/mL; NBT, Sigma-Aldrich). A549 and NCI-H460 cell lines (1×10^4 cells/well)
1331 were exposed to IC₅₀-TBB [A549 ($73 \mu\text{M} \pm 0.03$); NCI-H460 ($54 \mu\text{M} \pm 0.06$)] and incubated for
1332 24 h. Subsequently, the cells were washed with PBS, and 100 μL of NBT solution was added
1333 and incubated at 37 °C for 1 h. After centrifugation, the supernatant was discarded, and the
1334 formazan precipitate was dissolved by adding 60 μL of 2.0 M KOH and 60 μL of DMSO. The
1335 optical density was measured at 600 nm using a microplate reader (Multiskan GO Microplate
1336 Spectrophotometer, Thermo Scientific). Untreated cells served as the negative control, while
1337 treatment with phorbol myristate acetate (PMA) served as the positive control.

1338

1339 **2.6.7 Superoxide Dismutase (SOD)**

1340 For the determination of SOD activity, A549 and NCI-H460 cell lines (1×10^5 cells/well)
1341 were seeded in 6-wells plate, treated with the IC₅₀-TBB [A549 ($73 \mu\text{M} \pm 0.03$); NCI-H460 (54
1342 $\mu\text{M} \pm 0.06$)], and incubated for 24 h at 37 °C. Thereafter, the cells were lysed according to
1343 Concato-Lopes et al., (2023). The method used for assaying SOD activity was as described by
1344 Marklund and Marklund (1974), based on the enzyme's ability to inhibit auto-oxidation of
1345 pyrogallol. Total protein was determined on the cell lysate by the Lowry method (Lowry et al.,
1346 1951). The data were normalized as mg protein/mL. Untreated cells were used as negative
1347 control.

1348

1349 **2.6.8 Reduced glutathione (GSH)**

1350 To assess the degree of oxidative stress, the levels of reduced L-glutathione (GSH) were
1351 measured. Cells were seeded in 6-well plates at a density of 1×10^5 cells per well and treated with
1352 IC₅₀-TBB concentrations [A549 ($73 \mu\text{M} \pm 0.03$); NCI-H460 ($54 \mu\text{M} \pm 0.06$)]. Following a 24-h
1353 incubation period, GSH quantification was carried out following the methodology outlined by
1354 Rahman et al., (2006). Concentrations were determined by reference to a standard curve
1355 generated using serial dilutions of GSH. Untreated cells served as the negative control.

1356

1357 **2.6.9 Determination of Mitochondrial Membrane Potential ($\Delta\Psi_m$)**

1358 The evaluation of $\Delta\Psi_m$ was conducted through cell labeling with tetramethyl rhodamine-
1359 ethyl ester (TMRE) (Sigma-Aldrich, St. Louis, MO, USA). A549 and NCI-H460 cell lines (1×10^4
1360 cells/well) were exposed to IC₅₀-TBB concentrations [A549 ($73 \mu\text{M} \pm 0.03$); NCI-H460 ($54 \mu\text{M}$
1361 ± 0.06)] for 24 h, and the procedure followed the methodology outlined by Concato et al., (2020).
1362 Untreated cells served as the negative control, while carbonyl cyanide m-chlorophenylhydrazone
1363 (CCCP) ($100 \mu\text{M}$) was employed as the positive control. Data were expressed in arbitrary units.

1364

1365 **2.6.10 Detection of lipid droplets (LD)**

1366 A549 and NCI-H460 cells (1×10^4 cells/well) were treated with IC₅₀-TBB [A549 ($73 \mu\text{M}$
1367 ± 0.03); NCI-H460 ($54 \mu\text{M} \pm 0.06$)] for 24 h and analyzes performed by staining with Nile Red
1368 (NR) (Sigma-Aldrich) as described by Concato et al., (2020). Untreated cells were used as a
1369 negative control, and PBS treatment was used as a positive control. Data were normalized and
1370 expressed in arbitrary units.

1371

1372 **2.6.11 Determination of the formation of autophagic vacuoles**

1373 For the assessment of the formation of autophagic vacuoles, cells were labeled using
1374 monodansylcadaverine (MDC, Sigma-Aldrich). A549 and NCI-H460 cells (1×10^4 cells/well)
1375 were exposed to IC₅₀-TBB [A549 ($73 \mu\text{M} \pm 0.03$); NCI-H460 ($54 \mu\text{M} \pm 0.06$)] for 24 h. The
1376 assay procedure closely followed the methodology outlined by Concato et al., (2020). Untreated
1377 cells were used as a negative control, and PBS treatment was used as a positive control. Data
1378 were normalized and expressed in arbitrary units.

1379

1380 **2.6.12 Cell apoptosis assay**

1381 A549 and NCI-H460 cells (1×10^6 cells/well) were treated with IC₅₀-TBB [A549 ($73 \mu\text{M}$
1382 ± 0.03); NCI-H460 ($54 \mu\text{M} \pm 0.06$)] for 24 h and the analysis was performed using Annexin V
1383 and iodete propidium (PI) (Millipore, Billerica, MA, USA) labeling. The assay was achieved as
1384 described by Concato et al., (2020).

1385

1386 **2.6.13 Caspase-3/7 Assay**

1387 A549 and NCI-H460 cells (1×10^6 cells/well) were treated with IC₅₀-TBB [A549 ($73 \mu\text{M}$
1388 ± 0.03); NCI-H460 ($54 \mu\text{M} \pm 0.06$)] for 24 h and caspase-3/7 protease activity was measured
1389 using the EnzChek Caspase-3/7 Assay Kit (Invitrogen). The assay was achieved as described by
1390 Concato-Lopes et al., (2023). Data were normalized and expressed in arbitrary units.

1391

1392 2.7 Statistical analysis

1393 The data are presented as the mean \pm standard error of the mean (SEM). Three
1394 independent experiments were conducted, each with duplicate datasets, and subjected to analysis
1395 using GraphPad Prism 8 statistical software (GraphPad Software, Inc., USA, 500.288).
1396 Significant distinctions between groups were determined through t-tests or one-way ANOVA,
1397 followed by Tukey's test for multiple comparisons. A result was regarded statistically significant
1398 if the $p < 0.05$.

1399

1400 3 RESULTS

1401 3.6 Chemical and drug-likeness properties of TBB

1402 To validate the identification of the isolated compound (TBB; 495 mg), a thorough
1403 analysis was conducted using ^1H and ^{13}C NMR spectroscopy (**Supp. Tables 1-2**), which closely
1404 matched the values reported by (Ferreira et al., 1994). The m.p. of TBB was 218.52 °C For
1405 predicting the potential suitability of the molecule as an oral medication, Lipinski's (Ro5) and
1406 Veber rules were employed as reliable parameters. The results of the *in-silico* analysis of the
1407 TBB molecule indicated full compliance with the established rules, with no violation (**Table 1**).
1408 Theoretical predictions evaluated by the ADMET program revealed specific characteristics of
1409 TBB based on its chemical structure. Notably, the compound demonstrated a propensity for
1410 mitochondrial accumulation, high absorption in the human intestine, and was identified as a
1411 substrate and inhibitor of permeability glycoprotein. Importantly, TBB does not inhibit the
1412 human ether-a-go-go related gene (hERG), lacks carcinogenic components in its structure, does
1413 not induce toxicity according to the Ames test, and is non-carcinogenic. Furthermore, in
1414 accordance with theoretical predictions, TBB exhibited the potential to penetrate the blood-brain
1415 barrier and displayed interactions with several cytochrome P450 (CYP) isoforms, specifically
1416 CYP1A2, CYP2C9, CYP2D6, and CYP2C19. Also presented low CYP inhibitory promiscuity,
1417 *i.e.*, it does not alter the general functionality of CYP450 (**Table 2**) (Cheng et al., 2012; Yang et
1418 al., 2019).

1419

1420 **Supp. Table 1:** ^1H NMR data (400 MHz, CDCl_3 , J in Hz, τ_{H} in ppm)

Position	Ferreira et al. (1994)	This work
	τ_{H} (ppm)	TBB τ_{H} (ppm), m , J (Hz)
1	4.59 (<i>dd</i> , 10.5, 4.2)	4.59 (<i>dd</i> , 10.6, 4.2)

2	1.7 - 1.5	1.8 - 1.5
3	1.7 - 1.6	1.7 - 1.6
5	1.88 (<i>d</i> , 3.0)	1.88 (<i>d</i> , 2.9)
6	5.98 (<i>dl</i> , 3.0)	5.95 (<i>dl</i> , 3.0)
7	3.30 (<i>m</i>)	3.16 (<i>m</i>)
8	4.91 (<i>dd</i> , 8.1; 4.5)	4.88 (<i>dd</i> , 8.0, 4.5)
9	5.24 (<i>d</i> , 4.5)	5.21 (<i>d</i> , 4.5)
13 α	6.26 (<i>d</i> , 3.8)	6.23 (<i>d</i> , 3.8)
13 β	5.65 (<i>d</i> , 3.5)	5.66 (<i>d</i> , 3.5)
14	1.38 (<i>s</i>)	1.36 (<i>s</i>)
15	1.37 (<i>s</i>)	1.34 (<i>s</i>)
2'	2.6 (<i>spt</i> , 7.0)	2.58 (<i>spt</i>)
3'	1.23 (<i>d</i> , 7.0)	1.22 (<i>d</i> , 7.0)
4'	1.22 (<i>d</i> , 7.0)	1.20 (<i>d</i> , 7.0)
2''	1.99 (<i>s</i>)	2.02 (<i>s</i>)
2'''	1.96 (<i>s</i>)	1.93 (<i>s</i>)

(*dd*) double doublet, (*d*) doublet, (*dl*) doublet large, (*m*) multiplets, (*s*) singlet, (*spt*) septet.

