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- 4 **Running title:** Necroptosis in prostate cancer

6 20(S)-ginsenoside Rg3 induced the necroptosis of prostate cancer cells via ROS 7 overproduction

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Necroptosis is a programmed form of necrosis and compounds inducing necroptosis may contribute 25 to cancer treatment. 20(S)-ginsenoside Rg3 is a natural compound extracted from ginseng, which 26 exhibited a broad-spectrum of antitumor activity. In the present study, the potential role of 20(S)-27 ginsenoside Rg3 in inducing necroptosis in prostate cancer cells was evaluated. 20(S)-ginsenoside 28 Rg3 inhibited the proliferation of prostate cancer cells and upregulated the expression of necroptotic 29 proteins such as receptor-interacting serine/threonine-protein kinase 1 (RIPK1), RIPK3, and their 30 downstream mixed lineage kinase domain-like protein (MLKL). Pretreatment with the selective 31 RIPK1 inhibitor necrostatin-1 (Nec-1) partially reversed the inhibitory effect of 20(S)-ginsenoside 32 33 Rg3 on prostate cancer cell proliferation. 20(S)-ginsenoside Rg3 led to the accumulation of reactive oxygen species (ROS) and the regulation of autophagy in cancer cells. Scavenging ROS with N-34 acetyl-L-cysteine (NAC) antagonized the regulatory effects of 20(S)-ginsenoside Rg3 on cell 35 autophagy and necroptotic proteins expression. Moreover, 20(S)-ginsenoside Rg3 exhibited an 36 antitumor effect in a prostate cancer xenograft mouse model in which it upregulated the expression 37 of RIPK1, RIPK3, MLKL and led to a decrease in tumor weight, as well as an increase in necrotic 38 areas in tumor tissue. In conclusion, our study showed that 20(S)-ginsenoside Rg3 might induce 39 necroptosis in prostate cancer in vitro and in vivo via the ROS/autophagy signaling pathway. 40

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Key words: autophagy; 20(S)-ginsenoside Rg3; necroptosis; prostate cancer; reactive oxygen
 species

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46 Prostate cancer remains an important health issue since it was the second most frequent cancer and47 the fifth leading cause of cancer death among men worldwide in 2022 [1]. Prostate cancer is

48 initially sensitive to anti-androgen therapy. However, during the course of the treatment, prostate

49 cancer often becomes refractory to hormone therapy and develops into castration-resistant prostate

cancer (CRPC), which is also resistant to chemotherapy [2]. Therefore, there is no satisfactory
treatment against CRPC and it is urgent to develop new therapeutic options.

Necroptosis is a form of programmed cell death that resembles necrosis and depends on a unique 52 molecular pathway different from apoptosis. The formation of a RIPK1-RIPK3 necrosome through 53 auto- and trans-phosphorylation of RIPK1 and RIPK3 happens at the initiation of necroptosis [3]. 54 55 This is followed by MLKL activation via phosphorylation and translocation to the plasma membrane which causes cell death. Given the resistance of cancer cells to apoptosis, the induction 56 57 of necroptosis may be an alternative strategy for cancer therapy [4, 5]. Previous studies indicated that RIPK3 expression was often silenced in cancer cells due to genomic methylation near the 58 transcriptional start site of RIPK3. This repressed necroptosis-induced cell death suggests that 59 inducing necroptosis may be beneficial for cancer treatment [6]. In prostate cancer tissues, the 60 expression of RIPK3 was elevated at early stages of cancer, while it was repressed at late stages, 61 and negatively correlated with tumor size and prostate-specific antigen (PSA) levels [7]. Another 62 study reported the down-regulation of RIPK3 in prostate cancer cell lines and clinical prostate 63 tumor samples. Low expression of RIPK3 was closely related to tumor metastasis and poorer 64 survival, while up-regulation of RIPK3 could induce necroptosis and alleviate prostate cancer 65 progression [8]. Moreover, sirtuin 3 (SIRT3) and SIRT6 were increased in prostate cancer tissues 66 and this induced prostate cancer progression by inhibiting RIPK3-mediated necroptosis and innate 67 immune response [9]. Several natural compounds have been reported to induce necroptosis and may 68 be beneficial in the treatment of prostate cancer. Ophiopogonin D, a compound extracted from 69 Ophiopogon japonicus, exhibited an antiproliferative activity in androgen-dependent prostate 70 71 cancer cells via the induction of a RIPK1- and MLKL-dependent necroptosis [10]. Arctigenin, a natural product isolated from Arctium lappa, also induced necroptosis in acidity-tolerant prostate 72 cancer cells via ROS-mediated mitochondrial damage and the upregulation of cell communication 73 74 network factor 1 (CCN1) [11]. Hong et al reported that the rosin derivative IDOAMP increased the phosphorylation of RIPK1, RIPK3, MLKL and activated necroptosis in prostate cancer cells [12]. 75 Moreover, a recent report indicated that emodin induced necroptosis in prostate cancer cells via the 76 regulation of the mitochondrial fission pathway [13]. These studies highlight a potential strategy for 77 78 prostate cancer treatment via agent-induced necroptosis.

