



## p38 $\gamma$ and p38 $\delta$ as biomarkers in the interplay of colon cancer and inflammatory bowel diseases

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*Total number of authors:*  
16

*Published in:*  
Cancer Communications

*Link to article, DOI:*  
[10.1002/cac2.12331](https://doi.org/10.1002/cac2.12331)

*Publication date:*  
2022

*Document Version*  
Publisher's PDF, also known as Version of record

[Link back to DTU Orbit](#)

### *Citation (APA):*

Fajardo, P., Taskova, M., Martín-Serrano, M. A., Hansen, J., Slott, S., Jakobsen, A. K., Wibom, M-L., Salegi, B., Muñoz, A., Barbachano, A., Sharma, A., Gubatan, J. M., Habtezion, A., Sanz-Ezquerro, J. J., Astakhova, K., & Cuenda, A. (2022). p38 $\gamma$  and p38 $\delta$  as biomarkers in the interplay of colon cancer and inflammatory bowel diseases. *Cancer Communications*, 42, 897-901. <https://doi.org/10.1002/cac2.12331>

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## LETTER TO THE EDITOR

# p38 $\gamma$ and p38 $\delta$ as biomarkers in the interplay of colon cancer and inflammatory bowel diseases

Dear Editor,

Colorectal cancer (CRC) is the second leading cause of cancer death according to the World Health Organization. Patients with inflammatory bowel disease (IBD), ulcerative colitis (UC) or Crohn's disease (CD) are at increased risk of developing colitis-associated CRC (CAC) [1]; however, our understanding of the inflammation-cancer interplay at the molecular level is still limited.

p38 mitogen-activated protein kinases (MAPKs) (p38 $\alpha$ , p38 $\beta$ , p38 $\gamma$  and p38 $\delta$ ) modulate the inflammatory response which contribute to CAC development [2]. p38 $\alpha$  (called p38) has been a therapeutic target in CD treatment, but these treatments have not progressed beyond phase I/II clinical trials [2]. Interestingly, the less studied p38 $\gamma$  and p38 $\delta$  (p38 $\gamma$ /p38 $\delta$ ) are essential in the immune response and regulate inflammatory molecule production [2–6]. Moreover, the importance of p38 $\gamma$ /p38 $\delta$  in CRC development has been established in vitro and in an azoxymethane-dextran sodium sulfate (AOM-DSS)-induced CAC model using p38 $\gamma$ /p38 $\delta$ -knockout mice [2, 5]. However, their implication in human CRC and IBD is not yet well defined.

Here, we report that p38 $\delta$  mRNA and protein levels were markedly decreased, whereas p38 $\gamma$  phosphorylation was increased, in AOM-DSS-treated wild-type (WT) mice (Figure 1A, Supplementary Figure S1). Additionally, we performed a comparative analysis of p38 $\gamma$ /p38 $\delta$  mRNA expression in human CRC samples, using The Cancer Genome Atlas (TCGA) data and Gene Expression Omnibus (GEO) dataset (GSE39582). p38 $\gamma$  expression was significantly increased in primary colorectal tumors compared to normal colon tissues, whereas p38 $\delta$  was markedly decreased (Figure 1B, Supplementary Figure S2A–B). High p38 $\gamma$  mRNA expression, but not p38 $\delta$ , was associated with shorter overall survival of CRC patients

(Figure 1C). p38 $\gamma$ /p38 $\delta$  gene expression in CRC samples was not affected by patient gender or tumor stage (Supplementary Figure S2C–D). Additionally, p38 $\gamma$ /p38 $\delta$  genes showed no systematic loss-of-function mutations (Supplementary Figure S2E), in agreement with findings in other cancers [7].

