

1 **Sirtuin 3 silencing improves oxaliplatin efficacy through acetylation of MnSOD in**
2 **colon cancer**

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17 **Running head:** SIRT3 silencing improves oxaliplatin efficacy

18

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20

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30

31 **Abstract**

32

33 Sirtuin 3 (SIRT3) is the major mitochondria deacetylase and regulates ROS levels by
34 targeting several key proteins, such as those involved in mitochondrial function and
35 antioxidant defenses. This way, SIRT3 balances ROS production and scavenging and
36 promotes cell survival. The aim of this study was to analyze the effect of SIRT3
37 silencing on the antioxidant response in SW620 colon cancer cell line, and whether this
38 intervention could improve efficacy of oxaliplatin, a common drug used to treat colon
39 cancer. For this purpose, we obtained stable clones of SW620 with SIRT3 knockdown
40 and determined parameters such as ROS levels and ROS production, levels of several
41 antioxidant enzymes, cell viability and apoptosis. Results showed that after SIRT3
42 silencing, both ROS levels and production were increased, and antioxidant enzymes
43 gene expression was significantly reduced. Furthermore, manganese superoxide
44 dismutase levels and enzymatic activity were reduced. Combination of SIRT3
45 knockdown with oxaliplatin treatment further increased ROS production and
46 apoptosis, reducing cell viability. Finally, survival curves on colon cancer patients
47 suggested that SIRT3 expression is related to a poorer prognosis. In conclusion, SIRT3
48 could be a target for colon cancer, since it regulates the antioxidant response and
49 improves the efficacy of oxaliplatin treatment.

50

51

52 **Introduction**

53 Colorectal cancer is the third most common malignancy diagnosed worldwide and the
54 fourth leading cause of death by cancer (Favoriti et al. 2016; Arnold et al. 2017). It has
55 been reported that reactive oxygen species (ROS) and oxidative stress may play an
56 important role during tumorigenesis and cancer progression, including colon cancer
57 (Perše 2013; Galadari et al. 2017).

58 Cancer cells often show enhanced ROS production and accumulation compared to
59 normal cells. These features confer cancer cells advantages in cell growth, survival and
60 resistance to chemotherapy (Sullivan and Chandel 2014; Galadari et al. 2017).
61 However, excessive levels of ROS lead to growth arrest and apoptosis (Hussain et al.
62 2003; Sainz et al. 2012). Indeed, some of the current anticancer therapies act
63 increasing ROS levels, and therefore, it is important to consider the effective regulation
64 of oxidative stress for tumor progression and response to therapy (Nogueira and Hay
65 2013; Lee et al. 2014).

66 Sirtuins are a family of seven proteins with NAD⁺-dependent deacetylase activity and
67 are involved in the regulation of several processes, such as metabolism, cellular
68 proliferation and response to stress (Michan and Sinclair 2007; Torrens-Mas et al.
69 2017). Sirtuin 3 (SIRT3), the major mitochondrial deacetylase, plays a crucial role in
70 modulating ROS production and scavenging (Finley and Haigis 2012; Alhazzazi et al.
71 2013; Torrens-Mas et al. 2017). One of the main targets of SIRT3 is the manganese
72 superoxide dismutase or MnSOD, which detoxifies the radicals generated in the
73 mitochondrial respiratory chain (Tao et al. 2010; Ozden et al. 2011; Bause and Haigis
74 2013).

75 Thus, SIRT3 could function as a protective mechanism in normal cells; however, its
76 activity could also protect cancer cells from the deleterious effects of excessive
77 oxidative stress. Different studies have found a significant correlation between SIRT3
78 levels and progression of several types of cancer, including colon cancer (Liu et al.
79 2014; Torrens-Mas et al. 2017). Furthermore, MnSOD has also been reported to
80 enhance progression, invasion and metastasis in several types of cancer (Hempel et al.
81 2011).

82 The aim of this study was to analyze whether SIRT3 silencing in colon cancer cells could
83 increase oxidative stress through regulation of MnSOD and other antioxidant enzymes,
84 and thus increase cell death. For this purpose, a shRNA against SIRT3 was transfected
85 into SW620 colon cancer cells to generate a stable cell line with SIRT3 knockdown.
86 Parameters such as ROS levels, MnSOD levels, cell viability and apoptosis were
87 analysed.

