

1 **Methylglyoxal induces nuclear accumulation of p53 and**
2 **γH2AX in normal and cancer cells**

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26 **Abstract**

27 The formation and accumulation of methylglyoxal (MGO) is associated with age-related
28 diseases such as diabetes, cancer and neurodegenerative disorders. MGO is the major
29 precursor of non-enzymatic glycation of macromolecules affecting their function and
30 structure. We now show for the first time that MGO stress not only led to cellular aging
31 responses like DNA double-strand breaks indicated by an accumulation of γ H2AX in the
32 nucleus. We also observed an immediate increase of Ser15 phosphorylated p53 in the nucleus
33 of MGO treated cell lines which will change the cellular expression pattern with adverse
34 effects on the cell cycle and other cellular functions not necessarily related to aging.

35 **Introduction**

36 The formation and accumulation of methylglyoxal (MGO), the most potent glycating agent in
37 humans, is associated with age-related diseases such as diabetes, obesity, atherosclerosis,
38 cancer and neurodegenerative disorders [1]. Methylglyoxal is mainly formed by the non-
39 enzymatic degradation of triose phosphates, glyceraldehyde-3-phosphate (G3P) and
40 dihydroxyacetone-phosphate (DHAP), as a byproduct of glycolysis. It takes place in all cells
41 and organisms. [2, 3]. The actual concentration of MGO in the cell depends on several factors
42 like the rate of detoxification by glyoxalases, the phosphate pool of the cell and the rate of
43 influx of carbon sources [4].

44 MGO is the major precursor of non-enzymatic glycation of proteins, lipids and DNA,
45 subsequently leading to the formation of a heterogeneous group of molecules, collectively
46 called AGEs (advanced glycation endproducts) [1, 3, 5]. The glycation reactions can cause

47 the formation of complexes and irreversible adducts of these macromolecules affecting their
48 function and structure. Accumulation of AGEs compromises cellular processes resulting in
49 mitochondrial dysfunction, genomic instability, loss of proteostasis, inflammaging and
50 cellular senescence [6]. Moreover, other studies revealed that elevated MGO levels may also
51 have beneficial effects in cancer and lifespan [7, 8].

52 Tumor cells differ from non-tumor cells concerning their high glycolytic rates under
53 anaerobic conditions (Warburg effect). This inefficient energy production, cause a high rate
54 of glucose uptake and glycolysis in tumors resulting in elevated MGO levels [9]. Loarca *et al.*
55 reported that treatment of hepatocellular carcinoma (HCC) cell lines with MGO promote the
56 localization of p53 into the nucleus whereas total cellular p53 levels are not altered [10]. p53
57 is a well-known tumor suppressor that is mutated in many tumor cells [11]. It functions as a
58 transcription factor, which is involved in the regulation of the cell cycle, DNA repair and
59 apoptosis. Upon cellular stress such as DNA damage, hypoxia or viral infection p53 is
60 activated and stabilized by different post-translational modifications interfering with its
61 degradation [12]. It was shown that DNA damage induces phosphorylation of p53 on Ser15
62 [13]. This impairs the binding to MDM2, promoting the activation of p53 and thereby leading
63 to cell cycle arrest or apoptosis [14, 15].

64 Using several tumor and non-tumor cell lines we showed that MGO stress led to DNA
65 double-strand breaks, and an immediate increase of Ser15 phosphorylated p53 in the nucleus.
66 This upregulation of nuclear phospho-p53 seems not accompanied with an increase in
67 apoptosis rate but will likely activate p53 dependent signaling pathways and thereby change
68 the cellular expression pattern with adverse effects on the cell cycle and other cellular
69 functions. The application of MGO as a compound to mimic increased glycation in cells is a

70 widely accepted method particular in the aging research field. Therefore, we suggest
71 monitoring phosphorylated p53 and γ H2AX in MGO treated cells in age related issues to
72 exclude p53 signaling interfering with age response.

73 **Materials and methods**

74 **Reagents**

75 Methylglyoxal (MGO) was purchased from Sigma-Aldrich (catalog no. M0252). Primary
76 antibodies used were anti-phospho-p53 (Cell Signalling, catalog no. 9286), anti-p53 (Santa
77 Cruz), anti-phospho-Histone H2A.X (Ser139) clone JBW301 (Millipore, catalog no. 05-636),
78 anti-cleaved caspase3 (5A1E) (Cell Signalling, catalog no.9664) and anti-PARP (Cell
79 Signalling, catalog no.9542). F-actin was visualized with Atto 546 Phalloidin (Sigma-
80 Aldrich). Alexa Fluor-conjugated and IRDye® (800CW, 680RD) secondary antibodies were
81 from Thermo Fisher Scientific or Li-Cor (Bad Homburg, Germany), respectively. DNA was
82 stained with DAPI (Sigma-Aldrich).

83 **Cell Culture and Methylglyoxal (MGO) treatment**

84 The breast cancer cell line MDA-MB-468 was cultured at 37°C and 5% CO₂ in RPMI
85 medium supplemented with 10% (v/v) fetal calf serum (FCS) and Antibiotic-Antimycotic
86 (Thermo Fisher Scientific, Schwerte, Germany). The keratinocyte cell line HaCaT, cervical
87 adenocarcinoma Hela cells and neuroblastoma SH-SY5Y cells were cultured at 37°C and 5%
88 CO₂ in DMEM high glucose medium supplemented with 10% (v/v) fetal calf serum (FCS),
89 10% sodium pyruvate and Antibiotic-Antimycotic (Thermo Fisher Scientific). Primary
90 human fibroblasts were cultured at 37°C and 5% CO₂ in Dulbecco's Modified Eagle Medium

91 (DMEM)/Hams F12 supplemented with 8% (v/v) fetal calf serum (FCS), 2% Ultrosor G
92 (PALL Life Sciences) and 2 mM L-glutamine (Thermo Fisher Scientific).

93 In all experiments Methylglyoxal (MGO) (Sigma-Aldrich) was diluted to 0.1 mM, 0.5 mM or
94 1 mM and the cells were incubated for 1h and 18h respectively.

