

1 **For the submission:**

2

3

4 **FOXO3 upregulates and activates GSDME to trigger myeloma cell pyroptosis**

5

6 Yaner Wang<sup>1,2,#</sup>, Yali Wang<sup>1,3,#</sup>, Yaoli Cui<sup>1,#</sup>, Yuanming He<sup>1</sup>, Ye Yang<sup>4</sup>, Wen Zhou<sup>5</sup>,  
7 Longlong Liu<sup>1</sup>, Hua Wang<sup>6</sup>, Mo Liu<sup>7</sup>, Yongqiang Wei<sup>8</sup>, Zhenqian Huang<sup>1</sup>, Xiaolei Wei<sup>8</sup>,  
8 Xinliang Mao<sup>1,2,4\*</sup>

9

10 <sup>1</sup> Department of Hematology, The Key Laboratory of Advanced Interdisciplinary  
11 Studies, The First Affiliated Hospital of Guangzhou Medical University; Guangdong  
12 Provincial Key Laboratory of Protein Modification and Degradation, School of Basic  
13 Medical Sciences, Guangzhou Medical University, Guangzhou 511436, P. R. China

14 <sup>2</sup> Institute of Clinical Pharmacology, Science and Technology Innovation Center,  
15 Guangzhou University of Chinese Medicine, Guangzhou, 510405, China

16 <sup>3</sup> Department of Clinical Pharmacology, College of Pharmaceutical Sciences,  
17 Guangzhou Medical University, Guangzhou 511436, P. R. China

18 <sup>4</sup> School of Medicine & Holistic Integrative Medicine, Nanjing University of  
19 Traditional Medicine, Nanjing 210023, China

20 <sup>5</sup> Cancer Institute, Central China University, Changsha, 410078, P. R. China

21 <sup>6</sup> Department of Hematology, Southern Medical University, Guangzhou, 510515, P. R.  
22 China

23 <sup>7</sup> State Key Laboratory of Oncology in South China, Guangdong Provincial Clinical  
24 Research Center for Cancer, Sun Yat-sen University Cancer Center, Guangzhou, 510060,  
25 China

26 <sup>8</sup>Sino-French Hoffmann Institute, Guangzhou Medical University, Guangzhou 511436,

27 P. R. China

28

29 **Supplemental Table S1. Specific primers for FOXO3 cloning**

<b>Primers for FOXO3 cloning</b>	
Forward	5'-TCTAGAATGGACTACAAGGATGACGA-3'
Reverse	5'-GCGGCCGCTCAGCCTGGCACCCAGCTCT-3'
<b>Specific primers the construction of Tet-On FOXO3</b>	
Forward	5'-CCTACCCTCGTAAAGAATTCGCCACCATGGCAGAGGCACCG GCTTCCCCG-3'
Reverse	5'-AGGGGAGGTGGTCTGGATCCTCACTTGTCATCGTCATCCTTG TAATC-3'

30

31 **Supplemental Table S2. Specific primers for RT-PCR for FOXO3, BAX, BCL2,**  
32 **BCL2A1, BCL2L1, BNIP3, BNIP3L, BNIP1, MCL1 and GAPDH**

<b>Primers for FOXO3</b>	
Forward	5'- GCGTTGCGTGCCCTACTT -3'
Reverse	5'- CACTGACTGTGCTGGCGTTA-3'
<b>Primers for GSDME</b>	
Forward	5'- CCTCACCTTGGCGATGT -3'
Reverse	5'- GGTCTGGATGCCCACGAT -3'
<b>Primers for GSDMD</b>	
Forward	5'- GACAACGTGTACGTGGTGACTG -3'
Reverse	5'- TGGAAGGTCCTCTGCTTCTTAT -3'
<b>Primers for BAX</b>	
Forward	5'- CCCGAGAGGTCTTTTCCGAG-3'
Reverse	5'- CCAGCCCATGATGGTTCTGAT-3'
<b>Primers for BCL2</b>	
Forward	5'- GGTGGGGTCATGTGTGTGG-3'
Reverse	5'- CGGTCAGGTACTCAGTCATCC-3'
<b>Primers for BCL2A1</b>	
Forward	5'-TACAGGCTGGCTCAGGACTAT -3'
Reverse	5'-CGCAACATTTTGTAGCACTCTG-3'
<b>Primers for BCL2L1</b>	
Forward	5'-GAGCTGGTGGTTGACTTTCTC-3'
Reverse	5'-TCCATCTCCGATTGAGTCCCT-3'