88

89 **Materials and Methods**

90 *Reagents*

91 Dulbecco's Modified Eagle's medium (DMEM) high glucose was purchased from GIBCO
92 (Paisley, UK). Oxaliplatin ([SP-4-2-(1R-trans)]-(1,2-Cyclohexanediamine-N,N')[ethanedioate(2--
93)-O,O'] platinum or OXA) was obtained from Sigma-Aldrich (St. Louis, MO, USA). Routine
94 chemicals were supplied by Sigma-Aldrich (St. Louis, MO, USA), Roche (Barcelona, Spain),
95 Panreac (Barcelona, Spain) and Bio-Rad Laboratories (Hercules, CA, USA).

96 *Cell culture and stable transfection*

97 Human colon cancer cell line SW620 was purchased from American Type Culture
98 Collection (ATCC; Manassas, VA, USA) and maintained in DMEM supplemented with 10% (v/v)
99 heat-inactivated fetal bovine serum (FBS) and 1% (v/v) penicillin and streptomycin at 37 °C
100 with 5% CO₂.

101 Scrambled shRNA (TR20003) and shRNA targeting SIRT3 (TR309432) were purchased
102 from Origene (Rockville, MD, USA). cDNA clones were amplified in *Escherichia coli* DH5αF'
103 competent cells (Life Technologies, Paisley, UK) and isolated with MaxiPrep isolation kit (Life
104 Technologies, Paisley, UK). SW620 were seeded in 6-well plates and the next day were
105 transfected with either one of the plasmids and with Lipofectamine 2000 reagent (Life
106 Technologies, Thermo Fisher Scientific, Waltham, MA, USA) according to the manufacturer's
107 protocol. Two days after transfection, cells were then selected with 4 µg/mL puromycin for 14
108 days. Multiple monoclonal cell lines were isolated and checked for SIRT3 levels. The selected clones
109 were maintained with a mild selective pressure of 2 µg/mL puromycin for all subsequent
110 experiments.

111 *Measurement of ROS levels by flow cytometry*

112 Cells were seeded in 6-well plates at a density of 6×10^5 cells/well. The day after, cells
113 were trypsinized, harvested into cytometer tubes and centrifuged at 16000 rpm for 5 min.
114 Pellet was resuspended in 500 µL PBS with 10 µM 2',7'-dichlorofluorescein diacetate (DCFDA).
115 After incubation for 15 min in the dark, cells were analyzed immediately using an Epics XL flow
116 cytometer (Beckman–Coulter, Miami, FL, USA). The green fluorescence was measured using
117 the FL-1 setting (log mode) after the cell debris was electronically gated out. Ten thousand
118 events were acquired and analysis was performed with WinMidi software.

119 *RT-PCR*

120 Cells were seeded in 6-well plates at a density of 6×10^5 cells/well. Total RNA was
121 isolated from cultured cells using TRI Reagent® (Sigma-Aldrich, St. Louis, MO, USA) following
122 the manufacturer's protocol and then quantified using a BioSpec-nano spectrophotometer
123 (Shimadzu Biotech, Kyoto, Japan) set at 260 nm. For each sample, 1 µg of the total RNA was
124 reverse transcribed to cDNA at 42°C for 60 min with 25 U MuLV reverse transcriptase in a 10
125 µL volume of retrotranscription reaction mixture containing 10 mM Tris-HCl (pH 9.0), 50 mM
126 KCl, 0.1% Triton X-100, 2.5 mM MgCl₂, 2.5 µM random hexamers, 10 U RNase inhibitor, and

127 500 μ M each dNTP. Each cDNA was diluted 1/10 and aliquots were frozen (-20°C) until the PCR
128 reactions were carried out.

129 PCR was performed using SYBR Green technology on a LightCycler 480 System II rapid
130 thermal cycler (Roche Diagnostics, Basel, Switzerland). The genes, primers and temperatures
131 for the annealing step are specified in Table 1. Total reaction volume was 10 μ L, containing 7.5
132 μ L Lightcycler[®] 480 SYBR Green I Master (containing 0.5 μ M of the sense and antisense specific
133 primers) and 2.5 μ L of the cDNA template. The amplification program consisted of a
134 preincubation step for denaturation of the template cDNA (5 min, 95°C), followed by 45 cycles
135 consisting of a denaturation step (10 s, 95°C), an annealing step (10 s, temperature depending
136 on primers; listed in Table 1), and an elongation step (12 s, 72°C min). A negative control
137 lacking cDNA template was run in each assay.

