

# Panx1 knockout promotes preneoplastic aberrant crypt foci development in a chemically induced model of mouse colon carcinogenesis

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## Abstract

Colorectal cancer, which is the third leading cause of cancer-related deaths worldwide, is a multistep disease, featuring preneoplastic aberrant crypt foci (ACF) as the early morphological manifestation. The roles of hemichannel-forming transmembrane Pannexin 1 (Panx1) protein have not been investigated in the context of colon carcinogenesis yet, although it has contrasting roles in other cancer types. Thus, this study was conducted to examine the effects of Panx1 knockout (Panx1<sup>-/-</sup>) on the early events of chemically induced colon carcinogenesis in mouse. Wild type (WT) and Panx1<sup>-/-</sup> female C57BL/6J mice were submitted to a chemically induced model of colon carcinogenesis by receiving six intraperitoneal administrations of 1,2-dimethylhydrazine (DMH) carcinogen. Animals were euthanized 8 h (week 7) or 30 weeks (week 37) after the last DMH administration in order to evaluate sub-acute colon toxicity outcomes or the burden of ACF, respectively. At week 7, Panx1 genetic ablation increased DMH-induced genotoxicity in peripheral blood cells, malondialdehyde levels in the colon, and apoptosis (cleaved caspase-3) in colonic crypts. Of note, at week 37, Panx1<sup>-/-</sup> animals showed an increase in aberrant crypts (AC), ACF mean number, and ACF multiplicity (AC per ACF) by 56%, 57% and 20%, respectively. In essence, our findings indicate that Panx1 genetic ablation promotes preneoplastic ACF development during chemically induced mouse colon carcinogenesis, and a protective role of Panx1 is postulated.

## KEYWORDS

aberrant crypt foci, C57BL/6J mouse, colon carcinogenesis, dimethylhydrazine

## 1 | INTRODUCTION

Colorectal cancer (CC) is the third leading cause of cancer-related deaths worldwide, affecting both men and women.<sup>1</sup> (Sung et al., 2021). This disease, which is predominantly sporadic (up to 70% of cases), is closely related to smoking, alcohol intake, obesity, sedentary lifestyle, and a diet rich in processed foods, red meat, and sugary drinks.<sup>2–4</sup> Although new treatments have improved CC-related outcomes mainly in countries with high human development index (HDI), the 5-year relative survival remains at 64.7%, and it is up to 10% for patients with metastatic disease.<sup>3,5</sup> CC develops under the emergence and accumulation of a series of morphological, genetic or epigenetic changes in the epithelial cells. During this multi-step process, the aberrant crypt foci (ACF) are the earliest histologic alterations to arise. Morphologically, these lesions organize into clusters of differentially stained crypts with an abnormally thick epithelial lining, and oval to slit-shaped crypt openings. A percentage of these lesions also gather key molecular alterations, such as KRAS mutations and  $\beta$ -catenin mutation/accumulation.<sup>6,7</sup> Both clinical and preclinical assays have documented an association between the presence of ACF and risk for CC development. For these reasons, ACF are considered putative preneoplastic lesions in the human colonic mucosa, although their progression towards CC may depend on multiple features.<sup>6,7</sup> In short-to-medium term chemically induced colon carcinogenesis bioassays in rodents, ACF have already proved their predictive value for the prevention, treatment or progression to CC.<sup>8–11</sup> Although some ACF may gather molecular features that support them as CC precursors, the landscape of molecular alterations within these lesions is still not fully unveiled and the search for molecular targets is warranted.

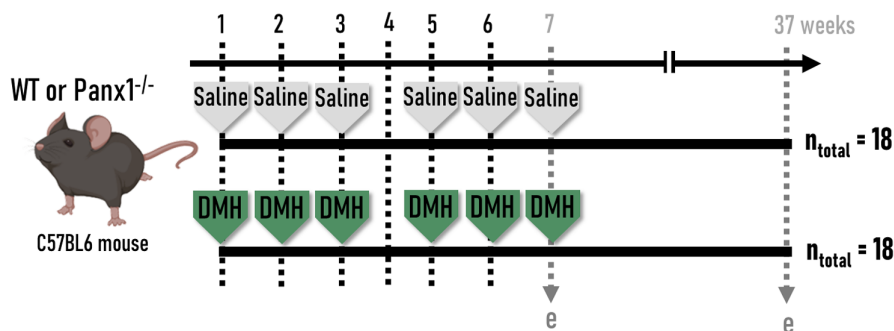
In this background, the roles of Pannexin 1 (Panx1) protein, and Panx1 hemichannels, have not yet been investigated. The family of three paralog transmembrane proteins called Pannexins (Panx1, Panx2, and Panx3), constitute channels that connect the intra and extracellular environments, enabling the bidirectional exchange of essential small molecules (e.g., ions and ATP).<sup>12,13</sup> Panx1 is the most commonly expressed protein of this family, although Panx family tissue distribution ranges from ubiquitous to very restricted, being paralog-dependent, cell type-specific, and developmentally regulated. Panx1 has been implicated in a myriad of physiological and pathophysiological processes.<sup>12–14</sup> There is little evidence for Panx1 distribution in the normal colon. Diezmos et al.<sup>15</sup> showed that Panx1 is broadly immuno-expressed in all layers of this organ, including the cytoplasm and cell membranes of the enterocytes and goblet cells of the colonic mucosa. In contrast, Panx1 channels are proposed