<b>Primers for BNIP3</b>	
Forward	5'-CAGGGCTCCTGGGTAGAACT -3'
Reverse	5'-CTACTCCGTCCAGACTCATGC-3'
<b>Primers for BNIP3L</b>	
Forward	5'-ATGTCGTCCCACCTAGTCGAG-3'
Reverse	5'-TGAGGATGGTACGTGTTCCAG-3'
<b>Primers for BNIPL</b>	
Forward	5'-GAGTCTGACTAAGGGGCCTG-3'
Reverse	5'-CTCCGAGTCTGAAGGTGTCT-3'
<b>Primers for MCL1</b>	
Forward	5'- GTAATAACACCAGTACGGACGG-3'
Reverse	5'- CCACAAACCCATCCTTGGAAG-3'
<b>Primers for GAPDH</b>	
Forward	5'- GGAAGATGGTGATGGGATT -3'
Reverse	5'- AACGGATTTGGTCGTATTG -3'

**Supplemental Table S3: Primers for qRT-PCR of FOXO3, GSDME, and GAPDH**

<b>Primers for FOXO3 (5'-3')</b>	
Forward	5'-CGGACAAACGGCTCACTCT-3'
Reverse	5'-GGACCCGCATGAATCGACTAT-3'
<b>Primers for GSDME (5'-3')</b>	
Forward	5'-ACATGCAGGTTCGAGGAGAAGT-3'
Reverse	5'-TCAATGACACCGTAGGCAATG-3'
<b>Primers for GAPDH (5'-3')</b>	
Forward	5'-GGAGCGAGATCCCTCCAAAAT-3'
Reverse	5'-GGCTGTTGTCATACTTCTCATGG-3'

**Supplemental Table S4: Primers for knockdown of FOXO3, BNIPL, and Bcl-xL**

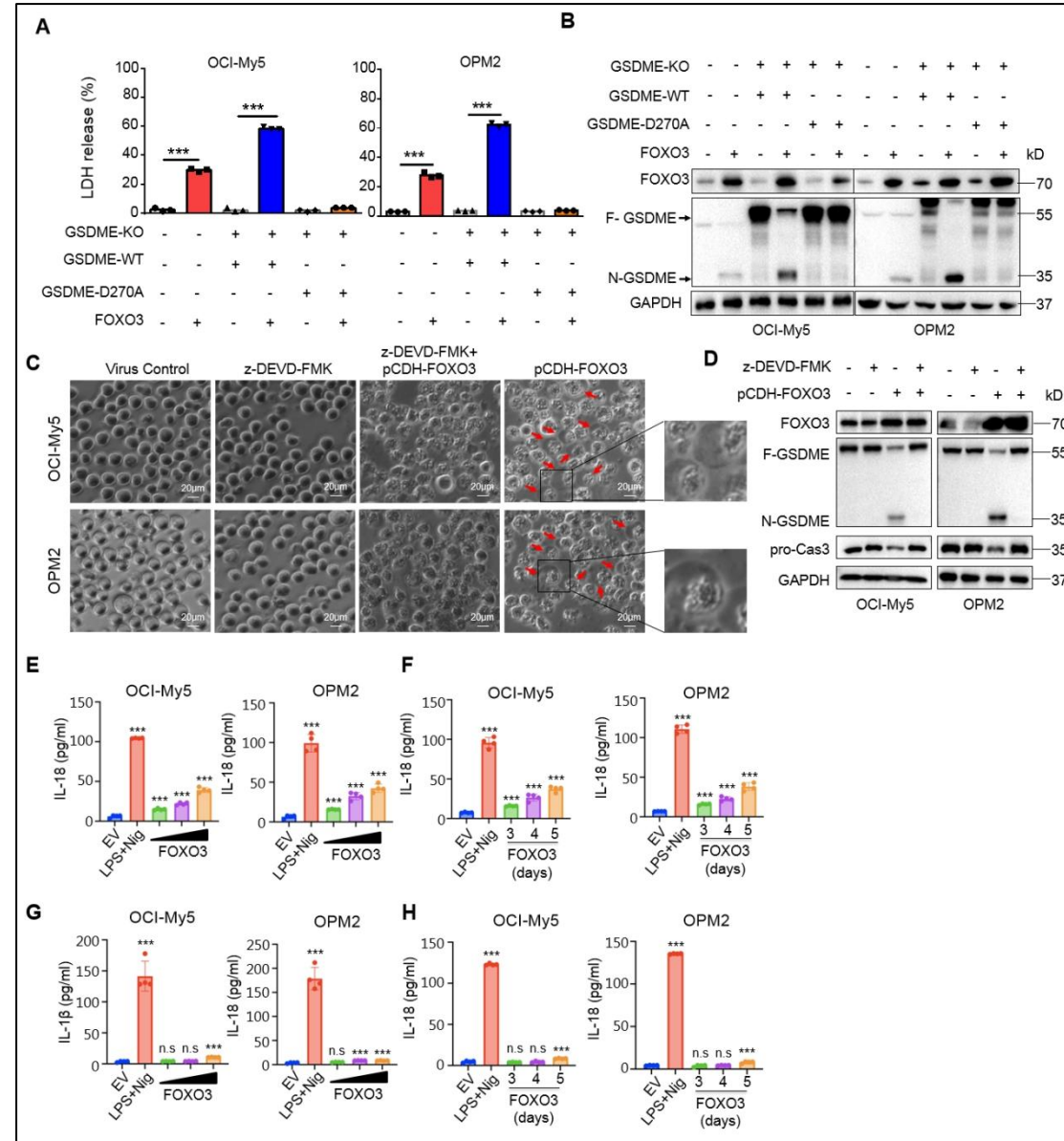
<b>Specific knockdown sequence for FOXO3</b>	
shFOXO3#1	5'-CCTACCCTCGTAAAGAATTCGCCACCATGGCAGAGGCA CCGGCTTCCCCG-3'
shFOXO3#2	5'-CGCTCGAAGTGGAGCTGGACCTTCAAGAGAGGTCCAG CTCCACTTCGAGCG