138 The Ct values of the real-time PCR were analyzed, taking into account the efficiency of
139 the reaction and referring these results to the total DNA amount, using the GenEx Standard
140 Software (Multi-DAnalises, Sweden).

141 *Enzymatic activities*

142 Cells were seeded in 6-well plates at a density of 6×10^5 cells/well. Cells were
143 harvested by scraping them into 200 μ L of STE buffer (250 mM sucrose, 3.59 mM Trizma-Base,
144 16.4 Tris-HCl pH 7.4, 2mM EDTA, 40 mM KCl). Then, cells were disrupted by sonication at 40%
145 amplitude for 10 seconds three times (VibraCell 75185) and centrifuged at 600xg for 10 min at
146 4°C to remove cell debris. Protein content (supernatant) was determined with a bicinchoninic
147 acid (BCA) protein assay kit (Pierce, Bonn, Germany) and the enzymatic assays were performed
148 immediately after.

149 MnSOD (SOD, EC 1.15.1.1) activity was determined by following the reduction of
150 cytochrome c by measuring the absorbance at 550 nm, as described by Quick et al (2000), and
151 adding 1 mM KCN in the reaction mixture to inhibit the activity from cytoplasmatic CuZnSOD.

152 *Western blot analysis*

153 Cells were seeded in 6-well plates at a density of 6×10^5 cells/well. Cells were harvested
154 by scraping them into 200 μ L of RIPA buffer (50 mM Tris-HCl pH 7.5, 150 mM NaCl, 0.1% SDS,
155 0.5% deoxycholate, 1% Triton X-100, 1 mM EDTA, 0.01 mM leupeptin, 0.01 M pepstatin, 2 mM
156 PMSF, 1 mM NaF and 1 mM Na_3VO_4) and disrupted by sonication at 40% amplitude for 10

157 seconds three times. Samples were then centrifuged at 14000 $\times g$ for 10 min at 4 °C and
158 protein content (supernatant) was determined with a bicinchoninic acid (BCA) protein assay kit
159 (Pierce, Bonn, Germany).

160 Twenty μg of protein from cell lysate were separated on 12% SDS–PAGE gels and
161 electrotransferred to 0.22 μm nitrocellulose membranes using the Trans-blot Turbo transfer
162 system (Bio-Rad). Membranes were blocked with 5% non-fat powdered milk in Tris-buffered
163 saline-Tween (TBS with 0.05% Tween-20) for 1h. Antisera against SIRT3 (Cell Signaling
164 Technology Inc, Danvers, MA), acetylated MnSOD (acetyl K63; Abcam, Cambridge, UK) GAPDH
165 and MnSOD (Santa Cruz Biotechnology, CA, USA) were used as primary antibodies. Protein
166 bands were visualized by Immun-Star® Western C® Kit reagent (Bio-Rad) Western blotting
167 detection systems. The chemiluminescence signal was captured with a Chemidoc XRS
168 densitometer (Bio-Rad) and results were analyzed with Quantity One Software (Bio-Rad).

169 *Cell viability assay*

170 Cells were plated in 96-well plates at a density of 7×10^4 cells/well. Cell viability was
171 determined by crystal violet assay. Briefly, cells were stained with 0.5% (p/v) crystal violet in
172 30% (v/v) acetic acid for 10 min. After washing in distilled water, 100 μL of methanol were
173 added to solubilize the dye and absorbance *was measured at* 595 nm using a PowerWave XS
174 Microplate Spectrophotometer (BioTek Instruments, Inc.).

175 *Fluorometric measurement of ROS production*

176 Cells were plated in 96-well plates at a density of 7×10^4 cells/well and treated with 5
177 μM OXA for 48 h. To test the effect of some ROS scavengers, cells were pre-treated for 3h with
178 250 μM ascorbic acid, and 400 μM N-acetyl-cysteine (NAC) was added at the same time of
179 OXA. To measure ROS production, the Amplex® Red Hydrogen Peroxide/Peroxidase Assay Kit
180 (Molecular Probes, Eugene, Oregon, USA) was used. Briefly, 50 μM Amplex red reagent and 0.1
181 U/mL horseradish peroxidase were diluted in Krebs-Ringer phosphate buffer (145 mM NaCl,
182 4.86 mM KCl, 0.54 mM CaCl₂, 1.22 mM MgSO₄, 5.5 mM glucose, 5.7 mM sodium phosphate,
183 pH 7.4) and the reaction mixture was added to cells. Fluorescence measurement was recorded
184 at times 0, 15, 30 and 60 minutes. An FLx800 microplate fluorescence reader (Bio-Tek
185 Winooski, Vermont, USA) was used, set at excitation and emission wavelengths of 571 and
186 585, respectively. Values were normalized per number of viable cells determined by crystal
187 violet assay.