79 20(S)-ginsenoside Rg3 is clinically used against several cancers and displays a remarkable 80 antitumor activity via multiple mechanisms [14, 15]. Several studies reported that 20(S)-81 ginsenoside Rg3 inhibited the proliferation and migration of prostate cancer cells as well as 82 enhanced the susceptibility of prostate cancer cells to docetaxel [16-18]. Our previous work 83 indicated that 20(S)-ginsenoside Rg3 upregulated ROS levels and induced cell cycle arrest to inhibit the proliferation of PC3, a human CRPC cell line [19]. The regulatory role of 20(S)-ginsenoside Rg3 was also reported in the prostatic microenvironment with an inhibition of senescence in prostate stromal cells through down-regulation of interleukin 8 (IL-8) expression [20]. In this study, we explored the *in vitro* and *in vivo* potential effects and underlying mechanisms of 20(S)ginsenoside Rg3 on the regulation of necroptosis in prostate cancer cells.

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90 Materials and methods

Cell lines and reagents. The prostate cancer cell lines 22RV1 and PC3 were purchased from 91 American type culture collection (ATCC, Manassas, VA, USA) and cultured in RPMI1640 medium 92 (Sigma, St Louis, MO, USA) supplemented with 10% fetal bovine serum (FBS, Lonsera Science 93 SRL, Canelones, Uruguay). 20(S)-ginsenoside Rg3 (CAS: 14197-60-5) was purchased from 94 Weikeqi Biological Technology Col, Ltd (Chengdu, Sichuan, China) and dissolved in dimethyl 95 sulfoxide (DMSO, purchased from Solarbio, Beijing, China). N-acetyl-L-cysteine (NAC) was 96 purchased from Beyotime institute of biotechnology (Shanghai, China), and Nec-1 from Med Chem 97 Express (Monmouth Junction, NJ, USA). 98

Analysis of online databases. The Genotype-Tissue Expression (GTEx) and The Cancer Genome Atlas (TCGA) databases were analyzed with an online tool (www.xiantaozi.com) to compare the expression of *RIPK1*, *RIPK3* and *MLKL* in normal prostate and prostate tumor samples (100 normal tissues, 496 tumor tissues and 52 para-cancerous tissues). Wilcoxon rank sum test and paired sample *t*-test were applied for statistical analysis. Disease specific survival analysis and receiver operating characteristic curve (ROC) analysis were also operated with the same tool. The diagnostic efficiencies of *RIPK1*, *RIPK3* and *RIPK3* in prostate tumor was compared with DeLong's test.