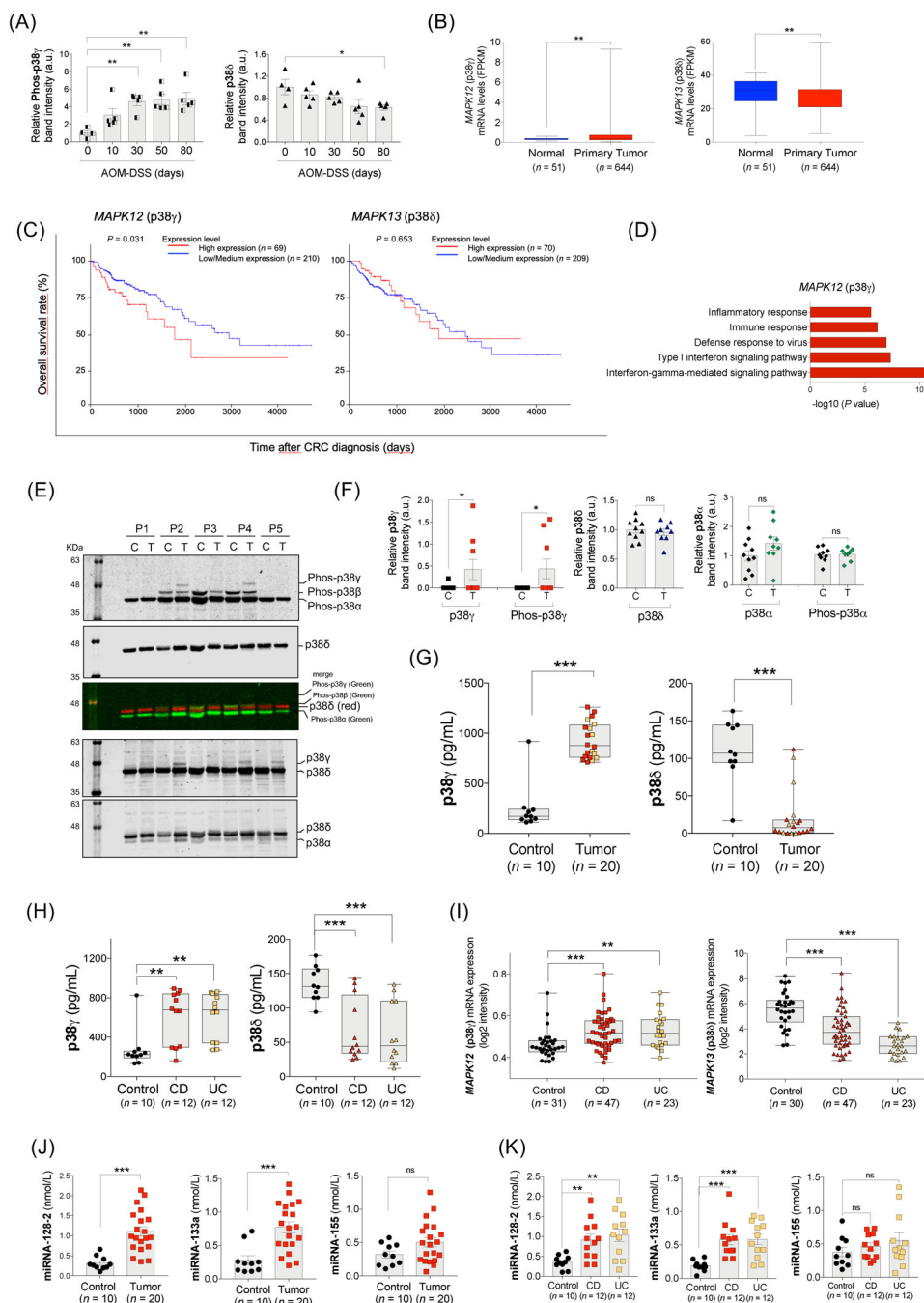
To understand the role of p38 $\gamma$ /p38 $\delta$  in CRC, we analyzed the correlation of their expression with other genes. Using the UALCAN portal and Gene Ontology (GO) enrichment analysis, we found that p38 $\gamma$  was positively correlated with genes whose biological functions are related to the immune and inflammatory response (Figure 1D, Supplementary Figure S3A–C, Supplementary Tables S1–S4). p38 $\gamma$  expression was positively correlated with matrix metalloproteinase 9 (*MMP9*), tissue inhibitor of metalloproteinase 1 (*TIMP1*) and interleukin-6 (*IL-6*) (Supplementary Figure S3D), which were overexpressed in CRC (Supplementary Figure S3E). *IL-6* is a marker of colitis that may precede CRC development. These results suggest that p38 $\gamma$  plays a role in the inflammatory tumor microenvironment which is associated to tumor formation and progression. This agrees with the observation that p38 $\gamma$  is activated in response to inflammatory stimuli, leading to the production of mediators, such as cytokines and other genes, which regulate cell proliferation, death or transformation [2–4]. On the other hand, the expression of p38 $\delta$  mRNA in CRC was positively correlated with genes involved in cell division and RNA/DNA processing, and negatively with *MMP9* and *TIMP1* (Supplementary Figure S3F–J).