188 *Apoptosis fluorometric assay*

189 Apoptosis was measured fluorometrically using Annexin V staining. Briefly, after
190 cytotoxic treatment, cells were fixed with 2% paraformaldehyde (BD Biosciences, NJ, USA) in
191 PBS for 30 minutes at room temperature. After washing cells twice with PBS, Annexin
192 V/AlexaFluor350 (Life Technologies, Thermo Fisher Scientific, Waltham, MA, USA) in annexin
193 binding buffer (10 mM HEPES/NaOH pH 7.4, 140 mM NaOH and 2.5 mM CaCl₂) was added for
194 10 minutes in the dark at room temperature. Cells were rinsed twice with annexin binding
195 buffer and fluorescence was measured. An FLx800 microplate fluorescence reader (Bio-Tek
196 Winooski, Vermont, USA) was used and excitation and emission wavelengths were set at 346
197 and 442 nm. Values were normalized per number of viable cells determined by crystal violet
198 assay.

199 *Kaplan-Meier survival curves*

200 Overall survival was assessed for 562 colon cancer patients from the data set
201 GSE39582 (Marisa et al. 2013). Survival curves were generated using Kaplan-Meier analysis
202 and assessed with Breslow test to check for statistical significance. Patients were split into high
203 SIRT3 expression levels and low SIRT3 expression levels groups by selecting the 50% top and
204 bottom expression patterns. The analysis was performed including all patients and evaluating
205 overall survival and relapse-free survival, and the Breslow test was used to check for statistical
206 significance.

207 *Statistical analysis*

208 The Statistical Program for the Social Sciences software for Windows (SPSS, version 21.0;
209 SPSS Inc, Chicago, IL) was used for all statistical analyses. Results are presented as mean values
210 ± standard error of the mean (SEM) from six independent experiments. The effects of SIRT3
211 silencing were assessed using the Student's t-test. The effects produced by cytotoxic agent
212 oxaliplatin in combination with SIRT3 silencing were assessed by a two-way analysis of
213 variance (ANOVA), and a post-hoc Student's t-test was performed when combinatory effects
214 were found. Statistical significance was set at P<0.05.

215

216 **Results**

217 *SIRT3 levels after stable knockdown*

218 SIRT3 knockdown by shRNA transfection was determined by measuring both mRNA
219 and protein levels in the selected clone as shown in Figure 1. Indeed, SW620
220 transfected cells showed a 69% decrease in SIRT3 mRNA levels (Figure 1A) and a 47%
221 decrease in protein levels (Figure 1B). In Figure 1C are shown representative bands of
222 the Western Blot.

223 To further check that SIRT3 knockdown was functional, ROS levels were measured by
224 flow cytometry. As expected, ROS levels were increased by 78% after SIRT3 silencing,
225 as shown in Figure 2.

226 *Antioxidant gene expression was decreased in SIRT3-knockdown cells*

227 Expression of several antioxidant genes was evaluated, including transcriptional factors
228 such as NRF2 and Foxo3a, antioxidant enzymes such as superoxide dismutases and
229 catalase, and enzymes involved in the glutathione and peroxiredoxin/thioredoxin
230 systems. Figure 3 shows that all of the genes analyzed showed a significant decrease in
231 their mRNA levels with SIRT3 knockdown.

232 *SIRT3 knockdown compromises MnSOD expression and activity*

233 The effects of SIRT3 knockdown on MnSOD, one of SIRT3 main targets, were further
234 analyzed. First, both levels and acetylation of MnSOD were evaluated by Western Blot.
235 As shown in Figure 4A, total levels of MnSOD were slightly decrease by SIRT3
236 knockdown. However, acetylated-MnSOD was significantly increased by 50% with
237 SIRT3 knockdown, as shown in Figure 4B. Thus, as shown in Figure 4C, the ratio
238 acetylated-MnSOD/total MnSOD was significantly higher in cells with SIRT3 silencing.
239 Representative bands of Western Blots are shown in Figure 4D. As a result, MnSOD
240 activity dropped in these cells to 55% of control values, as seen in Figure 4E.

