



Contents lists available at ScienceDirect

Experimental and Molecular Pathology

journal homepage: www.elsevier.com/locate/yexmp

Heme oxygenase-1 has antitumoral effects in colorectal cancer: Involvement of p53

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ARTICLE INFO

Article history:

Received 8 September 2014

Accepted 12 September 2014

Available online xxxxx

Keywords:

Heme oxygenase-1

Colorectal cancer

p53

Survival

Immunohistochemistry

ABSTRACT

The expression of heme oxygenase-1 (HO-1) has been shown to be up-regulated in colorectal cancer (CRC), but the role it plays in this cancer type has not yet been addressed. The aims of this study have been to analyze HO-1 expression in human invasive CRC, evaluate its correlation with clinical and histo-pathological parameters and to investigate the mechanisms through which the enzyme influences tumor progression. We confirmed that HO-1 was over-expressed in human invasive CRC and found that the expression of the enzyme was associated with a longer overall survival time. In addition, we observed in a chemically-induced CRC animal model that total and nuclear HO-1 expression increases with tumor progression. Our investigation of the mechanisms involved in HO-1 action in CRC demonstrates that the protein reduces cell viability through induction of cell cycle arrest and apoptosis and, importantly, that a functional p53 tumor suppressor protein is required for these effects. This reduction in cell viability is accompanied by modulation of the levels of p21, p27, and cyclin D1 and by modulation of Akt and PKC pathways. Altogether, our results demonstrate an antitumoral role of HO-1 and points to the importance of p53 status in this antitumor activity.

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1. Introduction

Colorectal cancer (CRC) is the third most commonly diagnosed cancer in males and the second in females (Ferlay et al., 2010). Histo-pathological staging from microscopic examination of tumor tissue is the standard for diagnosis and prognosis, as well as being the main guide for the choice of treatment (Compton and Greene, 2004). However, tumors with the same histo-pathological classification may display significant differences in progression and response to treatment (Liefers and Tollenaar, 2002). This reflects the need to identify new molecular markers that help in the diagnosis, prognosis and treatment of the disease.

Heme oxygenase (HO) is a microsomal enzyme catalyzing the first rate-limiting step in heme degradation, leading to the formation of equimolar quantities of carbon monoxide, biliverdin and free iron. HO-1, the inducible 32-kDa isoform, is an ubiquitous heat shock protein (HSP32) (Maines and Gibbs, 2005) that can be induced in response to cellular stress, oxidative stimuli and hypoxia, an important process

frequently occurring during tumor growth. An increasing body of evidence indicates that HO-1 may play an important role in cancer. Indeed, HO-1 was reported to be up-regulated in rat, mouse and human tumors (Jozkowicz et al., 2007; Was et al., 2006), although the significance of this up-regulation is not clear. In this regard, we have recently demonstrated that HO-1 protein is overexpressed and correlates with clinical parameters in head and neck squamous cell carcinoma (Gandini et al., 2012), glioma (Gandini et al., 2014) and in non-small cell lung cancer (Degese et al., 2012) and that the nuclear localization of the protein associates with tumor progression (Gandini et al., 2012). Several groups have studied the expression of the enzyme in intestinal diseases such as colitis (Berberat et al., 2005; Takagi et al., 2008), inflammation (Barton et al., 2003) and inflammatory bowel disease (Paul et al., 2005). In both the normal intestinal physiology and intestinal diseases, HO-1 has been shown to be increased in response to oxidative stress (Degese et al., 2012) and to play an important role in mucosal protection by scavenging free radicals and reducing inflammation (Berberat et al., 2005). To our knowledge, there are only three reports in the literature showing the expression of HO-1 in human CRC samples (Becker et al., 2007; Kang et al., 2012; Yin et al., 2014).

All this background suggested the need to address the role of HO-1 in CRC and for this purpose we evaluated enzyme expression in human CRC tissues and analyzed its correlation with clinic-pathological features. Additionally, and since the function of HO-1 in CRC biology is

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far from being completely understood, we have also begun to study the mechanisms through which the aforementioned protein influences colorectal tumor progression.

2. Materials and methods

2.1. Patients and tissue specimens

83 human CRC samples were retrieved from the Hospital Regional Italiano (Bahía Blanca, Argentina) with institutional approval. These samples corresponded to primary tumors obtained by surgical resection of invasive CRC patients. The staging of CRC was classified using the seventh edition of the International Union Against Cancer Tumor-Node-Metastasis (TNM) staging system. Data were obtained on diagnosis, treatment response, course and follow-up, gender, primary tumor site, date of surgery, date of death or last contact, date of relapse, site of relapse, obstruction and/or intestinal perforation at diagnosis and pre-surgical carcinoembryonic antigen (CEA). From the primary tumor biopsy we obtained tumor size, invasion to the intestinal wall, regional lymph node metastasis, vascular or perineural invasion, grade of differentiation, K-ras status and expression of EGFR-1 receptor. Additionally, a cohort of 15 samples was retrieved from private pathology laboratories. H&E staining was performed on each sample and the slides further re-evaluated by a pathologist. A series of 5- μ m sections were cut and transferred onto histological glass slides.

2.2. Immunohistochemistry (IHC)

The tissues obtained were fixed for 24 h in 10% formalin and embedded in paraffin using standard procedures. Then the slides were treated as previously described (Facchinetti et al., 2010; Gandini et al., 2012). Sections were then incubated overnight at 4 °C with primary rabbit anti-HO-1 antibody (SPA-896, Streetsgen; 1:100) followed by incubation for 30 min with diluted biotinylated secondary antibody and then incubation with Vectastain ABC Reagent (Vector Laboratories Inc.). For negative controls, the primary antibody was omitted. Diaminobenzidine/ H_2O_2 was used as substrate for the immunoperoxidase reaction and the tissues were lightly counterstained with hematoxylin (Harris), dehydrated through grade ethanol and xylene and mounted with Permount (Fischer Scientific) for analysis by bright-field microscopy.

2.3. Evaluation of staining intensity and statistical analysis

Human immunostained sections were scored semiquantitatively based upon the proportion of tumor cells stained and the staining intensity, by using the semi-quantitative immunoreactive score (IRS), as previously described (Gandini et al., 2012, 2014). To estimate the discriminative value of the IRS for HO-1 expression in CRC, receiver operating characteristic (ROC) curves were plotted and the corresponding areas under the curves (AUCs) were compared using various possible cut-off values, as already described (Gandini et al., 2014). In the animal model, the percentage of HO-1 expression was analyzed by counting cells in 10 random fields (400 \times). To study HO-1 nuclear staining in human samples and in the animal model, the total percentage of nuclei positive for HO-1 was assessed. Samples that had more than 10% of stained cells were considered positive. The software Graph Pad Prism 5 was used for the collection, processing and statistical analysis of all data. The statistical significance of HO-1 expression between groups was determined by the two-tailed χ^2 test or Mann-Whitney *U* test. *p* values of less than 0.05 indicated a significant result.

2.4. Animal model

Wistar male rats (*N* = 31) aged 8 weeks old were used. 28 of them were injected intramuscularly with 1,2-dimethylhydrazine (DMH) 20 mg/kg once a week for 8 weeks. At progressive stages of

the development of tumors, animals were anesthetized and sacrificed to collect the necessary samples (polyps, adenocarcinomas and signet-ring cell carcinomas). Normal tissue was collected from animals that belonged to the same colony but were not treated with DMH (*n* = 3), in order to perform comparative studies with normal mucosa. Samples were taken for fixing in 10% formalin for immunohistochemistry. The statistical significance of HO-1 expression between groups was determined by the two-tailed χ^2 test or Mann-Whitney *U* test. *p* values of less than 0.05 indicated a significant result.

2.5. Cell lines

Human CRC cell lines HCT116, HCT116 p53 $-/-$, HT29 and LoVo were maintained at 37 °C in a humidified incubator with 5% $CO_2/95\%$ air atmosphere in DMEM and F-12K Medium (Sigma) supplemented with 10% (v/v) FBS (Gibco), L-glutamine (5 mM, Gibco), penicillin (Gibco, 100 U/ml) and streptomycin (Gibco, 100 μ g/ml).