to have critical roles in colonic inflammation, as ATP released from these channels may activate purinergic P2X7 receptors (P2X7R), promoting inflammation by paracrine signalling. Indeed, the chemical blockade of Panx1 channels hindered cytokine-induced inflammation in CC Caco-2 cells and in normal cultured human colon strips.<sup>15</sup> In relation to carcinogenesis, Panx1 overexpression in breast and gastric tumour cells was positively associated with epithelial–mesenchymal transition phenotype-related outcomes, which were reversed by genetic knock-down.<sup>16,17</sup> In cohorts of breast, gastric, and liver cancers, Panx1 overexpression was also positively correlated with poor prognosis and/or tumour inflammation.<sup>17–19</sup> In contrast to these tumour-promoting outcomes, Panx1 was demonstrated to interact with liver X $\beta$  (LX $\beta$ ) receptors, leading to an increased ATP release and subsequent P2X7R activation, which induced pyroptosis in human and rodent CC cell lines, acting as a tumour suppressor protein.<sup>20</sup> This Panx1-dependent mechanism was also confirmed in a CT26 xenograft mouse model.<sup>20</sup> In the light of the available findings, the roles of Panx1 in the early stages of colon carcinogenesis are still unknown. Thus, the present study was conducted to examine the effects of Panx1 knockout (Panx1<sup>-/-</sup>) on the early events of chemically induced colon carcinogenesis in mouse. In contrast to other gastrointestinal cancers, our preclinical findings indicate that Panx1 deletion promotes preneoplastic ACF development, and a protective property of this protein in early colon carcinogenesis should be investigated further.

## 2 | METHODS

### 2.1 | Experimental design

The global Panx1<sup>-/-</sup> mouse strain (CMV-Cre/Panx1) in C57BL/6J background was generated as previously described.<sup>21</sup> All animals were obtained from the School of Veterinary Medicine and Animal Science of the University of São Paulo (USP), and kept in Botucatu Medical School of the São Paulo State (UNESP). After a 2-week acclimatization period, 6-week-old female wild type (WT) and female Panx1<sup>-/-</sup> C57BL/6J mice ( $n=18$  animals/genotype) received six intraperitoneal (i.p.) administrations of 1,2-dimethylhydrazine colon carcinogen (DMH, 20 mg/kg body weight),<sup>22,23</sup> divided in two cycles of three administrations with a rest week between them, or saline 0.9% vehicle, as shown in Figure 1. At week 7, in order to evaluate the sub-acute toxicity outcomes caused by DMH, animals (6 mice/group/genotype) were euthanized 8 h after the last administration, as this timepoint encompasses the peak of DMH-induced apoptosis.<sup>24</sup> At week 37, the remaining animals (12 mice/group/genotype) were



**FIGURE 1** Experimental design. Saline: 1 × NaCl 0.9% i.p. administration; DMH: 1 × 20 mg/kg/b.wt. i.p. administration; e: euthanasia;  $n_{\text{total}}$ : total number of mice per genotype.

euthanized 30 weeks after the last administration, allowing for screening of the earliest microscopic lesions induced by DMH, the preneoplastic ACF.<sup>25</sup> The euthanasia procedures were performed by exsanguination under ketamine/xylazine anaesthesia (100/16 mg/kg body weight, i.p.). The animals received diet and water ad libitum and were kept in polypropylene cages (up to 4 animals/cage) in a room maintained at  $22 \pm 2^\circ\text{C}$ ,  $55 \pm 10\%$  humidity, and with a 12-h light/dark cycle (light between 07:00 a.m. and 07:00 p.m.). All procedures described herein were approved by the Ethics Committee on Animal Use of the São Paulo State University (UNESP) (#569210721).