There are few databases containing comparative studies of protein expression in tumor versus normal tissue, and those available did not analyze p38 $\gamma$ /p38 $\delta$ . However, in the Human Protein Atlas portal, we found immunohistochemical staining of p38 $\gamma$ /p38 $\delta$  in CRC patients' tumor sections (Supplementary Figure S4A). To further determine p38 $\gamma$ /p38 $\delta$  protein expression and phosphorylation (which indicates protein activation) in CRC, we compared human colon organoids derived from CRC and normal colon tissues (Supplementary Table S5) by Western blotting

**Abbreviations:** AOM, azoxymethane; CAC, colitis-associated colorectal cancer; CD, Crohn's disease; CRC, colorectal cancer; DSS, dextran sodium sulfate; IBD, inflammatory bowel disease; MAPK, mitogen-activated protein kinase; miR, microRNAs; p38 $\gamma$ /p38 $\delta$ , p38 $\gamma$  and p38 $\delta$ ; UC, ulcerative colitis; WT, Wild type.

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**FIGURE 1** p38 $\gamma$  and p38 $\delta$  regulation in CRC and IBD. **(A)** WT mice were treated with AOM and DSS as described in Supplementary Figure S1A. Colon extracts (50  $\mu$ g) from AOM-DSS-treated WT mice were immunoblotted with antibodies to phosphorylated p38MAPKs and total p38 $\alpha$ , p38 $\gamma$  and p38 $\delta$ , as shown in Supplementary Figure S1D. Phospho-p38 $\gamma$  and p38 $\delta$  bands were analyzed and quantified. **(B)** *MAPK12* (p38 $\gamma$ ) and *MAPK13* (p38 $\delta$ ) mRNA expression levels in normal and primary CRC tumor tissues from the TCGA database. **(C)** Survival analysis (Kaplan-Meier plot with log-rank test) of CRC patients from the TCGA dataset (in UALCAN portal) stratified by *MAPK12* (p38 $\gamma$ ) and *MAPK13* (p38 $\delta$ ) mRNA expression levels. **(D)** Enrichment analysis of GO biological processes of the genes whose expression positively correlates with *MAPK12* (p38 $\gamma$ ). **(E)** Organoid lysates (20  $\mu$ g) from tumors (T) or normal tissues (C) from CRC patients (P1-5) were immunoblotted sequentially with the following antibodies, from top to bottom: 1) anti-phos-p38MAPK: secondary anti-rabbit green (this antibody recognizes all phospho-p38MAPK isoforms); 2) anti-p38 $\delta$ : secondary anti-sheep red (merge image of these two blots is shown in the figure); 3) anti-p38 $\gamma$ : secondary anti-sheep red (total p38 $\delta$  can be also visualized), and 4) anti-p38 $\alpha$ : secondary anti-rabbit red (total p38 $\delta$  can be also visualized). Blots were analyzed using the Odyssey infrared imaging system. The identity of the isoforms was determined, as in Supplementary Figure S1E-F, by their different mobility in the SDS-PAGE gel (to distinguish p38 $\gamma$  from p38 $\alpha$ , p38 $\delta$  and p38 $\beta$ ) and by using simultaneously specific antibodies for a particular p38MAPK isoform (in red) and anti-phospho-p38MAPK antibody (in green). After analysis of the blots in the Odyssey infrared imaging system, phospho-p38MAPK bands should be yellow when colors were merged. **(F)** Total and phospho-protein

(Figure 1E, Supplementary Figure S4B). Quantitative analysis revealed that p38 $\gamma$  was expressed and phosphorylated in 30% of the tumors examined, while p38 $\delta$  expression was similar in tumor and normal tissues and was not phosphorylated (Figure 1F). These data indicate that protein expression and activation of p38 $\gamma$ , but not p38 $\delta$ , is upregulated in CRC patients. An alternative method to evaluate protein expression changes in CRC is a blood-based test. We then measured p38 $\gamma$ /p38 $\delta$  protein levels, as well as p38MAPK phosphorylation, in plasma samples from healthy donors and CRC patients (Supplementary Table S6) by enzyme-linked immunosorbent assay (ELISA). We found that p38 $\gamma$  levels in plasma samples from CRC patients were markedly increased compared with healthy donors (Figure 1G), which was associated with increased levels of the inflammatory molecules IL-6, C-C motif chemokine ligand 5 (CCL5), tumor necrosis factor  $\alpha$  (TNF $\alpha$ ), Interferon stimulated gene 15 (ISG15) and anti-neutrophil cytoplasmic antibodies (ANCA) (Supplementary Figure S4C-D). In contrast, p38 $\delta$  levels were significantly decreased in plasma samples from CRC patients (Figure 1G). Phosphorylated p38MAPKs were decreased in CRC samples (Supplementary Figure S4E). These data agree with those observed for the p38 $\gamma$ /p38 $\delta$  mRNA expression in CRC patients and those found for p38 $\gamma$  in human tissue-derived organoids. The minimal changes in p38 $\delta$  protein observed in organoids could be due to that, although organoids replicate characteristics of tumor source, they do not contain non-tumor cells that could affect p38 $\delta$  expression, accounting for the difference observed.