95 **Immunofluorescence and Microscopy**

96 For immunofluorescence analysis, 8×10^4 cells (12-well plates) were grown on glass
97 coverslips, treated with 0.5 mM or 1 mM MGO for 1h and 18h respectively, washed twice
98 with PBS, fixed with 3.7% formaldehyde in PBS for 15 min, permeabilized with 0.2% (v/v)
99 Triton X-100 in PBS for 10 min and blocked in 10% FCS/1% BSA/0.05% Triton X-100 (v/v)
100 in PBS for 30 min. Primary antibodies were diluted 1:200 and incubated for 1 h at room
101 temperature. Alexa-conjugated secondary antibodies were diluted 1:200 and applied for 1 h at
102 room temperature. Samples were covered with ProLong Gold antifade reagent (Life
103 Technologies) and imaged using an Apotome-containing Axio Observer.Z1 (Zeiss, Jena,
104 Germany) equipped with a 63x oil objective and a monochrome Axiocam MRm camera.
105 Representative images are shown.

106 **Apoptosis assay**

107 2×10^5 cells were subjected to the Caspase-3/CPP32 Fluorometric Assay Kit (BioVision)
108 according to the manufacturer's protocol. Absorbance at 505 nm was measured after 2 h of
109 incubation at 37°C using a microplate reader. Quantitative results were calculated from at
110 least three independent biological experiments.

111 **Western Blot**

112 Cells were washed twice with ice-cold phosphate-buffered saline and lysed in RIPA buffer
113 (50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 2 mM EDTA, 1% (v/v) Triton X-100, 0.1% SDS,
114 Complete EDTA-free protease inhibitor cocktail) for 30 min on ice. The proteins were
115 separated on SDS-PAGE, transferred to polyvinylidene fluoride (PVDF) membranes (Merck
116 Millipore), incubated with indicated primary antibodies used at 1:200–1:5000 dilutions and
117 incubated overnight at 4°C. Fluorophore-labeled secondary antibodies used at 1:10.000
118 dilutions (LI-COR, Bad Homburg, Germany) were incubated for 1 h at room temperature.
119 The fluorescence signals were detected with ODYSSEY CLx (LI-COR) and quantified by the
120 associated software. Quantitative results were calculated from at least three independent
121 biological experiments.

122 **Results and Discussion**

123 In this study we analyzed the impact of the glycating agent Methylglyoxal on the
124 phosphorylation of p53 at Ser-15 and its subsequent translocation to the nucleus. We
125 compared different non-tumor and tumor cell lines to figure out if this response to MGO is
126 rather a more general than a cell-type specific (depending on the genetic background) answer.
127 Here we treated non-tumor cell lines (HaCaT keratinocytes, primary fibroblasts) and tumor
128 cells (cervical adenocarcinoma Hela cells, breast cancer MDA-MB-468 cells and
129 neuroblastoma SH-SY5Y cells) with concentrations of MGO between 0.1 mM and 1 mM
130 which are generally used in this research field. The localization of Ser-15 phosphorylated p53
131 was analyzed 1 h and 18 h after treatment in the above mentioned cell lines. In non-tumor
132 cells (HaCaT cells) as well as in tumor cells (MDA-MB-468 cells) MGO treatment caused a
133 significant increase of Ser15 phosphorylated p53 in the nucleus (Fig 1A-D). We found that

134 MGO increased nuclear phospho-p53 levels up to ~67 fold in HaCaT cells and up to ~1.8
135 fold in MDA-MB-468 cells (Fig 2). However, in Hela cells and primary human fibroblasts,
136 MGO treatment had no effect on the localization of phospho-p53 (S1 Fig; data not shown)
137 and in SH-SY5Y cells the nuclear localization of phospho-p53 is already very high without
138 MGO treatment (data not shown).

139 **Fig 1. Analysis of the expression and localization of phospho-p53 in HaCaT cells and**
140 **breast cancer MDA-MB-468 cells after treatment with MGO.** Representative images of
141 HaCaT (A,B), MDA-MB-468 cells (C,D) treated with indicated MGO concentrations for 1h
142 (A,C) and 18 h (B,D). The cells were stained for phospho-p53 together with phalloidin (F-
143 actin). Merged images show phospho-p53 (green) and F-actin (red). Scale bars, 20 μ m.

144 **Fig 2. MGO treatment results in an increase in nuclear localized phospho-p53 in**
145 **HaCaT and MDA-MB-468 cells.** Percentage of cells showing nuclear localization of
146 phospho-p53 normalized to the total cell number. All error bars represent \pm SEM, $n \geq 2$, $N > 50$.
147 All *P*-values were calculated using an unpaired two-sample Student's *t*-test ($*P \leq 0.05$,
148 $**P \leq 0.01$).

149 It was shown previously that MGO is a cytotoxic agent, which can lead to apoptosis in cancer
150 cells through the generation of ROS [16], DNA modification and DNA-protein crosslink
151 [17]. To answer the question if DNA damage esp. DSBs induced by MGO is a prerequisite
152 for the increase in nuclear localization of phospho-p53 we determined the phosphorylation
153 level of histone H2AX in HaCaT cells and human primary fibroblasts upon MGO treatment.
154 Phosphorylated H2AX (γ H2AX) specifically binds to damaged DNA and can be visualized
155 via immunofluorescence staining. Upon binding of γ H2AX to damaged DNA several proteins

156 are recruited for instance BRCA1 and p53 binding protein1 to induce DNA repair [18]. We
157 could demonstrate that MGO treatment already after 1 h increased nuclear γ H2AX
158 accumulation in HaCaT cells and human primary fibroblasts (Fig 3 and 4). Since in human
159 primary fibroblasts, MGO treatment does not lead to nuclear p53 translocation, DNA double
160 strand breaks seem to be not sufficient to trigger nuclear localization of phospho-p53.

161 **Fig 3. Analysis of DNA double-strand breaks and apoptosis rate in HaCaT cells after**
162 **treatment with MGO.** Representative images of HaCaT cells treated with indicated MGO
163 concentrations for 1h (A) and 18h (B). The cells were stained for γ H2AX together with
164 cleaved caspase. Merged images show γ H2AX (green), cleaved caspase (red) and DNA
165 (blue). Scale bars, 20 μ m. C. Apoptosis rate in HaCaT cells by measuring caspase-3 activity
166 using the Caspase-3/ CPP32 Fluorometric Assay Kit. All error bars represent \pm SEM, n=3. D.
167 Representative western blot of PARP in total cell lysates treated with indicated MGO
168 concentrations for 18h. GAPDH was used as a loading control. Quantification of PARP
169 protein levels normalized to GAPDH. All error bars represent \pm SEM, n=3.