1421

1422

1423

1424

Supp. Table 2: ^{13}C NMR data (100 MHz, CDCl_3 , $^{\text{TM}}$ in ppm)

Position	Ferreira et al. (1994)	This work
	$^{\text{TM}}_{\text{C}}$ (ppm)	TBB $^{\text{TM}}_{\text{C}}$ (ppm)
1	72.9	73.0
2	23.7	24.0
3	41.3	41.6
4	70.6	70.6
5	43.1	43.4
6	67.9	68.0
7	44.5	44.9
8	71.9	72.1
9	70.4	71.0

10	41.3	41.3
11	133.9	134.1
12	168.9	169.2
13	118.9	119.2
14	14.0	14.3
15	26.1	26.4
1'	175.9	176.2
2'	34.1	34.4
3'	18.2	18.4
4'	18.7	19.0
1''	170.3	170.6
2''	20.0	20.2
1'''	169.3	169.4
2'''	20.8	21.0

1425

1426 **Table 1:** Molecular properties of trilobolide-6-*O*-isobutyrate (TBB) based on Lipinski and
1427 Veber's criteria.

		Limits	TBB
Lipinski's rules	MW	≤500	452.50
	miLog P	≤5	1.72
	H-Acc	≤10	9
	H-Don	≤5	1
Veber's rules	RB	≤10	7
	tPSA	≤140°Å ²	125.45
	H-Acc + H-Don	≤12	10
N° of Violations		None	

1428 **MW:** molecular weight; **miLog P:** Log of partition-coefficient; **H-Acc:** number of Hydrogen bond acceptor; **H-**
1429 **Don:** number of Hydrogen bond donor; **RB:** number of rotatable bonds; **tPSA:** molecular polar surface area.
1430

1431 **Table 2:** Prediction of pharmacological parameters evaluating TBB using ADMETSar toolbox.

ADMET Properties	TBB*
------------------	------

Subcellular localization	Mitochondria
Blood-Brain Barrier permeant	High
Human Intestinal Absorption	High
Caco-2 Permeability	Non
P-glycoprotein Substrate	Yes
P-glycoprotein Inhibitor	Yes
CYP450 2C9 Substrate	Non
CYP450 2D6 Substrate	Non
CYP450 3A4 Substrate	Yes
CYP450 1A2 Inhibitor	Non
CYP450 2C9 Inhibitor	Non
CYP450 2D6 Inhibitor	Non
CYP450 2C19 Inhibitor	Non
CYP450 3A4 Inhibitor	Yes
CYP Inhibitory Promiscuity	Low
hERG	Non
AMES Toxicity	Non
Carcinogens	Non
Carcinogenicity	Non
Predicted aqueous solubility	Soluble

hERG, human Ether-a-go-go-Related Gene Inhibition. * (Cheng et al., 2012; Yang et al., 2019).

1432
1433
1434

1435 **3.7 TBB exhibits antiproliferative activity in A549 and NCI-H460 tumor cells and low**
1436 **toxicity in epithelial, alveolar macrophage, monocytic and blood red cells.**

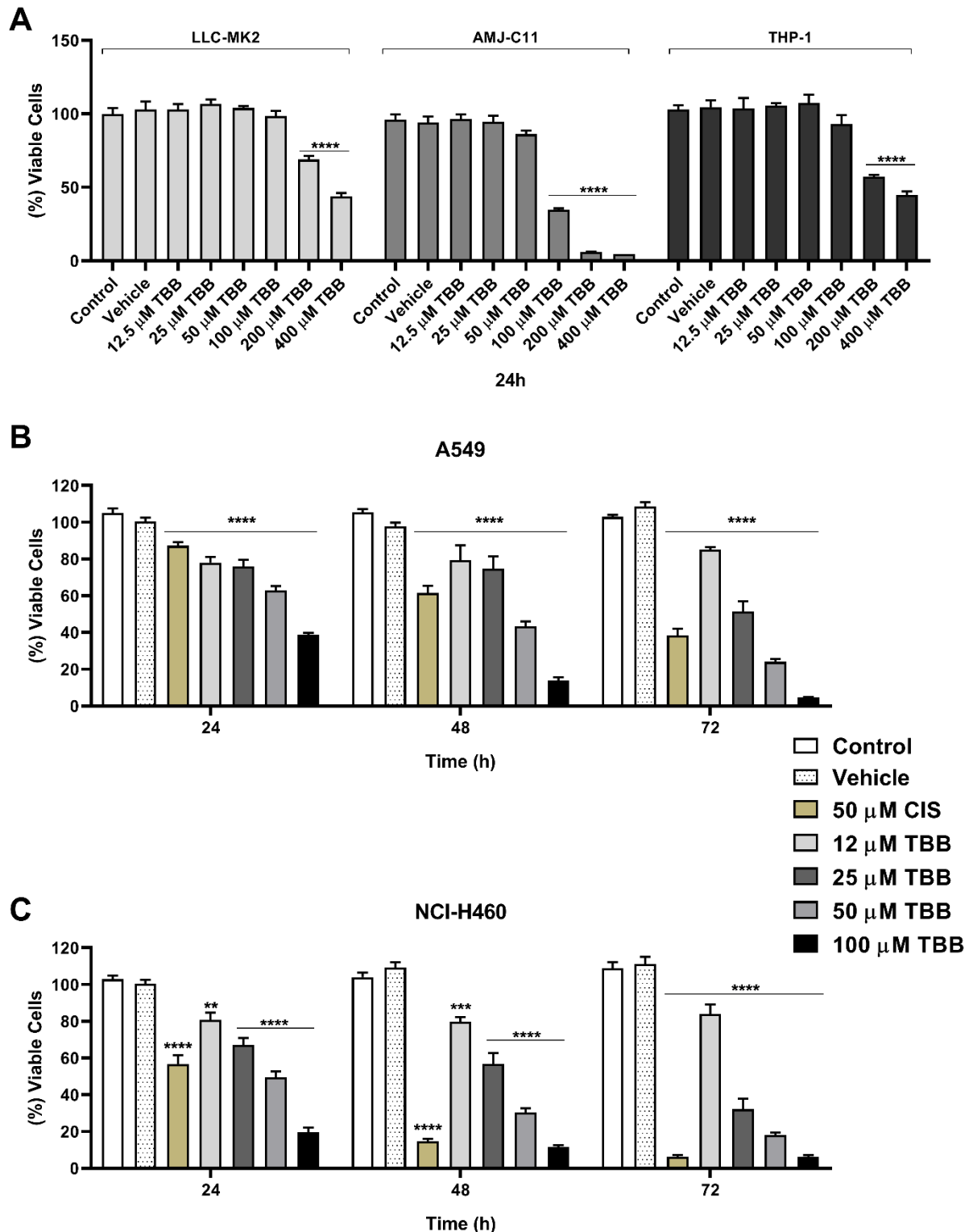
1437 The potential cytotoxicity of TBB was evaluated in epithelial, alveolar macrophage,
1438 monocytic, blood red cells and lung tumoral cell lines. Initially, we investigated whether
1439 treatment with TBB altered the viability of renal epithelial cells (LLC-MK2 from monkey),
1440 mouse alveolar macrophage (AMJ2-C11), human-derived leukemia monocytes (THP-1), and
1441 sheep erythrocytes employing the MTT and hemolytic assays. AMJ2-C11 cell line showed
1442 toxicity above 100 μ M TBB ($p < 0.0001$), while LLC-MK2 and THP-1 showed toxicity above
1443 200 μ M TBB (both $p < 0.0001$), when compared to control (**Fig. 2A**). Data from the hemolytic
1444 assay demonstrated that TBB did not cause hemolysis of sheep erythrocytes at any of the
1445 concentrations tested (3-200 μ M), while CIS (12-200 μ M) caused hemolysis at 50, 100 and 200

1446 μM ($p < 0.001$, $p < 0.01$, $p < 0.0001$, respectively) (data not shown), confirming the low toxicity
1447 of TBB for these red blood cells.

1448 The effect of TBB on the lung cancer cell lines A549 and NCI-H460 was investigated for
1449 24, 48 and 72 h (**Fig. 2B-C**). Treatment of A549 cells with TBB for 24 h at a concentration of
1450 100 μM was able to reduce the viability of the cells by 61% ($p < 0.0001$). Treatment with TBB
1451 at concentrations of 50 μM and 100 μM for 48 h reduced cell viability by more than 50% ($p <$
1452 0.0001), while at 72 h, we observed that concentrations of 25, 50, and 100 μM reduced cell
1453 viability by 51.35%, 78.85%, and 98.30%, respectively ($p < 0.0001$) (**Fig. 2B**). Further, in the
1454 NCI-H460 cells treated with TBB at concentrations of 50 and 100 μM for 24 h, cell viability was
1455 reduced by 53.44% and 83.38%, respectively ($p < 0.0001$). At 48 h, concentrations of TBB of
1456 25, 50 and 100 μM caused reduced cell viabilities of 46.01%, 72.48%, and 91.34%, respectively
1457 ($p < 0.0001$), while at 72 h, we observed that treatment with 25, 50, and 100 μM TBB caused
1458 reductions of 70.73%, 84.82%, and 96.64%, respectively ($p < 0.0001$) (**Fig. 2C**). The vehicle
1459 (DMSO 0.06%) exhibited no toxic effect at any of the time-points tested, and CIS (50 μM) as a
1460 positive control, inhibited cell viability in both lung cancer lines.

1461 The IC_{50} values of TBB were determined for LLC-MK2, THP-1, and AMJ2-C11 cells
1462 following a 24-h treatment period (**Table 3**). In the case of the tumor cells, A549 and NCI-H460,
1463 IC_{50} values were calculated at 24, 48, and 72 h (**Table 3 and Suppl. Table 3**). The TSI was also
1464 calculated, where a favorable TSI > 1.0 indicates a drug with efficacy against tumor cells greater
1465 than toxicity to normal cells. Our data showed TSI values >1 for both tumor cells relative to
1466 epithelial, monocytic and alveolar macrophage cells (**Table 3**).