106 CCK8 assays. 22RV1 and PC3 cells were seeded into 24-well plates $(1 \times 10^4 \text{ cells/well})$ and 107 treated with vehicle (DMSO, final concentration 0.1%), 50 or 100 μ M 20(S)-ginsenoside Rg3 for 0, 108 24, 48 or 72 h. Then, a CCK8 kit (US Everbright INC, Suzhou, China) was used to evaluate the 109 proliferation of cells according to the manufacturer instructions. The results are expressed as the 110 mean±SD of three independent experiments.

Flow cytometry assays. 22RV1 and PC3 cells were cultured into 6-well plate $(1 \times 10^5 \text{ cells/well})$ and treated with DMSO (final concentration 0.1%) or 100 μ M 20(S)-ginsenoside Rg3 for 48 h. Cells were digested with 0.25% trypsin (Solarbio) to detach from the plate and then a FITC-Annexin V and PI apoptosis kit (US Everbright) was used to label suspended cells according to the manufacturer instructions. Cell apoptosis and necrosis were analyzed by flow cytometry (Guava easyCyte b-2L, Millipore, Billerica, MT, USA). The proportion of cell subtypes are expressed as the mean±SD of three independent experiments. To evaluate the ROS level, 22RV1 and PC3 cells were cultured into 6-well plates $(1 \times 10^5 \text{ cells/well})$ and treated with DMSO (final concentration 0.1%) or 100 μ M 20(S)-ginsenoside Rg3 for 48 h, and subsequently labeled with 2,7dichlorodihydrofluorescein diacetate (DCFH-DA, Beyotime). Then flow cytometry assays were performed to evaluate the ROS levels in cells and the assays were performed three times independently.

Western blot analysis. 22RV1 and PC3 cells were cultured into 6-well plate (1×10^5 cells/well) 123 and then treated with DMSO (final concentration 0.1%) or 100 µM 20(S)-ginsenoside Rg3 for 48 h. 124 Total proteins were extracted with radio immunoprecipitation assay (RIPA) buffer (CWBIO, 125 Beijing, China) supplemented with 1 mM phenylmethanesulfonylfluoride (PMSF, Solarbio). The 126 protein concentration was quantified using the BCA method (CWBIO). Then, western blot analysis 127 was performed using the following antibodies (Cell Signaling Technology, Danvers, MT, USA): 128 RIPK1 (#3493, dilution 1:1000), RIPK3 (#13526, dilution 1:1000), MLKL (#14993, dilution 129 1:1000) and the HRP-linked second antibody (anti-rabbit, #7074, dilution 1:2500). LC3 (#bs-8878R, 130 dilution 1:1000) and p62 antibodies (#bs-55207R, dilution 1:1000) were purchased from Bioss 131 (Beijing, China). Caspase 3 antibody (#ab13847, diluted at 1:1000) was purchased from Abcam 132 expression was normalized to glyceraldehyde-3-phosphate (Cambridge, UK). Protein 133 dehydrogenase (GAPDH, #AB0037, Abways, dilution 1:12000). All the experiments were 134 performed three times independently. Quantitative analyses of the results were performed using 135 Image J software. 136

Nec-1 and NAC pretreatment. 22RV1 and PC3 cells were cultured into 6-well plates (1×10^5) 137 cells/well) and then pretreated with 10 mM NAC for 1 h, followed by a treatment with DMSO (final 138 concentration 0.1%) or 100 µM 20(S)-ginsenoside Rg3 for 48 h to extract total proteins for western 139 blot analyses. 22RV1 and PC3 cells were also cultured into 24-well plate (1×10^4 cells/well) and 140 pretreated with 100 µM Nec-1 for 1 h, followed by a treatment with DMSO (final concentration 141 0.1%) or 100 µM 20(S)-ginsenoside Rg3 for 72 h to measure cell proliferation using a CCK8 kit. 142 The experiments were performed independently for three times and the results of CCK8 assays are 143 expressed as the mean±SD. 144