2.6. Flow cytometry

For cell cycle analysis staining with propidium iodide (PI, Sigma) was used. 1×10^6 cells HCT116 and HCT116 p53 $-/-$ were seeded. The cells were synchronized by deprivation of fetal bovine serum for 24 h. Then they were treated with hemin and vehicle (100 μ M) for 24 h. The cells were trypsinized, fixed with ice-cold 70% ethanol, stained with PI, and analyzed for DNA content by FACScan flow cytometry (Becton Dickinson, Germany). Data were analyzed by Cell Quest software (Becton Dickinson). The percentage of apoptotic cells was measured by flow cytometry following Annexin V (FL1-H) and PI (FL2-H) labeling. All these experiments were carried out in triplicate and were repeated twice.

2.7. Transient transfections

We used two expression plasmids for HO-1 over-expression. One encoding the native form of the protein fused to enhanced green fluorescent protein (pEGFP-HO-1) was kindly donated by Dr. Phyllis A. Dennery from the Children's Hospital of Philadelphia, University of Philadelphia, USA. The other, pcDNA3-HO-1 was gently donated by Dr. Elba Vazquez (Universidad de Buenos Aires). Also, a pcDNA3-p53, kindly donated by Adriana De Siervi (Universidad de Buenos Aires) was employed to introduce p53 in the HCT116 p53 $-/-$ cell line. The transfection procedure was performed by using Lipoaectamine (Invitrogen, CA, USA) according to the manufacturer's instructions.

2.8. Cellular viability assays

HCT116, HCT116 p53 $-/-$ and HT29 CRC cell lines were seeded in 96 well plates. 48 h later, they were treated with an inductor or an inhibitor of the activity of HO-1 (hemin and tin dichloride (IV) protoporphyrin (IX) (SnPP), respectively) at different doses and different time points after which the cells were incubated with WST-1 cell proliferation reagent (Roche) and further counted manually using a hemocytometer (Becton Dickinson, Germany).

Alternatively, both HCT116 and HCT116 p53 $-/-$ cell lines were seeded in 96 well plates and 72 h later the HCT116 p53 $-/-$ cells were transfected as previously described. 24 h post-transfection the cells were treated with hemin (100 μ M) and vehicle and 96 h later the cells were incubated with WST-1 and manually counted.

2.9. Cell migration

Cell migration was measured by a "wound healing" assay as previously described (Petit et al., 2000). Cells were seeded and further treated for 24 h with vehicle or hemin (100 μ M).

189 2.10. Cell lysis and western blot analysis

190 The entire procedure was performed as already described (Facchinetti
191 et al., 2010). HCT116, HCT116 p53 $-/-$ and HT29 cells were grown for
192 48 h. Following cell line treatment with hemin (20, 40 and 100 μ M),
193 SnPP (2.5 and 10 mM) and vehicle (DMSO), or cell line transfection as de-
194 tailed previously, the cells were scraped and proteins were quantified
195 using the Bradford method (Bradford, 1976). The lysates were electro-
196 phoresed and the blots were incubated with primary rabbit polyclonal
197 anti-HO-1 antibody (SPA-896, Stressgen Bioreagents, Canada), mouse
198 polyclonal anti-p27/Kip1 (BD Transduction Laboratories™), mouse poly-
199 clonal anti-p21 (BD Pharmingen™), rabbit monoclonal anti-cyclin D1
200 (SP4, Thermo Scientific, Inc.), rabbit polyclonals anti-p53 (sc-6243,
201 Santa Cruz Biotechnology), anti-bax (sc-493, Santa Cruz Biotechnology,
202 Inc.), anti-PKC β I (sc-209, Santa Cruz Biotechnology, Inc.), anti-PKC β II
203 (sc-210, Santa Cruz Biotechnology) and goat polyclonal anti-actin
204 (sc-1615, Santa Cruz Biotechnology). The blots were finally washed
205 with PBS-T buffer, incubated with secondary horseradish peroxidase
206 conjugated antibody (Santa Cruz Biotechnology) and the reaction
207 was detected by chemiluminescence amplified (ECL, GE Healthcare
208 UK Limited).

209 2.11. Immunofluorescence

210 HCT116 and HCT116 p53 $-/-$ cells were plated on sterile glass
211 coverslips. The latter were transfected with pcDNA3-p53 and pcDNA3

plasmids. 48 h later they were treated with hemin and vehicle 212
(100 μ M) and then were fixed with 4% paraformaldehyde for 10 min 213
and permeabilized with 0.1% Triton X-100 (Sigma), followed by incuba- 214
tion with a blocking solution (1% BSA in PBS). Anti-HO-1 (SPA-896 215
Stressgen, dilution: 1:100) and anti-p53 (sc-6243, Santa Cruz Biotech- 216
nology, dilution: 1:100) were used. After incubation with primary anti- 217
body, cells were incubated with Alexa 566 fluoro-conjugated antibodies 218
(Molecular Probes, Invitrogen). Nuclei were stained with DAPI and then 219
mounted on slides with Prolong Gold anti-fade reagent (Invitrogen). 220
Images were captured with a Nikon Eclipse E600 fluorescence micro- 221
scope equipped with a Nikon camera. Counting of 200 cells in 10 ran- 222
dom fields (400 \times) was done in order to study the proportion of cells 223
containing HO-1 and p53 expression. 224

225 3. Results

226 3.1. Validation of the antibody for immunohistochemical analyses and HO-1
227 expression in human CRC biopsies

228 Immunospecificity of the antibody for HO-1 in human CRC was first
229 tested by western blot of protein lysates of human CRC cell lines HT29
230 and HCT116 (Fig. 1A). The antibody exclusively recognized one band
231 representing a protein with a mobility corresponding to the molecular
232 weight of HO-1 (32 kDa). This antibody was then used for further studies
233 of HO-1 expression in human CRC samples. We subsequently performed
234 immunohistochemical staining in 98 surgically-resected biopsies of

