Original Article

Novel CHOP activator LGH00168 induces necroptosis in A549 human lung cancer cells via ROS-mediated ER stress and NF-κB inhibition

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Aim: C/EBP homologous protein (CHOP) is a transcription factor that is activated at multiple levels during ER stress and plays an important role in ER stress-induced apoptosis. In this study we identified a novel CHOP activator, and further investigated its potential to be a therapeutic agent for human lung cancer.

Methods: HEK293-CHOP-luc reporter cells were used in high-throughput screening (HTS) to identify CHOP activators. The cytotoxicity against cancer cells in vitro was measured with MTT assay. The anticancer effects were further examined in A549 human non-small cell lung cancer xenograft mice. The mechanisms underlying CHOP activation were analyzed using luciferase assays, and the anticancer mechanisms were elucidated in A549 cells.

Results: From chemical libraries of 50,000 compounds, LGH00168 was identified as a CHOP activator, which showed cytotoxic activities against a panel of 9 cancer cell lines with an average IC₅₀ value of 3.26 µmol/L. Moreover, administration of LGH00168 significantly suppressed tumor growth in A549 xenograft bearing mice. LGH00168 activated CHOP promoter via AARE1 and AP1 elements, increased DR5 expression, decreased Bcl-2 expression, and inhibited the NF-κB pathway. Treatment of A549 cells with LGH00168 (10 µmol/L) did not induce apoptosis, but lead to RIP1-dependent necroptosis, accompanied by cell swelling, plasma membrane rupture, lysosomal membrane permeabilization, MMP collapse and caspase 8 inhibition. Furthermore, LGH00168 (10 and 20 µmol/L) dose-dependently induced mito-ROS production in A549 cells, which was reversed by the ROS scavenger N-acetyl-L-cysteine (NAC, 10 mmol/L). Moreover, NAC significantly diminished LGH00168-induced CHOP activation, NF-κB inhibition and necroptosis in A549 cells.

Conclusion: LGH00168 is a CHOP activator that inhibits A549 cell growth in vitro and lung tumor growth in vivo.

Keywords: CHOP; LGH00168; ER stress; NF-κB; necroptosis; RIP1; ROS; non-small cell lung cancer A549 cells; cancer xenograft mice

Introduction

Cancer is a leading cause of morbidity and mortality worldwide, with approximately 14 million new cases and 8.2 million cancer-related deaths worldwide in 2012. The lung was one of the most common sites of cancer diagnosed in 2012 (16.7% of the total among men and 8.7% of the total among women)⁵, with non-small cell lung cancer (NSCLC) accounting for 75% of all cases of lung cancer⁶; this statistic shows that NSCLC is an attractive disease for improved chemotherapy⁷–⁹.

The endoplasmic reticulum (ER) is a membrane-bound and structurally intricate organelle present in all eukaryotic cells; it is the primary site for the synthesis and folding of membrane and secretory proteins, and it also functions as storage for membrane lipids and internal calcium. Perturbations of ER homeostasis, a condition called ER stress, can be induced by homeostatic alterations that cause protein misfolding. These include redox status changes, nutrient deprivation, glycosylation changes, pathogen infection, and folding-defective protein expression⁴. ER stress activates the unfolded protein

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response (UPR), thereby re-establishing homeostasis in the ER, whereas extended ER stress activates apoptotic or necrotic pathways in damaged cells\(^{8-11}\). Several UPR effectors, including X-Box Binding Protein 1 (XBPs)\(^{12}\), Glucose-Regulated Protein, 78 kDa (GRP78/Bip)\(^{13,14}\) and CCAAT-enhancer-binding protein homologous protein (CHOP)\(^{15}\), are commonly used as ER stress indicators. CHOP is a transcription factor that is activated at multiple levels during ER stress\(^{16}\). CHOP expression is kept very low in non-stressed cells but is strongly stimulated upon acute ER stress through IRE1-, PERK- and ATF6-dependent transcriptional induction. CHOP overexpression promotes apoptosis in several cell lines, while CHOP\(^{-/-}\) cells are resistant to ER stress-induced apoptosis\(^{17,18}\). This finding suggests that CHOP plays an important role in ER stress-induced apoptosis. The downregulation of the anti-apoptotic protein Bcl-2 and the activation of the pro-apoptotic proteins Bim, Puma, Bax and DR5 are considered to contribute to CHOP-mediated apoptosis\(^{17-19}\). Moreover, the induction of caspase-independent necrosis by ER stress inducers can be blocked by the RIP1 inhibitor necrostatin-1 (Nec-1), indicating that ER stress can also induce necroptosis\(^{20}\).