170 **Fig 4. Analysis of DNA double-strand breaks and apoptosis rate in primary human**
171 **fibroblasts after treatment with MGO.** A. Representative images of primary human
172 fibroblasts treated with indicated MGO concentrations for 1h. The cells were stained for
173 γ H2AX together with cleaved caspase. Merged images show γ H2AX (green), cleaved
174 caspase (red) and DNA (blue). Scale bars, 20 μ m. B. Apoptosis rate in primary human
175 fibroblasts after treatment with indicated MGO concentrations for 1 h by measuring caspase-
176 3 activity using the Caspase-3/ CPP32 Fluorometric Assay Kit. All error bars represent \pm
177 SEM, n=3.

178 High levels of activated p53 may drive cells into an apoptotic fate. Thus we asked whether
179 the number of apoptotic cells is altered upon MGO treatment. To this end, expression and
180 activity of Caspase3 and PARP cleavage was investigated. We could demonstrate that upon
181 treatment of HaCaT cells and human primary fibroblasts with MGO the apoptotic rate was
182 not significantly changed (Fig 3A-C, Fig 4). Also PARP as one of the main cleavage target of
183 caspase3 in vivo ([19]) was not affected (Fig 3D).

184 p53 is activated upon many different stress factors influencing important cellular processes
185 either positively or negatively depending on the cellular context [20, 21]. Therefore, p53 is
186 thought to be required for the maintenance of cellular and particularly genomic integrity. This is
187 supported by the finding, that 50% of human tumors carry loss of function mutations in the
188 p53 gene [11, 22-25]. 95% of these detected mutations are located in the DNA binding
189 domain of p53 [26]. In the cell lines we used in this study MGO treatment leads to different
190 p53 responses. Only in HaCat and MDA-MB-468 cells MGO treatment causes increased
191 accumulation of Ser15 phosphorylated p53 in the nucleus. Interestingly in these cell lines the
192 p53 gene carries mutations in the DNA binding domain altering its ability to bind to p53
193 response elements and thereby changing its transcriptional activity. The MDA-MB-468 cell
194 line is hemizygous for a mutated p53 gene (R273C) [11, 27]. Whereas the spontaneously
195 immortalized skin keratinocyte cell line, HaCaT, possesses three point mutations (H179Y,
196 D281G, R282W) of p53 on both alleles [28]. In contrast to that, HeLa cells and probably also
197 primary fibroblasts which show no change in the nuclear localization of Ser15
198 phosphorylated p53 possess exclusively wild type alleles of p53. In these cells it is supposed
199 that other mechanisms may lead to a carcinogenic inactivation of p53, for instance HPV
200 infection for cervical carcinoma or increase in protein stability in neuroblastoma [29, 30].

201 In summary, in this study we showed for the first time that MGO stress leads to an immediate
202 increase of Ser15 phosphorylated p53 in the nucleus of several cell lines. This upregulation of
203 nuclear phospho-p53 seems not accompanied with an increase in apoptosis rate but will likely
204 activate p53 dependent signaling pathways. This would fit with recently published data
205 showing an activation of the p53 pathway upon MGO treatment in human umbilical vein
206 endothelial cells (HUVECs) [31]. Moreover, it was shown that the total amount of nuclear
207 p53 is increased in hepatocellular carcinoma cells upon MGO treatment. However, the type
208 of posttranslational modification of p53 was not analyzed. The authors correlate the nuclear
209 increase of p53 upon MGO treatment with a decreased migration, invasion and adhesion
210 phenotype in these cells [10].

211 The application of MGO as a compound to mimic increased glycation in cells is a widely
212 accepted method particular in the aging research field. Nevertheless, besides the enormous
213 progress in understanding AGE related phenotypes and diseases in more detail some critical
214 aspects of MGO treatment arises. It is important that the used concentrations of MGO are
215 comparable to the actual physiological concentration in this specific tissue and blood
216 respectively [1]. By using UPLC-MS/MS plasma levels of MGO in healthy individuals have
217 been estimated at ~60-250 nM, cellular levels ranges from ~1-5 μ M MGO [1, 32, 33]. In cell
218 culture model systems often near toxic concentrations of MGO ranging from 50 μ M up to 2
219 mM MGO are applied. For instance, HUVEC cells were treated with up to 800 μ M MGO to
220 copy injuries in endothelial cells typically seen in diabetes patients knowing that these
221 concentrations do not reflect the plasma concentrations of MGO even in diabetes patients
222 [31]. The authors argued that the in vivo situation is rather a complex mixture of compounds
223 acting in synergy than a single compound leading to this phenotype [31]. In addition, the

224 application of MGO in cell culture is different compared to the in vivo situation and thereby
225 MGO can act more direct on the cells. To exclude unphysiological responses of MGO
226 treatment in general, we suggest establishing a protocol for each cell line utilized, which
227 includes the analysis of the Ser-15 phosphorylated p53 and γ H2AX status to exclude
228 undesirably effects of MGO which would otherwise overlap with aging mechanisms (Fig 5).

229 **Fig 5. Model of MGO induced stress response.** MGO stress leads to DNA double-strand
230 breaks, and an immediate increase of phosphorylated p53 in the nucleus of MGO treated cell
231 lines. The subsequent activation of p53 dependent signaling pathways is supposed to overlap
232 with the aging response in these cells. We suggest monitoring phosphorylated p53 and
233 γ H2AX in MGO treated cells to exclude undesirable effects of MGO interfering with aging
234 response.

235

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238

239 **References**

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313 **Supporting information**

314 **S1 Fig. Analysis of the expression and localization of phospho-p53 in primary human**
315 **fibroblasts after treatment with MGO.** Representative images of primary human

316 fibroblasts treated with indicated MGO concentrations for 1h (A) and 18 h (B). The cells
317 were stained for phospho-p53 together with phalloidin (F-actin). Merged images show
318 phospho-p53 (green) and F-actin (red). Scale bars, 20 μ m.