<b>Specific knockdownsequence for Bcl-xL</b>	
shBcl-xL #1	5'-GCTCACTCTTCAGTCGGAAAT-3'
shBcl-xL #2	5'-GTGGAACCTCTATGGGAACAAT-3'
<b>Specific knockdownsequence for BNIPL</b>	
shBNIPL#1	5'-GCTGGACAGTGGACATGAATTCTCGAGAATTCATGTCC ACTGTCCAGC-3'
shBNIPL#2	5'-GCTGGATACGTCAGTGTTACCTTCAAGAGAGGTAACAC TGACGTATCCAGC-3'

39 **Supplementary Table S5. The primers for the regulatory region of GSDME**

<b>GSDME (1~2000): P0</b>	
Forward	5'-TGCAGGTGCCAGAACATTTCTCTATCGATAGGTACCCCCA AAGCACCCCCAAGGCTGG-3'
Reverse	5'-AGGACACCTTCTTATTGCTTCTACTCATAACTAAAAGGTCT GCGGAGATTC-3'
<b>GSDME (-500~-1): P4</b>	
Forward	5'-TGCAGGTGCCAGAACATTTCTCTATCGATAGGTACCCCCA AA-3'
Reverse	5'-TGGTGGCTTTACCAACAGTACCGGAATGCCAAGCTTCATA AC-3'
<b>GSDME (-1000~-501): P3</b>	
Forward	5'-TGCAGGTGCCAGAACATTTCTCTATCGATAGGTACC-3'
Reverse	5'-TGGTGGCTTTACCAACAGTACCGGAATGCCAAGCTT-3'
<b>GSDME (-1501~-1001): P2</b>	
Forward	5'-TGCAGGTGCCAGAACATTTCTCTATCGATAGGTACC-3'
Reverse	5'-TGGTGGCTTTACCAACAGTACCGGAATGCCAAGCTT-3'
<b>GSDME (-2000~-1500): P1</b>	
Forward	5'-TGCAGGTGCCAGAACATTTCTCTATCGATAGGTACC-3'
Reverse	5'-TGGTGGCTTTACCAACAGTACCGGAATGCCAAGCTT-3'
<b>GSDME-ΔFRE1 (Δ1)</b>	
Forward	5'-CAATTTCTGCCTCCAAAGGAAGAATAAAAGGCAGAAATG AAATCCACAGG-3'
Reverse	5'-CTGTGGATTTCATTTCTGCCTTTTATTCTTCCTTTGGAGGC AGAAATTGG-3'
<b>GSDME-ΔFRE2 (Δ2)</b>	
Forward	5'-ATAGGTACCTATAGACTGTATTCTTATAAGAGAAAAGCAA ATGTTACGAAG-3'