At the necropsy, the large intestine was removed, opened longitudinally, and gently rinsed with 0.9% saline to remove residual bowel contents, and the colon length was then measured (cm). At week 7, distal colon samples (1–2 cm) were collected, snap-frozen in liquid nitrogen, and stored at  $-80^\circ\text{C}$  for biochemical analysis. Other distal colon specimens were fixed in methacarn solution (60% methanol, 30% chloroform, and 10% acetic acid) for 12 h ( $4^\circ\text{C}$ ), transferred to 70% alcohol, and embedded in paraffin. At week 37, the colon was removed and the whole sample was fixed flat in 10% formalin solution for 24 h and transferred to 70% alcohol for ACF analysis. Distal colon was sampled as they display a higher DMH-induced DNA adduct and higher incidence of (pre)neoplastic alterations than middle/proximal colon.<sup>23</sup>

## 2.2 | Comet assay

Peripheral blood samples from the retro-orbital plexus were obtained 4 h after the last DMH administration (week 7) for alkaline single-cell gel electrophoresis assay.<sup>26</sup> Blood samples were mixed with low-melting point agarose (100  $\mu\text{L}$  0.75% in PBS, Invitrogen), spread on slides pre-coated with normal point agarose (1.5% in PBS, Invitrogen), and coverslipped. Following agarose solidification ( $4^\circ\text{C}$  for 5 min), coverslips were carefully removed, and slides were incubated with cold lysis solution (2.5 M NaCl, 100 mM  $\text{Na}_2\text{EDTA}$ , 10 mM Tris-HCl, 1% Triton X-100 and 10% DMSO, pH 10) overnight, at

$4^\circ\text{C}$ . Subsequently, the slides were immersed in fresh cold alkaline electrophoresis buffer (300 mM NaOH, 1 mM  $\text{Na}_2\text{EDTA}$ , pH > 13) for 20 min. Electrophoresis was conducted for 20 min at 1 V/cm (300 mA). The slides were neutralized with 0.4 M Tris (pH 7.5), dehydrated in 100% ethanol, and stained with SYBR Gold solution (1:10,000) (Invitrogen). Fifty randomly selected nucleoids were counted per slide (duplicate) in an epi-fluorescence microscope (Olympus BX-50) using Comet Assay IV software (Perceptive Instruments). Tail intensity (% of DNA in comet tail) was calculated.

## 2.3 | ACF screening

Flat-fixed colon specimens were stained with 1% methylene blue solution and topographically evaluated for ACF occurrence, considering that these lesions feature larger, darker, thicker, and “slit-like” crypts.<sup>25</sup> Then, the ACF number/ $\text{cm}^2$  (mean number of ACF per colon length), aberrant crypt (AC)/ $\text{cm}^2$  (mean number of AC per colon length), and ACF multiplicity (mean number of AC per ACF) were calculated for each group.<sup>25</sup>

## 2.4 | Immunohistochemistry

Distal colon sections (5  $\mu\text{m}$ ) were subjected to antigen retrieval in 0.01 M citrate buffer (pH 6.0, 5 min.,  $120^\circ\text{C}$ ) in a Pascal Pressure Chamber (Dako Cytomation). Following endogenous peroxidase blockade with 10%  $\text{H}_2\text{O}_2$  in phosphate-buffered saline (PBS) (15 min), the slides were treated with skimmed milk (60 min) and incubated with primary antibodies directed against Ki-67 (i.e., cell proliferation, MA5-14520, 1:100 dilution, Thermo Fisher), cleaved caspase-3 (i.e., apoptosis, PA577887, 1:50 dilution, Thermo Fisher, EUA), phosphorylated-H2AX (i.e., DNA damage marker, MA5-27753, 1:200, Thermo Scientific) or  $\beta$ -catenin (ab32572, E247, 1:400 dilution, Abcam) in a humidified chamber (overnight,  $4^\circ\text{C}$ ). Slides were incubated with a one-step horseradish peroxidase polymer (EasyPath - Erviegas) (20 min). The reaction was visualized

with 3-diaminobenzidine (DAB) chromogen (Sigma Aldrich), and counterstained with Harris' haematoxylin. Semiquantitative analysis for Ki-67, cleaved caspase-3, and H2AX were calculated in 15 randomly selected crypts/animal/section (distal colon). Both positive and negative epithelial cells were counted, and results were expressed as the percentage (%) of Ki-67+, cleaved caspase-3+, or H2AX+ positive cells per crypt. The analyses were performed by using Image J software (National Institutes of Health).