241 *SIRT3 silencing affects response to oxaliplatin treatment*

242 Effects of SIRT3 knockdown on cell viability, ROS production and apoptosis were
243 evaluated alone and in combination with the cytotoxic treatment oxaliplatin. SIRT3
244 silencing reduced cell viability only by 17%, while oxaliplatin treatment decreased it by
245 45%, as shown in Figure 5A. The combination of SIRT3 knockdown and oxaliplatin
246 treatment resulted in a 51% decrease in cell viability.

247 Figure 5B shows that ROS production is significantly higher after SIRT3 silencing and
248 oxaliplatin treatment, 29% and 101% respectively. Combination of both resulted in a
249 147% increase in ROS production.

250 To check whether programmed cell death was also affected by SIRT3 silencing, an
251 apoptosis assay was performed. As shown in Figure 5C, SIRT3 knockdown increased
252 annexin V fluorescence by 112%, while oxaliplatin treatment significantly increased it
253 by 342%. Both SIRT3 silencing and cytotoxic treatment resulted in a 657% increase in
254 the apoptosis marker.

255 A treatment with different ROS scavengers was performed to prove a causal
256 relationship between increased ROS production and lower cell viability. As seen in
257 Figure 6A, cell viability in SIRT3-silenced cells was restored after OXA and NAC or
258 ascorbic acid treatment, reaching comparable or even higher values to control cells
259 after OXA treatment. Furthermore, ROS production was lowered by the addition of
260 NAC or ascorbic acid in SIRT3-silenced cells after OXA treatment, as shown in Figure
261 6B.

262 *SIRT3 expression is related to a poorer prognosis in colon cancer patients*

263 SIRT3 expression was analyzed as a possible prognostic factor in colon cancer patients
264 using the publicly available data set described under the Kaplan-Meier survival curves
265 section of Materials and Methods. Figure 7A and 7B represent the overall survival (OS)
266 and relapse free survival (RFS) in colon cancer patients expressing high or low levels of
267 SIRT3, respectively. Although RFS did not reach a significant difference between both
268 groups ($P < 0.07$), OS was significantly lower in those patients with higher expression of
269 SIRT3 ($P < 0.009$).

270

271 **Discussion**

272 In this study, we have shown that SIRT3 silencing resulted in a downregulation of
273 several antioxidant genes, as well as a reduction in MnSOD protein levels and activity.
274 Furthermore, SIRT3 knockdown lead to higher ROS levels and ROS production, which

275 was accompanied by lower cell viability and an increase in apoptosis. Combination of
276 SIRT3 silencing and cytotoxic treatment with oxaliplatin further increased ROS
277 production and apoptosis. Finally, we show here, using bioinformatic tools, that high
278 SIRT3 expression could be related to a poorer prognosis for colon cancer patients.

279 Growing evidence suggests that free radicals and oxidative stress play an important
280 role in the development and progression of colon cancer (Perše 2013). In fact, it has
281 been reported that human colorectal tumors present high levels of ROS and markers of
282 oxidative stress such as catalase, glutathione peroxidase or MnSOD (Skrzycki et al.
283 2009; Perše 2013). However, the role of ROS remains controversial. A light increase in
284 ROS production and mild oxidative stress may be beneficial for cell survival, while
285 rising ROS levels above a certain threshold may trigger cell death (Trachootham et al.
286 2008; Panieri and Santoro 2016; Galadari et al. 2017).

287 SIRT3 is considered a crucial protein against oxidative stress, since it deacetylates and
288 activates several proteins related to mitochondrial function and antioxidant enzymes
289 (Alhazzazi et al. 2011a; Torrens-Mas et al. 2017). In this regard, we observed an
290 increase in ROS levels and ROS production after SIRT3 knockdown, as has been widely
291 described before (Kong et al. 2010; Finley et al. 2011; Finley and Haigis 2012; Kwon et
292 al. 2015; Torrens-Mas et al. 2016). This increase has been attributed to the
293 hyperacetylation, and thus inactivation, of SIRT3 targets, mainly MnSOD, which is one
294 of the most important antioxidant enzymes of the cell (Ozden et al. 2011; Tao et al.
295 2014; Zou et al. 2016).