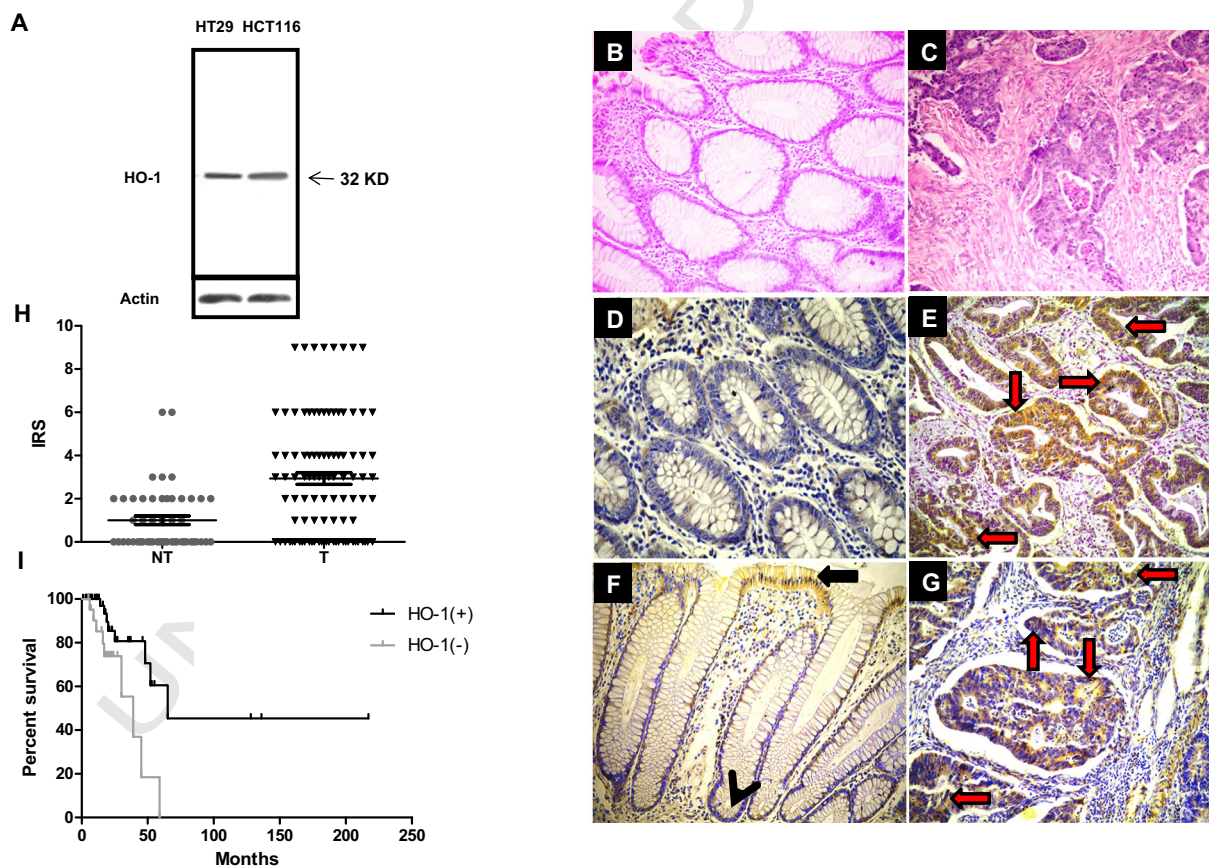


Fig. 1. Specific detection of HO-1 in human colorectal biopsies and association of HO-1 expression with longer overall survival times in patients with invasive CRC. Western blot analysis of HT29 and HCT116 cell lines (A) probed with anti HO-1 (SPA-896). Hematoxylin and eosin staining of histologically normal tissue adjacent to the tumor (B) and of tumor tissue (C) (40 \times). HO-1 expression in non-malignant adjacent tissues (D and F) (100 and 40 \times) and in their respective tumor tissues (E and G) (100 \times). Black arrow: HO-1 immunoreactivity observed in apical cells of the crypt within the non-malignant epithelia. Arrowhead: basal cells lacking HO-1 staining. Red arrows: High HO-1 expression in tumor tissues. H. Dot plot showing the levels (IRS) of HO-1 in tumor tissues (T) and in adjacent non-malignant tissues (NT). Immunohistochemical analysis revealed differences in the levels of HO-1 expression between adjacent non-tumor tissues (NT) and tumor tissues (T) ($p = 0.0001$, Mann Whitney test). I. Kaplan Meier survival plots. Patients with positive HO-1 expression in their tumors present a longer OS time than patients displaying negative HO-1 expression ($N = 74$; $p = 0.002$, log-rank test). (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

invasive CRC. Staining with hematoxylin & eosin to each slide was first done to study histopathological characters in order to confirm the diagnosis (representative samples: Fig. 1B and C). Thirty two of the samples contained histologically-normal tissues (non-malignant epithelia) adjacent to the tumor tissues. We then proceeded with the immunohistochemistry for HO-1 as described in the Materials and methods section. HO-1 immunoreactivity was observed in apical cells of the crypt within the non-malignant epithelia (Fig. 1F, black arrow), and basal cells showed no HO-1 staining (Fig. 1F, red arrow). Tumor specimens showed higher rates of expression (69.3%, 68/98; Fig. 1E and G) than their respective surrounding non-malignant tissues (12.5%, 4/32; $p = 0.04$; Fig. 1D and F and Supplementary Table 1). The analysis of HO-1 expression levels also showed differences between tumor (T) and adjacent non-malignant tissues (NT) (median IRS: 3 versus median IRS: 1, respectively; $p = 0.0001$; Fig. 1H).

3.2. HO-1 is associated with increased overall survival time of invasive CRC patients

We subsequently studied the correlation between HO-1 protein expression and several clinic-pathological parameters important for CRC prognosis such as gender, tumor location, differentiation grade, mitotic index, nuclear index, lymph node involvement, the presence of metastasis, K-ras status, pre-surgical CEA levels and EGFR expression (Table 1). This analysis revealed a significant correlation between HO-1 positive expression and wild-type K-ras status ($p = 0.04$) and normal CEA levels ($p = 0.04$, χ^2 test).

Our next step was to examine whether HO-1 could be considered a prognostic factor in invasive CRC, and for this purpose we analyzed if HO-1 protein expression was associated with patient overall survival time. For this analysis and in order to avoid the problems of multiple cut-point selection, ROC curve analysis was performed to determine a reasonable cut-off point of HO-1 in the CRC samples. The best HO-1 cut-off point for overall survival (OS) was a score ≥ 1 (sensitivity: 61.11 and specificity: 69.09; AUC = 0.703; 95% confidence interval: 0.585–0.804; $p = 0.0018$). In brief, patients with a cut-off score ≥ 1 were regarded as HO-1 positive and those with a score < 1 were regarded as HO-1 negative.

Interestingly, we found an association between tumor HO-1 positivity and a longer patients' survival time (Fig. 1I; $p = 0.002$, log rank test). The patients whose tumors presented HO-1 positive expression had a median survival time of 65 months whereas the patients presenting negative expression of the protein had a median survival time of 39 months.

3.3. An increase in HO-1 expression and nuclear localization is associated with malignant progression in a CRC animal model

In order to gain insight into the significance of HO-1 in CRC, we assessed the expression of HO-1 during the progression of a chemically-induced CRC model. For this purpose, we isolated normal epithelial tissues (Fig. 2A and E), polyps (Fig. 2B and F), adenocarcinoma tissues (Fig. 2C and G) and signet-ring cell carcinoma tissues (Fig. 2D and H) that were obtained at different times of disease progression. Signet-ring cell carcinomas have been characterized as very malignant entities (Nissan et al., 1999; Bradford, 1976). We found positive staining in 2/10 (20%) of the normal tissue, 3/3 (100%) of the polyps, 12/12 (100%) of adenocarcinoma specimens and 3/3 (100%) of signet-ring cell carcinoma samples. Protein expression showed similar immune-staining pattern and sub-cellular localization to those observed in the human samples. Polyps ($p = 0.008$), adenocarcinoma ($p < 0.001$) and signet cell carcinoma ($p = 0.009$) showed higher rates of HO-1 protein expression than normal epithelia (Fig. 2I). HO-1 has been originally described as a microsomal enzyme. However an increasing number of reports (Sacca et al., 2007; Gueron et al., 2009; Birrane et al., 2013; Yin et al., 2014) including ours (Gandini et al., 2012, 2014) showed nuclear localization of the

Table 1
Summary of patient characteristics and HO-1 expression and correlation with clinic-pathological features.

	Cases	HO-1 positive n (%)	HO-1 negative n (%)	<i>p</i>
<i>Gender</i>				
Female	36	20(55.5)	16(44.5)	0.400
Male	39	25(64.1)	14(35.9)	
Not available	8			
<i>Tumor location</i>				
Colon	68	45(66.2)	23(33.8)	0.900
Rectus	15	10(66.7)	5(33.3)	
Not available	–			
<i>Differentiation grade</i>				
I	20	17(85)	3(15)	0.058
II	53	30(56.6)	23(43.4)	
III	5	4(80)	1(20)	
Not available	5			
<i>Mitotic index</i>				
I	20	16(80)	4(20)	0.100
II	32	18(56.25)	14(43.75)	
III	2	1(50)	1(50)	
Not available	29			
<i>Nuclear index</i>				
I	8	5(62.5)	3(37.5)	0.200
II	41	23(56.10)	18(43.9)	
III	4	4(100)	0(0)	
Not available	30			
<i>Lymph node involvement</i>				
No	17	12(70.6)	5(29.4)	0.400
Yes	46	28(60.87)	18(39.13)	
Not available	12			
<i>K-ras status</i>				
Wild type	55	36(65.45)	19(34.55)	0.040
Mutated	16	6(37.5)	10(62.5)	
Not available	12			
<i>CEA levels</i>				
Normal	20	15(70.6)	5(29.4)	0.040
High	26	12(46.2)	14(53.8)	
Not available	40			
<i>EGFR1 positivity</i>				
1+	17	8(47)	9(53)	0.100
2+	13	10(77)	3(23)	
3+	5	4(80)	1(20)	
Not available	48			

χ^2 test was used.

enzyme in some tissues and cell types and demonstrated an association of nuclear localization with the progression of the disease (Gandini et al., 2012). Therefore we studied HO-1 sub-cellular localization in tissues from this animal model of CRC. Nuclear staining in 3/3 normal tissues (mean percentage of stained nuclei of all the positive samples: $7.3 \pm 2\%$), 3/3 of the polyps ($15 \pm 3\%$; $p = 0.035$), 8/12 of adenocarcinoma samples ($15 \pm 5\%$; $p = 0.019$) and 3/3 of signet cell carcinoma ($25.6 \pm 3\%$; $p = 0.002$) was observed, thus showing that HO-1 nuclear localization rates increase with disease progression ($p < 0.0001$, ANOVA; Fig. 2J).