## 2.5 | Biochemical analysis

Total proteins were extracted from 50 to 100 mg of distal colon samples in KCl buffer (1.15%) and were quantified by Bradford's method. Total proteins were used to determine malondialdehyde (MDA) levels,<sup>27</sup> and activities of superoxide dismutase (SOD)<sup>28</sup> and catalase.<sup>29</sup> For MDA, the proteins were precipitated in a 10% trichloroacetic acid solution, centrifuged (3000×g, for 5 min) and the supernatant will be removed. Thiobarbituric acid (TBA) was added in a proportion of 0.67% (1:1) and the samples were heated to 100°C (15 min). MDA reacted with TBA in a 1:2 ratio (MDA:TBA). SOD activity was measured based on the inhibition of superoxide radical reaction with pyrogallol. The catalase activity was determined in a 30% H<sub>2</sub>O<sub>2</sub> PBS solution. Absorbance was measured at 535 (MDA), 420 (SOD) or 240 (catalase) nm in a microplate reader (Spectra Max 190, Molecular Devices).

## 2.6 | Statistical analysis

All the aforementioned analyses were performed blind. Pairwise comparisons were evaluated by the Student's *t*-test. Other data were analysed using one-way analysis of variance (ANOVA) or Kruskal-Wallis and a posteriori Tukey test. Data presentation and the number of replicates are detailed in the footnotes and captions. The aforementioned analyses were performed on Prism GraphPad software (V4.03, GraphPad Software). The level of significance was set at  $p < .05$ .

## 3 | RESULTS

### 3.1 | General findings

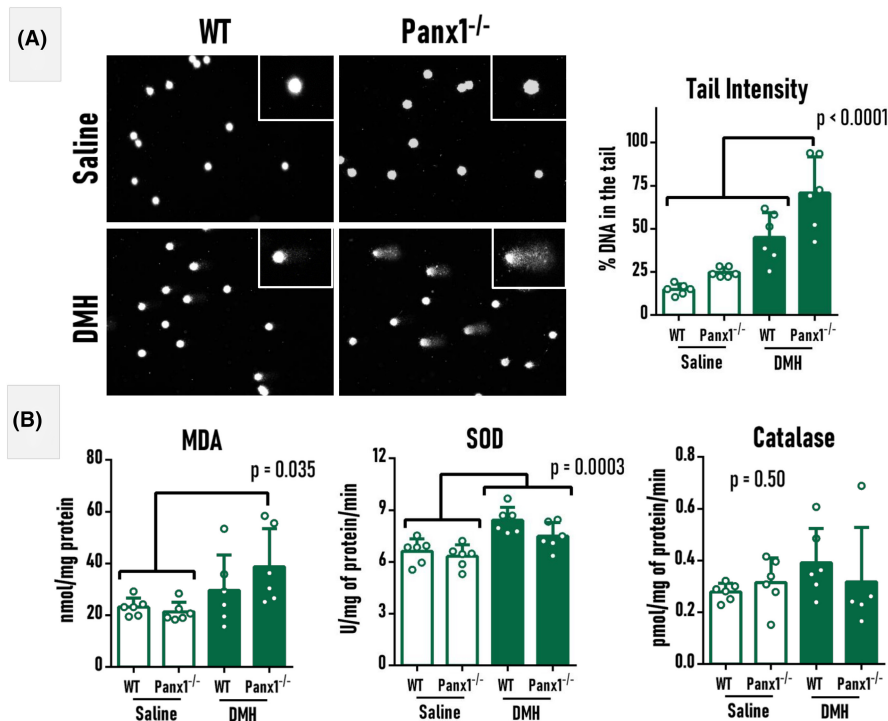
At week 7, groups showed no significant differences in initial and final body weights, absolute and relative liver weights (Table 1). At week 37, Panx1<sup>-/-</sup> animals exposed to DMH had decreased final body weight compared to WT counterparts ( $p = .0021$ ). DMH-treated Panx1<sup>-/-</sup> mice also displayed diminished absolute and relative liver weights compared to all groups ( $p < .0001$ , for both) (Table 1). Colon length remained unaltered within the groups at both timepoints. Saline-treated Panx1<sup>-/-</sup> animals showed no alterations compared to WT counterparts. Animals presented similar diet intake in both timepoints analysed (data not shown).

**TABLE 1** Implications of Pannexin 1 knockout on body weight, liver weight, and colon length during 1,2-dimethylhydrazine-induced colon carcinogenesis.