296 However, we have shown here that gene expression of several antioxidant enzymes
297 was also affected by SIRT3 silencing. The downregulation of these antioxidant genes is
298 presumably due to a reduction of transcriptional activity of FOXO3a, which is a target
299 of SIRT3 and regulates the expression of crucial proteins involved in ROS scavenging
300 and mitochondrial integrity, such as MnSOD and catalase (Sundaresan et al. 2009;
301 Bause and Haigis 2013; Tseng et al. 2013; Rangarajan et al. 2015). Thus, SIRT3 silencing
302 prevents FOXO3a deacetylation and leads to a reduction in the activity of this
303 transcription factor.

304 To further study the effect of SIRT3 knockdown on antioxidant enzymes, we measured
305 protein levels of MnSOD and acetylated MnSOD. As expected, the acetylation level of
306 MnSOD was higher after SIRT3 silencing. Moreover, total protein levels of MnSOD
307 were significantly reduced and the ratio acetylated MnSOD/total MnSOD almost
308 doubled in cells with SIRT3 silencing. A previous study by our group also showed a
309 reduction in MnSOD protein levels and other targets after treatment of MCF-7 breast
310 cancer cell line with siRNA against SIRT3 (Torrens-Mas et al. 2016). The increase in
311 acetylation in MnSOD resulted in a reduction of its enzymatic activity, as has also been
312 described before (Tao et al. 2010; Ozden et al. 2011).

313 On the other hand, SIRT3 knockdown also reduced cell viability, which occurred with
314 an increase in apoptosis. This may be due to the increase in ROS production that can
315 lead to oxidative damage and cell death (Alhazzazi et al. 2011b; Papa and Germain
316 2014). These results are agreement with several studies that refer to SIRT3 as an
317 oncogene, since SIRT3 limits ROS production and increases cellular resistance to
318 oxidative stress, thus promoting cell proliferation and avoiding apoptosis (L. Zhang et
319 al. 2013; Wang et al. 2015). However, some studies describe SIRT3 as a tumor
320 suppressor in some types of cancer, such as breast cancer (Buler et al. 2012), ovarian
321 cancer (Dong et al. 2016), hepatocellular carcinoma (B. Zhang et al. 2013; Liu et al.
322 2017) or B cell malignancies (Yu et al. 2016). In these reports, SIRT3 reduces cell
323 proliferation of cancer cells and limits their metabolic reprogramming. Furthermore,
324 patients with higher SIRT3 expression present a good outcome and an increased
325 overall survival. These studies suggest a dual role for SIRT3 in cancer, which may
326 depend on cellular type and cellular context (Torrens-Mas et al. 2017).

327 Finally, we observed that oxaliplatin increased apoptosis and ROS production, and
328 when combined with SIRT3 knockdown, these parameters were further increased. As
329 expected, the addition of ROS scavengers, NAC and ascorbic acid, reduced ROS
330 production after OXA treatment. Accordingly, combination of OXA treatment and ROS
331 scavengers produced a recovery in cell viability, reducing the cytotoxic effect of the
332 increased ROS produced by OXA treatment.

333 Previous studies have shown that SIRT3 knockdown sensitizes cells to cytotoxic
334 treatments and reduces cell proliferation (Alhazzazi et al. 2011b; George et al. 2016;
335 Torrens-Mas et al. 2016). In this regard, our results show that SIRT3 silencing enhances
336 the effectivity of oxaliplatin through increasing ROS production to excessive levels
337 incompatible with cell viability. Moreover, when SIRT3 is silenced we observed an
338 inability to induce the antioxidant defenses and counteract the harmful effects of ROS,
339 which leads to an increase in the apoptosis process.

340 Taken together, these results support the role of SIRT3 as a crucial protein in limiting
341 ROS production and promoting cell survival by preservation of proper mitochondrial
342 function and integrity (Kim et al. 2010; Park et al. 2011). Furthermore, the survival
343 analysis presented here suggests that when SIRT3 expression is high, tumor cells can
344 balance ROS production activating the antioxidant response and could be more
345 resistant to oxidative stress and even cytotoxic therapy (Papa and Germain 2014;
346 Galadari et al. 2017). This way, patients with high expression of SIRT3 show a poorer
347 survival.