1467



1468

1469 **Figure 2: TBB exhibited superior efficacy against tumor cells and low toxicity to epithelial,**
 1470 **monocytic, and alveolar macrophage.** Cell lines LLC-MK2, THP-1, and AMJ2-C11 (A)
 1471 treated with TBB at different concentrations (12.5-400 μ M) for 24 h and analyzed by the MTT
 1472 assay. Lung cancer cell lines A549 (B) and NCI-H460 (C) treated with TBB (12 - 100 μ M) for
 1473 24 h, were also evaluated by MTT. Values are expressed as mean \pm SEM. Significant difference

1474 with $**p \leq 0.01$, $***p \leq 0.001$, and $****p \leq 0.0001$ vs. control (untreated cells); vehicle (DMSO
 1475 0.06%); CIS (cisplatin, 50 μM).

1476

1477 **Table 3**

1478 IC_{50} of cell lines: LLC-MK2, THP-1 and AMJ2-C11, and TSI of lung cancer cells A549 and
 1479 NCI-H460 following treatment with trilobolide-6-*O*-isobutyrate (TBB).

Compound	LLC-MK2 IC_{50} μM ($\pm\text{SEM}$)	THP-1 IC_{50} μM ($\pm\text{SEM}$)	AMJ-C11 IC_{50} μM ($\pm\text{SEM}$)	A549 IC_{50} μM ($\pm\text{SEM}$)	NCI-H460 IC_{50} μM ($\pm\text{SEM}$)
TBB	335 μM (± 0.01)	305 μM (± 0.04)	81 μM (± 0.03)	73 μM (± 0.06)	54 μM (± 0.03)
TSI A549	5	4	1	-	-
TSI NCI-H460	6.2	6	1.5	-	-

1480 Rhesus monkey kidney epithelial cells (LLC-MK2), human leukemia monocytic cell line (THP-1),
 1481 alveolar macrophage murine (AMJ2-C11), lung adenocarcinoma (A549) and large cell lung cancer (NCI-
 1482 H460). SEM – standard error of the mean.

1483

1484 **Supp. Table 3:** Half-maximal inhibitory concentration (IC_{50}) in lung cancer cells at 24-72 h after
 1485 treatment with TBB.

Cell line	IC_{50} ($\mu\text{M} \pm \text{SEM}$)		
	Time of treatment (h)		
	24	48	72
A549	73 (± 0.06)	47 (± 0.03)	27 (± 0.02)
NCI-H460	54 (± 0.03)	30 (± 0.03)	21 (± 0.02)

1486

SEM – standard error of the mean.

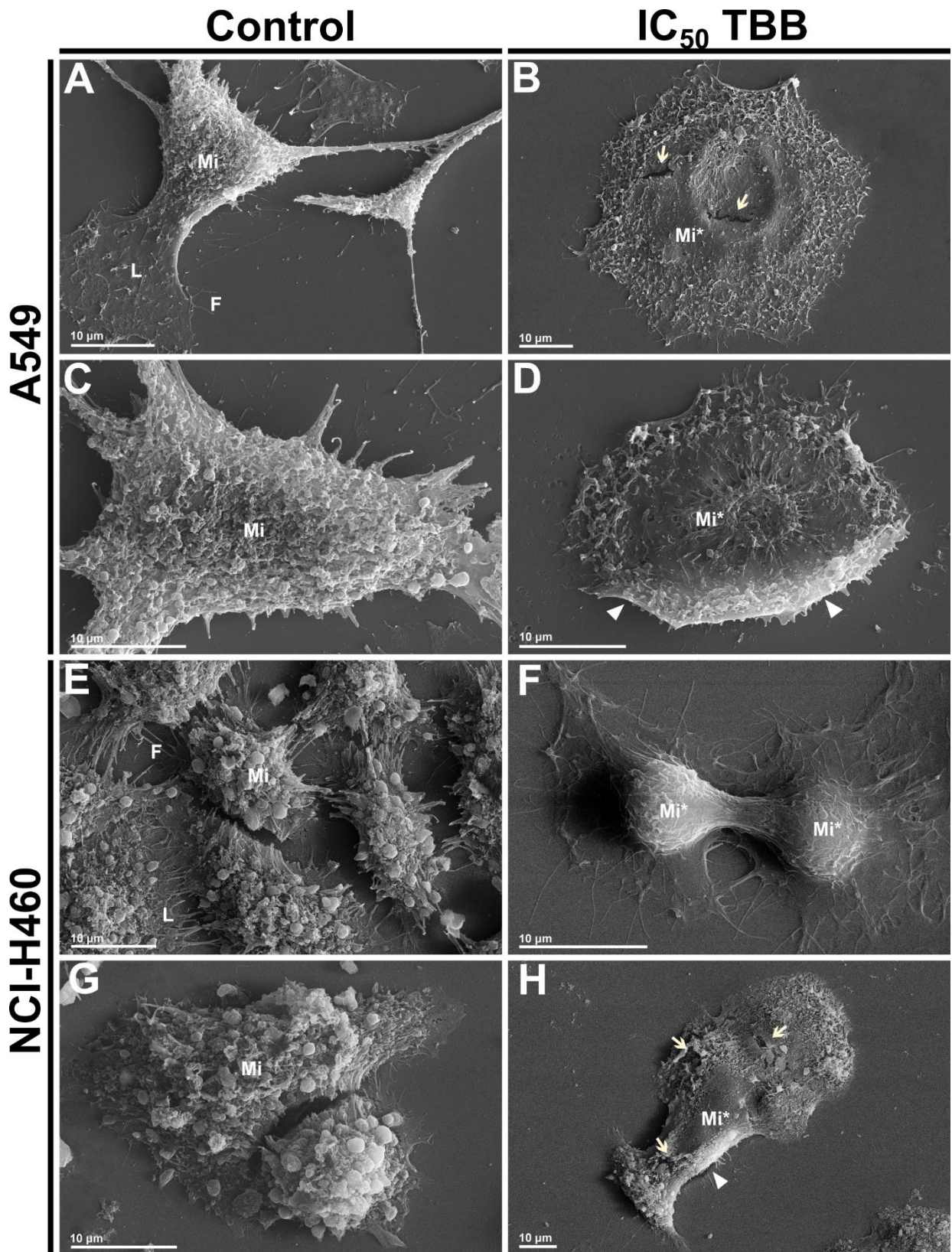
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1488 **3.8 TBB induces morphological and ultrastructural changes in lung cancer cell lines**
 1489 **treated with TBB**

1490 Having observed the inhibitory effect of TBB on the proliferation of the lung cancer cells,
 1491 a comprehensive investigation was conducted to assess its impact on the morphological and
 1492 ultrastructural aspects of A549 and NCI-H460 cells using SEM and TEM. SEM analysis revealed
 1493 distinct morphological changes: untreated A549 cells (**Fig. 3A, C**) exhibited typical
 1494 characteristics, such as strong adhesion to the glass coverslips, presence of an intact cell
 1495 membrane, the presence of lamellipodia (L), filopodia (F) and microvilli (Mi) across the cell

1496 surface, and well-preserved cellular structures. Following treatment with IC₅₀-TBB, these cells
1497 displayed altered morphologies, characterized by a transition to rounded shapes, reduced
1498 lamellipodia, irregularities on the cell surface (arrowhead), diminished or absent microvilli
1499 (Mi*), and evidence of nuclear and cytoplasmic disruptions (arrow) (**Fig. 3B, D**). Similarly,
1500 untreated NCI-H460 cells showed a normal appearance consistent with the features (**Fig. 3E, G**).
1501 Upon treatment with TBB, NCI-H460 cells also showed reduced or absent microvilli (Mi*),
1502 irregularities on the cell surface (arrowhead), and along with nuclear and cytoplasmic disruptions
1503 (arrow) (**Fig. 3F, H**).

1504 Ultrastructural evaluation by TEM showed that the control groups of A549 and NCI-
1505 H460 cells presented a homogeneous cytoplasm with regular organelles (mitochondria and
1506 endoplasmic reticulum) and without the presence of vacuole-like structures, indicating whole cell
1507 integrity (**Fig. 4A, C and E, G**). After treatment with TBB, A549 showed the absence of
1508 microvilli (arrowhead), changes in mitochondrial structures such as swelling and enlargement of
1509 these organelles (m*). Accumulation of lipid storage droplets (*) and autophagic vacuoles
1510 (arrow) were also observed. In NCI-H460 cells, we observed reduced microvilli (Mi*),
1511 accumulation of lipid storage droplets (*) and autophagic vacuoles (arrow) following treatment
1512 with TBB. Furthermore, the presence of substances inside the vacuoles can be observed (**Fig.**
1513 **4B, D and F, H**).