Mice xenograft model. The protocol for *in vivo* studies was approved by the Experimental Animal Ethics Committee of the Institute of Radiation Medicine, Chinese Academy of Medical Sciences (Approval No. IRM-DWLL-2019157). Sixteen Balb/c nude mice (male, 6 weeks old) were purchased from Huafukang Biotechnology Company (Beijing, China). The mice were acclimated in specific pathogen-free units for 1 week, followed by subcutaneous injection of 5×10^6 PC3 cells on the right side of the armpit. Mice were normally fed for another 2 weeks and divided into two groups: control group (eight mice) and ginsenoside Rg3 treatment group (eight mice). Daily gavage with phosphate buffer saline (PBS) was performed in the control group and daily intragastric gavage with 20(S)-ginsenoside Rg3 (25 mg/kg body weight) was performed in the treatment group for 2 weeks. Body weight and tumor volumes were monitored every two days. The volume of tumors was calculated with the formula: $V=1/2 \times \log diameter \times (\text{short diameter})^2$. At the end of the treatment, mice were sacrificed and the xenografts were collected for hematoxylin-eosin (H&E) staining, immunofluorescent (IF) staining and western blot analysis.

H&E and IF staining. Each tissue sample was separated in two parts. One part was embedded in 158 paraffin and the other one was frozen. The paraffin sections were used to perform H&E staining and 159 the pictures were taken using an optical microscope (Leica, Wetzlar, Germany) at 100, 200 and 160 400× magnification. The frozen sections were blocked in PBS supplemented with 10% FBS. RIPK1 161 (bs-5805R, dilution 1:200), RIPK3 (bs-3551R, dilution 1:200) and MLKL (bsm-33339M, dilution 162 1:200) antibodies were purchased from Bioss and the secondary antibodies (CoraLite594-163 conjugated recombinant rabbit anti-mouse IgG and CoraLite488-conjugated goat anti-rabbit IgG) 164 were both purchased from Proteintech. 4',6-Diamidino-2-phenylindole (DAPI, Solarbio) was used 165 to label the nucleus. Pictures were taken with a fluorescence microscope (Leica DMi8) at 200× 166 167 magnification.

168 Statistical analysis. All statistical analyses were performed using SPSS software (version 23.0; 169 IBM Corp., Armonk, USA). Data are presented as the mean \pm SD. *t*-Tests were used to analyze the 170 data between two groups. One-way ANOVA and Turkey's post hoc correction was performed when 171 there were multiple groups. P < 0.05 indicates statistical significance.

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173 **Results**

The expression of necroptotic genes was down-regulated in prostate cancer compared to 174 **normal prostate samples.** Necroptosis may contribute to prevent the progression of carcinogenesis 175 176 since the expression of its key mediator RIPK3 is decreased in several types of cancer cells. To validate the potential effect of necroptosis in prostate cancer, the expression of RIPK1, RIPK3 and 177 MLKL was explored in normal prostate and cancer tissues with data obtained through the GTEx and 178 TCGA databases. A higher expression of RIPK1 was observed in normal tissues compared to 179 prostate cancer tissues. Moreover, the expression of RIPK3 and MLKL was also down-regulated in 180 prostate cancer samples compared to either normal tissues or corresponding para-cancerous tissues 181 182 (Figures 1A, 1B). In consideration of these results, the diagnostic potential of RIPK1, RIPK3 and MLKL for prostate cancer was compared using a ROC analysis and results indicated that RIPK3 183 184 (AUC=0.903) had a higher diagnostic efficiency for prostate cancer compared to RIPK1

(AUC=0.685) and *MLKL* (AUC=0.854) (Figure 1C). Furthermore, survival analysis was performed
and lower expression of *RIPK3* was correlated with poor disease specific survival (Figure 1D).