348 In conclusion, SIRT3 silencing in colon cancer cells not only affects the activity of
349 antioxidant enzymes, but also its expression at the mRNA level, which may reduce the
350 antioxidant capacity of the cell and thus increasing cell death. In this regard, SIRT3
351 knockdown could be an adjuvant treatment for colon cancer, enhancing the effect of
352 cytotoxic treatments such as oxaliplatin, which rely on increasing ROS production to
353 induce cell death.

354

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495 **Figure legends**

496 **Figure 1.** SIRT3 levels after shRNA SIRT3 transfection and clonal selection. (A) SIRT3 mRNA levels after
497 SIRT3 silencing. (B) SIRT3 protein levels measured by Western Blot after SIRT3 silencing. (C)
498 Representative bands of SIRT3 Western Blot and GAPDH as a loading control. Values are expressed as
499 mean \pm SEM (n=6) and normalized to control value. Cells transfected with control shRNA are shown in
500 white, and cells with SIRT3 shRNA are shown in black. * indicates a significant difference between
501 control and SIRT3-silencing cells (P<0.05).

502 **Figure 2.** SIRT3 silencing increased ROS levels measured by flow cytometry. Intracellular levels of ROS
503 determined by DCF fluorescence. Values are expressed as mean \pm SEM (n=6) and normalized as
504 percentage of the control value. * indicates a significant difference between control and SIRT3-silencing
505 cells (P<0.05).

506 **Figure 3.** SIRT3 silencing reduced antioxidant gene expression. NRF2: nuclear respiratory factor 2;
507 Foxo3a: Forkhead box O3; MnSOD: manganese superoxide dismutase; CuZnSOD: copper-zinc superoxide
508 dismutase; CAT: catalase; Gpx: glutathione peroxidase; GRd: glutathione reductase; Prx2-6:
509 peroxiredoxine 2-6; Trx1-2: thioredoxin 1-2; TrxR2: thioredoxin reductase 2. Values are expressed as
510 mean \pm SEM (n=6) and values of control cells were set at 1. * indicates a significant difference between
511 control and SIRT3-silencing cells (P<0.05).

512 **Figure 4.** SIRT3 silencing increased acetylated-MnSOD/total-MnSOD ratio and reduced MnSOD
513 enzymatic activity. (A) Total protein levels of MnSOD as measured by Western Blot. (B) Acetylated-
514 MnSOD levels measured by Western Blot. (C) Ratio of acetylated-MnSOD/total-MnSOD. (D)
515 Representative bands of Western Blots against total MnSOD and acetylated MnSOD. (E) MnSOD
516 enzymatic activity. Values are expressed as mean \pm SEM (n=6) and normalized as percentage of the
517 control value. * indicates a significant difference between control and SIRT3-silencing cells (P<0.05).

518 **Figure 5.** SIRT3 silencing increased oxaliplatin efficacy by increasing ROS production and apoptosis. (A)
519 Cell viability after SIRT3 silencing and oxaliplatin treatment measured by crystal violet. (B) ROS
520 production measured fluorometrically after SIRT3 silencing and oxaliplatin treatment. (C) Annexin V
521 fluorescence, an apoptosis marker, measured after SIRT3 silencing and oxaliplatin treatment. Values are
522 expressed as mean \pm SEM (n=6) and normalized as percentage of the control value. S indicates a SIRT3-
523 silencing effect, T indicates treatment effect, and SxT indicates a combinatory effect of both. A Student's
524 t test was performed when combinatory effects were observed, where: * indicates a significant
525 difference between control and SIRT3-silencing cells and # indicates statistical difference between
526 vehicle- and oxaliplatin-treated cells (P<0.05).

527 **Figure 6.** ROS scavengers reduce ROS production and recover cell viability after SIRT3 silencing and OXA
528 treatment. (A) Cell viability after SIRT3 silencing and oxaliplatin, NAC and ascorbic acid treatment
529 measured by crystal violet assay. (B) ROS production was measured fluorometrically after SIRT3

530 silencing and oxaliplatin, NAC and ascorbic acid treatment. Values are expressed as mean \pm SEM (n=6)
531 and normalized as percentage of the control value. * indicates a significant difference between control
532 and SIRT3-silenced cells (P<0.05).

533 **Figure 7.** Kaplan-Meier survival curves showed that high SIRT3 expression is correlated to a lower overall
534 survival for colon cancer patients. (A) Overall survival of colon cancer patients. (B) Relapse-free survival
535 of colon cancer patients.