1514

1515

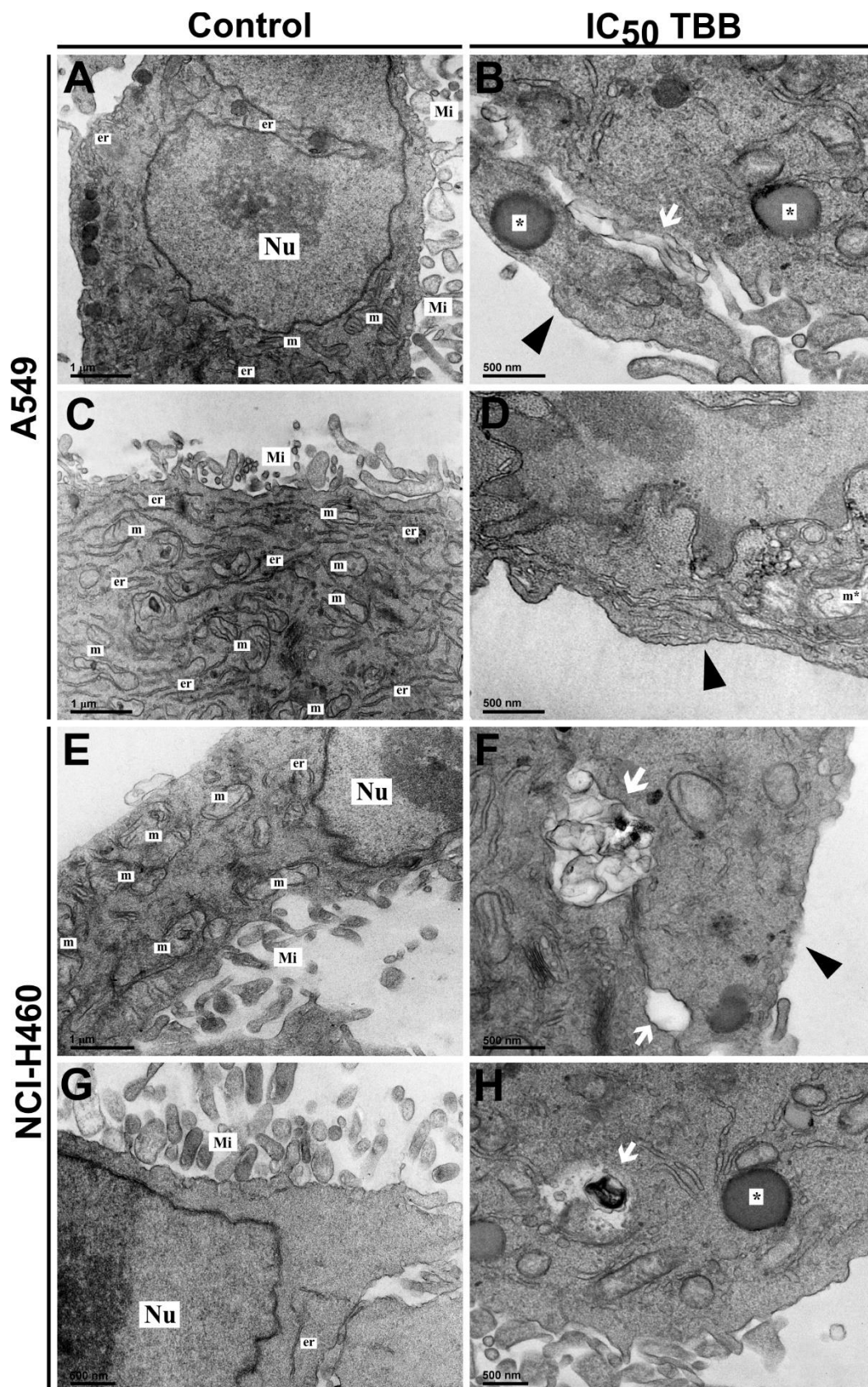
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Figure 3: Effect of TBB treatment on the morphology of lung cancer cell lines. Morphological changes observed by scanning electron microscopy after treatment with TBB for 24 h. **(A and C)** Control A549 cell line, untreated cells; **(B and D)** A549 cells treated with IC₅₀ of TBB (73 μM). **(E and G)** Control NCI-H460 cell line, untreated cells; **(F and H)** NCI-H460

1519 cells treated with IC₅₀ of TBB. (54 μM). (Mi) microvilli, (L) lamellipodia and (F) filopodia,
 1520 altered microvilli (Mi*), (arrowhead) lamellipodia and filopodia alterations, (arrow) cell
 1521 disruption. Scale bars = 10 μm (A-I).

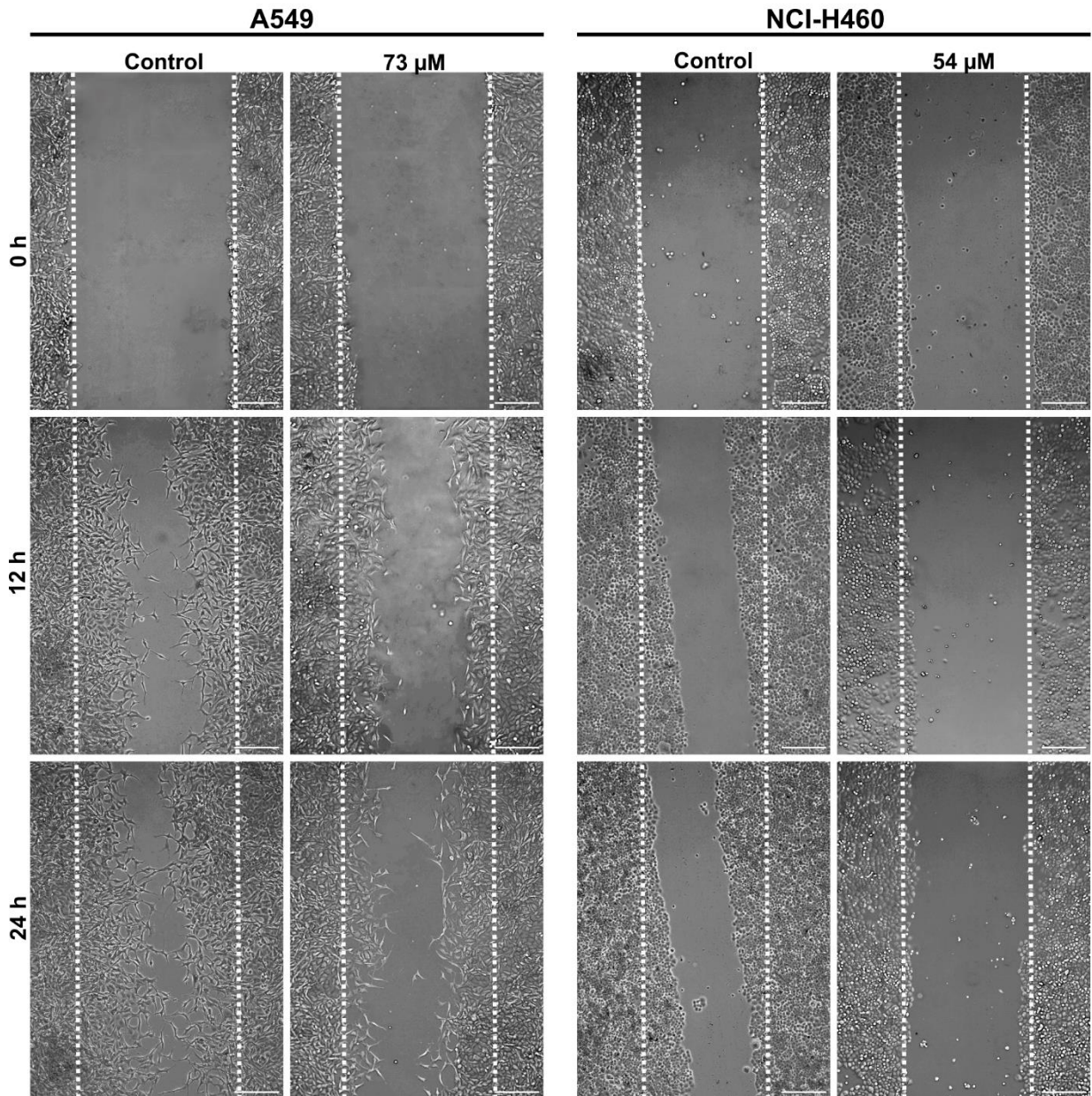
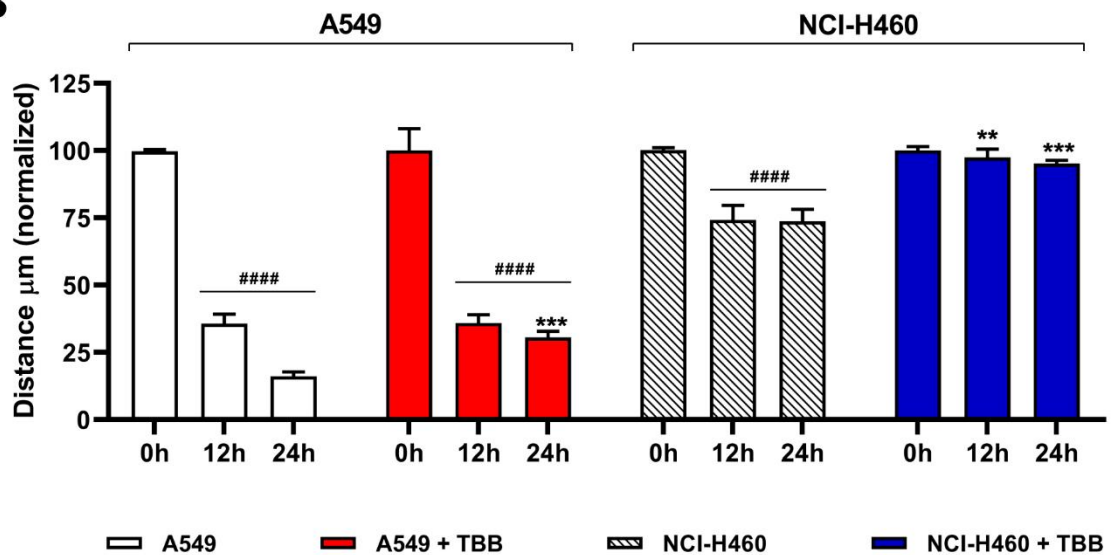


1523 **Figure 4: Ultrastructural changes in lung cancer cell lines treated with TBB for 24 h and**
1524 **analyzed by transmission electron microscopy. (A and C, A549; E and G, NCI-H460)**
1525 **untreated cells; (B and D, A549; F and H, NCI-H460) cells treated with IC₅₀ TBB (73 and 54**
1526 **μM, respectively); (m) mitochondria; (Gc) Golgi complex; (er) endoplasmic reticulum; (m*)**
1527 **mitochondrial swelling; (Nu) nucleus; (arrow) cytoplasmic autophagic vacuole; (arrowhead)**
1528 **absence of microvilli; (*) lipid droplets storage areas. Scale bars = 1 μm (A, C), 600 nm (B) and**
1529 **500 nm (D-F).**

1530

1531 **3.9 TBB reduces migratory capacity in A549 and NCI-H460 cell lines.**

1532 A wound healing assay was performed to evaluate migration of lung cancer cells after
1533 treatment with TBB (**Fig. 5**). Cell movements at the wound margins were photographed
1534 immediately at the start of scratching and at regular time intervals (0, 12, and 24 h) (**Fig. 5A**).
1535 Thereafter quantification, the control group of A549 cells showed an increase in migration in
1536 relation to the elapsed time, while the TBB-treated group showed a reduction in migration time
1537 at 24 h ($p \leq 0.001$) compared to the untreated group (**Fig. 5B**). A549 cells also showed a
1538 significant difference between the time points analyzed, with an increase in migration at 12 and
1539 24 h ($p \leq 0.0001$) compared to 0 h. For NCI-H460 cells, the migration effect was even more
1540 pronounced. The untreated group also showed an increase in migratory capacity, but the treated
1541 group showed a discrepant reduction at 12 h ($p \leq 0.01$) and 24 h ($p \leq 0.001$) (**Fig. 5B**). It is
1542 noteworthy that there were no discernible variations in the temporal outcomes among the TBB-
1543 treated group for this cell line, elucidating the compound's impact during the initial phases of
1544 treatment.