20(S)-ginsenoside Rg3 induced the expression of necroptotic genes in prostate cancer cells. 187 Our previous research indicated that 20(S)-ginsenoside Rg3 induced cell cycle arrest in PC3 cells 188 [19]. In the present study, the inhibitory effects of 20(S)-ginsenoside Rg3 on cell proliferation were 189 190 observed both in 22RV1 and PC3 cells. The cells were treated with various doses of 20(S)ginsenoside Rg3 (0, 50 and 100 µM) for 24, 48 and 72 h. CCK8 assays confirmed that 50 and 100 191 192 µM 20(S)-ginsenoside Rg3 inhibited the proliferation of 22RV1 cells at all three time points (Figure 2A). In PC3 cells, 100 µM 20(S)-ginsenoside Rg3 inhibited cell proliferation at 24, 48 and 72 h, 193 while 50 μ M 20(S)-ginsenoside Rg3 did not exhibit the inhibitory effects on cell proliferation at 72 194 h (Figure 2B). The percentage of apoptotic and necrotic cells in 22RV1 and PC3 cells treated with 195 100 µM 20(S)-ginsenoside Rg3 for 48 h was also evaluated by flow cytometry with a FITC-196 Annexin V and PI apoptosis assay (Figure 2C). 20(S)-ginsenoside Rg3 increased the proportion of 197 late apoptotic/necrotic (PI⁺/FITC⁺) cells and decreased the proportion of viable cells (PI⁻/FITC⁻) in 198 both 22RV1 and PC3 cells (Figures 2D, 2E). Treatment with 100 µM 20(S)-ginsenoside Rg3 for 48 199 h did not regulate the expression and cleavage of caspase 3 both in 22RV1 and PC3 cells, 200 suggesting that 20(S)-ginsenoside Rg3 may induce necroptosis rather than apoptosis in prostate 201 cancer cells (Figure 2F). To explore the potential regulatory effects of 20(S)-ginsenoside Rg3 on 202 cell necroptosis, western blots were performed and the results showed that treatment with 100 μ M 203 20(S)-ginsenoside Rg3 for 48 h led to an increase in the expression of RIPK1, RIPK3 and MLKL, 204 suggesting that 20(S)-ginsenoside Rg3 might induce necroptosis in prostate cancer cells (Figures 205 3A, 3B). Pretreatment with the 100 µM RIPK1 inhibitor Nec-1 for 1 h at least partially reversed the 206 inhibitory role of 20(S)-ginsenoside Rg3 on the proliferation of prostate cancer cells (Figure 3C). 207

20(S)-ginsenoside Rg3 blocked the autophagy flux and up-regulated the expression of 208 209 necroptotic proteins in prostate cancer cells via ROS accumulation. ROS plays an important role in the modulation of cell fate. Our previous report indicated that 20(S)-ginsenoside Rg3 210 211 elevated ROS level in PC3 cells in a dose-dependent manner [19]. In the present study, the accumulation of ROS was observed both in 22RV1 cells and PC3 cells in the treatment with 100 212 213 µM 20(S)-ginsenoside Rg3 for 48 h using the DCFH-DA dye followed by flow cytometry analysis (Figure 4A). Since there is a crosstalk between cell autophagy and necroptosis, the regulatory effect 214 of 20(S)-ginsenoside Rg3 on cell autophagy was also explored. The lipidation of LC3 and the 215 protein expression of p62 were both increased by 20(S)-ginsenoside Rg3 (100 µM for 48 h) in 216 217 22RV1 and PC3 cells, suggesting that 20(S)-ginsenoside Rg3 inhibited the late autophagy and decreased the autophagy flux in prostate cancer cells (Figures 4B, 4C). NAC, a ROS scavenger, 218

reversed the regulatory role of 20(S)-ginsenoside Rg3 in LC3 lipidation and p62 protein expression, indicating that 20(S)-ginsenoside Rg3 blocked the autophagy flux via the accumulation of cellular ROS (Figures 4D, 4E). Moreover, pretreatment with 10 mM NAC for 1 h also blocked the upregulation of RIPK1, RIPK3 and MLKL protein expression in 22RV1 and PC3 cells induced by 100 μ M 20(S)-ginsenoside Rg3, suggesting that the compound might increase necroptosis via the accumulation of ROS and subsequent blockage of autophagy flux (Figures 4F, 4G).

