

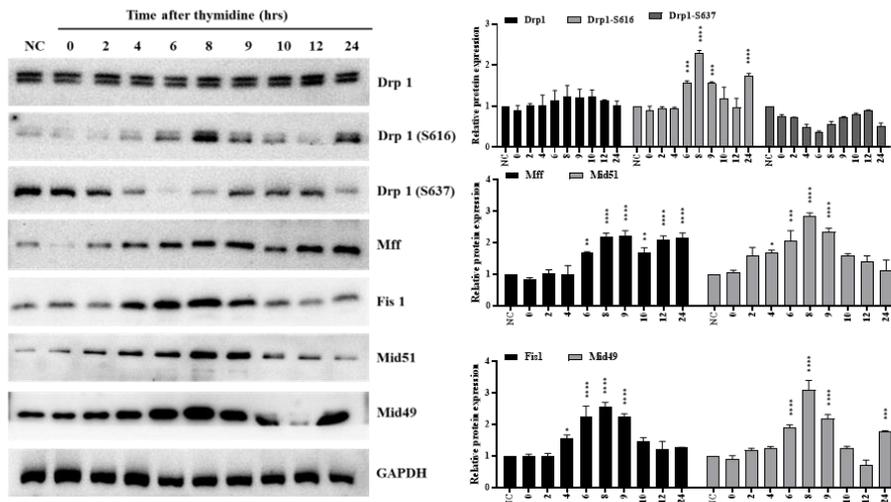
## **Supplemental information**

### **The phospho-Drp1-637/Fis1/Mid49 cluster modulates mitochondrial periphery fission signature responses to mitotic arrest**

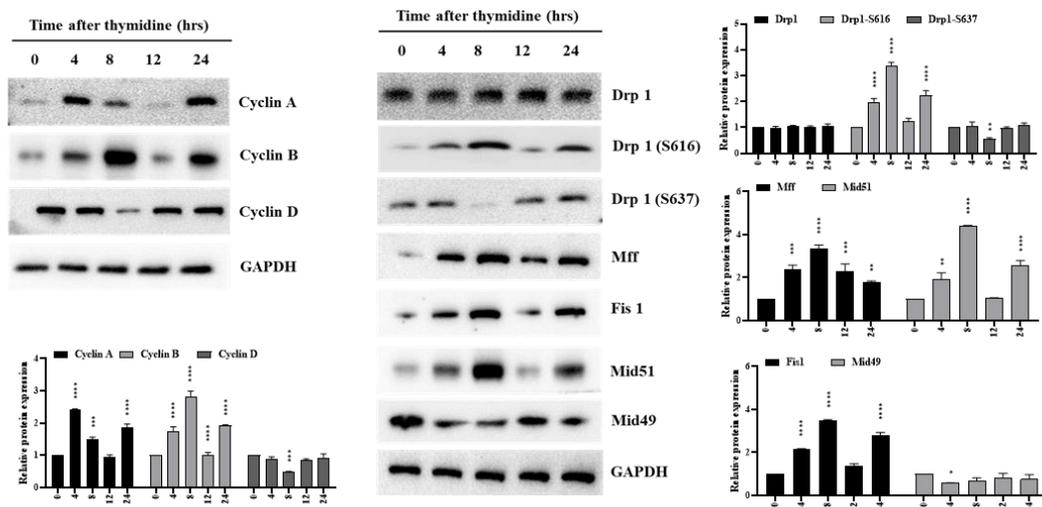
**Huey-Jiun Ko, Cheng-Yu Tsai, Thi-Huong Nguyen, Tai-Shan Cheng, Chi-Ying F. Huang, Nian-Siou Wu, Chun-Li Su, Po-Yu Tsai, Shean-Jaw Chiou, Ann-Shung Lieu, Joon-Khim Loh, Yi-Ren Hong**

1 **Figure S1**

A



B



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3 **Figure S1. Dynamics of Drp1 Phosphorylations during Cell Cycle Progression.** (A)