A**B**

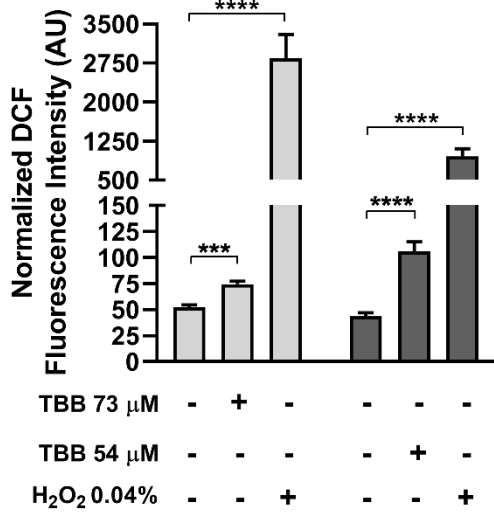
1546 **Figure 5: TBB treatment inhibited lung cancer cell line's migration.** The scratch assay
1547 procedure was performed to observe the effect of TBB treatment on cell migration over time (0,
1548 12 and 24 h) (**A**). The distance of the free surface of these treated cells (**B**) was quantified. Values
1549 represent the mean \pm SEM of three different experiments performed in triplicate. Significant
1550 difference versus control ** ($p \leq 0.01$); *** ($p \leq 0.001$). Significant difference inters groups #####
1551 ($p < 0.0001$). Scale bar = 200 μ m.

1553 **3.10 TBB promotes oxidative stress in lung cancer cells by increasing free radicals and** 1554 **reducing the activity of *in-situ* antioxidant enzymes.**

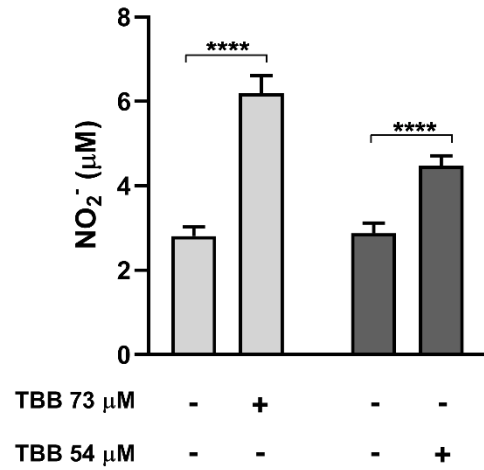
1555 As treatment of lung cancer cells with TBB demonstrated antiproliferative activity and
1556 halted cell migration, we investigated whether TBB would act as a pro- or antioxidant in these
1557 cells. Therefore, in evaluating total ROS production after 24 h, treatment with TBB showed high
1558 levels of these reactive species in the A549 (22.23 AU \pm 4.15) and NCI-H460 (62.61 AU \pm 9.52)
1559 (**Fig. 6A**) cell lines. One of the reactive oxygen species produced intracellularly mainly by
1560 mitochondria is $O_2^{\cdot-}$. When evaluating whether TBB would alter the production of these radicals,
1561 we found that the treatment promoted a significant increase in both A549 ($p = 0.0075$) and NCI-
1562 H460 ($p < 0.0001$) (**Fig. 6C**) cell lines. Reactive nitrogen species also have a highly potent pro-
1563 oxidant capacity. When evaluating the levels of NO after treatment with TBB, both A549 and
1564 NCI-H460 showed an increase of levels ($p \leq 0.0001$) compared to the control groups (**Fig. 6B**).

1565 Parameters related to the antioxidant profile were also altered by TBB-treatment of the
1566 cells. The data showed a decrease in SOD activity and level of GSH for the A549 ($p = 0.0086$, p
1567 = 0.0009, respectively) and NCI-H460 ($p = 0.0445$, $p = 0.0002$, respectively) cell lines (**Fig. 6D-**
1568 **E**). Overall, these data suggest that TBB acts by increasing oxidative stress in these cancer cells.

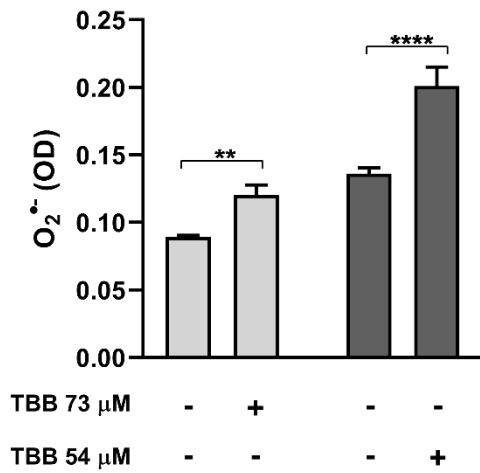
A Reactive Oxygen Species



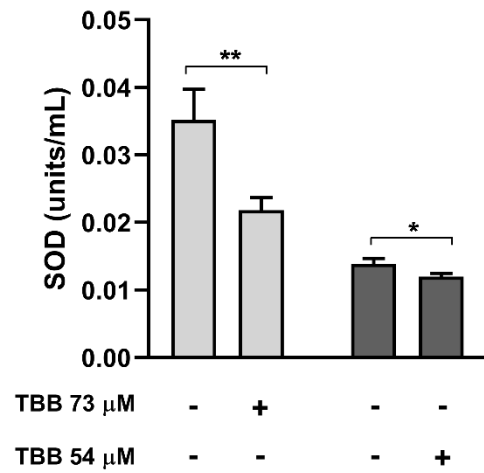
B Nitric Oxide



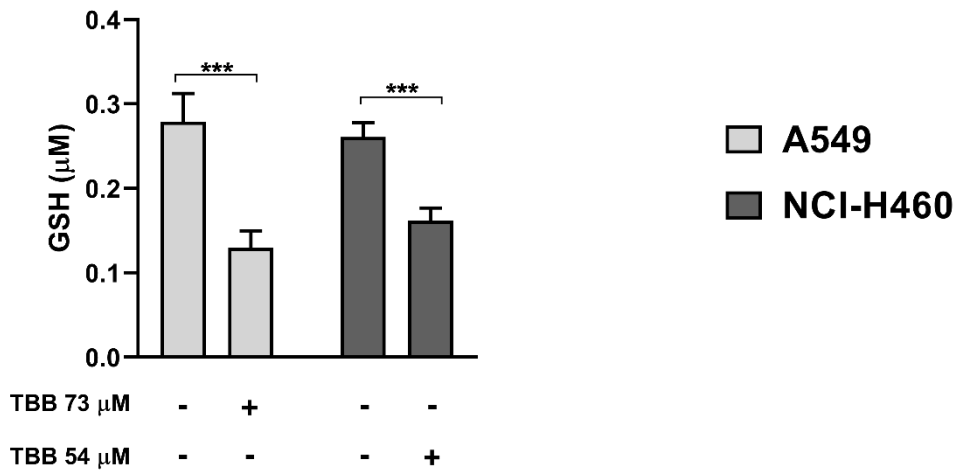
C Superoxide Anion Radical



D Superoxide Dismutase



E Reduced L-glutathione



1570 **Figure 6: The TBB treatment of lung cancer cells trigger oxidative stress, while decreasing**
1571 **levels of antioxidant molecules.** Both A549 and NCI-H460 cells were exposed to TBB at IC₅₀
1572 concentrations (73 μM for A549, and 54 μM for NCI-H460) for a duration of 24 h. To measure
1573 the extent of oxidative stress, ROS levels were assessed using the H₂DCFDA probe (A). Nitric
1574 oxide (NO) levels were quantified using the Griess method (B). Additionally, superoxide radical
1575 anion (O₂^{•-}) levels were determined through the NBT assay (C). The activity of superoxide
1576 dismutase (SOD) (D), and the reduced levels of glutathione (GSH) (E) were evaluated in these
1577 lung cancer cell lines. Data are expressed as mean ± SEM of three independent experiments
1578 performed in triplicate. * ($p < 0.05$); ** ($p \leq 0.01$); *** ($p \leq 0.001$); **** ($p \leq 0.0001$) vs. control.
1579

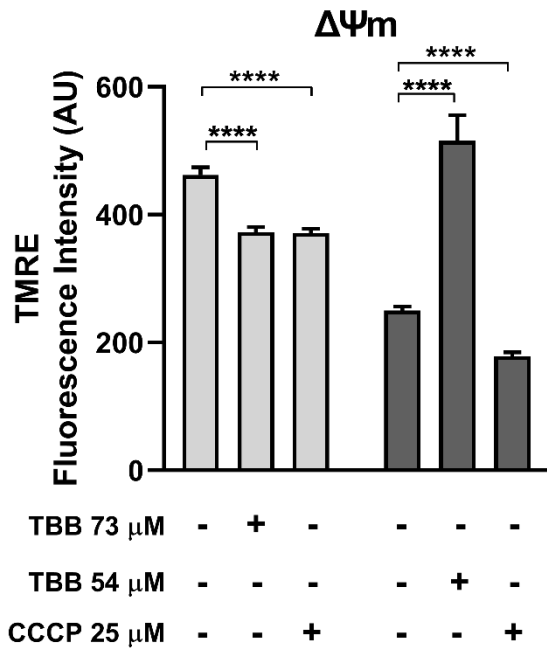
1580 3.11 TBB causes metabolic stress in A549 and NCI-H460 cancer cells

1581 Cellular metabolism is essential for the maintenance and acquisition of energy. We
1582 evaluated the $\Delta\Psi_m$ using the TMRE assay and found that treatment with TBB in A549 cells
1583 caused depolarization of $\Delta\Psi_m$, with a decrease in the total fluorescence intensity of TMRE
1584 compared to the control ($p < 0.0001$) (Fig. 7A), indicating a loss of organelle integrity. In NCI-
1585 H460 cells, however, the opposite effect was obtained, i.e., treatment with TBB caused
1586 hyperpolarization of $\Delta\Psi_m$, and induced an increase in TMRE-mitochondria complexation ($p <$
1587 0.0001) (Fig. 7A), indicating an increase in function and damage in this organelle.