Apoptosis and necrosis are two distinct cell death pathways\(^{20}\). Apoptosis is highly regulated programmed cell death that is morphologically characterized by cytoplasmic shrinkage, chromatin condensation, phosphatidylserine externalization, DNA fragmentation, and the formation of apoptotic bodies without a loss of plasma membrane integrity\(^{21}\). By contrast, necrosis is viewed as unprogrammed cell death caused by external factors, such as trauma or infection. Necrosis is characterized by the swelling of cells, the early breakdown of plasma membranes, and the induction of inflammatory responses resulting from released cellular content\(^{23}\). Recently a new form of regulated necrosis, called necroptosis, has been recognized as an alternative form of programmed cell death\(^{25}\). Necroptosis is generally triggered by ligands of the death receptor family when caspase-dependent apoptotic pathways are inhibited\(^{24}\). Death receptors are members of the TNFR superfamily and are major mediators of cell death\(^{25}\). The activation of death receptors by their respective ligands recruits and activates caspase 8 at the death-inducing signaling complex (DISC) and triggers apoptosis in the absence of the NF-kB survival pathway\(^{26}\). Under caspase 8 inhibition, necroptosis can be stimulated via an RIP1 kinase and the activation of RIP3 by an RIP1 kinase and MLKL\(^{27-30}\). Inhibition of an RIP1 kinase by a specific RIP1 kinase inhibitor Nec-1 blocks the initiation of necroptosis\(^{30}\). Alternatively, reactive oxygen species (ROS) also play important roles in necroptosis. Antioxidant treatment can reduce ROS levels and also inhibit cell death in some but not all cell types\(^{32,33}\), suggesting that ROS is an executioner of necroptosis in a cell type-dependent manner. Here, we stably introduced a CHOP luciferase reporter into a cell line as a tool to identify ER stress inducers for lung cancer therapy in humans. Using high-throughput screening (HTS), we evaluated one series of analogs for their CHOP-inducing activities; of these analogs, the active compound LGH00168 was further examined as a CHOP activator. The NSCLC cell line A549 is a well-studied lung adenocarcinoma cell line that has been used in xenograft models, including orthotopic xenograft models, of NSCLC to study primary tumor growth and metastasis\(^{24}\). Further investigation of the anti-cancer mechanism was performed in A549 human lung cancer cells. LGH00168 inhibited A549 cell growth in vitro and A549 lung tumor growth in vivo. Mechanistic experiments demonstrated that LGH00168 induces necroptosis via ROS production.

Materials and methods

Cell culture

All of the cell lines mentioned in this paper were obtained from the Cell Bank (Chinese Academy of Sciences, Shanghai, China). The HeLa human cervical cancer cell line and the MCF-7 human breast cancer cell line were cultured in DMEM (high glucose) containing 10% FBS. The MDA-MB-231 human breast cancer cell line, COLO 205 human colon cancer cell line and NCI-H1299 human lung cancer cell line were cultured in RPMI-1640 medium containing 10% FBS. The HCT116 human colon cancer cell line and SK-OV-3 human ovarian cancer cell line were cultured in McCoy’s 5A modified medium containing 10% FBS. The A549 human lung cancer cell line was cultured in Ham’s F12 medium containing 10% FBS. These cells were maintained in a humidified atmosphere of 5% CO\(_2\) and 95% air at 37°C.