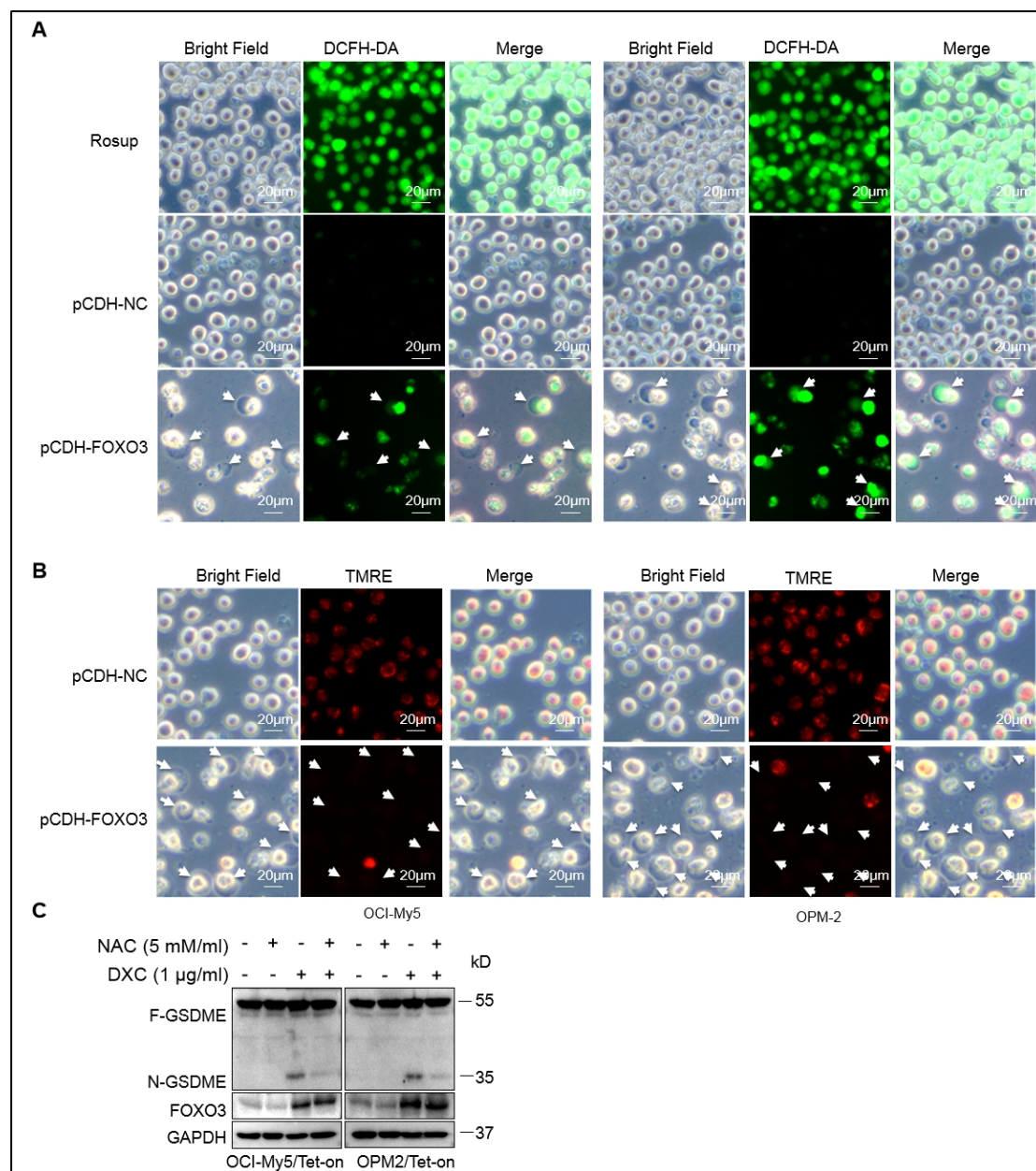
Reverse	5'-TGGTGGCTTTACCAACAGTACCGGAATGCCAAGCTTGGGG GCTGCATGCAC-3'
<b>GSDME-ΔFRE1ΔFRE2 (GSDME-Δ1Δ2)</b>	
Forward	5'-GATAGGTACCTATAGACTGTATTCTTATAAGAGAAAAGCA AATGTTACGAAGAAAAAC-3'
Reverse	5'-GCCTTTTATTCTTCCTTTGGAGGCAGAAATTGGGCATAAGA CAATAC-3'