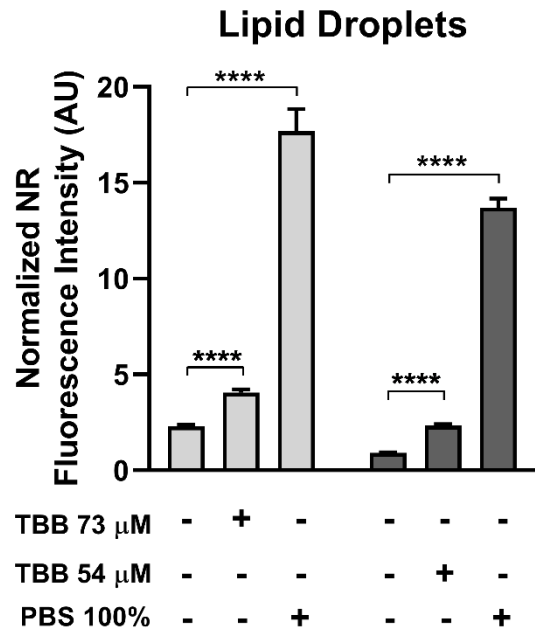
1588 Considering the observed mitochondrial alterations, our investigation then assessed the
1589 formation of lipid droplets (LDs) and autophagic vacuoles in the TBB-exposed cells, with the
1590 objective to substantiate the presence of metabolic and/or energy imbalances induced by TBB
1591 treatment of the lung cancer cells. Nile Red (NR) fluorescence was utilized since it serves as a
1592 reliable indicator of LDs. The results clearly demonstrated a substantial increase in LD formation
1593 in both A549 ($p < 0.0001$) and NCI-H460 ($p < 0.0001$) cells following treatment with TBB, as
1594 compared to the control group (Fig. 7B). This observation supports the premise of altered cellular
1595 metabolism and energy dynamics induced by treatment with TBB.

1596 The presence of autophagic vacuoles was identified using monodansylcadaverine (MDC)
1597 fluorescence. Notably, we observed a significant elevation in fluorescence intensity in A549
1598 (58.68 AU ± 4.12) and NCI-H460 (82.29 AU ± 2.06) cells post-TBB treatment (Fig. 7C). These
1599 findings align with the results presented in figure 4, where TBB induced marked changes in the
1600 cellular environment, indicative of an adaptive process aimed at maintaining cell viability.

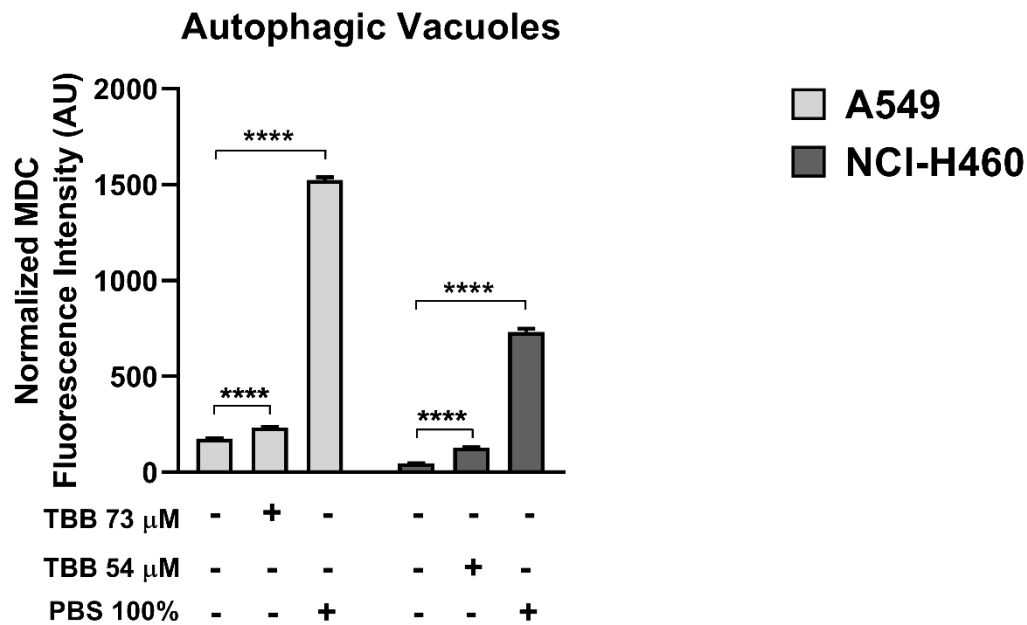
A



B



C



1601

1602 **Figure 7: TBB treatment alters metabolism in lung cancer cells.** A549 and NCI-H460 cells

1603 were treated with IC₅₀-TBB (73 and 54 μM , respectively) for 24 h. (A) $\Delta\Psi_m$, (B), lipid droplets

1604 and (C) autophagic vacuoles were quantified by fluorescence. Data are expressed as mean \pm SEM

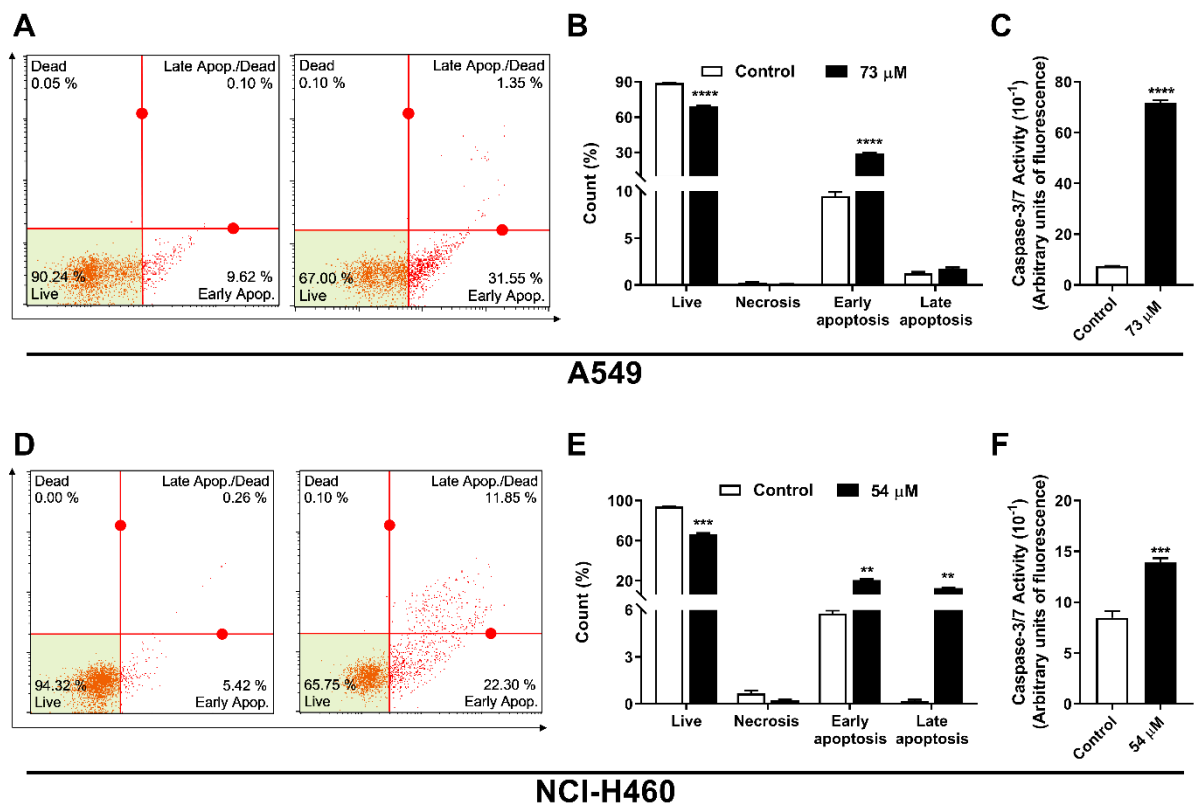
1605 of three independent experiments, each performed in triplicate. **** ($p < 0.0001$) vs. control.