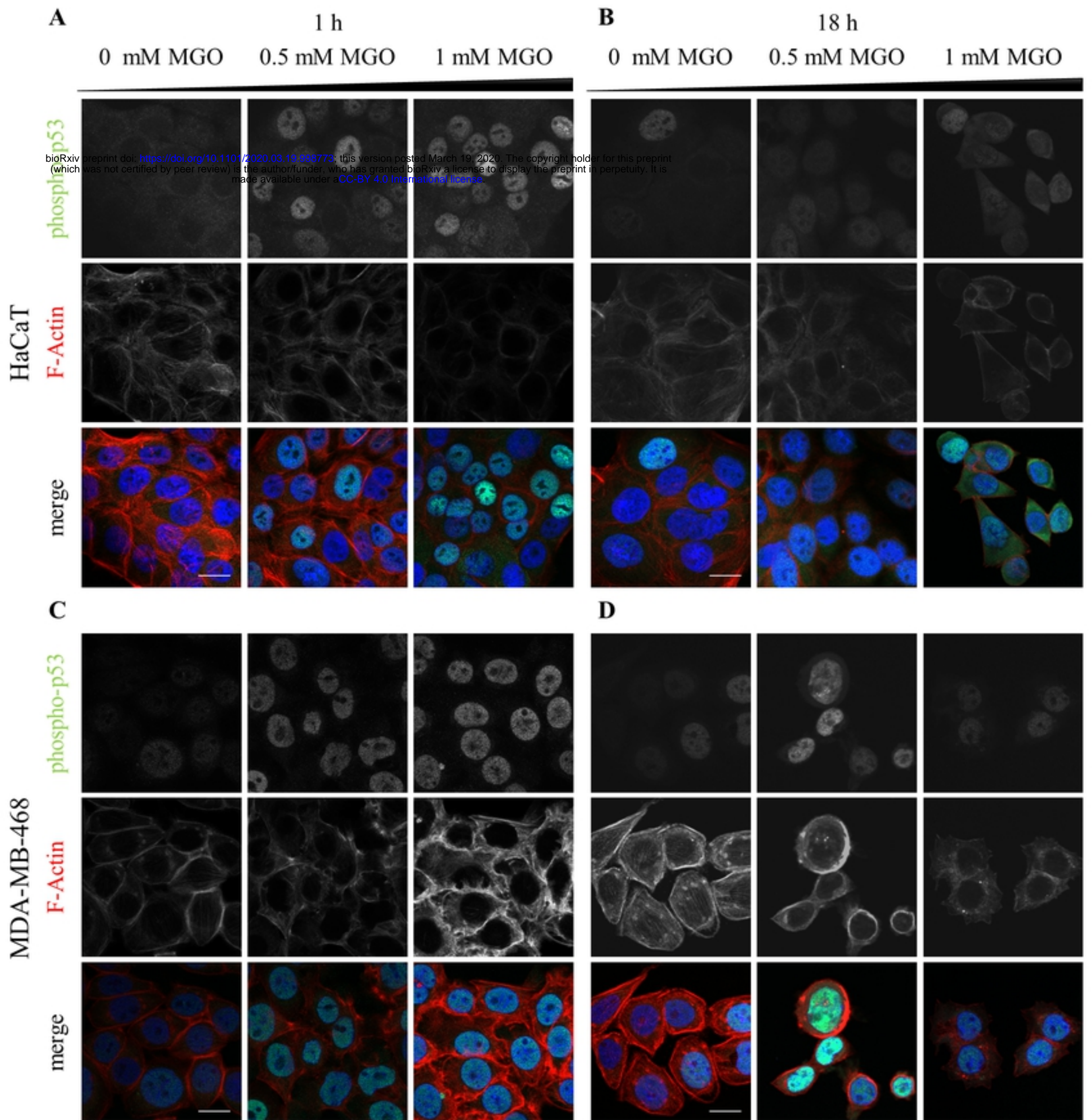


Fig 1

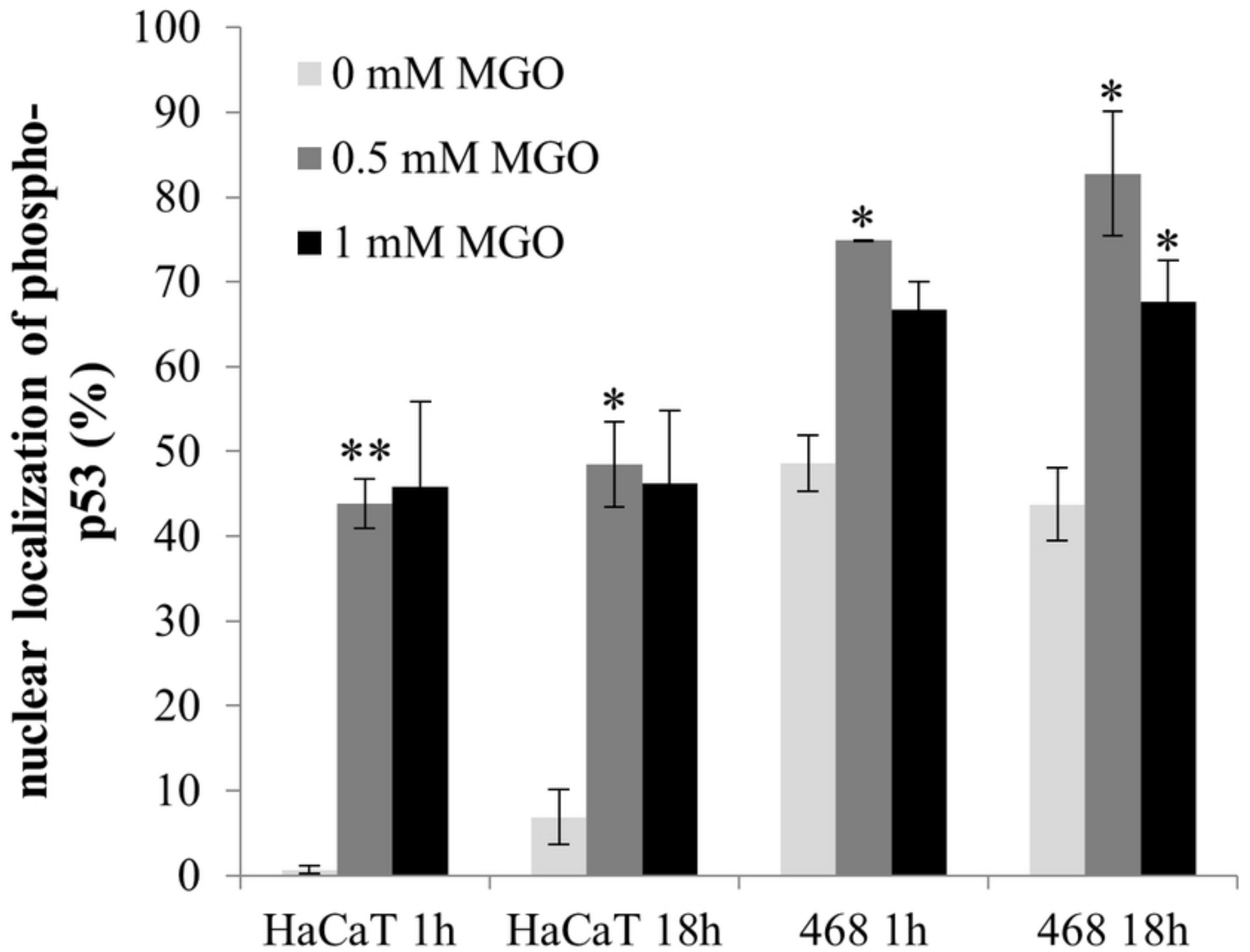