- 20(S)-ginsenoside Rg3 up-regulated the expression of necroptotic proteins in PC3 mice 225 xenografts. To evaluate the *in vivo* effect of 20(S)-ginsenoside Rg3, a PC3 mouse xenograft model 226 was established by subcutaneous injection of PC3 cells in the right side of the armpit. Two weeks 227 later, daily gavage with PBS or 20(S)-ginsenoside Rg3 (25 mg per kg body weight) was performed 228 for additional 2 weeks. Then, the mice were sacrificed, and the volume and weight of the tumors 229 were measured. No obvious differences were observed for the volume of tumors between the 230 control and 20(S)-ginsenoside Rg3-treated groups though tumors in the latter looked smaller 231 (Figures 5A, 5B). However, the weights of tumors decreased after 20(S)-ginsenoside Rg3 treatment 232 (Figure 5C). HE staining was performed to detect the histological features of the tumors and the 233 results showed that 20(S)-ginsenoside Rg3 increased a necrosis-like phenotype in the tissues with 234 more ruptured nuclei and disintegrated cells (Figure 5D). IF staining was also performed and the 235 results showed that 20(S)-ginsenoside Rg3 led to an upregulation of the expression of RIPK1, 236 RIPK3 and MLKL in tumor tissues (Figure 5E). Finally, protein expression of RIPK1, RIPK3 and 237 MLKL was upregulated in xenografts isolated from the 20(S)-ginsenoside Rg3-treated group 238 compared to the control (Figures 5F, 5G). 239
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241 Discussion

Necroptosis is recognized as a programmed necrotic cell death and is triggered by several intrinsic 242 243 factors, such as tumor necrosis factor (TNF) and interferon (IFN) [21]. Although the formation of the necrosome, which is composed of RIPK1 and RIPK3 has been considered as a feature of 244 necroptosis, the actual mechanism is much more complicated. In addition to forming the necrosome, 245 RIPK1 can also mediate RIPK1-dependent apoptosis and the decision to which way the cells die 246 may be determined by mitogen-activated protein kinase 7 (MAP3K7, also known as TAK1) [22]. 247 RIPK3 was reported to mediate RIPK1-independent necroptosis via its interaction with TNFRSF1A 248 249 associated via death domain (TRADD) to activate the RIPK3-MLKL signaling pathway [23]. Recent research disclosed a new regulatory mechanism of necroptosis in which linear ubiquitin 250 251 chain assembly complex (LUBAC) was identified as a novel checkpoint for necroptosis. LUBAC and it mediated M1 poly-ubiquitination modification promoted MLKL membrane accumulation and 252

subsequent cell necroptosis in human cells without affecting the phosphorylation of RIPK1, RIPK3
and MLKL or necrosome formation [24]. In the present study, although the expression of RIPK1,
RIPK3 and MLKL was up-regulated in prostate cancer cells, the possible necroptotic mechanism
induced by 20(S)-ginsenoside Rg3 still needs further exploration.