1606

1607 **3.12 TBB induces apoptosis-like cell death and caspase-3 activation**

1608 Cell staining with Annexin V, which binds to phosphatidylserine when externalized, and
 1609 PI, which diffuses through permeable membranes and binds DNA. This labeling technique
 1610 allows us to identify the mechanism of cell death induced by IC₅₀-TBB treatment in the lung
 1611 cancer cells. We differentiated the cell death mechanism into necrotic (PI+), apoptotic (Annexin
 1612 V+) or late apoptotic (co-staining V+/PI+) processes. Our data revealed that A549 cells treated
 1613 with TBB increased V+ labeling (31.55% ± 0.93) compared to the control group, while PI or co-
 1614 staining labeling were not different from the control (Fig. 8A, B). In NCI-H460 cells, treatment
 1615 with TBB was able to increase the percentage of annexin V+ cells (22.30 ± 1.17) and Annexin
 1616 V+/PI+ double positive cells (11.85% ± 0.40), and PI-stained cells did not differ from the control
 1617 group (Fig. 8D, E).

1618 In support of these data, we found that IC₅₀-TBB treatment also induced an increase in
 1619 caspase 3/7 activity in both cancer cell lines (Fig. 8C, F). A549 and NCI-H460 cells showed an
 1620 increase in fluorescence intensity of 64.37% (± 1.30) (Fig. 8C) and 50% (± 0.70) (Fig. 8F),
 1621 respectively, compared to the control. These data indicate a predominance of apoptotic-type cell
 1622 death processes resulting from TBB treatment.



1623
 1624 **Figure 8: TBB treatment induces apoptosis via increased caspase 3/7 activity in lung cancer**
 1625 **cells.** A549 and NCI-H460 cell lines treated with IC₅₀ of TBB for 24 h and evaluated in dot plots

1626 of cytometry, the induction of viable cells, apoptosis, and necrosis (A, D). Quantitative analysis
1627 of the dot plots of both cell lines (B, E). Caspase 3/7 was quantified by fluorescence for A549
1628 (C) and NCI-H460 (F) cells. Values are expressed as mean \pm standard deviation. Cytometry data
1629 are expressed as % vs. control; and mean \pm standard deviation. $**p \leq 0.01$; $***p \leq 0.001$; $****p$
1630 < 0.0001 vs. control.

1631

1632 4 DISCUSSION

1633 Chemotherapy consists of a group of drugs, either alone or in combination, which
1634 represents a therapeutic strategy aimed at inhibiting cell growth as the standard treatment for
1635 non-small cell lung cancer. The FDA has compiled a list of approved drugs for combating
1636 NSCLC, but both the treatments and the disease itself can lead to several adverse effects. These
1637 side effects can vary significantly among patients, even when they are administered the same
1638 drug. This variability primarily arises from the substantial heterogeneity observed in NSCLC
1639 patients (Majeed et al., 2021; Araghi et al., 2023). Consequently, there remains an ongoing
1640 imperative for the development of novel, or newer, therapies to address and mitigate this complex
1641 condition (Zappa and Mousa, 2016).

1642 Within this framework, we perform preliminaries study and offers novel insights into the
1643 mechanism underlying the *in vitro* action of trilobolide-6-*O*-isobutyrate (TBB) on NSCLC cells,
1644 particularly the A549 and NCI-H460 cell lines. Initially, we presented the characterization and
1645 obtention of the compound, in addition, revealed that TBB meets the critical criteria for drug-
1646 likeness *in silico*. Our findings revealed that TBB reduced cell viability in the A549 and NCI-
1647 H460 cell lines while exhibiting low toxicity in epithelial cells, alveolar macrophages, and blood
1648 cells. Treatment with TBB reduced migration ability and induced morphological and
1649 ultrastructural changes. These observed outcomes were accompanied by an increase in oxidative
1650 stress, leading to an adaptive response aimed at mitigating damage and evading cell death.
1651 However, TBB ultimately induced cell death through apoptosis-like processes in both cell lines,
1652 activating caspase 3/7.

1653 Sesquiterpene lactones (SLs) are naturally occurring products found in *Asteraceae* plants.
1654 They hold significant promise for advancing human health due to history of traditional medicine
1655 use, and the biomolecular diversity for drug discovery (Chadwick et al., 2013). They can serve
1656 as valuable pharmaceutical agents with promising applications in oncotherapy due to their
1657 anticancer properties (Liang et al., 2008; Chan et al., 2016; Wang et al., 2017). One noteworthy
1658 compound is TBB, classified as a SL, which boasts advantageous pharmacological attributes. *In*
1659 *silico* studies have consistently demonstrated its adherence to Lipinski and Veber's rules,

1660 exhibiting excellent parameters of drug-likeness predictions, such as bioavailability, metabolic
1661 stability, and intestinal absorption without any violations (Lipinski, 2004; Veber et al., 2002).
1662 Moreover, the structural characteristics of TBB demonstrated favorable calculation of ADMET
1663 (Absorption, Distribution, Metabolism, Excretion, and Toxicity) properties, including a high
1664 potential for oral bioavailability (Cheng et al., 2012). It's important to note that unfavorable or
1665 poor ADMET properties often represent a significant challenge for drug candidates during
1666 clinical trials (van de Waterbeemd and Gifford, 2003).

1667 The primary objective of chemotherapy is to achieve a high degree of selective
1668 cytotoxicity against cancer cells while minimizing harm to healthy cells. Following the
1669 administration of chemotherapy drugs, such as docetaxel, cisplatin, and fluorouracil, several
1670 issues arise. These include immunosuppression, leading to conditions like monocytopenia, which
1671 involves a reduction in the number of monocytes in peripheral blood. Additionally, renal toxicity
1672 becomes evident, as the kidney is susceptible to damage from drugs due to its vital role in the
1673 elimination of ingested, injected, or administered chemicals, toxic metabolites, and medications
1674 (Poprac et al., 2017; Griffin et al., 2019; Patysheva et al., 2022).

1675 Furthermore, during-chemotherapy, hematological toxicity is a significant concern. This
1676 complication is a leading cause of treatment discontinuation, primarily due to its adverse effects
1677 on red blood cells, causing anemia, white blood cells, leading to neutropenia or granulocytopenia,
1678 and platelets, resulting in thrombocytopenia. These conditions can pose a significant risk to the
1679 patient's well-being (Ouyang et al., 2013). Therefore, it is crucial to note that treatment with TBB
1680 showed low toxicity towards renal epithelial cells, monocytic cells, and red blood cells. This
1681 suggests a promising potential for reduced side effects and a more favorable outcome for patients
1682 undergoing this treatment.

1683 The TSI evaluates the cytotoxicity of a compound with respect to tumor and normal cells.
1684 A high value (>1) indicates greater selectivity for tumor cells (Badisa et al., 2009; Lica et al.,
1685 2021). The present study demonstrated for the first time the anticancer activity of TBB against
1686 A549 and NCI-H460 cell lines. Our results revealed that TBB inhibited proliferation of the lung
1687 tumor cells with low toxicity to epithelial cells, monocytic and alveolar macrophage, with TSI
1688 values >1 . Furthermore, we observed a decrease in cell migration, a crucial factor in metastasis
1689 (Stuelten et al., 2018), along with significant morphological and ultrastructural alterations
1690 indicating the hallmarks of apoptosis-induced cell death and loss of mobility in both A549 and
1691 NCI-H460 cancer cells.

1692 In the study conducted by Zhou et al. (2021), they demonstrated the effectiveness of TBB,
1693 isolated from *S. trilobata*, against various cell lines of hepatocellular carcinoma (HCC), including

1694 Huh7, SK-Hep-1, HepG2, and Bel7402. Their study assessed multiple time points and revealed
1695 that TBB treatment significantly inhibited tumorigenesis. Building upon the findings regarding
1696 TBB's action on HCC cells, the primary focus of our investigation was to explore the mechanistic
1697 aspects of cell death induced by TBB in NSCLC cells.

1698 Generally, cancer cells have elevated levels of ROS compared to normal cells. The
1699 presence of small amounts of ROS in cells promotes cancer development. However, when ROS
1700 production exceeds the toxic threshold, this can be associated with mitochondrial dysfunction
1701 and can trigger apoptosis in cancer cells. Therefore, a highly effective approach used by cytotoxic
1702 agents to induce cell death, is to generate oxygen-containing species. Our results demonstrated
1703 that TBB was effective in increasing total ROS and superoxide anion radical production beyond
1704 the threshold in both lung cancer cells. In addition, these results were accompanied by increased
1705 NO levels, while decreased levels of SOD and GSH in both A549 and NCI-H460 cell lines after
1706 treatment with TBB. The literature database has highlighted the dual activities exhibited by NO,
1707 which are contingent upon cellular response, redox status, and concentration of NO (Kolb, 2000).
1708 Physiologically (nM range) relevant levels of NO have been observed to inhibit apoptosis in
1709 diverse cell types, including lung cancer cells. However, at higher concentrations (μM range),
1710 NO is known to induce apoptosis through mechanisms involving oxidative stress and time-
1711 dependent activation of caspases 1, 3 and 6 (Mintz et al., 2021; Umansky et al., 2000).

1712 The cellular defense against oxidative stress is a fundamental aspect of maintaining
1713 cellular homeostasis and preventing cellular death. The enzymes SOD and GSH play pivotal
1714 roles in orchestrating the antioxidant response within cells. SOD catalyzes $\text{O}_2^{\cdot-}$ into less harmful
1715 species like hydrogen peroxide (H_2O_2), thereby mitigating the initial burst of oxidative damage.
1716 Subsequently, GSH participates in detoxifying H_2O_2 through the action of glutathione
1717 peroxidases, preventing the formation of highly reactive hydroxyl radicals ($\cdot\text{OH}$). Together, SOD
1718 and GSH act synergistically, forming an essential frontline defense mechanism against oxidative
1719 stress, and their proper regulation and activity are crucial for maintaining cellular viability
1720 (Poprac et al., 2017). Consequently, our findings indicate that TBB employs a strategy of
1721 promoting oxidative stress in tumor cells in two ways: (1) increasing the production of reactive
1722 oxygen and nitrogen species and (2) reducing the levels of antioxidant enzymes. This reinforces
1723 its potential to combat the proliferation of lung cancer cells.

