


## Original Research

# The gene expression alterations in chronic hypoxic PANC-1 pancreatic cancer cell line

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### Abstract

**Aim:** Cancer cells divide excessively, resulting in overpopulation and hypoxia. Tumor hypoxia drives tumor development and therapeutic resistance. Hypoxia dominates pancreatic tumor microenvironments. This study aimed to ascertain alterations in gene expression linked to chronic hypoxia in the pancreatic cancer cell (PANC-1) line. **Methods:** PANC-1 had eight-hour hypoxic events with oxygen levels below 1%. 40 episodes were exposed three times a week. Real-time PCR arrays were used to analyze gene expression changes. This investigation compared cells treated with 20 and 40 hypoxia episodes to normoxic cells. MTT cell proliferation assays assessed hypoxic cell doxorubicin resistance. Wound-healing assays measured cell migration. 20 and 40 hypoxia exposures altered gene expression patterns significantly. **Results:** No alterations were observed in either stage, as most genes exhibited a resurgence after 40 episodes. Following exposure to 20 episodes of hypoxia, the expression levels of the genes HIF1AN, HMOX-1, and PKM were significantly increased by factors of 6.9, 4.5, and 3.4, respectively. The IC50 value of Doxorubicin in PANC-1 cells exhibited a 2.7-fold increase and an approximately 3.6-fold increase after 20 and 40 episodes, respectively, compared to normoxic cells. This study demonstrates that subjecting cells to extended durations of hypoxia results in distinct alterations in gene expression compared to those induced by short-term hypoxia, specifically lasting less than 72 hours. Furthermore, it sanctioned the facilitation of chemotherapy resistance through prolonged exposure. **Conclusions:** The study suggests that HIF1AN, HMOX-1, and PKM may serve as promising biomarkers for hypoxia in pancreatic cancer and contributors to the cellular response to prolonged hypoxia.

**Keywords:** Cancer, Cell Line, Gene Expression, Hyoxia, PANC-1

## INTRODUCTION

Pancreatic cancer is presently regarded as one of the most difficult forms of human malignancies. It continues to pose a significant unresolved public health challenge in the early years of the 21st century. Despite substantial reductions in cancer-related mortality in recent years, the prognosis for pancreatic

cancer has remained persistently unfavorable<sup>1</sup>. Pancreatic ductal adenocarcinoma (PDAC) constitutes the predominant form of pancreatic cancer, comprising approximately 85-90% of all pancreatic tumor cases<sup>2</sup>.

The PDAC is classified as the fourth most prevalent cause of mortality associated with cancer in the United States<sup>2</sup>. This malignancy is recognized as one of the most lethal among human diseases, exhibiting a mortality rate approaching nearly 100% within a short time<sup>2</sup>. It is one of the deadliest human malignancies, with a high mortality rate approaching 100%<sup>3</sup>. Pancreatic cancer is one of top ten leading cause of cancer death among Jordanian, accounting for 6.2 % of cancer deaths in 2012<sup>4</sup>.

Currently, surgery is the only curative treatment for PDAC. Nevertheless, only a minority of PDAC patients are considered surgical candidates due to late-stage diagnosis common with this disease<sup>5</sup>. Chemotherapy is a standard treatment option in managing all stages of pancreatic cancer. In 1997, Burris et al. conducted a study to evaluate the efficacy of gemcitabine as the primary treatment option for patients with advanced, unresectable pancreatic cancer<sup>6</sup>. The use of gemcitabine (Gemzar<sup>®</sup>) or the combinations of gemcitabine with other cancer drugs, such as cisplatin, or erlotinib, has confirmed efficacy in treating pancreatic cancer<sup>7</sup>. The potential role of radiotherapy as a component of adjuvant therapy in the postoperative regimen remains debatable<sup>8</sup>. Pancreatic cancer is well recognized as a very challenging disease on many fronts. It is one of the major aggressive human malignancies, clinically marked by local invasion, early metastasis, and resistance to

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standard chemotherapy<sup>1</sup>. Hypoxia is a characteristic feature of solid tumors and define as a reduction in the normal level of tissue oxygen tension<sup>9</sup>. Cells go through genetic and adaptive alterations, which permit such cells to survive and proliferate in a hypoxic environment. Tumor hypoxia has been recognized as a potential therapeutic problem because it causes resistance to chemo- and radiotherapy and promotes a more aggressive phenotype, increasing mutation rates, invasion, and metastasis<sup>10-14</sup>.

Genetic changes in the aforementioned observations have not yet been completely agreed<sup>15</sup>. Therefore, analyzing the modulated genes induced by hypoxia in cancer is greatly beneficial<sup>16</sup>. This study was designed to simulate actual hypoxic conditions in order to investigate the alterations in gene expression that occur in pancreatic cancer cells during prolonged exposure to hypoxia. The primary objective of this study is to elucidate novel biomarkers associated with tumor hypoxia. Our objective is to investigate the correlation between hypoxia and chemo-resistance and identify potential underlying mechanisms for this phenomenon.