- The relationship between ROS and necroptosis has been widely explored and abundant evidences 257 258 confirm they are positively correlated [25, 26]. Early research indicated that RIPK3 could activate several metabolic enzymes which induced aerobic respiration and oxidative respiration, resulting in 259 260 an increase in ROS production [27]. RIPK1 has been reported to inhibit the activity of the mitochondrial adenine-nucleotide translocase (ANT), which led to a decrease in ADP/ATP 261 exchange and ROS production [28]. Besides, many studies revealed the promoting role of ROS in 262 necroptosis. Previous research reported that ROS mediated the modification of RIPK1 cysteine 263 residues, which facilitated RIPK1 autophosphorylation and subsequent necrosome formation [29]. 264 Shikonin, a necroptosis inducer, promoted the overproduction of ROS in nasopharyngeal carcinoma 265 cells and glioma cells, which led to the upregulation of RIPK1 and RIPK3 expression, as well as the 266 induction of necroptosis [30, 31]. A recent study reported that the combined treatment between 267 resveratrol and docetaxel induced apoptosis and necroptosis in prostate cancer cells via ROS 268 production [32]. Another antiproliferative natural compound, curcumin, also induced prostate 269 cancer cell apoptosis and necroptosis, and the reduction of ROS levels could reverse the effects 270 induced by curcumin [33]. Our study also showed that 20(S)-ginsenoside Rg3 upregulated ROS 271 levels in prostate cancer cells, and scavenging ROS with NAC pretreatment antagonized 20(S)-272 ginsenoside Rg3-induced necroptotic protein expression. 273
- The crosstalk between ROS and cell autophagy was well reviewed in a recent report [34]. ROS 274 could transcriptionally and post-transcriptionally regulate cell autophagy, and in turn, autophagy 275 also regulated ROS levels through several pathways. In our study, 20(S)-ginsenoside Rg3-elevated 276 277 ROS levels induced changes in the autophagy flux in prostate cancer cells. Autophagy also plays an important role in the regulation of necroptosis [35]. An impaired autophagy flux contributed to the 278 279 induction of necroptosis [36]. In the prostate cancer cell line DU145, sorafenib induced the formation of ATG5-deficient autophagosomes and promoted the interaction between p62 and 280 281 RIPK1, subsequently triggering necroptosis [37]. Artepillin C (ArtC), a cinnamic acid derivative, induced apoptosis in 22RV1 cells. Co-treatment with ArtC and autophagy inhibitors not only 282 283 exacerbated apoptosis but also induced necroptosis, suggesting that the inhibition of autophagy may help trigger necroptosis in prostate cancer cells [38]. A previous report has indicated that 20(S)-284 285 ginsenoside Rg3 could inhibit autophagic flux in the late stages of autophagy and thus sensitized doxorubicin-induced cell death in hepatocellular carcinoma cell [39]. Our results were consistent 286

with the report and the autophagy flux was also blocked by 20(S)-ginsenoside Rg3 treatment with an increase in LC3 lipidation and p62 expression in prostate cancer cells, suggesting that the upregulation of p62 induced by 20(S)-ginsenoside Rg3 may contribute to the formation of the necrosome. However, the exact mechanism needs to be further explored.

The present study also evaluated the effect of 20(S)-ginsenoside Rg3 in PC3 mice xenografts. Results indicated that the treatment with 20(S)-ginsenoside Rg3 decreased tumor weight and increased necrotic areas in tumor tissues. Furthermore, 20(S)-ginsenoside Rg3 upregulated the expression of RIPK1, RIPK3 and MLKL in xenograft tissues. However, there was no noteworthy differences in tumor size between control and treated mice, which may be due to the short treatment time.

In conclusion, the present study showed that 20(S)-ginsenoside Rg3 induced the expression of necroptotic proteins in prostate cancer cells *in vitro* and *in vivo*. The underlying mechanism involved ROS accumulation and subsequent autophagy flux impairment.

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426 Figure Legends

[39]

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Figure 1. Analysis of necroptotic gene expression based on GTEx and TCGA databases. A) Expression of *RIPK1*, *RIPK3* and *MLKL* in normal prostate and prostate tumor tissues. B) Expression of *RIPK1*, *RIPK3* and *MLKL* in prostate tumor tissues and corresponding paracancerous tissues. C) ROC analysis of the diagnostic efficiency of the three genes in prostate cancer. D) Disease specific survival analysis of *RIPK3* expression in prostate cancer. PRAD, prostate adenocarcinoma. *p < 0.05, ***p < 0.001 vs. Normal