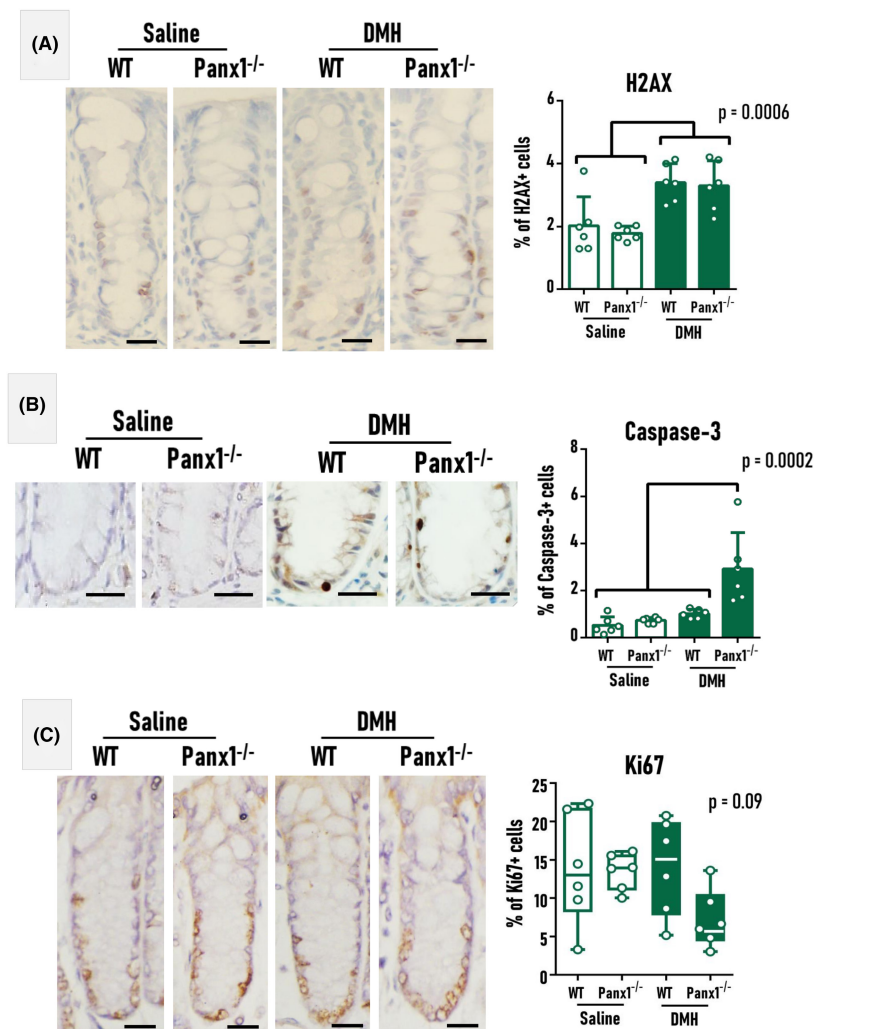
Parameters/Groups	Saline		DMH		p value
	WT	Panx1 <sup>-/-</sup>	WT	Panx1 <sup>-/-</sup>	
Weeks 0–7					
<i>n</i>	6	6	6	6	
Initial body weight (g)	18.10 ± 1.26	17.80 ± 0.61	17.85 ± 0.51	17.55 ± 0.95	.76
Final body weight (g)	19.48 ± 0.63	19.17 ± 0.86	20.83 ± 1.03	20.33 ± 0.78	.51
Absolute liver weight (g)	0.97 ± 0.10	1.05 ± 0.08	0.92 ± 0.11	1.05 ± 0.09	.07
Relative liver weight (%)	4.36 ± 0.30	4.90 ± 0.27	4.75 ± 0.23	5.18 ± 0.49	.055
Colon length (cm)	10.00 ± 0.56	9.66 ± 0.51	10.31 ± 0.76	9.17 ± 1.25	.13
Weeks 0–37					
<i>n</i>	12	12	12	12	
Initial body weight (g)	17.30 ± 0.92	17.33 ± 1.22	18.26 ± 0.69	17.96 ± 1.72	.15
Final body weight (g)	24.16 ± 2.65 ab	25.65 ± 2.09 a	23.50 ± 2.53 ab	22.03 ± 1.03 b	.0021
Absolute liver weight (g)	1.07 ± 0.19 a	1.07 ± 0.14 a	0.95 ± 0.14 a	0.73 ± 0.15 b	<.0001
Relative liver weight (%)	4.68 ± 0.60 a	4.20 ± 0.39 ab	4.03 ± 0.38 b	3.32 ± 0.68 c	<.0001
Colon length (cm)	9.95 ± 1.05	9.50 ± 0.95	9.93 ± 0.97	9.59 ± 0.63	.51

Note: WT: wild-type. *n* = number of mice/group/genotype. Data are presented as mean + standard deviation. Different letters correspond to statistical difference among groups by one-way ANOVA and a posteriori Tukey test ( $p < .05$ ). The *p* value in the tables refers to the ANOVA-derived analysis.