3.4. Nuclear localization of HO-1 is also observed in a CRC cell line and in human tissues

Since we had observed that nuclear localization was present and increased with tumor progression in the animal model of CRC, we also analyzed the presence of nuclear HO-1 by immunofluorescence in the human CRC cell line HCT116 and by immunohistochemistry in human CRC tissues. HO-1 expression was nuclear in HCT116 cells and increased with pharmacological activation (Fig. 3A). Additionally, all the adjacent non-malignant tissues that were positive for HO-1 showed cytoplasmic localization of the protein (100%) (Fig. 3B) whereas of the tumor samples

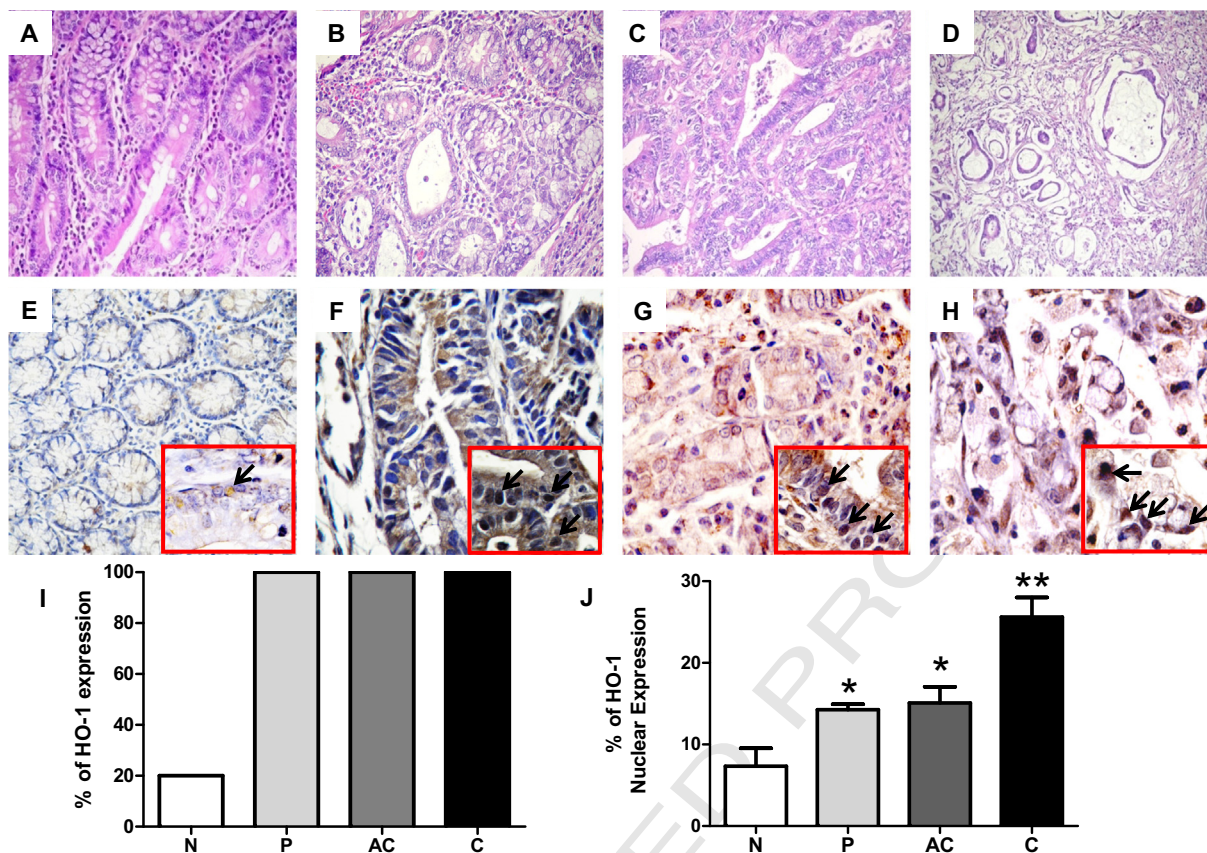


Fig. 2. HO-1 expression and nuclear localization increases with malignant progression in a chemically-induced animal model of CRC. Hematoxylin and eosin staining (A–D) (100 \times) and immunohistochemistry for HO-1 (E–H) (400 and 1000 \times) in non-malignant tissues (A and E), polyps (B and F), adenocarcinoma tissues (C and G) and signet-ring cell carcinoma tissues (D and H) obtained from the animal model. Black arrows show cytoplasmic (E) and nuclear (F–H) staining of HO-1. I. Analysis of HO-1 expression in non-malignant epithelia (N), polyps (P), adenocarcinoma (AC) and a signet-ring cell carcinoma (C). J. Graph showing the frequencies observed for the nuclear expression of HO-1 in non-malignant epithelia (N), polyps (P), adenocarcinoma (AC) and signet-ring cell carcinoma (C); * $p = 0.035$, * $p = 0.019$ and ** $p = 0.002$ compared to adjacent non-malignant tissues (N).

317 that were positive for HO-1, 61.8% (42/68) showed only cytoplasmic localization and 38.2% (26/68) showed both cytoplasmic and nuclear localization of the protein (Fig. 3C and D) ($p = 0.0001$, χ^2 test).

320 3.5. HO-1 decrease viability of cell lines that do not have alterations in the p53 gene

322 The HO-1 protein has been mainly described as having a pro-tumoral activity in most types of cancers. However, as already mentioned, an antitumoral role has been suggested in prostate (Gueron et al., 2009) and colorectal cancers (Becker et al., 2007; Kang et al., 2012). The results obtained in human CRC biopsies corroborate that HO-1 expression is increased in tumors and that it is associated with a better patient outcome, thus suggesting an antitumoral activity for HO-1 in CRC. Therefore, in order to further investigate the role of HO-1 in CRC progression, we first evaluated the involvement of HO-1 on the viability of the colorectal cancer cell line HCT116. For this purpose, we first performed time-response analyses for cell viability after activation of HO-1 with hemin (100 μ M) and observed that the number of cells decreased at 96 h of treatment ($p = 0.01$) (Fig. 4A). Taking this time point for future studies of cellular viability, we proceeded to perform dose–response studies with hemin and with the HO-1 inhibitor SnPP. A decrease in the viability of the cells was observed when doses of 40 μ M and 100 μ M of hemin were used (Fig. 4B). Contrariwise, an increase in cell number was observed when cells were treated with SnPP (10 μ M) (Fig. 4C). Since the tumor suppressor p53 is important for the regulation of cellular survival and is frequently des-regulated in CRC, we performed similar experiments using the cell line HCT116 p53 –/–

343 which lacks p53 protein. Interestingly, the effect of hemin and SnPP treatment on cellular survival was blunted (Fig. 4D and E). This difference observed between the two cell lines regarding their response to HO-1 up-regulation may be due to the presence of p53 acting through regulation of the expression of p21, an inhibitor of cyclin-dependent kinases (CDK's) and regulator of cell cycle progression. We therefore analyzed whether activation of HO-1 induced the expression of this and other proteins related to cell cycle arrest and cellular viability. As seen in Fig. 4F, modulation with hemin produced up-regulation of HO-1 in both cell lines (with and without p53), thus suggesting that the expression of HO-1 is independent of p53. Also hemin-up-regulation of HO-1 was accompanied with an increase in p21, p27 (Fig. 4G and H) and a decrease in cyclin D1 (not shown) in HCT116 cells. Instead, in the cell line lacking p53, overexpression of HO-1 was not accompanied by up-regulation of p21 or p27 (Fig. 4E and F). Time-response analyses performed with hemin revealed that p53 induction occurs simultaneously with HO-1 induction (Fig. 4I).