**Suppl. Fig. S1. GSDME is required for FOXO3-induced myeloma cell pyroptosis.**

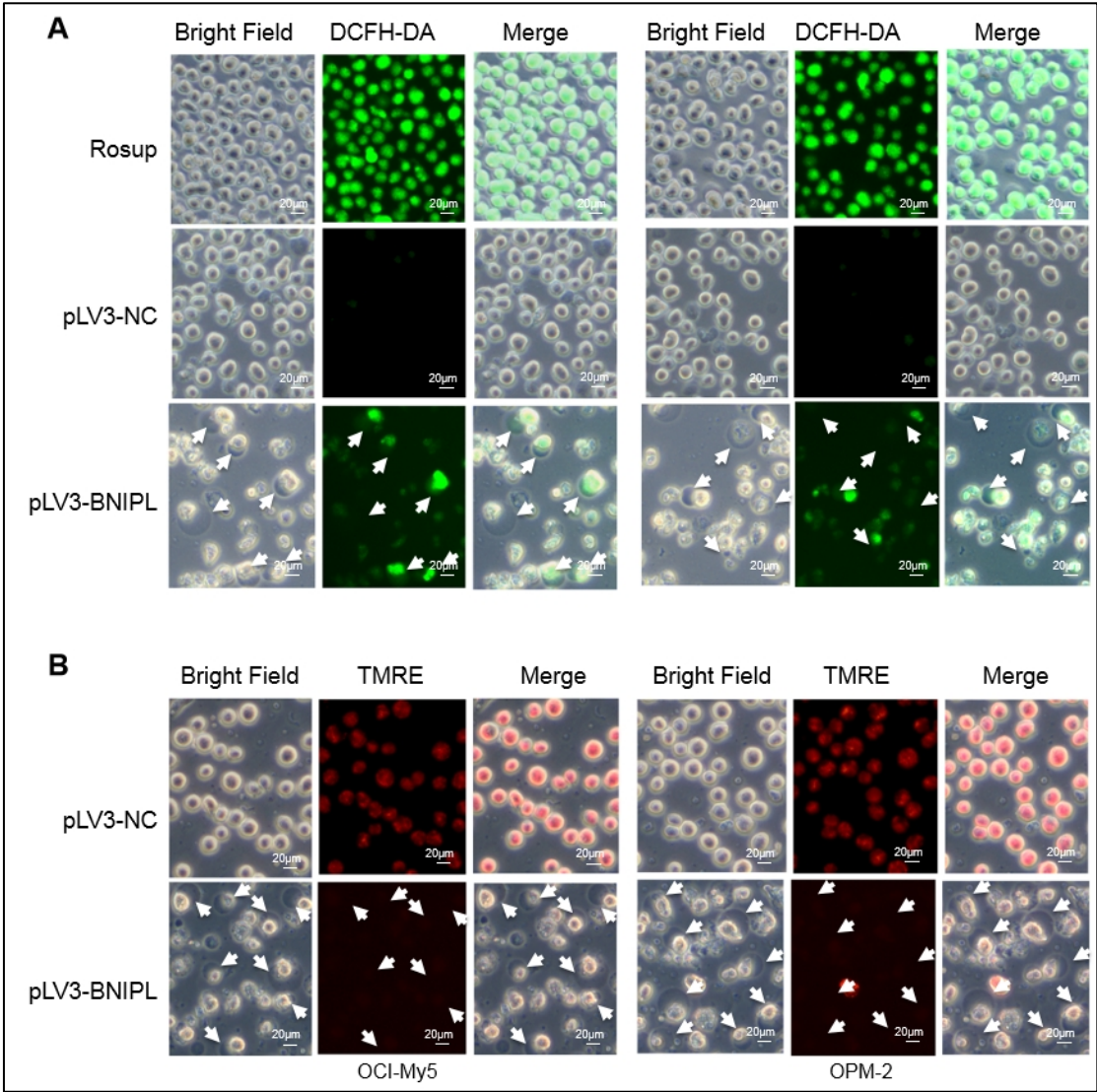
OCI-My5 and OPM2 were knocked out GSDME, followed by infection with FOXO3 along with wild-type or D270A mutant GSDME. A, LDH in culture media was measured. B, Cell

lysates were subjected to IB assays. C-D, OCI-My5 and OPM2 cells were infected with lentiviral FOXO3 for 72 h, followed by zDEVD-FMK treatment. Cell pyroptosis was subjected to Phase-contrast microscope analysis (C) and the cell lysates were subjected to WB for GSDME and Caspase-3 activation(D). E-F, MM cells were infected with lentiviral FOXO3 with increased titration (E) for 3 days or increased incubation days (F), followed by measurement of IL-18 by using an ELISA kit. LPS/Nigericin was used as a positive control. G-H, MM cells were infected with lentiviral FOXO3 with increased viral particles of infection (G) for 3 days or increased incubation days (H), followed by measurement of IL-1 $\beta$  by using an ELISA kit. LPS/Nigericin was used as a positive control.n.s., not significant. \*\*\*,  $P < 0.001$ .