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Figure 2. 20(S)-ginsenoside Rg3 inhibited cell proliferation and increased the proportion of 435 necroptotic cells in prostate cancer cells. A) CCK8 assays to evaluate the effects of 20(S)-436 ginsenoside Rg3 on the proliferation of 22RV1 cells. B) CCK8 assays to evaluate the effects of 437 20(S)-ginsenoside Rg3 on the proliferation of PC3 cells. C) Flow cytometry analysis of apoptotic 438 and necrotic cells in prostate cancer cells treated with DMSO or 100 µM 20(S)-ginsenoside Rg3 for 439 48 h. D, E) Quantitative analysis of flow cytometry results. F) Western blot assays for the 440 expression and cleavage of caspase 3 in 22RV1 and PC3 cells treated with 100 µM 20(S)-441 ginsenoside Rg3 for 48 h. p < 0.05, p < 0.01 vs. CTRL 442

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Figure 3. 20(S)-ginsenoside Rg3 up-regulated the expression of necroptotic proteins in prostate cancer cells. A) Western blot assays for the expression of necroptotic proteins in 22RV1 and PC3 cells treated with 100 μ M 20(S)-ginsenoside Rg3 for 48 h. B) Quantitative analysis of protein expression in 22RV1 and PC3 cells. C) CCK8 assays to explore the proliferation of 22RV1 and PC3 cells pretreated with 100 μ M Nec-1 for 1 h. **P*<0.05, ***P*<0.01 vs CTRL

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Figure 4. 20(S)-ginsenoside Rg3 blocked autophagy flux and up-regulated the expression of necroptotic proteins in prostate cancer cells via overproduction of ROS. A) Flowcytometry analysis of ROS levels in 22RV1 and PC3 cells treated with 100 μ M 20(S)-ginsenoside Rg3 for 48 h. B) Western blot showing LC3 lipidation and p62 expression in 22RV1 and PC3 cells treated with 50 or 100 μ M 20(S)-ginsenoside Rg3 for 48 h. C) Quantitative analysis of the lipidation of LC3 and the expression of p62 in 22RV1 and PC3 cells treated with different doses of 20(S)-ginsenoside Rg3. D) Western blotting showing the effects of 100 μ M 20(S)-ginsenoside Rg3 on LC3 lipidation and p62 457 expression in 22RV1 and PC3 cells pretreated with 10 mM NAC for 1 h. E) Quantitative analysis of 458 LC3 lipidation and p62 expression in 22RV1 and PC3 cells. F) Western blotting showing the effects 459 of 100 μ M 20(S)-ginsenoside Rg3 on the expression of necroptotic proteins in 22RV1 and PC3 460 cells pretreated with 10 mM NAC for 1 h. G) Quantitative analysis of necroptotic proteins 461 expression in 22RV1 and PC3 cells. C, CTRL; R, 20(S)-ginsenoside Rg3; N, NAC; N+R, 462 NAC+20(S)-ginsenoside Rg3. *p < 0.05, **p < 0.01 vs. CTRL

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Figure 5. 20(S)-ginsenoside Rg3 up-regulated necroptotic protein expression in a PC3 mouse xenograft model. A) Xenografts isolated from control mice and 20(S)-ginsenoside Rg3-treated mice. B) tumor volume in both control and ginsenoside Rg3-treated mice. C) Tumor weights from control and 20(S)-ginsenoside Rg3-treated mice. D) HE staining of tumor tissues isolated from control and 20(S)-ginsenoside Rg3-treated mice. E) IF staining analysis of the expression of RIPK1, RIPK3 and MLKL in xenograft tissues. F) Protein expression of RIPK1, RIPK3 and MLKL in tumor tissues. G)

470 Quantitation analysis of the results of western blot. *p < 0.05, **p < 0.01 vs. CTRL

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Fig. 2 Download full resolution image





Fig. 4 Download full resolution image