359 Since both HCT116 cell lines differ mainly in the presence or absence of p53, we inferred that this tumor suppressor might be involved in the effect of HO-1 on cell viability. Because of this, we additionally investigated the activation of HO-1 in two different CRC cell lines, the LoVo cell line that is wild type for p53 gene and the HT29 cell line that has a mutation at codon 273 of p53 gene. This mutation alters p53 sequences that are directly responsible for sequence-specific DNA binding, thus allowing the cell cycle to proceed unchecked (Oliver et al., 2010). We performed viability analyses after activation of HO-1 with hemin (100 μ M). The analysis performed in LoVo cells showed that activation of HO-1 resulted in decreased cell number ($p = 0.037$) at 96 h post-

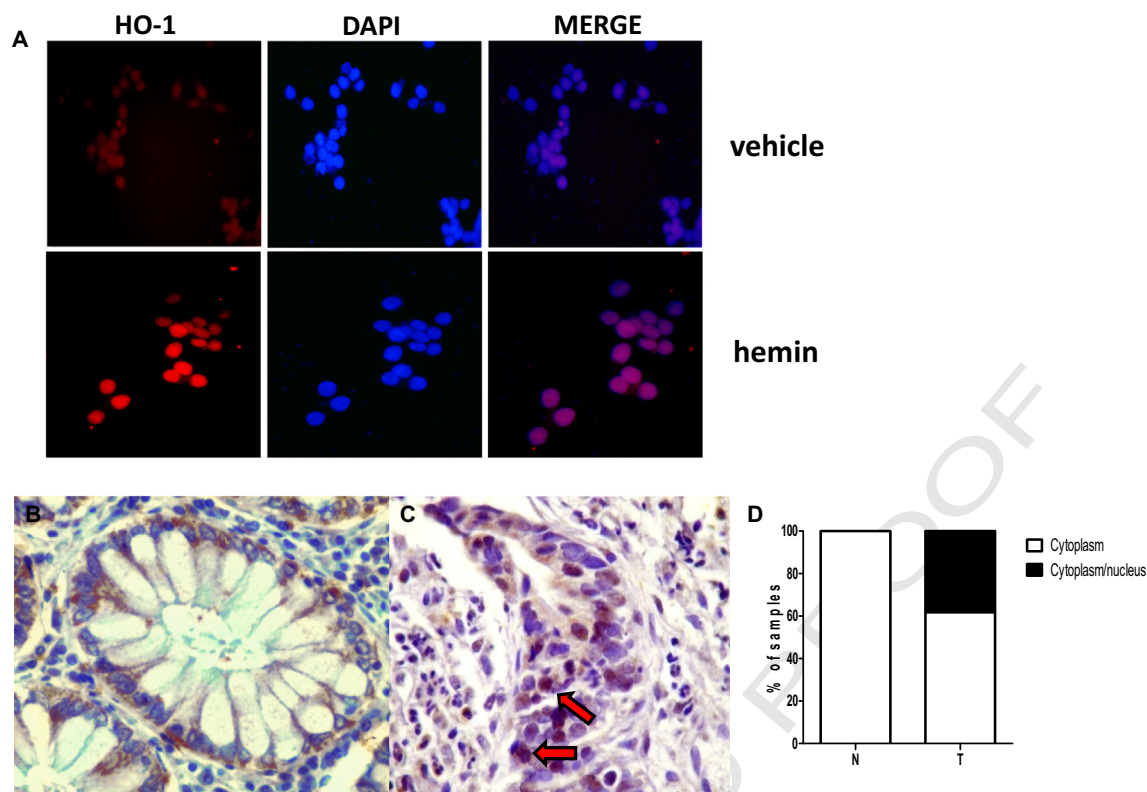


Fig. 3. Nuclear localization of HO-1 in HCT116 cell line and in human biopsies. A. HCT116 cells were treated with hemin (100 μ M) or vehicle for 24 h and immunofluorescence for HO-1 was carried out. DAPI counter-stain was performed. B. The absence of nuclear localization of HO-1 in adjacent non-malignant human tissue. C. Nuclear localization of HO-1 in human tumor assayed by immunohistochemical staining (1000 \times). Red arrows show nuclear staining. D. Graph depicting nuclear and cytoplasmic rates of HO-1 in human tumor. 61.8% of samples showed cytoplasmic expression, while 38.2% showed both cytoplasmic and nuclear staining ($p = 0.0001$, χ^2 test). (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

371 treatment (Fig. 5A). On the contrary, a time- and dose–response survival
 372 assay carried out in the HT29 cell line showed that the number of cells
 373 remained unchanged or slightly increased when treated with various
 374 doses of hemin at different time points (Fig. 5B). These results suggest
 375 that a functional p53 is necessary for HO-1-inhibition of cellular viability.
 376 This hypothesis was confirmed by re-expressing p53 tumor suppressor
 377 in HCT116 p53 $^{-/-}$ cell line by transient transfection of a pcDNA3 p53
 378 plasmid and further performing viability analyses. We detected that the
 379 cell lines with altered p53 neither responded to HO-1 modulation nor
 380 showed a slight increase in cell survival, as previously observed (Fig. 5C;
 381 $p = 0.361$). However, when p53 was re-introduced in cells, their viability
 382 decreased after hemin treatment (Fig. 5C; $p = 0.0047$) presenting a
 383 similar behavior to p53-containing HCT116 cells ($p = 0.0006$). The
 384 efficiency of the transfection was analyzed by western-blot (Fig. 5D) and
 385 also by immunofluorescence (not shown), with 10% of HCT116 p53 $^{-/-}$
 386 cells expressing p53 protein after transfection.

388 Since it has been demonstrated that HO-1 modulators display direct
 389 effects on some cellular processes that are not mediated by HO-1 activity
 390 (La et al., 2009) we proceeded to genetically over-express the enzyme
 391 in order to confirm the results obtained with pharmacological modulation
 392 of HO-1. We repeated the viability assay after genetic over-expression
 393 of HO-1 in the HCT116 and LoVo cell lines (Fig. 5E and F), obtaining
 394 similar results to those obtained by pharmacological over-expression
 395 of the protein ($p = 0.0017$ and $p = 0.0023$, respectively). The
 396 efficiency of the transfection was analyzed by immunofluorescence
 397 (50% of cells, data not shown).

398 The expression of cell survival-related proteins was analyzed by
 399 western-blot in HO-1-overexpressing HCT116 cells. Similar results
 400 were obtained than with the pharmacological induction of HO-1 (Fig. 5G).

3.6. The over-expression of HO-1 causes cell cycle arrest and apoptosis 401

402 We subsequently analyzed the possible mechanisms underlying
 403 HO-1 effects on cell viability. For this purpose, we studied if pharmaco-
 404 logic modulation of HO-1 induces cell cycle arrest by PI staining follow-
 405 ed by flow cytometry. As observed in Fig. 6A, hemin treatment induces a
 406 G_0/G_1 arrest in HCT116 cell line ($p = 0.004$). No G_0/G_1 arrest was ob-
 407 served in HCT116 p53 $^{-/-}$ cells although a G_2/M arrest was induced
 408 (Fig. 6B; $p < 0.05$).

409 Subsequently, we considered if HO-1 activation could also regulate
 410 apoptosis in HCT116 cells and for this purpose labeling with Annexin
 411 V-fluorescein isothiocyanate (FITC) was performed. As shown in Fig. 6C,
 412 activation of HO-1 induces an increase in Annexin V staining in these
 413 cells ($p = 0.01$), thus suggesting that apoptosis is involved in HO-1
 414 modulation of cellular viability.

415 In order to further analyze the mechanisms of HO-1-induced apo-
 416 ptosis we checked the expression of Bax, a pro-apoptotic protein that
 417 is known to be up-regulated by p53 (Chipuk et al., 2004). We observed
 418 an increase in the levels of this protein in the HCT116 cells when HO-1
 419 was pharmacologically induced (Fig. 6D).