Fig 2

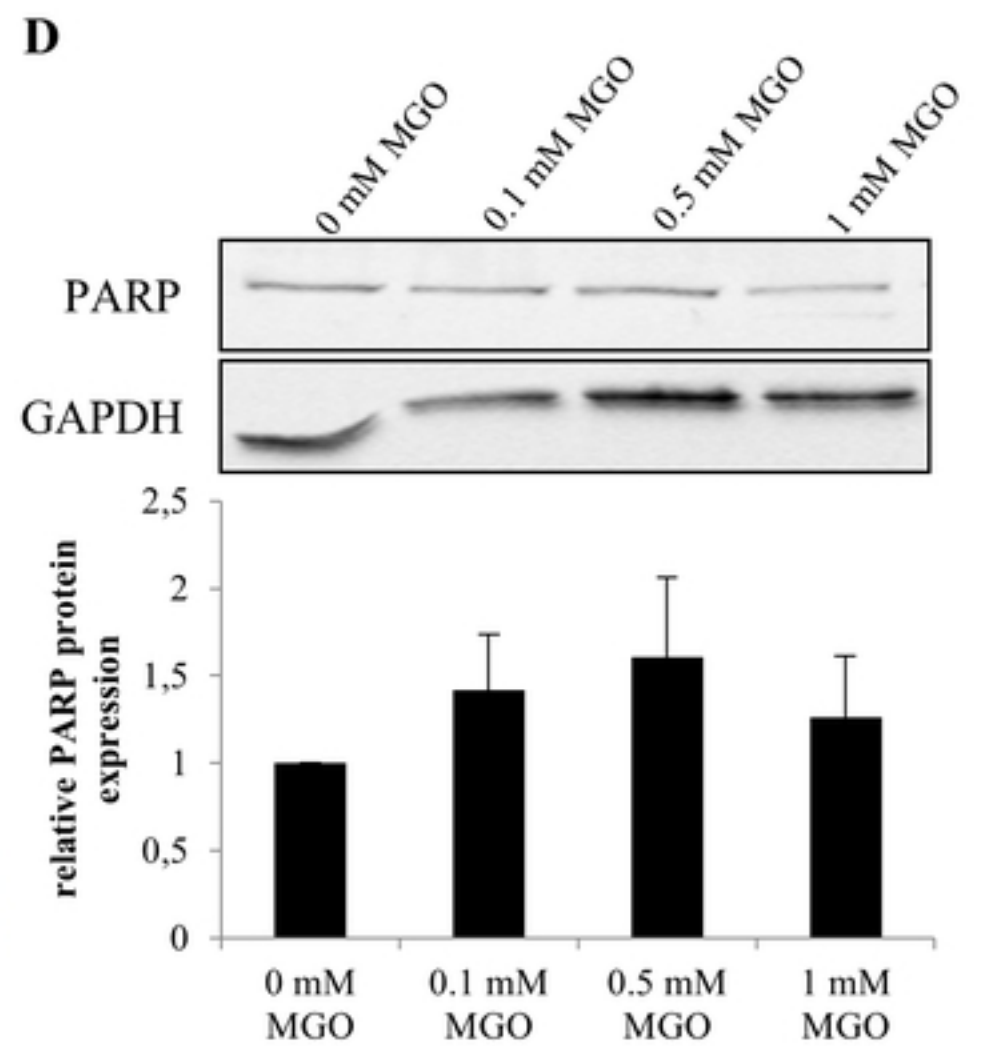
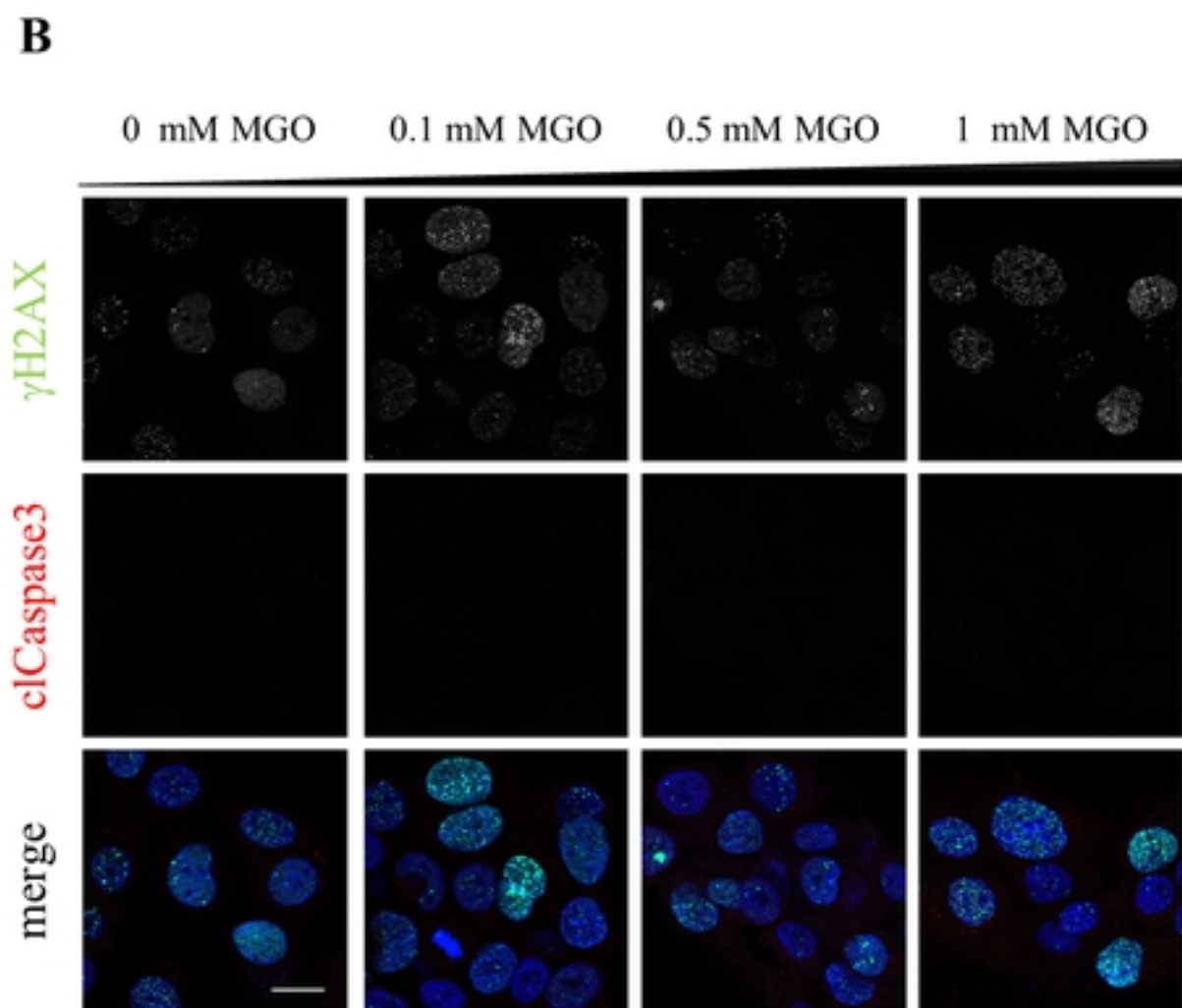