Reagents

LGH00168 was obtained from Specs (Delft, The Netherlands), DMEM media (high glucose; Cat #11995-065), RPMI-1640 media (Cat #31800022), MEM media (Cat #11575032), Ham’s F-12 media (Cat #21700-075), Hoechst 33342 (Cat #H3570), JC-1 Dye (Cat #T-3168), and MitoSOX\(^{\text{TM}}\) Red mitochondrial superoxide indicator (Cat #M36008) were obtained from Life Technologies (Grand Island, NY, USA). Hyclone FBS (Cat #SH30071.01) and ECL Western blotting detection reagents (Cat #RPN2108) were purchased from GE Healthcare Life Sciences (Logan, UT, USA). Penicillin/streptomycin (Cat #1540-122) was purchased from the Invitrogen Corporation (Carlsbad, CA, USA). Paclitaxel (Cat #T7402), methylthiazolylidiphenyl-tetrazolium bromide (Cat #M2003), sulforhodamine B (Cat #S1402), N-Acetyl-L-Cysteine (Cat #A7250), propidium iodide (Cat #P4170), Acridine Orange hemi (zinc chloride) salt (Cat #A6014), and McCoy’s 5A modified media (Cat #M4892) were purchased from Sigma-Aldrich (St Louis, MO, USA). A CellTiter-Glo\(^{\text{TM}}\) luminescent cell viability assay kit (Cat #G7573) and Steady-Glo\(^{\text{TM}}\) Luciferase Assay System (Cat #E2550) were purchased from Promega (Madison, WI, USA). An Annexin V-FITC/PI double staining kit (Cat #KGA108) was purchased from Kaiji Bio Co (Nanjing, China).

MTT cytotoxicity assay

An MTT viability assay was used to evaluate the anti-proliferative effects of LGH00168 on different cancer cell lines\(^{33}\). Briefly, cells were seeded into 96-well plates and cultured at 37°C. After 24 h, compounds at concentrations ranging from 0.63 μmol/L to 20 μmol/L were added at a dilution of...
Soft cells were collected, the mice were randomized into treatment groups (5 × L/2, De Novo) by proton trapping within the acidic vacuolar compartment. Cells were treated with LGH00168 for 4 h and analyzed with FCS Express V3 software (San Jose, CA, USA). Data were expressed as the mean ± SEM.

Anti-tumoral effect of LGH00168 in lung carcinoma-bearing mice

Tumor xenografts were established by the injection of A549 cells (5 × 10⁶ cells/animal, sc) into the armpit of athymic mice (National Rodent Laboratory Animal Resource, Shanghai, China). Tumor volume (cubic millimeters) was measured with calipers and calculated by the following formula: volume = W² × L / 2, where W is the short diameter, and L is the long diameter. When tumors reached a mean group size of approximately 110 mm³, the mice were randomized into treatment groups (5 mice per group), and treatment was commenced. The mice were administered with LGH00168 (5, 10, and 20 mg/kg, im), a vehicle (tea tree oil), or the positive control Paclitaxel (10 mg/kg) once every 2 d. Animal care was in accordance with institutional guidelines.

Propidium iodide/Hoechst 33342 (PI/HO) double staining

Plasma membrane permeability (PMP) and nuclear morphology were evaluated using PI/HO double staining as described previously with modifications as follows: A549 cells were cultured on glass coverslips and incubated with LGH00168 for 4 h. Next, the cells were stained with 40 mg/L of PI and 20 mg/L of Hoechst 33342 in 1× PBS for 15 min in the dark. After washing twice with 1× PBS, the slides were mounted, sealed and observed under an Olympus BX51 fluorescence microscope (Tokyo, Japan). Images were captured with an Olympus DP72 digital camera (Tokyo, Japan) and analyzed using Adobe Photoshop CS4 software (San Jose, CA, USA).