1724 Cellular responses to stress can trigger either the activation of cell death pathways, or an
1725 adaptive response that ensures cell survival. The adaptive response empowers cells to better
1726 withstand stress-induced damage. Throughout the adaptive process, cells undergo significant
1727 metabolic and physiological adjustments to safeguard against damage and maintain internal

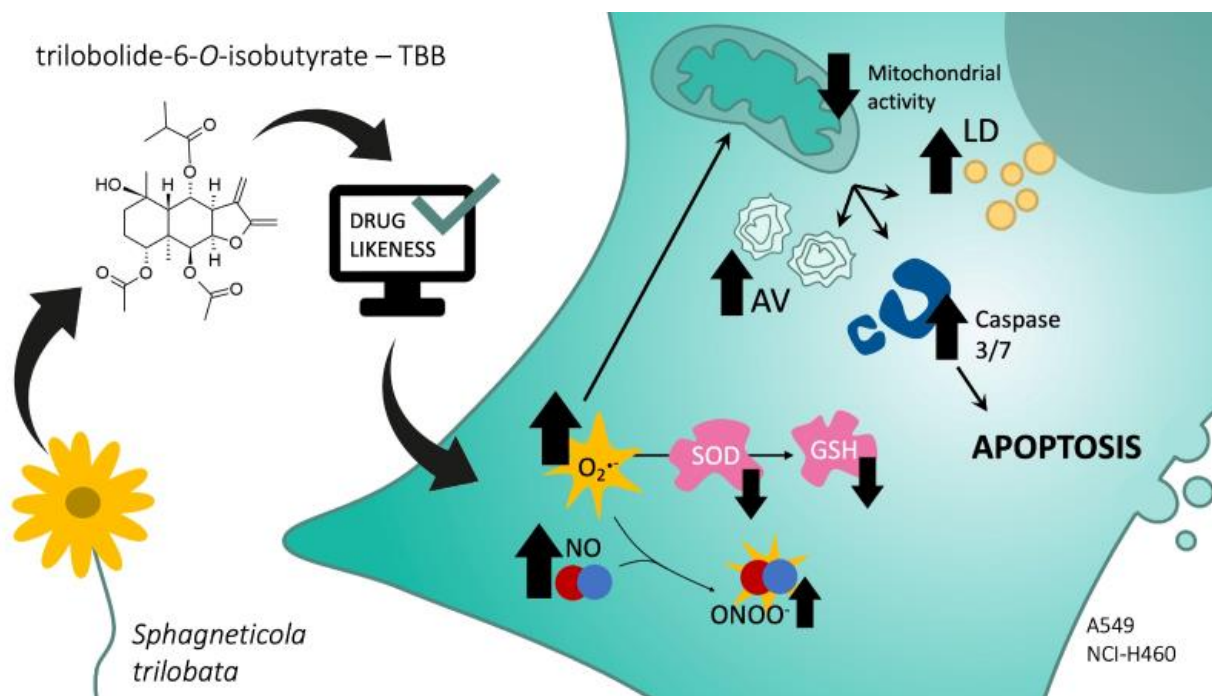
1728 balance (homeostasis). One of the primary mechanisms through which cells promote survival is
1729 by removing damaged organelles and recycling nutrients via the autophagic process (Chern and
1730 Tai, 2020; Labrie et al., 2022). However, when subjected to TBB treatment, A549 and NCI-
1731 H460 cells exhibited an increase of lipid droplets and autophagic vacuoles. Despite this, the
1732 treatment resulted in mitochondrial depolarization (MDP) and hyperpolarization (MHP) in the
1733 respective cancer cell lines, indicating functional loss and organelle damage that culminates in
1734 the loss of the electron transport chain and energy synthesis.

1735 Maintaining the integrity of mitochondria is of paramount importance for efficient energy
1736 production and overall cellular survival, especially when exposed to environmental stressors such
1737 as nutrient scarcity or genotoxic agents. Disruptions in mitochondrial membrane potential, as
1738 indicated by MDP and MHP, are critical events associated with cell death. These mitochondrial
1739 dysfunctions represent a crucial "point of no return" in cascade signaling leading to the loss of
1740 efficiency in the electron transport chain and reduction in energy synthesis. These events
1741 culminate in cell death, and the release of factors such as cytochrome c, into the cytosol (Ganta
1742 et al., 2017; Kroemer and Reed, 2000; Susin et al., 1997). Among the various mechanisms of cell
1743 death explored, two prominent pathways stand out: apoptosis, a programmed process, and
1744 necrosis, which can act as a precursor to several chronic illnesses, including cancer, due to
1745 initiation of an inflammatory cascade. Therefore, there is a distinct preference for therapeutic
1746 agents that can selectively induce apoptosis in cancer cells (Pfeffer and Singh, 2018).

1747 Apoptosis can be initiated by the intrinsic (mitochondrial) and extrinsic (death receptor)
1748 pathways, leading to the activation of effector caspases (caspase 3, -6, and -7) that trigger
1749 apoptotic death. DNA damage, oxidative and nitrosative stress, and cytotoxic agents, ultimately
1750 lead to permeabilization of the mitochondrial outer membrane and subsequently release
1751 cytochrome c, typically initiating the intrinsic pathway. The released cytochrome c leads to the
1752 formation of the apoptosome complex, which in turn activates effector caspases. Notable features
1753 of apoptotic cell death include some hallmarks such as chromatin condensation, nuclear
1754 fragmentation, mitotic cell rounding, reduced cell volume (pyknosis), membrane blebbing
1755 formation, ultrastructural modification of cytoplasmic organelles, loss of membrane integrity,
1756 and withdrawal of pseudopodia (Chaudhry et al., 2022; Gottlieb et al., 2003; Wong, 2011). In
1757 our study, we identified several features associated with apoptosis and, in support of these data,
1758 we also observed cell death such as apoptosis with 3/7 caspase activation. However, further
1759 experiments are required to substantiate this hypothesis.

1760 Thus, our data showed that cells treated with TBB exhibited morphological and
1761 ultrastructural changes indicative of cell damage and death. These changes indicated a profile

1762 favorable to oxidative stress, evidenced by increased levels of ROS and NO, potentially leading
 1763 to the production of highly reactive radicals, such as peroxynitrite and hydroxyls. Furthermore,
 1764 tumor cell lines exhibited adaptations that culminated in cell death through processes similar to
 1765 apoptosis and caspase 3/7 activation (**Fig. 9**).
 1766



1767
 1768 **Figure 9: Representative scheme of the antiproliferative action of TBB on human lung**
 1769 **tumor cell lines, A549 and NCI-H460.** Captions: LD, lipid droplets; NO, Nitric oxide; $O_2^{\bullet-}$
 1770 Superoxide anion radical; SOD, Superoxide dismutase; GSH, Reduced L-glutathione; AV,
 1771 Autophagic vacuoles; $ONOO^-$, Peroxynitrite.

1772
 1773 **CONCLUSION**

1774 This research provided for the first-time new insights into the *in vitro* activity of
 1775 trilobolide-6-O-isobutyrate in lung cancer cell lines, as well as its characterization and
 1776 predictions based on the *in-silico* study. This research has provided novel insights into the *in*
 1777 *vitro* activity of trilobolide-6-O-isobutyrate in lung cancer cell lines. TBB exerts a direct
 1778 inhibitory effect, leading to a reduction in cell proliferation and migration, accompanied by
 1779 notable morphological, ultrastructural, and biochemical alterations. Additionally, TBB induced
 1780 elevated levels of oxidative and metabolic stress, culminating in a process resembling apoptosis,
 1781 as evidenced by the activation of effector caspases. These findings, coupled with theoretical
 1782 assessments of drug-likeness analysis, position TBB as a promising candidate for further
 1783 exploration in cancer studies. Furthermore, this study represents a pioneering effort in unraveling

1784 the mechanisms underlying the *in vitro* treatment of lung cancer cells with TBB, and more studies
1785 need to be explored to develop as a potential lead in drug development in cancer treatment.

1786

1787 **Credit author statement**

1788 All persons who meet authorship criteria are listed as authors, and all authors certify that they
1789 have participated sufficiently in the work to take public responsibility for the content, including
1790 participation in the concept, design, analysis, writing, or revision of the manuscript. Furthermore,
1791 each author certifies that this material or similar material has not been and will not be submitted
1792 to or published in any other publication before its appearance in *Phytomedicine*.

1793

1794 **Author's contribution**

1795 **V.M.C.L.** Conceptualization; Formal analysis; Investigation; Resources; Methodology;
1796 Roles/Writing - original draft. **M.D.G. and F.T.P.:** Formal analysis; Investigation;
1797 Methodology; Roles/Writing - original draft. **M.B.D., E.M.S.C and B.T.S.B:** Investigation;
1798 Methodology; Roles/Writing - original draft. **AC.M.C.:** Formal analysis; Investigation;
1799 Methodology; Roles/Writing - original draft. **A.C.J.R.:** Formal analysis; Methodology. **T.F.S.:**
1800 Formal analysis; Methodology; Roles/Writing - original draft. **E.S.S.:** Investigation;
1801 Methodology. **R.L.N.M.:** Formal analysis; Investigation; Roles/Writing - original draft.
1802 **I.L.A.C.:** Investigation; Methodology. **I.C.C.:** Writing - review & editing. **D.B.L.:** Methodology
1803 and writing - review & editing. **N.S.A.:** Data curation; Formal analysis; Investigation; Resources;
1804 Methodology and revised the manuscript. **R.F.H.D.:** Data curation, formal analysis, and revised
1805 the manuscript. **M.S.M.:** supervised the study and revised the manuscript. **W.R.P.:** supervised
1806 the study and revised the manuscript. All authors read and approved the final manuscript. All
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1809

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1818

1819 **Ethics approval**

1820 This study was approved by the ethics committee (8286.2016.60 and approved in 1/6/2016) of
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1822

1823 **Competing interests**

1824 The authors declare that they have no competing interests.

1825

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