**FIGURE 2** Effects of pannexin 1 knockout (Panx1<sup>-/-</sup>) on (A) peripheral blood genotoxicity, (B) malondialdehyde (MDA) levels and antioxidant superoxide dismutase (SOD) and catalase activities in the colon during 1,2-dimethylhydrazine (DMH)-induced colon carcinogenesis in mice at week 7. WT: wild-type.  $n = 6$  mice/group/genotype. Data are presented as mean + standard deviation, and data points. Data were analysed by one-way ANOVA and a posteriori Tukey test ( $p < .05$ ). The  $p$  value in the graphs refers to the ANOVA-derived analysis.



**FIGURE 3** Effects of pannexin 1 knockout (Panx1<sup>-/-</sup>) on (A) colonocyte genotoxicity (H2AX), (B) crypt apoptosis (caspase-3), and (C) proliferation (Ki67) and during 1,2-dimethylhydrazine (DMH)-induced colon carcinogenesis in mice at week 7. Representative microscopic overviews (scale bar: 10  $\mu$ m) show that H2AX and Ki67 labelling comprised top, middle and/or bottom crypt compartments, while caspase-3 was restricted to bottom crypt compartment. WT: wild-type.  $n = 6$  mice/group/genotype. Data are presented as mean + standard deviation or box plots, and data points. Data were analysed by one-way ANOVA (H2AX and caspase-3) or Kruskal-Wallis (Ki-67) and a posteriori Tukey test ( $p < .05$ ). The  $p$  value in the graphs refers to the ANOVA-derived analysis.

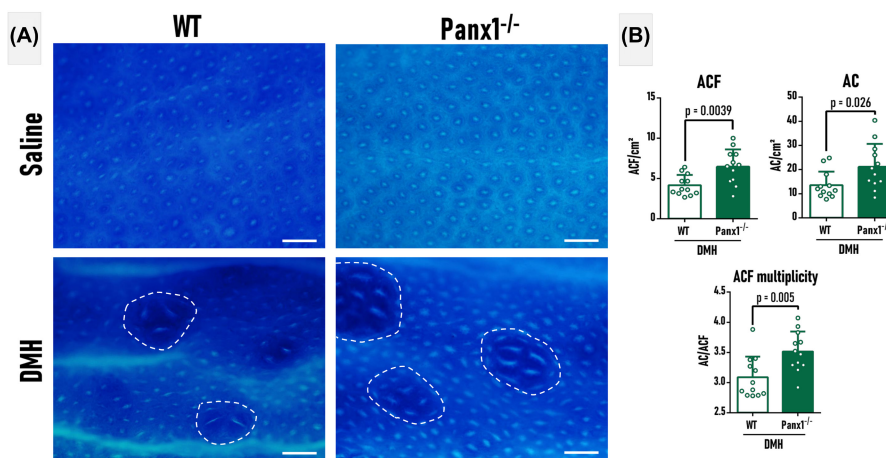
### 3.2 | Panx1 knockout promotes carcinogen-induced peripheral blood genotoxicity and lipid peroxidation in the colon at week 7

Four hours after the last carcinogen administration, we observed that Panx1 ablation increased the DMH-induced genotoxicity in nucleated peripheral blood cells, as inferred by comet assay ( $p < .0001$ ) (Figure 2A). Eight hours after the last carcinogen administration, although we found that Panx1 deletion resulted in no alterations in the DMH-mediated increase in the number of colonocytes positive for H2AX (Figure 3A) - an indicator of double-strand DNA breaks<sup>30</sup> - Panx1 knockout did result in higher DMH-induced MDA levels in the colon ( $p = .035$ ) (Figure 2B), which is an oxidative stress byproduct involved implicated in adduct formation, cytotoxicity and cell death.<sup>31</sup> These findings are in line

with the increase in cleaved caspase-3 immunolabelling index in the colonic crypts of DMH-exposure Panx1<sup>-/-</sup> mice ( $p = .0002$ ) (Figure 3B). The activities of SOD, and catalase (Figure 2B), and the proliferation of colonocytes (Ki67) (Figure 3C) remained unaltered by Panx1 genetic ablation.