Annexin V/PI analysis

The quantitative assessment of cell death was determined using an Annexin V/PI double staining kit according to the manufacturer’s instructions. Briefly, the cells were treated with LGH00168 for 4 h. Next, 5 × 10⁶ cells were collected, washed twice with cold 1× PBS, and stained with Annexin V and PI for 15 min at room temperature in the dark. Finally, the percentages of necrotic cells were measured using a BD FACSCalibur Flow Cytometer (BD Inc, San Jose, CA, USA) and analyzed with FCS Express V3 software (De Novo Software, Glendale, CA, USA).

Determination of lysosomal membrane permeabilization (LMP) using acridine orange/Hoechst33342 (AO/HO) staining

The metachromatic fluorophore AO is retained in its charged form (AOH⁻) by proton trapping within the acidic vacuolar compartment. When normal cells are excited by blue light, highly concentrated lysosomal AO emits an intense red granular fluorescence, and the nuclei and cytosol show weak diffuse green fluorescence. LMP results in reduced red fluorescence and bright green cytoplasmic and nuclear fluorescence. LMP induction by LGH00168 was observed using AO/HO double staining. Cells were treated with LGH00168 for 4 h and then incubated with 5 mg/L of AO for 10 min in the dark. Images were captured with an Olympus DP72 digital camera (Tokyo, Japan) and analyzed in Adobe Photoshop CS4 software (San Jose, CA, USA).

Caspase 8 activity evaluation

Caspase 8 activity was evaluated using the Caspase-Glo 8 assay system (Promega, Madison, WI, USA) according to the manufacturer’s instructions. Briefly, the cells were treated as described with LGH00168 before the addition of 1/2 volume of the assay reagent. The lysates were then incubated at room temperature for 1 h before luminescence was measured on an EnVision Multilabel Plate Reader (Perkin Elmer, Waltham, MA, USA).

Quantitative polymerase chain reaction (qPCR) detection of CHOP, Bcl-2, and DR5

qPCR was utilized to detect the CHOP, Bcl-2, and DR5 mRNA levels. Briefly, cells were treated with 10 µmol/L of LGH00168 for 0, 5, 24, and 48 h. Next, the total RNA was extracted using a TRizol reagent (Invitrogen, Carlsbad, CA, USA) according to the manufacturer’s instructions. GAPDH was used as an internal control. Complementary DNA generated by M-MLV Reverse Transcriptase (Promega, Madison, WI, USA) was analyzed by qPCR using SYBR® Premix Ex Taq II (TakaRa, Dalian, China). The sequences of the primers were as follows: human CHOP forward 5'-CTGCAAGAGTGCTTGCTCTCA-3' and reverse 5'-TGGTCTCCTGTTCTGTC-3'; human Bcl-2 forward 5'-ATGTTGGAGGAGCTTTGCA-3' and reverse 5'-ACAGTTCCACAAAGGCATCC-3'; human DR5 forward 5'-CCTATCGCTAGGATGACTTCTCC-3' and reverse 5'-CAGCAGAATGCATGTTCTC-3'. All of the reactions were performed in triplicate, and the cycle threshold (CT) value in each reaction well was recorded. The relative quantification of CHOP, Bcl-2, and DR5 mRNA expression was calculated using the 2⁻¹⁰ΔΔCT method.