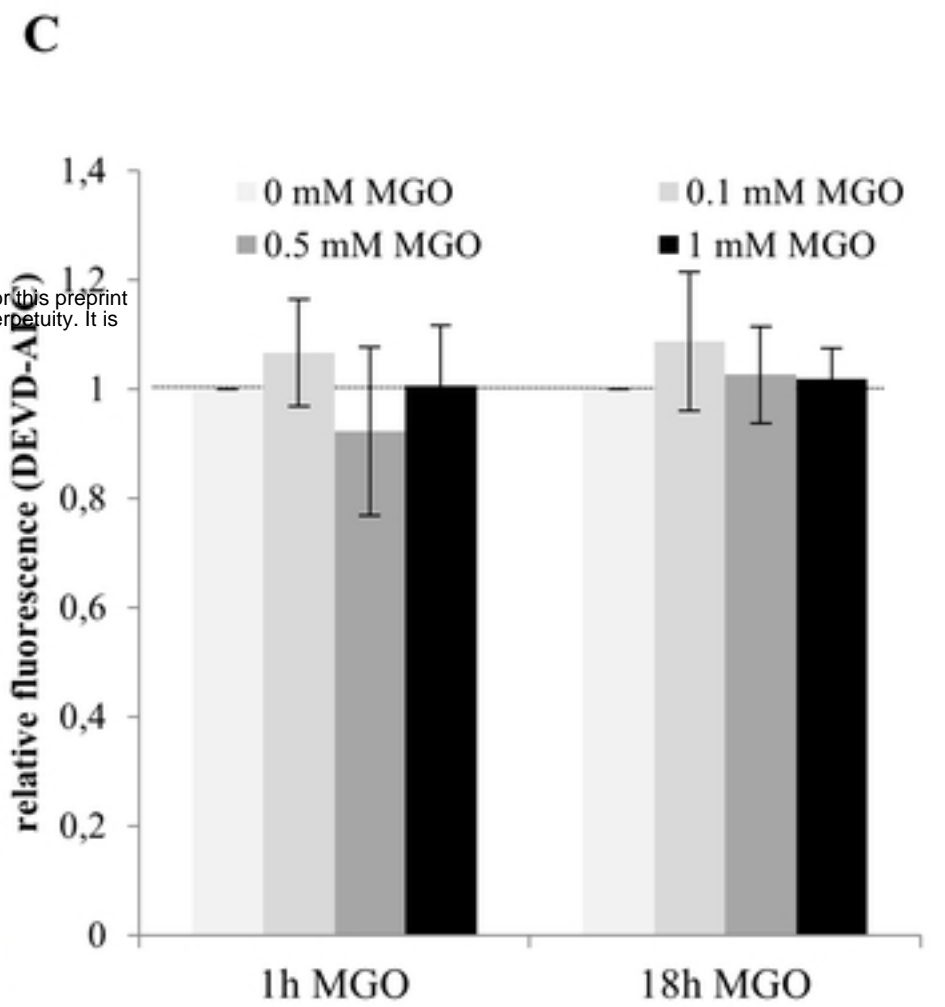
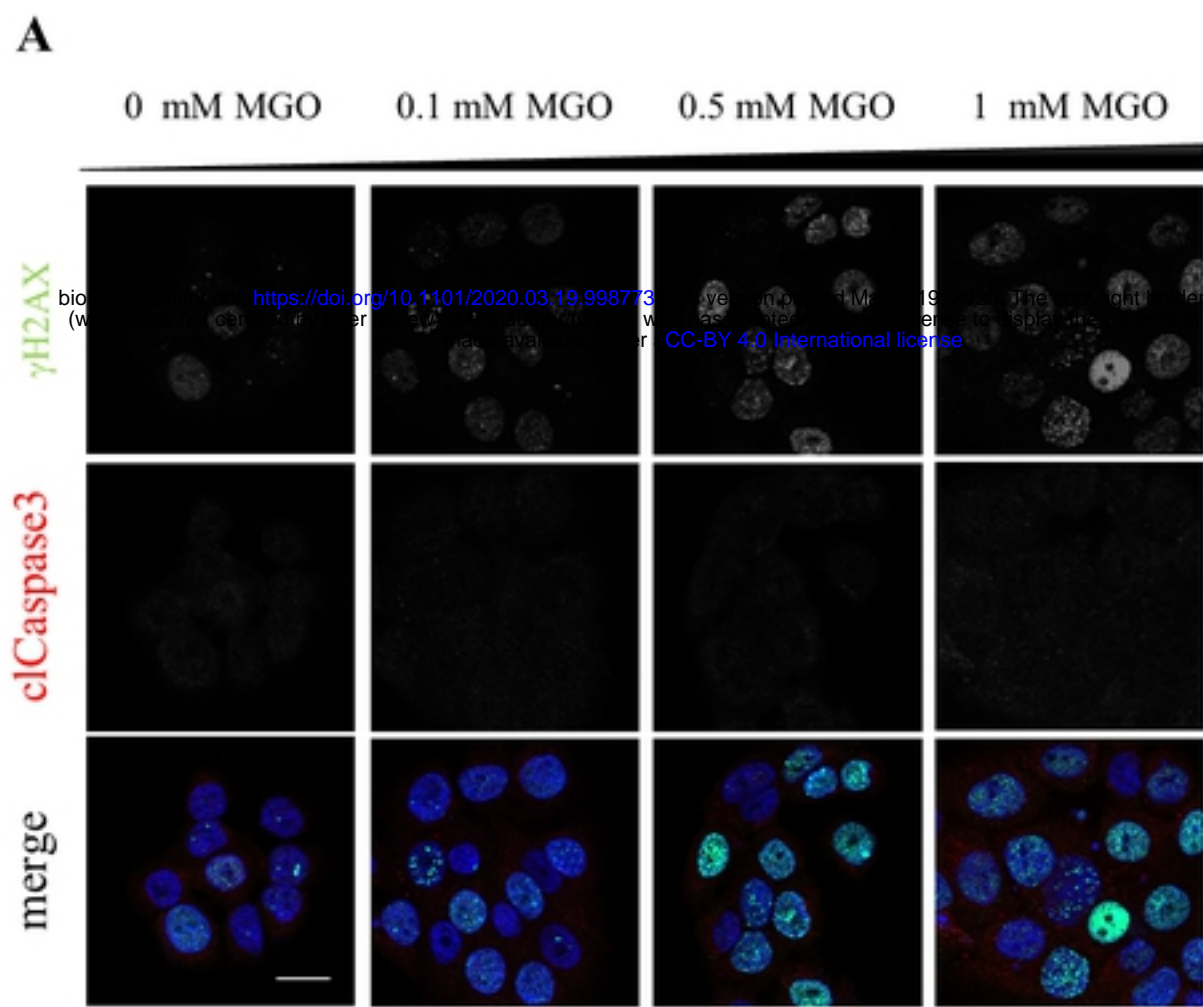