420 One common alteration in colon cancer is the hyper activation of the
 421 Akt and PKC (mainly β isoform) pathways. Therefore, we evaluated the
 422 activation state of Akt and expression of PKC following pharmacological
 423 modulation of HO-1. There was an increase in PKC β I levels in the paren-
 424 tal cell line HCT116 when HO-1 was activated. In turn, there was a de-
 425 crease of PKC β II following treatment with hemin, and an increase
 426 when SnPP was used (Supplementary Fig. 1). Importantly, a delay in
 427 the phosphorylation of Akt following HO-1 activation was observed in
 428 these cells (Fig. 6E). These results suggest that Akt and PKC β I pathways
 429 are involved in the effects observed on cellular survival.

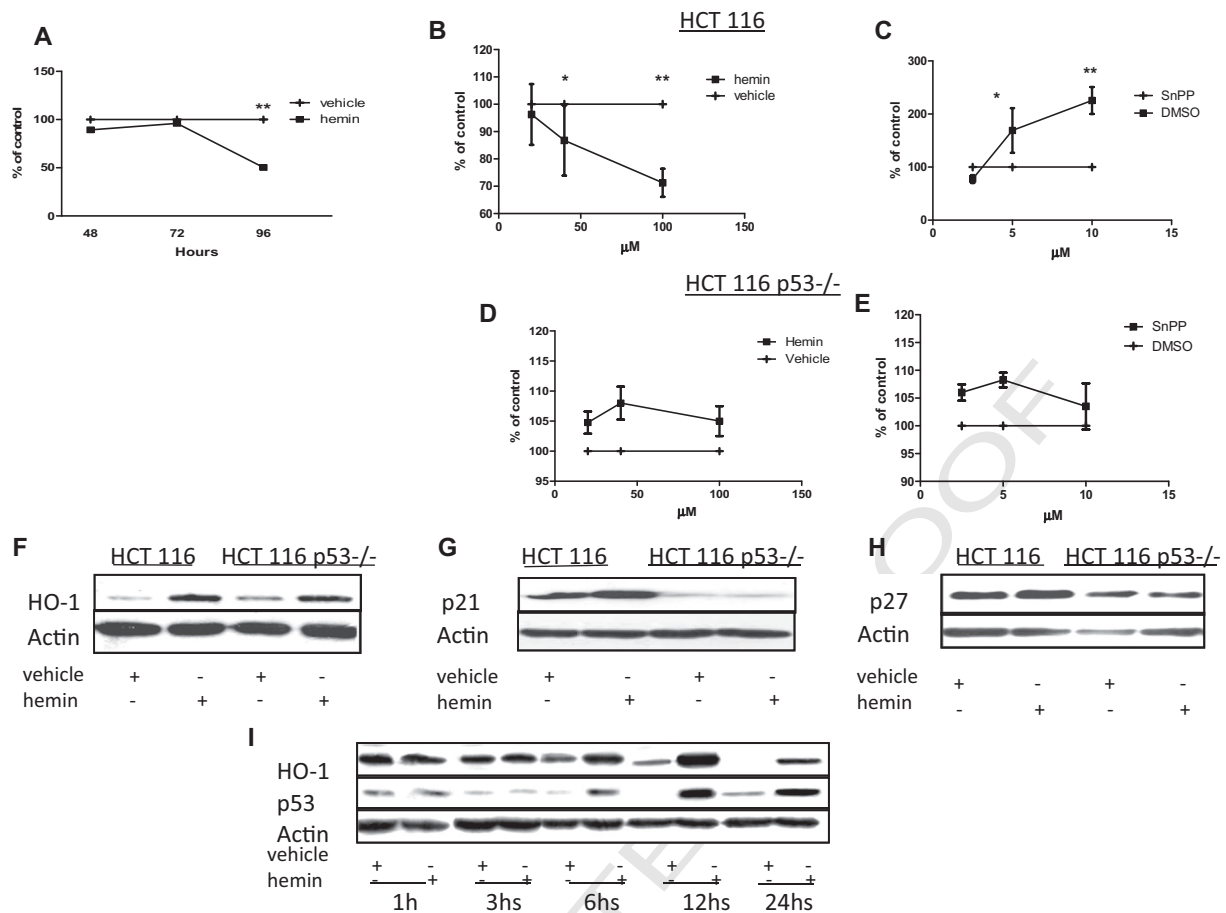


Fig. 4. Effect of pharmacological modulation of HO-1 on cell viability and on cell cycle-related protein expression in HCT116 and HCT116 p53^{-/-} cell lines. A. Time-response survival assay following hemin (100 μM) or vehicle treatment. The number of HCT116 cells decreased at 96 h. B–E. Dose-response viability assays following hemin (B and D) (20 μM, 40 μM or 100 μM) – or SnPP (C and E) (2 μM, 4 μM or 10 μM) – treatment in HCT116 (B and C) and HCT116 p53^{-/-} (D and E) cell lines. HCT116 and HCT116 p53^{-/-} cells were treated for 24 h with vehicle or hemin (100 μM) and protein lysates were electrophoresed and blotted against HO-1 (F), p21 (G) and p27 (H). I. HCT116 cells were treated for the times indicated, with vehicle or hemin (100 μM). The expression of HO-1 and p53 was analyzed by western-blotting. Actin was used as loading control.

3.7. HO-1 modulates cellular migration of HCT116 cell line

To evaluate the migratory capacity of HCT116 cells following HO-1 modulation, we used an in vitro scratch wound assay. Confluent monolayers of hemin-, vehicle- and SnPP-treated HCT116 cells were wounded. Wound closure was monitored every hour for 24 h as previously described. HCT116 cells treated with vehicle migrated and almost covered the wound by 24 h (uncovered wound area 60.28%), whereas in hemin-treated cells, a significant area of the wound (78.95%) remained uncovered over the same period. The inhibition of HO-1 with SnPP produced opposite results, that is the wound closed faster with SnPP than with the vehicle-treated cells (Supplementary Fig. 2). These results suggest that HO-1 is involved in modulating the migratory capacity of HCT116 cell line.

4. Discussion

In this study we have demonstrated that HO-1 is over-expressed in tumor epithelium of invasive CRC biopsies compared to their adjacent non-malignant epithelium. These results are in agreement with those obtained by our group in squamous cell carcinoma (Gandini et al., 2012), glioma (Gandini et al., 2014) and non-small cell lung cancer (Degese et al., 2012) where an up-regulation of HO-1 with tumor progression was found. Since induction of HO-1 is a fundamental cellular defense process against oxidative stress and other environmental insults, its increase in tumor cells may provide the first line of cellular

defense of cancer cells against these insults. This might explain the increase in HO-1 expression observed in many different tumors. Indeed, HO-1 over-expression has also been demonstrated in lymphosarcoma (Schacter and Kurz, 1982), prostate carcinoma (Sacca et al., 2007) brain tumors (Deininger et al., 2000; Hara et al., 1996) renal carcinoma (Goodman et al., 1997), hepatoma (Doi et al., 1999), melanoma (Torisu-Itakura et al., 2000), Kaposi sarcoma (McAllister et al., 2004) pancreatic cancer (Berberat et al., 2005) and in chronic myeloid leukemia (Mayerhofer et al., 2004).