**Suppl. Fig. S2. FOXO3 triggers ROS production and decreases mitochondrial membrane potential (MMP) in MM cells.** A, MM cells were treated the positive control Rosup or infected with FOXO3 lentivirus. Cells were then stained with DCFH-DA to measure ROS production (A) or stained with TMRE to measure MMP (B). Arrows indicated pyroptotic cells. C, OCI-My5 and OPM2 stably infected with Tet-On/FOXO3 lentivirus were pre-treated with N-acetyl cysteine (NAC) for 5 h, followed by treatment with doxycycline (DXC) for 48 h. Cell lysates were then prepared for IB assays.



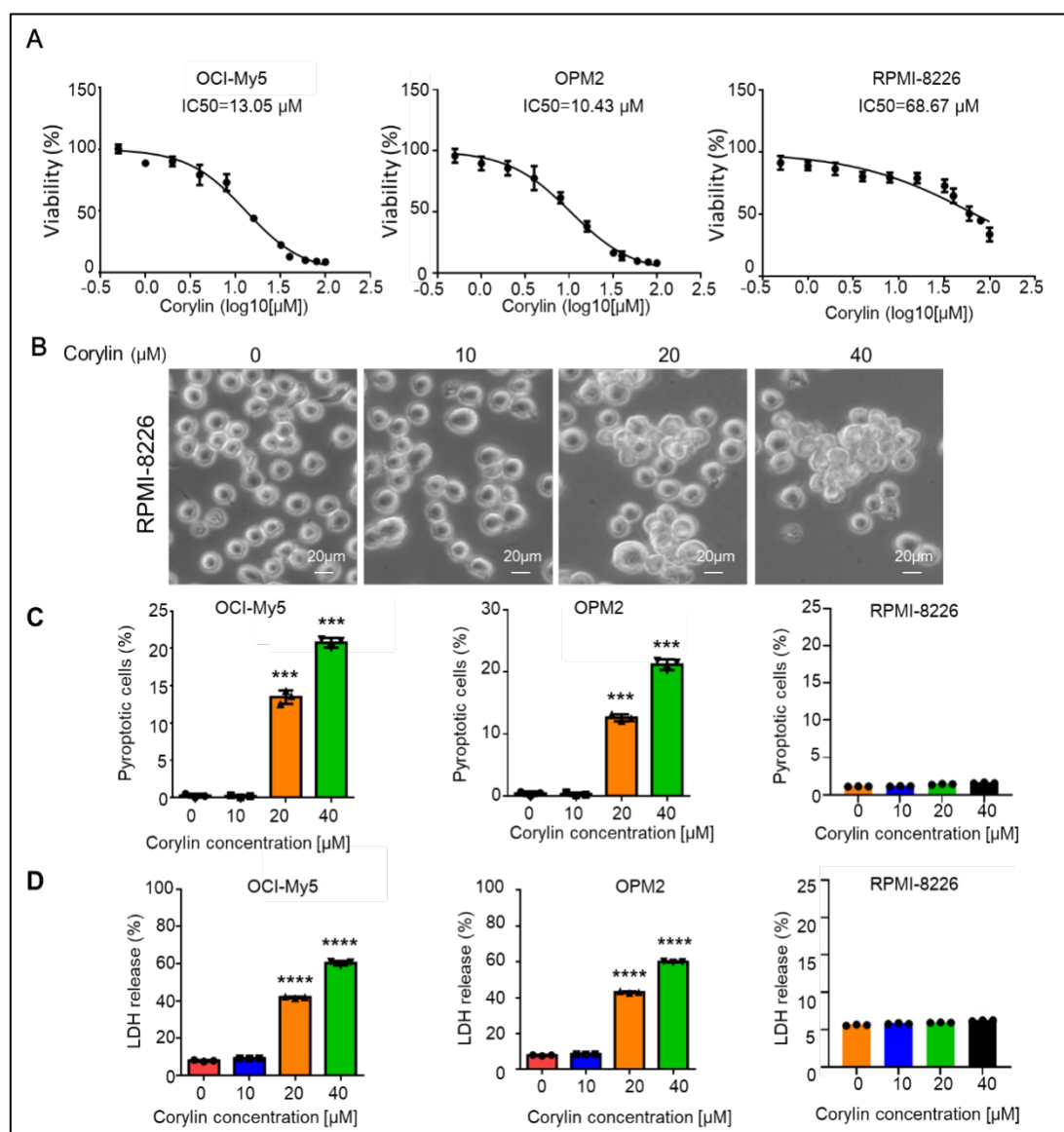


66

67        **Suppl. Fig. S3. BNIP1 overexpression elevated intracellular ROS levels and**  
68 **decreased MMP in MM cells.** A, MM cells were treated the positive control Rosup or  
69 infected with BNIP1 lentivirus. Cells were then stained with DCFH-DA to measure ROS  
70 production (A) or stained with TMRE to measure MMP (B). Arrows indicated pyroptotic  
71 cells.