Fig 3

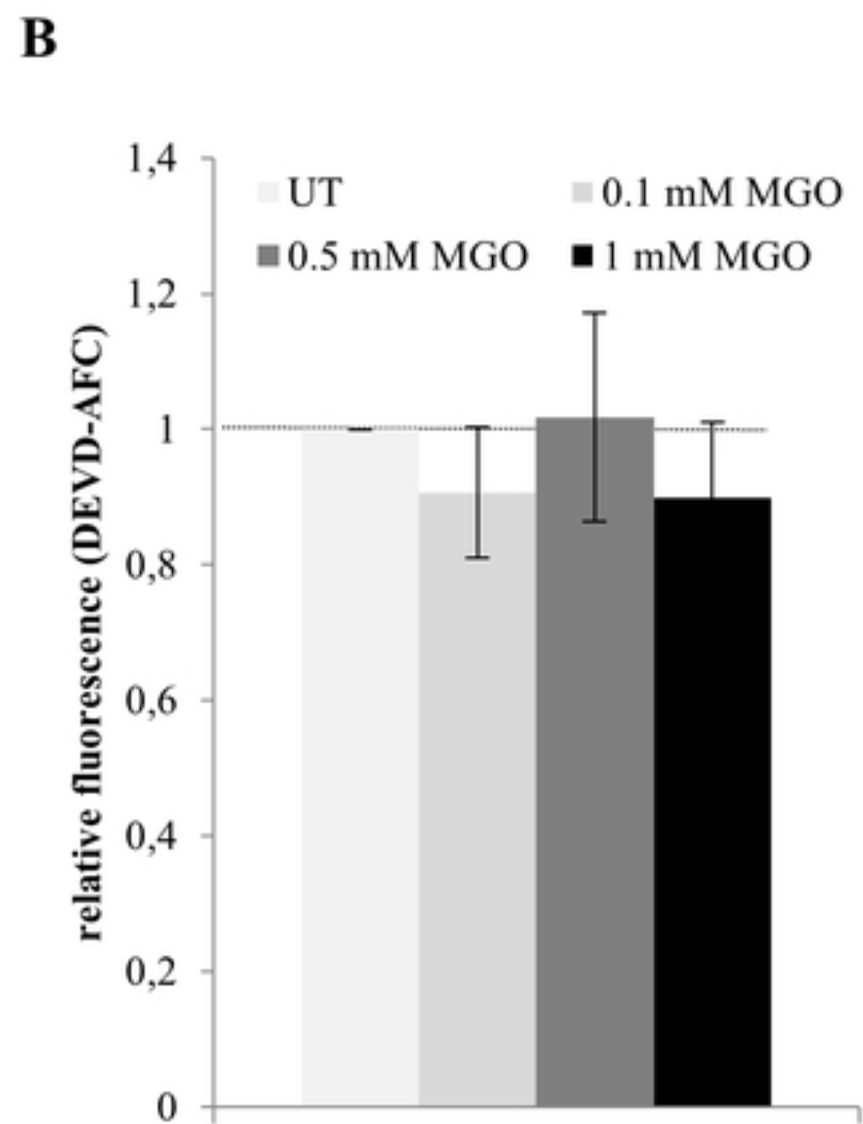
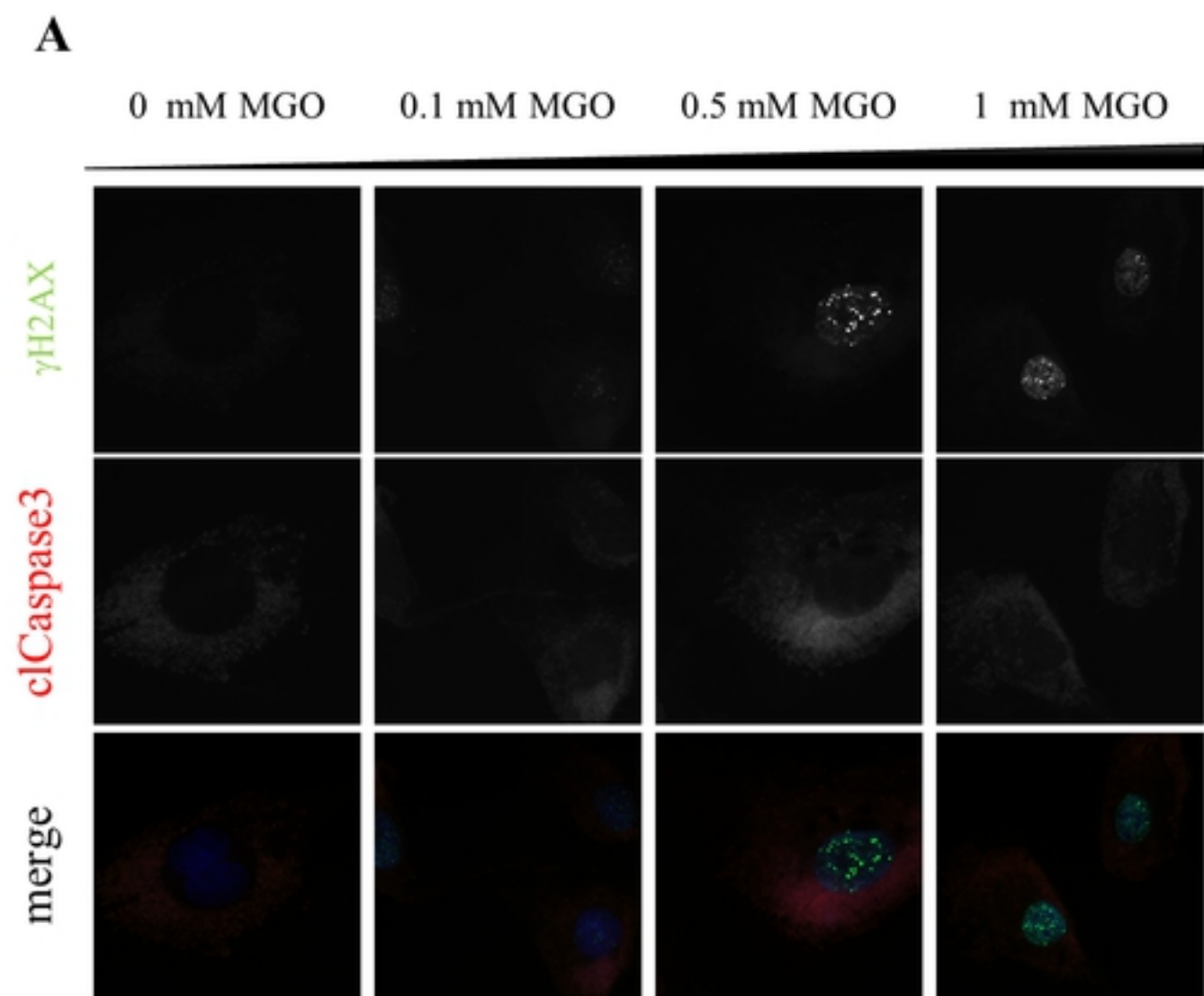