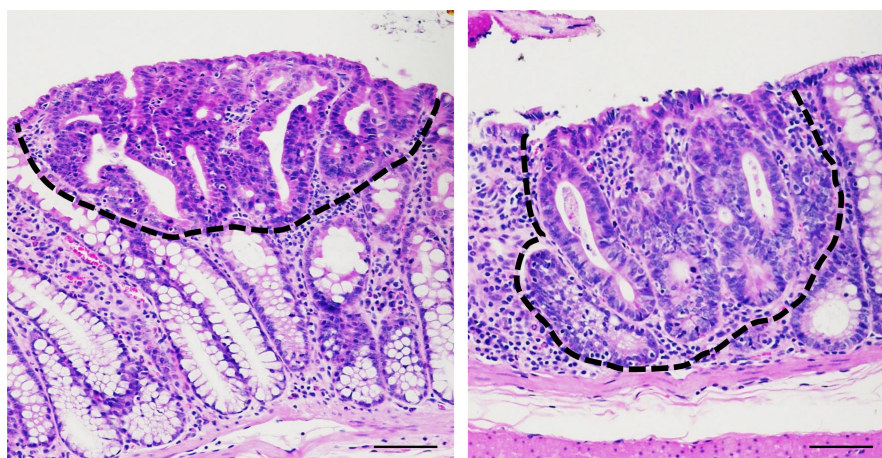
### 3.3 | Panx1 knockout increases preneoplastic ACF burden at week 37

Thirty weeks after the last carcinogen administration, Panx1<sup>-/-</sup> animals showed increased ACF ( $p = .0039$ ), AC ( $p = .026$ ) and ACF multiplicity ( $p = .005$ ) by 56%, 57% and 20%, respectively, compared to the WT counterpart (Figure 4). Representative microscopic overview of these lesions in haematoxylin and eosin (HE)-stained sections are also shown (Figure 5).



**FIGURE 4** Effects of pannexin 1 knockout (Panx1<sup>-/-</sup>) on aberrant crypt foci (ACF) burden during 1,2-dimethylhydrazine (DMH)-induced colon carcinogenesis in mice at week 37. (A) Representative photomicrographs of a whole mount of colon specimens stained with methylene blue (scale bar: 100 μm) - depicting the absence (non-initiated) or the occurrence of ACF (DMH-initiated mouse) (dotted lines). (B) Colon length, ACF, aberrant crypt (AC), and ACF multiplicity (AC/ACF) data in both genotypes.  $n = 12$  mice/group/genotype. WT: wild type. Data are presented as mean + standard deviation, and data points. Data were analysed by Student's  $t$ -test ( $p < .05$ ). The  $p$  value in the graphs refers to the ANOVA-derived analysis.

**FIGURE 5** Representative photomicrographs of dimethylhydrazine (DMH)-induced aberrant crypt foci (ACF) - featuring dysplastic phenotype - in haematoxylin and eosin (HE)-stained sections (dotted lines, scale bar: 50 μm).



## 4 | DISCUSSION

In the present investigation, we examined the modifying effects of Panx1 global ablation on the early events involved in chemically induced colon carcinogenesis in mouse, evaluating both (A) the sub-acute toxicity outcomes and (B) preneoplastic lesion burden. In brief, regarding the sub-acute effects, the genetic removal of Panx1 resulted in heightened genotoxicity induced by DMH in peripheral blood cells. It also led to elevated levels of MDA in the colon and increased apoptosis (cleaved caspase-3) in colonic crypts. Importantly, at the later timepoint, animals lacking Panx1 exhibited a decreased body weight and showed a 56% increase in mean number of aberrant ACF and a 20% increase in ACF multiplicity. Of note, the Panx1 KO did not exert any noticeable impact on all the measured parameters in mice treated with saline. Overall, our findings suggest that the genetic elimination of Panx1 fosters the formation of preneoplastic aberrant crypt foci (ACF) during chemically induced colon carcinogenesis in mice. Conversely, our observations propose a protective role of Panx1 in the early steps of this process. The genetically ablated Panx1 mice were used since these animals are functional knockouts that express a truncated form of active Panx1 subunits.<sup>21</sup> This model is considered a useful tool for biomedical research, including the investigation of Panx1 roles on gastrointestinal diseases.<sup>32–34</sup>