As all our data indicated that p38 $\gamma$ /p38 $\delta$ , particularly p38 $\gamma$ , played an essential role in colon inflammation and since CRC is associated to or can be caused by chronic colon inflammation, we examined the levels of p38 $\gamma$ /p38 $\delta$  in plasma samples from IBD (UC and CD) patients (Supplementary Table S7). p38 $\gamma$  protein levels in the plasma

samples from IBD patients were markedly increased compared to healthy donors, while the amounts of p38 $\delta$  and phosphorylated p38MAPKs were decreased (Figure 1H, Supplementary Figure S4F). We also assessed the mRNA expression of p38 $\gamma$ /p38 $\delta$  in IBD patients versus healthy controls using the GEO datasets (GSE179285; GSE166925) [8]. p38 $\gamma$  mRNA expression was significantly increased in UC and CD compared to healthy controls, whereas p38 $\delta$  was markedly decreased in UC and CD (Figure 1I, Supplementary Figure S4G). These results indicate that increased levels of p38 $\gamma$  protein in plasma and p38 $\gamma$  mRNA in colon samples as well as decreased levels of p38 $\delta$  mRNA may be good indicators of both CRC and IBD. These data and previous findings strongly suggest the p38 $\gamma$ 's role in linking inflammation and tumor development.

p38 $\gamma$ /p38 $\delta$  have a pro-oncogenic function in CRC by regulating the production of inflammatory molecules and miRNAs [2, 5, 9, 10]. We then examined their effect on the levels of miR-128-2, miR133a and miR-155, which are implicated in inflammation and cancer development [9, 10] and are detectable in plasma samples. miR-128-2 and miR133a were regulated by p38 $\gamma$  and p38 $\delta$  in a mouse model of colitis and CAC (Supplementary Figure S5A-D). In the plasma samples from IBD and CRC patients, miR128-2 and miR133a, but not miR-155, were increased compared to healthy donors (Figure 1J-K); however, only miR128-2 was correlated with p38 $\gamma$ /p38 $\delta$  expression levels (Supplementary Figure S5E-G). These results indicate a link between the expression of p38 $\gamma$ /p38 $\delta$  and molecules implicated in inflammation and cancer development, reinforcing the described interplay of CRC and colitis (Supplementary Figure S6).

This study showed that p38 $\gamma$ /p38 $\delta$  protein could be detected in human plasma. Given the high mortality of colon cancer, the easy access to early risk assessment would contribute to improved survival rates. Our work

bands of all organoid samples (9 organoids derived from tumor tissues and 10 from normal tissues) shown in panel (E) and Supplementary Figure S4B were quantified using ImageJ. As a control, p38 $\alpha$  expression and phosphorylation were examined, and they were similar in organoids from normal and tumor tissues. (G) p38 $\gamma$  and p38 $\delta$  expression in plasma samples from 20 colon cancer patients and 10 healthy controls were measured by ELISA. Black dots represent the data of healthy donors; red symbols represent the data of CRC patients without IBD; yellow symbols represent the data of CRC patients with IBD. (H) p38 $\gamma$  and p38 $\delta$  protein expression in the plasma samples from IBD (CD and UC) patients were measured by ELISA. (I) *MAPK12* (p38 $\gamma$ ) and *MAPK13* (p38 $\delta$ ) mRNA expression in IBD (UC and CD) patients and healthy controls using data from GEO (GSE179285). Colon tissues were collected from patients with UC; colon and ileum tissues were collected from patients with CD and healthy donors. (J, K) Expression of miRNAs in the plasma samples from colon cancer (J) and colitis patients (K). miRNA-128-2, miRNA-133a and miRNA-155 were measured by FLEET CTD analysis in plasma samples from 20 healthy donors (10 for CRC and 10 for IBD), 20 colon cancer patients, 12 UC and 12 CD patients. Data are shown as mean  $\pm$  SEM. Student's *t*-test was used for comparison. ns, not significant; \**P*  $\leq$  0.05; \*\**P*  $\leq$  0.01; \*\*\**P*  $\leq$  0.001. Abbreviations: AOM: azoxymethane; CD: Crohn's disease; CRC: colorectal cancer; DSS: dextran sodium sulfate; FPKM: fragments per kilobase of transcript per million mapped reads; ELISA: enzyme-linked immunosorbent assay; GEO: gene expression omnibus; GO: gene ontology; IBD: inflammatory bowel disease; MAPK: mitogen-activated protein kinase; miRNA: microRNAs; p38 $\gamma$ /p38 $\delta$ : p38 $\gamma$  and p38 $\delta$ ; SDS-PAGE: sodium dodecyl sulfate-polyacrylamide gel electrophoresis; SEM: standard error of the mean; TCGA: The Cancer Genome Atlas; UC: ulcerative colitis; WT: Wild type; au, arbitrary units