Western blot analysis

Cells were treated with LGH00168 and then lysed with a cell lysis buffer (1% NP-40, 150 mmol/L of NaCl, 20 mmol/L of Tris-HCl, 1 mmol/L of EDTA, 1 mmol/L of EGTA and complete protease inhibitor cocktail). The lysates were fractionated by SDS-PAGE, transferred to Hybond-C nitrocellulose membranes (GE Life Sciences, Pittsburgh, PA, USA) and immunoblotted as described previously. Immunoreactive bands were detected using ECL Western blotting detection reagents (Cat #RPN2108, GE Life Sciences, Pittsburgh, PA, USA).
Luciferase assay
HEK293-ERSE-luc, HEK293-AARE1-luc, HEK293-AARE2-luc, HEK293-AP1-luc, HEK293-NF-κB-luc, or HEK293-CHOP-luc reporter cells were seeded into 96-well plates and cultured at 37°C. After 24 h, different doses of drugs were added. After 24 h of treatment, luciferase activity was determined using the Promega Steady-Glo luciferase assay system according to the manufacturer’s instructions. The luminescent signal was recorded using the EnVision Multilabel Plate Reader (Perkin Elmer, Waltham, MA, USA). All of the luciferase assay experiments were repeated at least three times.

Measurement of the mitochondrial ROS (mito-ROS)
The mito-ROS level was examined by a MitoSOX Red mitochondrial superoxide indicator (Cat #M36008, Life technologies, Grand Island, NY, USA) according to the manufacturer’s instructions. Briefly, cells were treated with LGH00168 for 1 h, collected by trypsin treatment, washed twice with ice-cold 1× PBS, and then incubated with 5 μmol/L of MitoSOX for 20 min in the dark. The mito-ROS level was measured using a BD FACSCalibur Flow Cytometer and analyzed with FCS Express V3 software (De Novo Software, Glendale, CA, USA).

Measurement of the mitochondrial membrane potential (Δψm)
The Δψm level was measured using a JC-1 dye (Cat #1-T-3168, Life Technologies, Grand Island, NY, USA) according to a previous report with modifications. Briefly, cells were seeded into black-bottomed 96-well optical plates (Greiner Bio-One, Dusseldorf, Germany). After 24 h, LGH00168 was added, and the cells were incubated at 37°C for 1 h. JC-1 at a final concentration of 10 μg/mL was then added. The plates were incubated for another 20 min at 37°C, and each well was washed three times with a Krebs-Ringer phosphate HEPES buffer. Fluorescence was measured in a Flexstation II plate reader (Molecular Device, Union City, CA, USA) first at excitation and emission wavelengths of (Ex/Em) 530 nm/580 nm (“red”) and then at Ex/Em 485 nm/530 nm (“green”). The ratio of red to green reflects Δψm. The results were analyzed using GraphPad Prism version 5.02 (GraphPad Software Inc, La Jolla, CA, USA).

Measurement of the cellular ATP level
The cellular ATP level was detected using a CellTiter-Glo assay (Cat #G7573; Promega, Madison, WI, USA). Briefly, cells (5000 cells per well) were seeded into white-bottomed 96-well plates and cultured at 37°C. After 24 h, the cells were treated with LGH00168 for 4 h. Next, the plates were equilibrated at room temperature for 30 min, a CellTiter-Glo reagent was added, and the plates were shaken with an orbital shaker for 2 min. After incubation at room temperature for 10 min, the luminescent signal was recorded using the EnVision Multilabel Plate Reader (Perkin Elmer, Waltham, MA, USA).

Sulforhodamine B (SRB) assay
The SRB assay was used to evaluate the effect of NAC (N-Acetyl-L-Cysteine) on LGH00168-induced cell death. Briefly, cells were seeded into 96-well plates and cultured at 37°C. After 24 h, LGH00168 with final concentrations ranging from 20 μmol/L to 2.5 μmol/L at a dilution ratio of 1:2 with or without NAC (final concentration: 10 mmol/L) was added, and the cells were further incubated for 72 h. Next, the cells were fixed with 3.3% (w/v) trichloroacetic acid for 1 h at 4°C and stained with 0.4% SRB dissolved in 1% (v/v) acetic acid for 30 min. The unbound dye was removed by four washes with 1% acetic acid, and the protein-bound dye was extracted with 10 mmol/L of an unbuffered Tris base (pH 10.5) for 5 min. The optical density was determined at 510 nm using a VersaMax Microplate Reader (Molecular Devices, Sunnyvale, CA, USA). Experiments were performed in triplicate. IC50 values were derived from a nonlinear regression model (curvfit) based on a sigmoidal dose response curve (variable slope) and computed using GraphPad Prism version 5.02 (GraphPad Software Inc, La Jolla, CA, USA). Data were expressed as the mean±SEM.