DMH is an alkylating colon-specific complete procarcinogen. DMH is first metabolically activated by hepatic cytochrome P450 (CYP) enzymes, primarily CYP2E1 and CYP2A6, through N-demethylation. This, methyl diazonium, methyl ions and other reactive species are generated, reaching the colon via the bloodstream.<sup>8</sup> In the colon, DMH metabolites alkylate colonocytes, “initiating” them for colon carcinogenesis.<sup>8,23</sup> Furthermore, other secondary reactive metabolites may also be generated as DMH-derived radicals that react with membrane lipids, as MDA, which is a known marker of oxidative stress/lipid peroxidation.<sup>9,31</sup> MDA is formed through the process of lipid peroxidation. Lipid peroxidation occurs when reactive oxygen species (ROS), such as free radicals, attack and react with polyunsaturated fatty acids (PUFAs) present in cell membranes.<sup>31</sup> When evaluating the sub-acute toxicity outcomes at week 7, we observed that Panx1 deletion promoted the carcinogen-induced systemic genotoxicity, also increasing MDA levels in whole colon samples and the number of caspase-3 positive colonic epithelial cells. MDA is a highly reactive and cytotoxic molecule. It can form adducts with proteins, DNA, and other biomolecules, leading to cellular damage, dysfunction and death. Thus, at early this timepoint, the increased apoptosis in

epithelial colonic cells observed in Panx1 deletion is probably a response to the increased MDA levels. It is also important to point out that the increase in lipid peroxidation is clearly implicated on CC pathogenesis in humans and in preclinical models.<sup>35,36</sup> Panx1 hemichannels and oxidative stress have an intimate bidirectional relationship. Reactive species may negatively regulate Panx1 hemichannel activity, as nitric oxide (NO)-induced structural modification of Panx1 protein resulted in the inhibition of the hemichannel in human embryonic kidney cells.<sup>37</sup> On the other hand, the Panx1-mediated release of ATP activated P2Y/P2X receptors, subsequent G proteins, triggering the accumulation of NADPH oxidases-generated reactive species *in vitro*.<sup>38</sup> Although our findings may sound contrasting, most of the available findings mechanistically linking Panx1 and oxidative stress are restricted to *in vitro* bioassays in physiological contexts. Thus, further molecular investigations on Panx1 KO in the multi-stage colon carcinogenesis context are necessary.

At week 37, we observed that animals lacking Panx1 had decreased body and liver weights, and increased preneoplastic ACF burden. As diminished body and liver weights are known effects of DMH upon multiple administrations,<sup>39,40</sup> the findings indicate that Panx1 knockout may predispose mice to these deleterious toxicological outcomes. As ACF are the earliest morphological manifestation of colon carcinogenesis, the modulation of these classical parameters is predictive of the preventive or predisposing interventions. In both preclinical and clinical investigations, the occurrence of ACF positively correlates with colon tumorigenesis.<sup>6,7</sup> Thus, the findings indicate that Panx1 knockout may promote the early stages of DMH-induced colon carcinogenesis. The promotion of lipid peroxidation by Panx1<sup>-/-</sup> at week 7 is proposed to be partly involved to the enhanced ACF burden at week 37, although other predisposing mechanisms should be considered regardless of carcinogen-related ones. In general, recent findings from human tumours and several tumour-derived cell lines indicate that high Panx1 expression enables survival advantages to tumours, increasing their potential for progression.<sup>16–19,41</sup> Nonetheless, the attenuation of Panx1 does not have a common outcome in all cancers. In human oral cancer cells, the microRNA-mediated Panx1 downregulation is proposed as a chemotherapy resistance mechanism, as the inhibition of Panx1 hemichannels may negatively interfere with ATP discharge that contributes to chemotherapy-induced tumour cell apoptosis.<sup>42</sup> Panx1 is expressed in immune cells that infiltrate tumours and promote antitumor immunity.<sup>43</sup> In a certain way our findings are in line with Derangère et al.,<sup>20</sup> in the sense that Panx1 may have a tumour suppressor role of Panx1 in CC cells, although this study addresses



the progression (malignant) stage of colon carcinogenesis and not the preneoplastic step. Differently from these findings, on which Panx1 was associated with a pyroptosis mechanism by the activation of LXR $\beta$ /Panx1/NLRP3/caspase-1-pathway in CC cells, our increased caspase-3 labelling may be a response to the enhanced oxidative stress right after carcinogen exposure. It is noteworthy that preneoplastic stages – as ACF – are usually underappreciated in the available bioassays. As Panx1 (and hemichannels) may have a complex interplay with a myriad of cellular and non-cellular components of tumour microenvironment, the preventive, suppressive or promoting outcomes should consider late stage of disease, and others tumour types. Considering that saline-treated Panx1 ablated mice did not developed ACF, and did not spontaneously develop tumours in previous assays,<sup>41</sup> the suppressive role of this protein in carcinogenesis should not be overestimated as well.

In essence, our findings innovatively indicate that Panx1 genetic ablation promotes preneoplastic ACF development during early stage of chemically-induced mouse colon carcinogenesis. A protective role of Panx1 on colon carcinogenesis is postulated, although further investigation is required in late stages of this disease.

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#### CONFLICT OF INTEREST STATEMENT

The authors declare no conflict of interest.

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