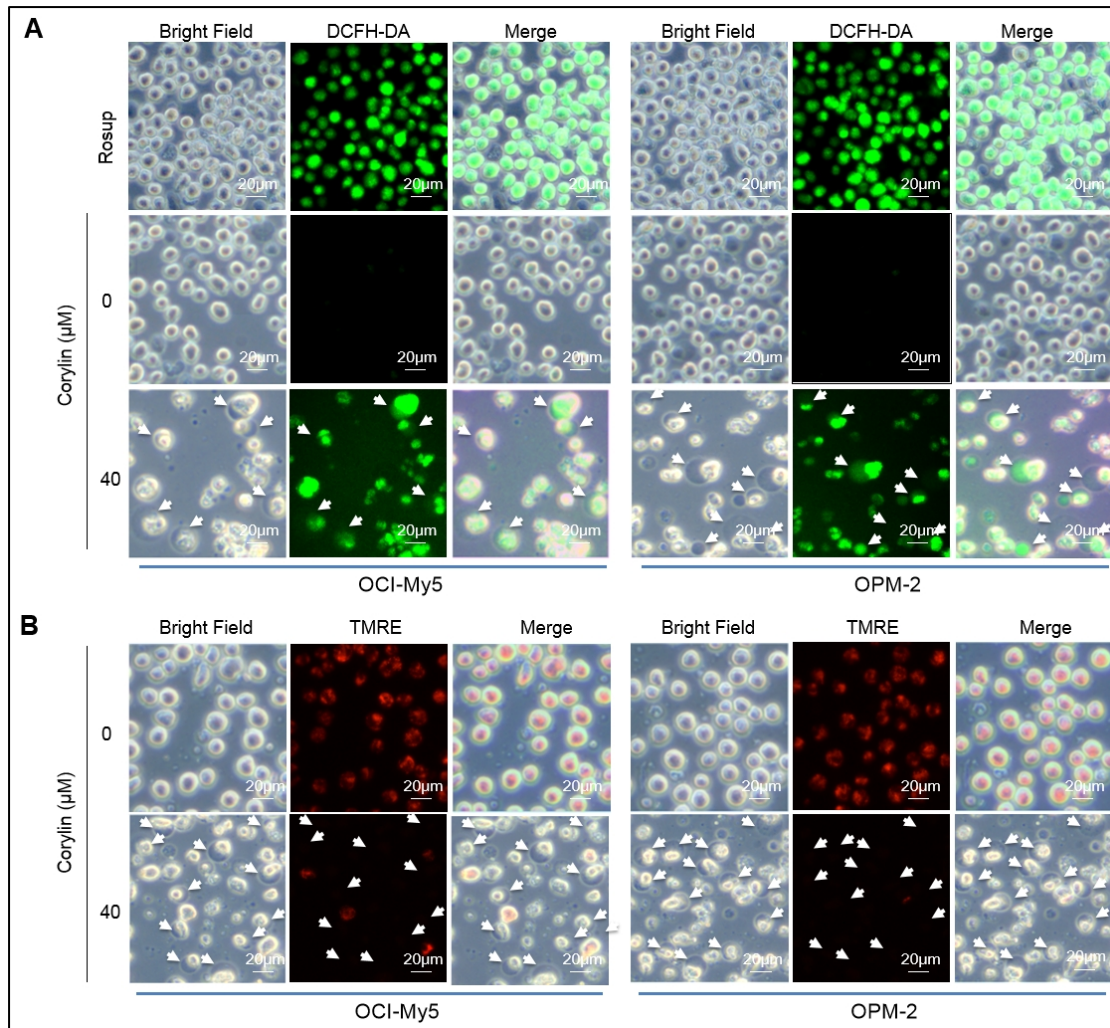
72





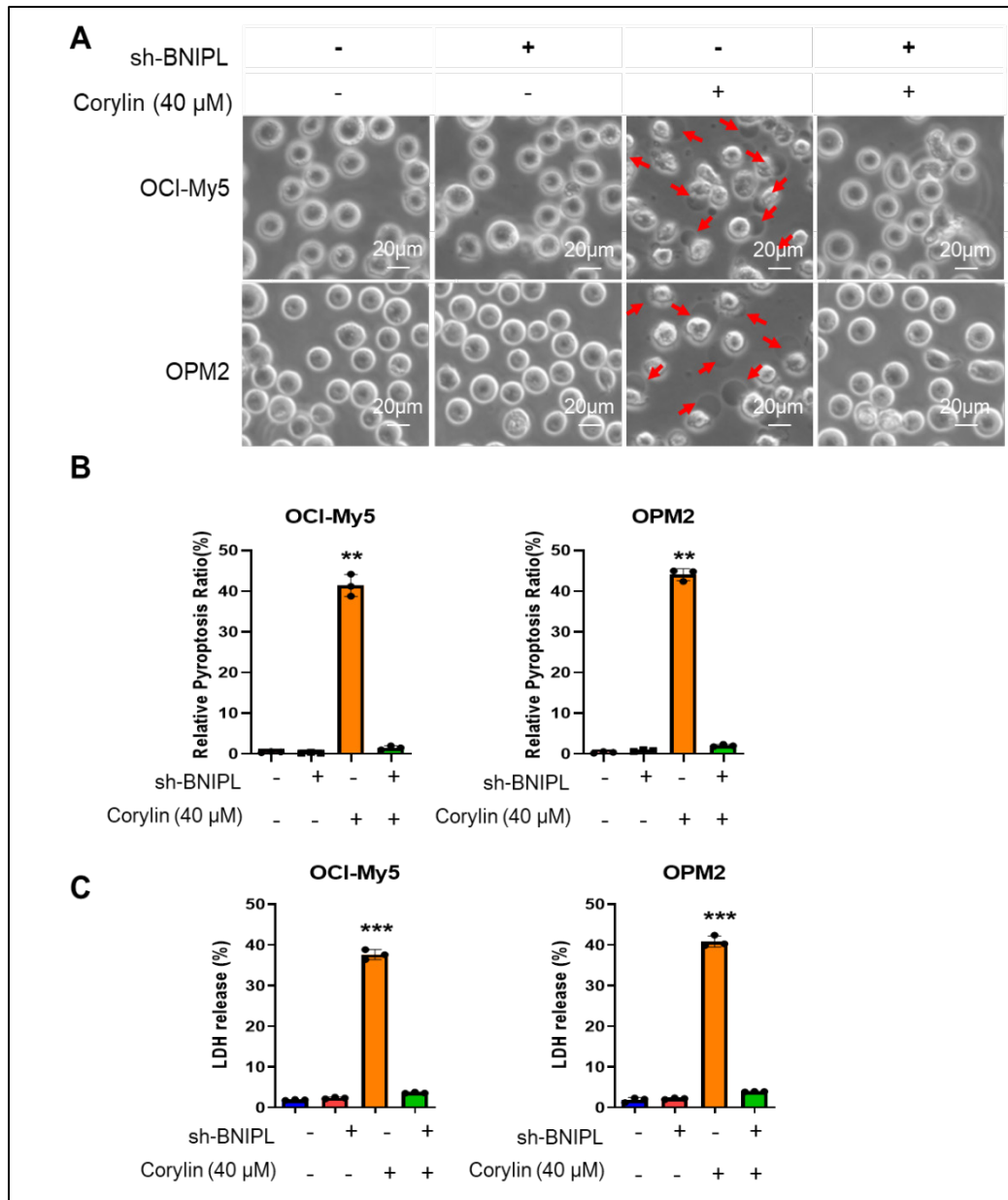
**Suppl. Fig. S4. Corylin induces MM cell pyroptosis in a GSDME-dependent manner.**

A, OCI-My5, OPM2 and RPMI-8226 cells were incubated with increased corylin for 48 h followed MTT assay. IC50 was calculated. B, RPMI-8226 cells were treated with increased corylin for 24 h, followed by Phase-contrast microscopy analysis. C, Statistic analysis of pyroptotic cells from corylin-treated MM cells. D, LDH release into culture media after cells were treated with corylin was measured. \*\*\*,  $P < 0.001$ ; \*\*\*\*,  $P < 0.0001$ .



**Suppl. Fig. S5. Corylin treatment elevates ROS levels and reduces MMP in MM cells.**

A-B, MM cells were treated the positive control Rosup or Corylin for 24 hrs. Cells were then stained with DCFH-DA to measure ROS production (A) or stained with TMRE to measure MMP (B). Arrows indicated pyroptotic cells.



**Suppl. Fig. S6. BNIPL is required for Corylin to induce MM cell pyroptosis.** A-C, MM

cells were knocked down BNIPL followed by incubation with increased corylin for 48 hrs. Cell

pyroptosis was analyzed by Phase-contrast microscopy (A), pyroptosis ratio (B) and LDH

measurement (C). \*\*\*,  $P < 0.001$ . Arrows indicated pyroptotic cells.