Fig 4

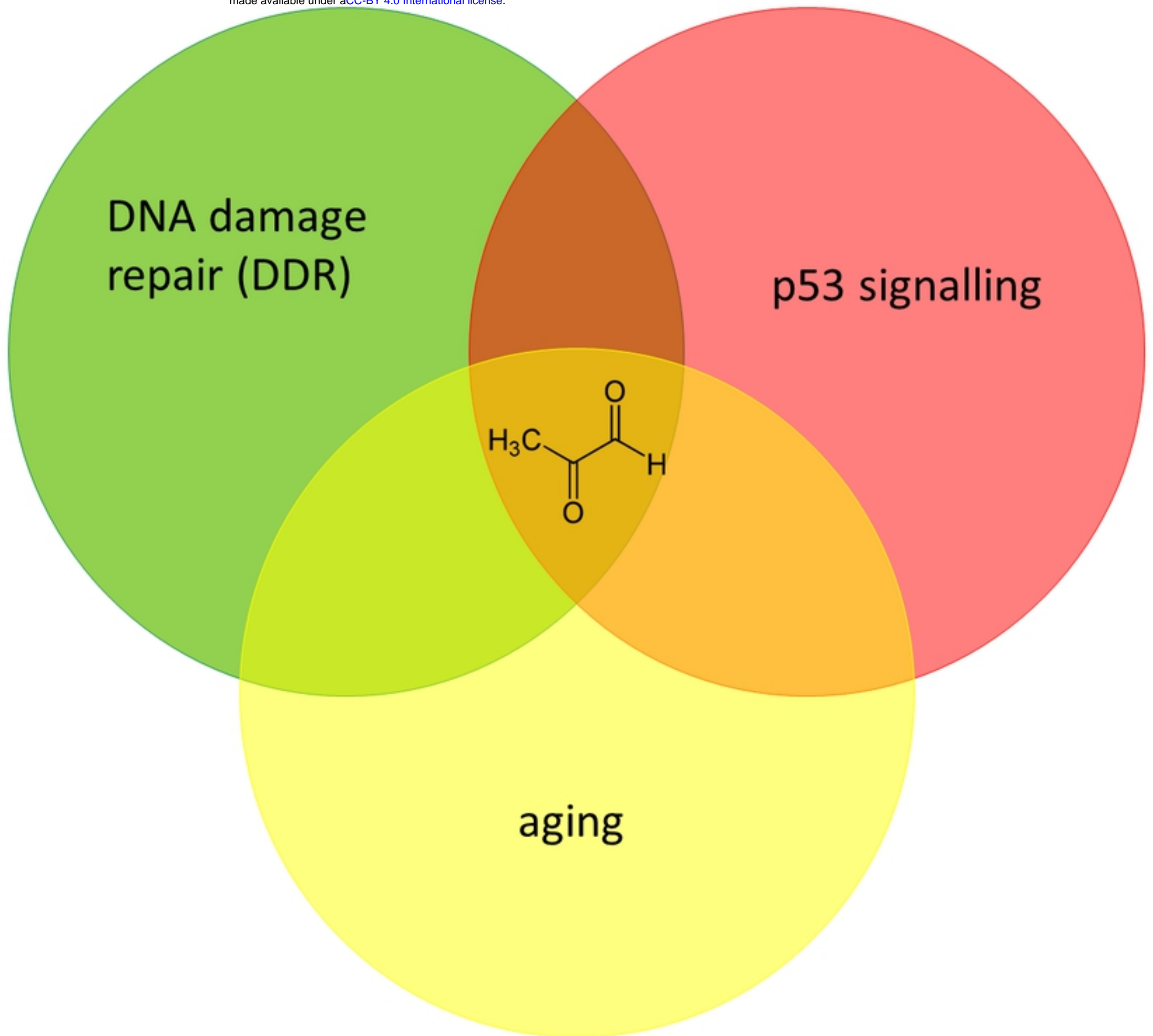


Fig 5