4 HeLa cell synchronization was achieved using a double-thymidine block, and the

5 expression levels of total Drp1, phospho-Drp1 (Ser616), phospho-Drp1 (Ser637), Mff,

6 Fis1, MiD49, and MiD51 were assessed at indicated time points after release. (B) HeLa

7 cells were synchronized through double thymidine treatment and subsequently released

8 for the specified durations. Expression levels of total Cyclin A, Cyclin B, Cyclin D,

9 Drp1, phospho-Drp1 (Ser616), phospho-Drp1 (Ser637), Mff, Fis1, MiD49, and MiD51

10 were analyzed at indicated time points after release. GAPDH served as an internal

11 loading control. The term "NC" (normal control) refers to samples without

12 synchronization or double-thymidine treatment.



15 **Figure S2. Effects of Mdi, Doxo, Oxa, and 5-Fu on Drp1 Phosphorylation and**  
16 **Mitochondrial Receptor Proteins.** (A) HeLa cells, synchronized through double-  
17 thymidine treatment, were treated with various concentrations of Mdi, Doxo, Oxa,  
18 and 5-Fu, then released for specified durations. Expression levels of total Drp1,  
19 phospho-Drp1 (Ser616/Ser637), Mff, Fis1, MiD49, and MiD51 were analyzed at 1, 4,  
20 8, and 12 hours after release by western blotting. (B) Cytoplasmic and mitochondrial  
21 fractions were separated to quantify total Drp1, phospho-Drp1 (Ser-616/Ser-637),  
22 Mff, Fis1, Mid49, and Mid51. VDAC1 served as an internal control, and subcellular  
23 purity was confirmed with GAPDH. "NC" denotes untreated samples, and "M"  
24 represents synchronized/mock-treated samples.

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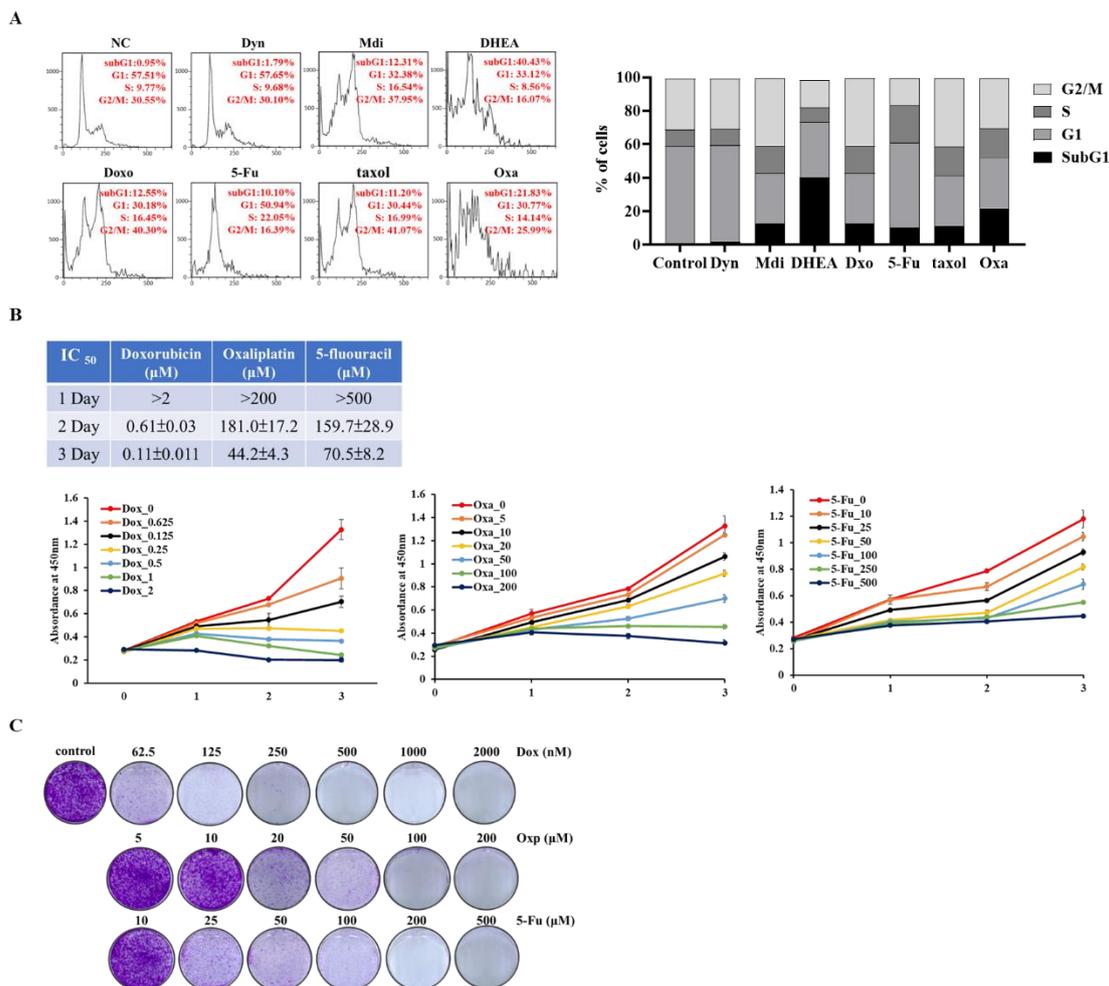
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38 **Figure S3**