NAC treatment
To evaluate the role of ROS in the anticancer activity of LGH00168 against A549 lung cancer cells, NAC (final concentration: 10 mmol/L) was used as ROS scavenger. mRNA levels of CHOP, protein levels of p-p65, p65, IκBα, and β-actin, levels of LMP, MMP, and cellular viability were evaluated in LGH00168-treated cells with or without NAC.

Statistical analysis
The statistical analyses were performed by paired two-tailed Student’s t-tests (GraphPad Prism version 5.02, GraphPad Software Inc, La Jolla, CA, USA). A level of P<0.05 was considered statistically significant. The data are presented as the mean±SEM.

Results
LGH00168 was identified as a CHOP activator that inhibited A549 cell growth in vitro and lung tumor growth in vivo
A high-throughput screening (HTS) assay was utilized to identify CHOP activators from 50,000 compounds from the chemical libraries of the Chinese National Center for Drug Screening (data not shown). LGH00168 was identified from among its analogs (Figure 1A) as a CHOP activator (Figure 1B) with good cytotoxic activity in vitro (Figure 1C) and in vivo (Figure 1D, 1E). LGH00168 showed good cytotoxic effects against nine cancer cell lines that were derived from six different tissues, with an average IC50 value of 3.26 μmol/L in vitro. Moreover, LGH00168 effectively inhibited the growth of A549 tumor xenografts in vivo compared with the control group.

LGH00168 induced CHOP activation and NF-κB inhibition
CHOP promoter-luciferase reporter assays demonstrated the activation of AARE1 and AP1 elements of the CHOP promoter by LGH00168 treatment (Figure 2A). qPCR results indicated increased DR5 expression and decreased Bcl-2 expression of two CHOP downstream targets in a time-dependent manner (Figure 2B). Moreover, a luciferase assay and Western blot
data indicated the inhibitory effect of LGH00168 on the NF-κB pathway (Figure 2C and 2D).

**LGH00168 induced RIP1-dependent necroptosis**

The mechanism of LGH00168-induced cell death was investigated by morphological observation, PI/HO double staining, and an Annexin V/PI assay. LGH00168 induced morphological changes and an increase in PI staining, which is a sign of plasma membrane rupture (Figure 3A). Moreover, an Annexin V/PI assay examination indicated that cell death was due to necrosis without apoptotic induction (Figure 3B). Additionally, the correspondent FSC signal decreased (Figure 3C), demonstrating that LGH00168-treated cells were dying. Moreover, severe LMP and MMP collapse were induced after 4 h of treatment (Figure 3D and 3E), and caspase 8 activity was inhibited (Figure 3F). PI staining and a cell viability assay were utilized to evaluate whether LGH00168-induced necrosis can be reversed by the RIP1 inhibitor Nec-1. As shown in Figure 3G and 3H, Nec-1 partially reversed PI staining and cell death, indicating that LGH00168 induced RIP1-dependent necroptosis.

**LGH00168 induced mito-ROS production**

Mitochondria are reported as the main site of ROS generation. Additionally, superoxide (O$_2^-$) is the primary mitochondrial ROS. We next examined whether O$_2^-$ was generated in mito-
chondria using the MitoSOX Red mitochondrial superoxide indicator, which only detects $O_2^{.-}$ generated in mitochondria. LGH00168 induced mito-ROS generation in a dose-dependent manner (Figure 4A), which was effectively reversed by the ROS scavenger NAC (Figure 4B).