Regarding human CRC, a previous report showed apical staining in crypts of normal colonic epithelia, similar to our observations, and higher expression rates in CRC than in colon adenoma samples (Becker et al., 2007). Contrary to our results, they reported lower staining in high grade tumors. Two recent reports also showed higher expression levels in tumors (Kang et al., 2012; Yin et al., 2014). To our knowledge, no works have followed up with the study of HO-1 expression in human CRC. Instead, there are several studies addressing HO-1 function in both the normal intestinal physiology and inflammatory intestinal diseases. Interestingly, HO-1 expression is usually increased in gastrointestinal inflammation and injury, processes associated with cancer progression; this up-regulation was shown in gastric ulcers (Guo et al., 2002), colitis (Wang et al., 2001), radiation enteritis (Giriş et al., 2006), inflammatory bowel disease (Paul et al., 2005) both in patients and animal models. This up-regulation of HO-1 has been shown to play a role in protecting from inflammation and oxidative injury in the gastrointestinal tract (Zhu et al., 2011), a role that may inhibit CRC tumor progression and/or may improve patient outcome. According to

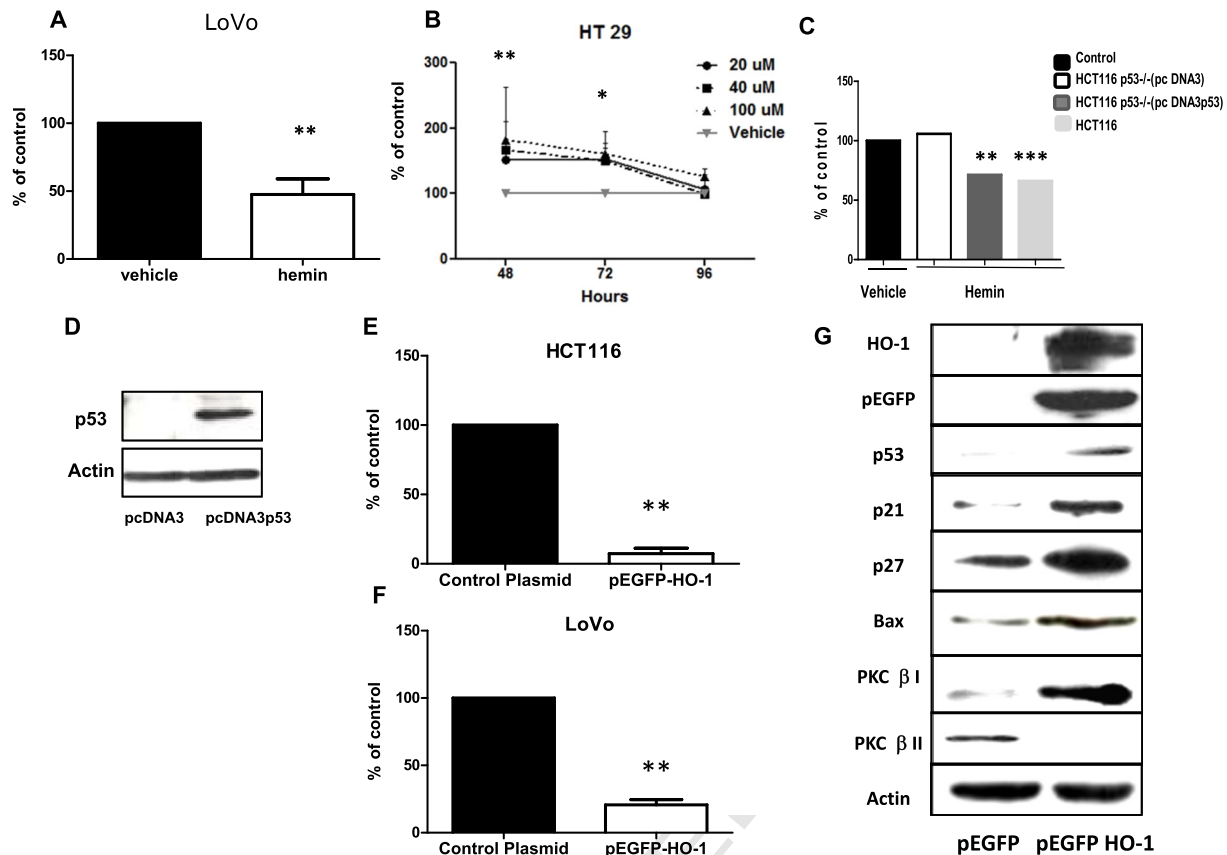


Fig. 5. Involvement of a functional p53 on HO-1 modulation of cell viability. **A.** Viability assay in the p53-wild type-containing LoVo cell line. Up-regulation of HO-1 with hemin (100 μ M) decreased cell number ($p = 0.037$; T test). **B.** Viability assay in the p53-mutated-containing HT29 cell line. Dose (20, 40 and 100 μ M) and time (48, 72 and 96 h)-response analysis. Hemin treatment leads to increased cell count at 48 and 72 hour treatment (** $p = 0.01$, * $p = 0.05$). **C.** Viability assays in HCT116 p53 $^{-/-}$ cell line transfected with pcDNA3 or pcDNA3 p53 and in HCT116 following hemin treatment (** $p = 0.0047$, *** $p = 0.0006$). **D.** Western blot analysis showing re-expression of p53 in the HCT116 p53 $^{-/-}$ cell line. Viability assays in the HCT116 (**E**) and LoVo (**F**) cell lines. Genetic over-expression of HO-1 decreased cell number in both cell lines (** $p = 0.0017$, ** $p = 0.0023$). **G.** Protein lysates of transiently transfected HCT116 cells were electrophoresed and blotted against HO-1, pEGFP, p53, p21, p27, Bax, PKC β I, and PKC β II. Actin was used as loading control.

480 this, we have also demonstrated that the expression of HO-1 in tumors
 481 is associated with increased overall survival of patients with CRC. These
 482 results are in agreement with previous studies of HO-1 expression in
 483 CRC where a correlation between HO-1 positivity and a better long
 484 term survival was found (Becker et al., 2007). However, they are con-
 485 trary to the majority of the tumor types analyzed in which HO-1 has
 486 been positively associated with tumor progression (Was et al., 2006).
 487 The exceptions to this pro-tumoral role for HO-1 have been the findings
 488 of Becker and col in CRC already mentioned (Becker et al., 2007) and the
 489 observations in tongue squamous cell carcinomas (Yanagawa et al.,
 490 2004), mammary tumors (Hill et al., 2005 and unpublished observa-
 491 tions from our laboratory) and prostate cancer (Gueron et al., 2009).

492 In order to further demonstrate that HO-1 expression increases with
 493 tumor progression, we also evaluated its expression in a CRC animal
 494 model. HO-1 up-regulation as well as an increase in the incidence of nu-
 495 clear localization were observed during tumor progression. Further-
 496 more, HO-1 nuclear expression was associated with less differentiated,
 497 more aggressive tumors in the animal model. This nuclear expression
 498 was also observed in human samples. These results are in agreement
 499 with the observations of Yin et al. (2014) where an increase in nuclear
 500 HO-1 was observed with advanced tumor stages. Nuclear localization
 501 was also reported by our group in human samples and in an animal
 502 model of squamous cell carcinoma (Gandini et al., 2012) and also
 503 agree with previous reports demonstrating an increase in HO-1 nuclear
 504 staining when oral epithelial dysplasias progress from moderate to
 505 severe (Lee et al., 2008). Similar results were also obtained in prostate

506 cancer in which HO-1 nuclear expression was found to be lower in ad-
 507 jacent non-malignant tissues than in prostate carcinoma ones (Sacca
 508 et al., 2007). Additionally, in support of a role of nuclear HO-1 in tumor
 509 progression, a recent report demonstrated that cigarette smoke induces
 510 nuclear translocation of HO-1, and this localization promotes vascular en-
 511 dothelial growth factor secretion, which favors prostate tumor progres-
 512 sion (Birrane et al., 2013). Although this novel nuclear localization of
 513 HO-1 has been demonstrated in some tissues, its significant role has not
 514 been completely addressed. In this regard, it was postulated that the nu-
 515 clear form of HO-1 may up-regulate genes that promote cytoprotection
 516 against oxidative stress (Lin et al., 2007). There is evidence showing that
 517 nuclear HO-1 could have a physiological role independent of its enzymatic
 518 activity. For example, gene transfection of the activity-lacking mutant
 519 HO-1 protects cells against oxidative stress (Busserolles et al., 2006). As
 520 the HO-1 structure does not show DNA-binding motifs it seems that
 521 this protein is not a typical transcription factor (Lin et al., 2007). However,
 522 HO-1, acting as a transcriptional co-regulator protein, may be able to
 523 modulate transcription factors, nuclear localization being thus neces-
 524 sary for these effects, and this explains its presence in the nuclear
 525 compartment.