indicates that p38 $\gamma$ /p38 $\delta$  could be useful biomarkers for CRC/IBD diagnostic blood-based tests for colitis and early-stage CRC. Additionally, our results support the role of p38 $\gamma$ /p38 $\delta$  in promoting tumor growth and give further evidence that one of the mechanisms by which p38 $\gamma$  promotes tumorigenesis is linked to its elevated protein expression and activation, thus increasing inflammation and providing an inflamed tumor environment that would promote tumor growth.

## DECLARATIONS

## ACKNOWLEDGMENTS

We thank our colleagues Tirso Pons and Lorena Madraner for informatic support. We thank Prof. MD Elizabeth D Mellins, Stanford University School of Medicine and Prof. N. Heegaard, SSI, Denmark, for providing us with healthy human controls for this study. We acknowledge S. Khatri, Technical University of Denmark, for technical assistance with performing ELISA and bead-based assay. We want to particularly acknowledge the patients and the Biobank IdiPAZ (PT20/00004) from the Spanish National Biobanks Network.

## CONFLICTS OF INTEREST

The authors declare no conflict of interest.

## FUNDING

This research was funded by the MCIN/AEI/10.13039/501100011033 (PID2019-108349RB-I00 and SAF2016-79792R) to AC and JJSE; Villum Foundation, grant no. 13152 to KA; by Agencia Estatal de Investigación (PID2019-104867RB-I00/AEI/10.13039/501100011033) and the Instituto de Salud Carlos III- Fondo Europeo de Desarrollo Regional (CIBERONC/CB16/12/00273 and ICI20/00057) to AM and AB. PF received MCIN FPI fellowship (BES-2017-080139).

## AUTHOR CONTRIBUTIONS

AC, KA and MT conceived and planned the experiments. PF, MT, MAMS, BS and AB carried out the experiments. PF, MT, MAMS, JH, SS, AM, AB, AS, JJSE, AH, JMG and AC contributed to sample collection and preparation. PF, MT, AKJ, MLW, JJSE, KA and AC contributed to the analysis and interpretation of the results. MT, AKJ, MLW, KA and AC contributed to writing the original draft; AC writing-Review and editing. All authors provided critical feedback, helped shape the research, analysis and manuscript, and agreed to the published version of the manuscript.

## ETHICS APPROVAL AND CONSENT TO PARTICIPATE

De-identified human colitis plasma samples were obtained from Stanford University (USA) under Stanford Institutional Review Board (IRB protocol 28427). Human colon cancer samples were provided by Danish Biobank RBGB Herlev, under EVK number 21015520. All the samples were fully anonymized prior to analyses. Samples from patients used for organoid cultures were provided by the Biobank IdiPAZ (PT20/00004) with the approval of the Ethics and Scientific Committees of La Paz University Hospital (HULP-PI-1425 and HULP-PI-3169). Normal tissue samples were obtained from the CRC-adjacent non-tumoral tissue. Informed consent was obtained from all participants. Mouse studies were conducted in specific pathogen-free conditions in accordance with European Union regulations and the regulation of local Centro Nacional de Biotecnología and Consejo Superior de Investigaciones Científicas bioethics committees and were approved by the Community of Madrid PROEX316/15 and PROEX289.2/20.

## CONSENT FOR PUBLICATION

Not applicable.

## DATA AVAILABILITY STATEMENT

Not applicable.

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