The ROS scavenger NAC blocked LGH00168-induced CHOP activation, NF-$\kappa$B inhibition, and necroptosis. NAC effectively blocked LGH00168-induced CHOP activation, NF-$\kappa$B inhibition, LMP, and MMP collapse (Figure 5A–5D), indicating that these effects were dependent on ROS generation. Moreover, NAC completely blocked LGH00168-induced cell death (Figure 5E), showing that necrototic cell death was dependent on ROS production.

**Discussion**

It has long been accepted that cell death can be classified into two types: apoptosis and necrosis. The activation of necrotic events was considered, until recently, to be promoted by extensive cellular trauma, whereas apoptosis was considered to be driven by a genetically controlled cascade of events initiated by minor insults or extracellular signals. Recent work has suggested that anti-cancer treatments also trigger other, non-apoptotic, cell death modalities (autophagy, mitotic catastrophe and necrosis). Additionally, some studies have found that necrosis induction enhances the anti-cancer efficacy for some chemo-resistant tumors.

Here, LGH00168 was identified as a novel CHOP activator that effectively inhibited the in vitro growth of nine cancer cell lines derived from six different tissues and that inhibited lung tumor growth in vivo. As previously reported, cells undergoing apoptosis exhibit cardinal features such as changes in cellular and nuclear morphology, a non-random phosphatidylserine inversion on the plasma membrane, and the formation of apoptotic bodies. Additionally, cells undergoing necrosis will change morphologically, showing swelling with a rupture of the plasma membrane. In this study, cells with typical apoptotic nuclei and cells in the early apoptotic stage were not observed; however, rupture of the plasma membrane was induced, suggesting that LGH00168 induces necrosis not apoptosis.

Necrosis is the consequence of extensive crosstalk between several biochemical and molecular events. Currently, there is no single well-described signaling cascade that defines necrotic cell death. As previously reported, necrosis is the result of the interplay between several signaling cascades, and it is thought to be regulated. There are several important mediators involved in the propagation and execution of necrotic cell death, including ER stress induction, NF-$\kappa$B inhibition, ROS, proteases, and ceramide. Here, we found that LGH00168 induced necrotic cell death accompanied by ER stress induction, NF-$\kappa$B inhibition, mito-ROS generation, severe LMP, and

**Figure 2.** LGH00168 activated CHOP and inhibited the NF-$\kappa$B pathway. (A) The luciferase assay indicated that CHOP was activated via the AARE1 and AP1 elements in HEK293 cells. **$P<0.01$** vs DMSO group. (B) qPCR indicated that LGH00168 activated CHOP and DR5 in a time-dependent manner and inhibited Bcl-2. (C) The luciferase assay indicated the inhibitory effect of LGH00168 on NF-$\kappa$B activity in HEK293 cells. (D) The Western blot assay indicated the inhibition of p-p65 and the accumulation of IκBα in LGH00168-treated cells.
the loss of Δψm, which might be the mediators of LGH00168-induced necrosis. As previously reported, an RIP1 kinase was found to initiate necroosome formation and trigger necroptosis, and we further utilized an RIP1 inhibitor, Nec-1, to evaluate the role of RIP1 in LGH00168-induced necrosis. Nec-1 effectively reversed LMP and necrotic cell death, indicating that LGH00168 induced RIP1-dependent necroptosis.

Mitochondria are the organelles that are central to many fundamental cellular processes, such as energy generation, and are an important source of ROS within most mammalian cells. We next examined whether LGH00168 induced the generation of ROS in mitochondria. Here, LGH00168 induced mito-ROS generation. The ROS scavenger NAC effectively eliminated excessive ROS and reversed the loss of MMP, the induction of LMP and necroptotic cell death, indicating that necroptosis was mainly induced through the production of ROS.