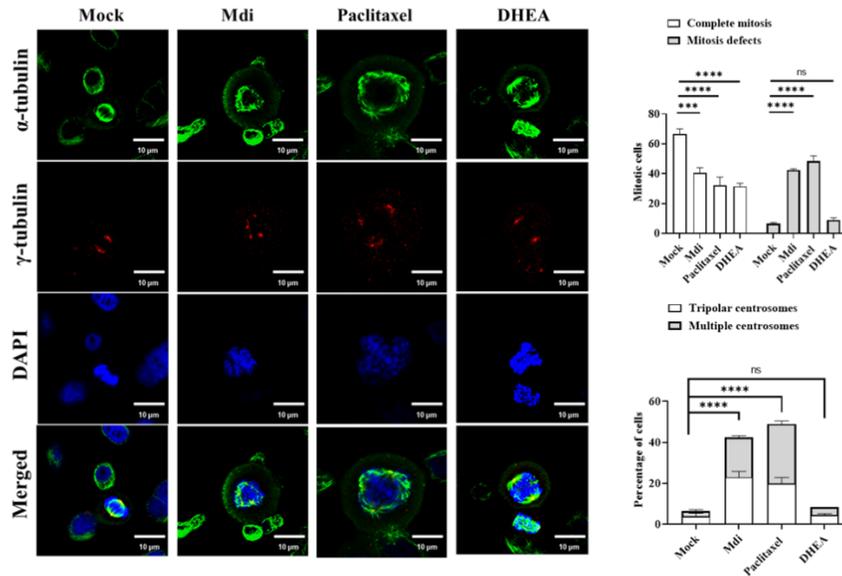


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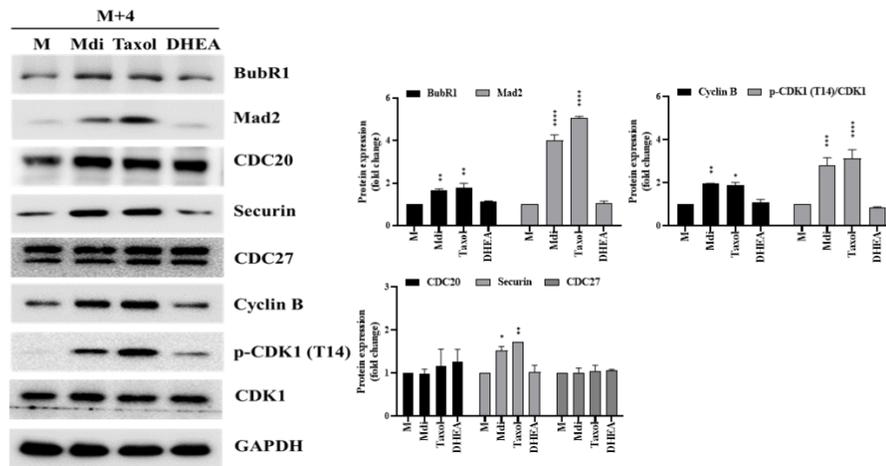
40 **Figure S3. Different drugs induce cell death and G2/M cell cycle arrest in HeLa**  
 41 **cells.** (A) Cell cycle analysis of HeLa cells following treatment with various drugs. The  
 42 histogram illustrates the percentages of cells in different cell cycle phases. The bar chart  
 43 below shows the distribution of cells across different cell cycle phases. (B) Assessment  
 44 of the effects of the different drugs on HeLa cell viability. HeLa cell viabilities were  
 45 evaluated after treating with Dox, Oxa, and 5-Fu at indicated concentrations from 24  
 46 hours to 72 hours. The right panel represents % cell viability versus the logarithm of  
 47 the concentration, and IC<sub>50</sub> values were obtained for Dox, Oxa, and 5-Fu. (C) The  
 48 proliferation of HeLa cells was suppressed by Dox, Oxa, and 5-FU. The ability of HeLa  
 49 cells to form colonies was assessed following treatment with Dox, Oxa, and 5-Fu.

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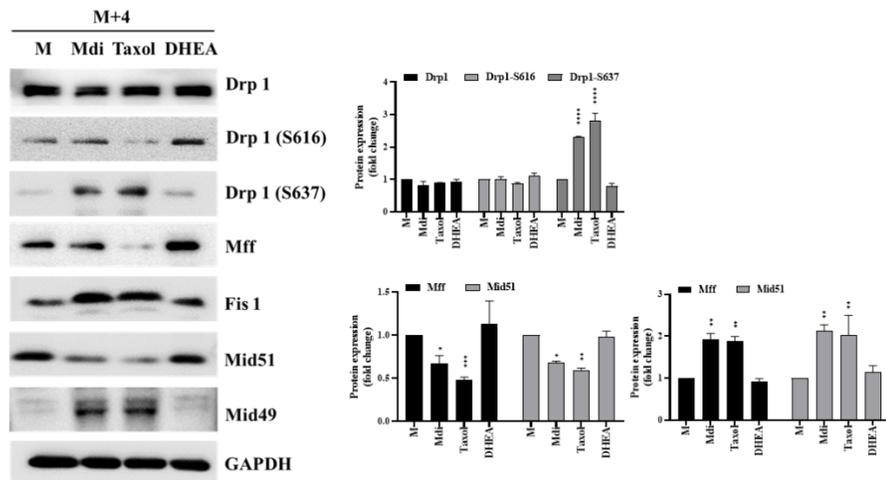
**A**



**B**



**C**



53 **Appendix Figure S4. Induced in nocodazole-treated mitotic HeLa cells are**  
54 **multiple spindle foci by taxol and DHEA. (A)** Confocal images depict microtubules  
55 (green), centrosomes (red), and DNA (blue) in HeLa cells after a 16-hour nocodazole  
56 treatment. (B) Impact of Mdi, Taxol and DHEA on the SAC during the M phase. (C)  
57 Dynamic changes in Drp1 phosphorylation and mitochondrial adaptor protein levels  
58 during treatment with Mdi, Taxol, and DHEA in nocodazole-treated mitotic HeLa  
59 cells. The scale bar is 10  $\mu$ m, and the cells were stained with anti- $\alpha$ -tubulin and anti- $\gamma$ -  
60 tubulin, with DNA stained using DAPI. Inhibition of multipolar spindle formation in  
61 HeLa cells by drug treatment is shown in panels C and F on the right. The percentage  
62 of cells exhibiting bipolar spindles and multiple spindle foci is presented. The data  
63 represent the mean  $\pm$  SE from three independent experiments, with approximately  
64 300–350 mitotic cells analyzed per experiment.

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