526 As already stated, most of the literature shows a pro-tumoral role of
 527 HO-1 (Was et al., 2006) with a few exceptions already mentioned. Since
 528 we observed an association of HO-1 with longer patients' survival time,
 529 we hypothesized that HO-1 expression in CRC cells could inhibit cellular
 530 survival. Our results demonstrate that HO-1 activation decreases cell sur-
 531 vival through cell cycle arrest and induction of apoptosis. Importantly,

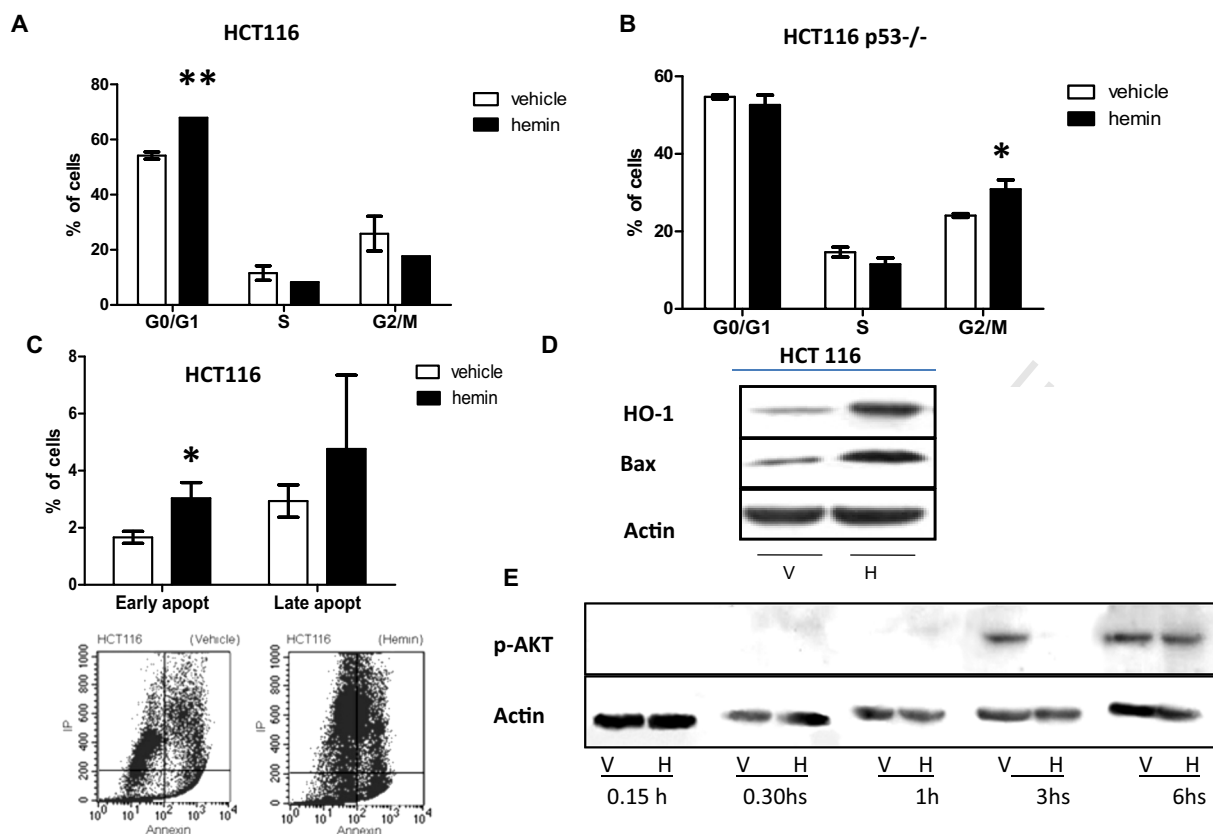


Fig. 6. HO-1 produces cell cycle arrest in G₀/G₁ and early apoptosis in HCT116 cells and modulates the expression of the pro-apoptotic protein Bax. A and B. Distribution of the phases of the cell cycle by determining the DNA content carried out by flow cytometry using IP staining. The histograms show the percentage of cells from each cell line, HCT116 (A) and HCT116 p53^{-/-} (B), located in different cell cycle phases after treatment with hemin and vehicle (100 μM). Cell arrest was observed in G₀/G₁ phase in those cells that have wild type p53 (***p* = 0.004) and G2/M in cells lacking p53 (**p* < 0.05). C. Detection of apoptosis in HCT116 cells. The cells were analyzed by flow cytometry using Annexin-V staining. Percentage of cells in early apoptosis (**p* = 0.01) and late apoptosis (*p* > 0.17) after treatment with hemin (100 μM) and vehicle. D. HCT116 cells were treated with vehicle (V) and hemin (H, 100 μM) for 24 h and HO-1 and Bax expression determined. Actin was used as loading control. E. The HCT116 cells were subjected to a 6-hour pretreatment with vehicle (V) and hemin (H); then they were maintained for 16 h in serum-free medium and subsequently treated for 15 min, 30 min, 1 h, 3 h and 6 h with vehicle (V) or hemin (H, 100 μM) in medium containing serum. The expression of phosphorylated Akt was analyzed by western blot. The results were normalized with actin.

p53 tumor suppressor protein is necessary for these effects, since HO-1 reduction in cell survival is blunted in the HCT116 cell line lacking p53 and in the HT29 which bears a mutated p53. On the other hand, previous work from other laboratories done with the p53-mutated Caco-2 cell line showed an anti-apoptotic activity for HO-1 (Busserolles et al., 2006). Altogether, these results support the hypothesis of HO-1 decreasing survival in CRC through wild type p53 up-regulation. p53 tumor suppressor gene is mutated in over 50% of human tumors and plays an important role in the response to genotoxic stress and hypoxia. The contradictory responses to HO-1 activation observed in different tumor cells might be explained by the different status of p53 that they carry. However, the relationship between HO-1 and p53 is not yet clear as there are some reports showing that p53 induces HO-1, and other ones demonstrating that HO-1 is upstream of p53. For example in the lymphoid organs of γ-irradiated mice, HO-1 was demonstrated to be a direct p53 target gene (Meiller et al., 2007). Nam et al. also demonstrated a role for p53 in promoting cellular survival through the activation of HO-1 (Nam and Sabapathy, 2011). On the other hand, Lee et al. (2008) have demonstrated that HO activity is involved in the regulation of p53 expression in a human retinal pigment epithelial cell line and Kim et al. (2014) showed that HO-1 is necessary for the up-regulation of p53 induced by 15d-PGJ₂ in breast cancer cells. Importantly, in non-small cell lung carcinoma HO-1 up-regulated p53 and this was accompanied by a reduction in proliferation, migration and angiogenic potential (Skrzypek et al., 2013). Additional investigations should be performed in order to further establish wild-type p53 as responsible for HO-1 effects on cellular survival in CRC.

Part of the observed effects upon HO-1 activation may be secondary to modulation of Akt and PKC β pathways. The role of PKC β is controversial and varies in different cell lines. According to the results obtained by Choi et al. (1990) PKC βI is associated with reduced tumorigenicity, whereas activation of PKC βII isoform plays a direct role in increasing colorectal cancer cell proliferation (Sauma et al., 1996). The role of Akt pathway in CRC is very well understood and modulates both cell survival and migration processes (Agarwal et al., 2013). Finally, we demonstrated that HO-1 activation reduces cell migration. This action could involve modulation of the matrix metalloproteinases (MMP)'s levels specially the gelatinases MMP-2 and MMP-9 that have important roles in tumor invasion, metastasis and angiogenesis in colorectal cancer (Tutton et al., 2003). In relation to this, there is evidence that indicates that HO-1 reduces the MMP-9 levels (Gueron et al., 2009).

In conclusion, we provide evidence that HO-1 is up-regulated during CRC progression and this overexpression is associated with an increase in the overall survival time of patients. We also provide evidence that demonstrates that the effects of HO-1 in CRC involve a decrease in cellular migration and survival and that the p53 tumor suppressor protein is necessary for this effect. Altogether these results point to an anti-tumoral role of HO-1 in wild-type-p53-bearing CRC.

Supplementary data to this article can be found online at <http://dx.doi.org/10.1016/j.yexmp.2014.09.012>.

Conflict of interest statement

The authors disclose no potential conflicts of interest.

Acknowledgments

This work was supported by grants from the CONICET, ANPCyT, the Secretaría Técnica de la Universidad Nacional del Sur, and from the University Medical Center of Groningen.

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