Necrosis is an irreversible inflammatory form of cell death, and it is also known that therapy-induced necrotic cell death initiates an immune response to tumor cells. However, whether the inflammation is associated with necrosis remains unclear, but therapeutics that target the regulators of

Figure 3. LGH00168 induced necroptosis in A549 cells. (A) PI/HO staining indicated the induction of cellular swelling and plasma membrane rupture. (B) The Annexin V/PI assay indicated the induction of necrosis. (C) The SSC/FSC parameters showed cell death of LGH00168-treated cells. (D) AO/EB staining indicated the induction of severe LMP. (E) JC-1 staining indicated the collapse of MMP in LGH00168-treated cells. (F) A caspase 8 activity assay indicated the inhibition of caspase 8 activity. (G) The RIP1 inhibitor Nec-1 partially reversed LGH00168-induced PMP. (H) The RIP1 inhibitor Nec-1 partially reversed LGH00168-induced necroptosis. *P<0.05, **P<0.01.
necrotic cell death are already in early-phase clinical trials\[^{45}\]. It has been reported that a cell can initiate its own demise by necrosis, which initiates both inflammatory and/or reparative responses in the host\[^{50}\]. Moreover, most chemotherapy drugs induce apoptosis, but some drugs can cause necrosis\[^{51, 52}\]. Additionally, drug resistance is often associated with an imbalance in pro-apoptotic and anti-apoptotic mediators, with the net result favoring cell survival. Here, we identified LGH00168 as a new necroptosis inducer. We propose that LGH00168 stimulates severe ER stress, NF-κB inhibition, ROS production, MMP loss and severe LMP induction, which finally lead to necroptosis in A549 cells. LGH00168, which shows good anti-cancer activity \textit{in vitro} and \textit{in vivo}, has potential as a new drug candidate and provides an ideal strategy for developing combination therapies to eliminate subpopulations of cells that are resistant to apoptotic cell death. Additionally, the further study of LGH00168 would provide more detailed information about its anti-cancer mechanism, which might be helpful for further optimization.

**Abbreviations**

CHOP, C/EBP homologous protein; GADD153, growth arrest- and DNA damage-inducible gene 153; HTS, high-throughput

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**Figure 4.** LGH00168 induced mito-ROS production in A549 cells. (A) LGH00168 induced mito-ROS production in a dose-dependent manner. (B) The ROS scavenger NAC (10 mmol/L) effectively blocked LGH00168-induced mito-ROS production.

**Figure 5.** The ROS scavenger NAC (10 mmol/L) blocked LGH00168-induced CHOP activation, NF-κB inhibition, and necroptosis in A549 cells. (A) NAC reversed LGH00168-induced CHOP activation. (B) NAC reversed LGH00168-induced NF-κB inhibition. (C) NAC blocked LGH00168-induced LMP. (D) NAC blocked LGH00168-induced MMP collapse. (E) NAC protected cells from LGH00168-induced necroptotic cell death. **P<0.01.
screening; MTT, methylthiazolyldiphenyl-tetrazolium bromide; CT, cycle threshold; ER, endoplasmic reticulum; UPR, unfolded protein response; XBPI, X-box binding protein 1; DISC, death-inducing signaling complex; NF-kB, nuclear factor κB; NSCLC, non-small cell lung cancer; PMP, plasma membrane permeabilization; LMP, lysosomal membrane permeabilization; MPP, mitochondrial membrane potential; AO, acridine orange; HO, Hoechst 33342; qPCR, quantitative polymerase chain reaction; RIP1, receptor-interacting protein 1; ΔψM, mitochondrial membrane potential; mito-ROS, mitochondrial ROS; NAC, N-Acetyl-L-Cysteine; SRB, sulforhodamine B; Nec-1, necrostatin-1.

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Author contribution
Yi-ming MA, Yan-min PENG, Qiong-hua ZHU, An-hui GAO, Bo CHAO, Qiao-jun HE, You-hong HU, Jia LI, and Yu-bo ZHOU conceived and designed the experiments; Yi-ming MA, Yan-min PENG, Qiong-hua ZHU, and An-hui GAO performed the experiments; Yi-ming MA, Yan-min PENG, Qiong-hua ZHU, and An-hui GAO analyzed the data; Yi-ming MA and Yu-bo ZHOU wrote the paper.

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