



p53 accumulation following cytokinesis failure in the absence of caspase-2

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Dear Editor

Caspase-2 is the most conserved member of the caspase family, but its physiological function(s) remains a matter of some debate [1]. Recent work suggests that caspase-2 deficiency in mice makes animals more susceptible to tumorigenesis in various models leading to the concept that caspase-2 is a tumor suppressor [1–3]. One of the key characteristics of caspase-2 knockout (KO) MEFs in culture and cells from caspase-2-deficient tumors is a high degree of chromosomal instability (CIN), suggesting that caspase-2 prevents accumulation of aneuploid cells and CIN [1–4]. Real time imaging of KO cells following prolonged spindle-assembly checkpoint arrest suggests that caspase-2 is required for the deletion of cells carrying mitotic aberrations and this requires the catalytic function of caspase-2 [5]. Another model that may explain increased CIN is suggested by Fava et al. [6]. Their data suggest that supernumerary centrosomes trigger PIDDosome (a caspase-2 activating platform containing Pidd and Raidd)-dependent caspase-2 activation, which then cleaves MDM2, resulting in p53 stabilization and p21-dependent cell cycle arrest [6]. Thus PIDDosome-mediated caspase-2 activation is predicted to be required to restrain polyploidization, preventing CIN, and therefore cancer. The main caveat with such a model is that, unlike *Casp2* KO animals, *Pidd* and *Raidd*

KO mice are not susceptible to enhanced tumorigenesis indicating that PIDDosome is not responsible for the tumor suppressor function of caspase-2 [7, 8].

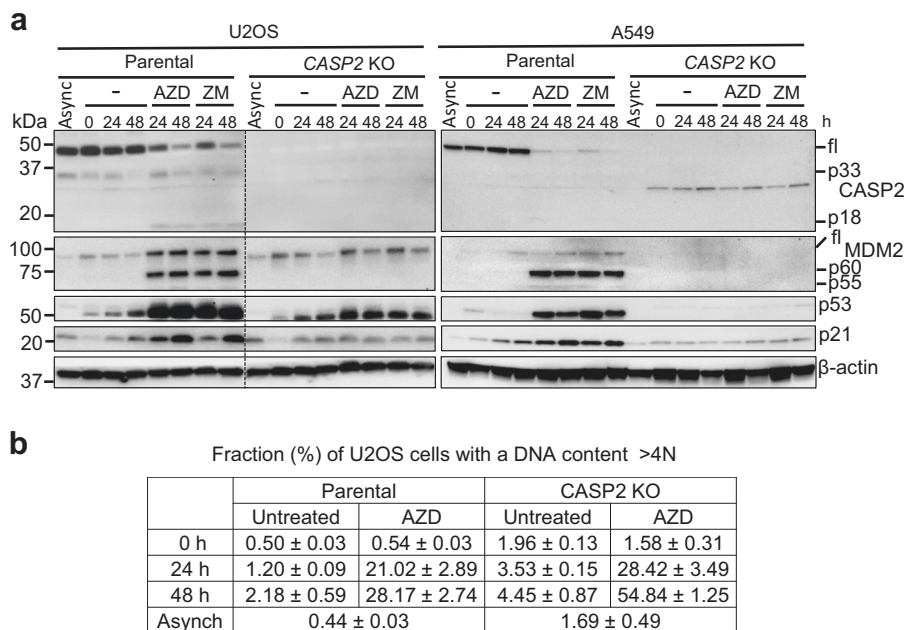
Most of the studies by Fava et al. used A549 lung adenocarcinoma cells, likely because it is a good in vitro model for cell cycle studies. To examine this pathway in other cell types, we generated *CASP2* KO U2OS (human osteosarcoma with wild-type p53) cells and compared them with A549 cells following treatment with the Aurora kinase B (AURKB) inhibitors, ZM447439 (as used by Fava et al.) or AZD1152-HQPA, to induce cytokinesis failure (Suppl. Fig. S1a). AZD1152-HQPA is >3000 times more specific toward AURKB compared with AURKA, while ZM447439 used by Fava et al. in most of their studies has lower selectivity [9]. The results confirm some of the observations reported by Fava et al., including MDM2 cleavage and increased levels of p53 and p21 in parental cells treated with the inhibitors (Fig. 1a). We did not observe any difference between the two AURKB inhibitors. Fava et al. [6] found that *CASP2* KO in A549 cells results in complete abrogation of MDM2 cleavage, p53 accumulation, and cell cycle arrest of tetraploid cells following cytokinesis failure. Our data are also consistent with these findings in *CASP2* KO A549 cells (Fig. 1a). However, we observed that *CASP2* KO U2OS cells, still showed increased p53 levels that appear to be independent of MDM2 cleavage in response to the AURKB inhibitors (Fig. 1a). The p53 response was not as robust as in the parental U2OS cells, as indicated by reduced p21 levels in the absence of caspase-2, and this has previously been documented [1, 4]. Interestingly, we also observed that MDM2 cleavage increases at 24 h and then decreases at 48 h following treatment in the non-synchronized A549, but not in U2OS cells (Suppl. Fig. S1b), which is consistent with previous data [6]. The data here, suggest different responses to AURKB inhibition in different cell types. From a functional perspective, DNA content analysis of *CASP2*KO U2OS cells treated with AZD1152-HQPA showed increased accumulation of polyploid (>4N) cells when compared to parental U2OS cells.

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Fig. 1 Aurora kinase B inhibition results in p53 accumulation in both parental and *CASP2* KO U2OS cells. U2OS and A549 cells were synchronized in G1/S by treatment with 2 mM thymidine for 24 h. After 3 h release, the cells were treated with DMSO or AURKB inhibitors (400 nM AZD1152-HQPA (AZD) or 2 μ M ZM447439 (ZM)) for 0, 24, and 48 h, followed by immunoblot and fluorescence-activated cell sorting (FACS) analyses. **a** Representative immunoblots of cell lysates from treated parental and *CASP2* KO cells. Antibodies used for immunoblotting are as indicated. **b** Percentage of U2OS parental and *CASP2* KO cells with polyploid (>4N) DNA content following AZD treatment. Async, asynchronous



However, these results are not as dramatic as reported for A549 cells. In particular, only marginal differences were observed at 24 h following treatment with AZD1152-HQPA (Fig. 1b). We found that U2OS cells are efficiently blocked in cytokinesis by AZD1152-HQPA, ruling out any possibility that the observations are affected by asynchronous cell populations (Suppl. Fig. S1c). Our data indicate that cytokinesis failure can trigger p53-mediated cell cycle arrest in the absence of caspase-2. This suggests the presence of alternative pathways and cell-type specific differences may drive caspase-2 responses to prevent polyploidization.

As AURKB has various functions in mitosis including mitotic condensation, spindle-assembly checkpoint and cytokinesis [10], inhibiting AURKB by drugs such as AZD1152-HQPA can cause not only cytokinesis failure but also prolonged mitosis [11] and DNA damage, that can induce a p53 response and lead to cell cycle arrest. This, in turn may also contribute to differential responses to these drugs in different cell types [12].

In conclusion, we found that MDM2 cleavage in response to cytokinesis failure is not essential for cell cycle arrest and p53 accumulation in all cell types. Importantly, p53-mediated cell cycle arrest can still occur in the complete absence of caspase-2.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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