## MATERIALS AND METHODS

### Cell culture

The Panc-1 human pancreatic cancer cell lines were purchased from the American Type Culture Collection (ATCC; USA), The Panc-1 cells were kept as an attached monolayer culture in the DMEM High Glucose (EuroClone, Via Figino, Italy), supplemented with 10% (v/v) heat-inactivated fetal bovine serum (FBS) (Biowest, South America), 2 mM L-glutamine, and 100 U/mL and 100 µg/mL penicillin– streptomycin (EuroClone, Italy). The cells were grown on 75 cm<sup>2</sup> attached types, filter-cap culture flasks (Membrane Solutions, USA). Exposure to hypoxia PANC-1 cells were exposed to an 8 hours of hypoxia (0.1% oxygen), three times a week for a total of 40 hypoxic episodes (continuous hypoxic PANC-1 cells). The experimental setup involved utilizing the AnaeroGen Compact system (Oxoid, UK), designed to generate an anaerobic atmosphere. This system was employed to create hypoxic conditions in the experimental environment<sup>17</sup>. The vented culture flasks were positioned within the plastic bags, while the paper sachets were opened and placed inside the bag. Subsequently, the bag was sealed using the plastic zip closure.

Throughout a five-month duration. In addition to administering hypoxic shots, the cells were subjected to incubation alongside their normoxic counterparts under normoxic conditions. The normoxia-exposed PANC-1 cells served as the control group.

### Cell proliferation assay

7000 cells were seeded into coated 96-well plate (Greiner, Germany). Each of the hypoxic and normoxic cell line was seeded in triplicates in plain medium for at least 18 h. Then the media were aspirated from the wells. Then, gradually decreasing concentrations between 0.01 µM and 500 µM of Doxorubicin were added in fresh media. The rest of assay was done according to<sup>14</sup>. The MTT proliferation assay

was done after the 20<sup>th</sup>, and 40<sup>th</sup> for hypoxic shots. As well, the MTT proliferation assay was done for the control PANC-1 cells which incubated under normoxic condition.

### Wound Healing Experiment

A total of 50,000 cells were seeded in each well of a 6-well plate, originating from three different sources: control Panc-1 cells, Panc-1 cells after 20 hypoxic shots, and Panc-1 cells after 40 hypoxic shots. The cells were cultured until they reached confluence. A simulated wound was generated by a linear abrasion using a yellow pipette tip. To limit the width of the scratch, a deliberate angle of approximately 30 degrees was maintained while manipulating the pipette tip. Subsequently, the media underwent a process of alteration and cleansing utilizing a phosphate buffer solution in order to eliminate the deceased cellular entities. The experiment was conducted in a hypoxic environment with a 1% oxygen concentration. Photographs were captured at three distinct time intervals: 0 hours, 24 hours, and 48 hours.

### RNA isolation and Real-time PCR

The RNA was extracted utilizing the RNeasy® Mini kit (Qiagen, Germany) in accordance with the instructions provided by the manufacturer.

The impact of hypoxia on the gene expression of the PANC-1 pancreatic cancer cell line was investigated utilizing a 96-well PCR array, specifically the RT2 Profiler PCR array (PAHS-032Z, Human Hypoxia Signaling Pathway PCR Array, Qiagen, USA). The array consists of 96-well plates containing a selection of primers for 84 genes identified as responsive to low oxygen levels, along with an additional set of 12 genes used for quality control purposes. The cDNA resulting from the First Strand Kit® reaction was mixed with RT2 SYBR® green master mix (Qiagen, USA) and nuclease free water (Biobasic, USA). Then, 20µl of the mix was placed in every well and the plate was subsequently centrifuged (Hettich, Germany) at 1000g for 1 minute to remove any air bubbles that could affect the results. CFX (Bio-Rad, USA) thermo-cycler performed the Real-time PCR; the program consisted of 95 °C for 10 min, followed by 40 cycles of 95 °C for 15.

## RESULTS

### Effect of hypoxia on PANC-1 cell line resistance to Doxorubicin

The investigation was conducted to verify the emergence of the hypoxic phenotype by assessing the resistance to Doxorubicin, a frequently employed chemotherapeutic agent. The IC<sub>50</sub>, which represents the concentration at which half of the maximal inhibitory effect is observed, was determined using the MTT cell proliferation assay, as described in the Materials and Methods section. The IC<sub>50</sub> value exhibited a 2.7-fold increase following 20 instances of hypoxic episodes in comparison to the normoxic cells. Following a series of 40 hypoxic episodes, the cells exhibited an approximately 3.6-fold increase in their IC<sub>50</sub> values when compared to the normoxic cells. Table 1 displays the IC<sub>50</sub> values of the cells measured at different time points.



**Table 1.** Half maximal inhibitory concentrations (IC50) of Doxorubicin in hypoxic and normoxic cells measured at different stages of the experiment.

Cell Lines	Control Cells (H0) IC50 in nM(STD)	Hypoxic Cells(H--) IC50 in nM (STD)	Hypoxic Cells IC50/ Control Cells IC50 (fold)
H20/comparable age H0	344 (± 0.034)	934(± 0.014)	2.7 (INCREASE )
H40/comparable age H0	369 (±0.011)	1244(± 0.07)	3.6 (INCREASE)

Gene Expression Changes in H20-PANC-1 Cancer Cell Line

The changes in the gene expression of PANC-1 cells that were induced by hypoxia could be potentially used as biomarkers for hypoxia in pancreatic cells. RT-PCR was performed using a hypoxia-signaling pathway PCR array. An arbitrary cut-off point of 2 folds was chosen so that the changes identified would be substantial. When comparing cells exposed to 20 episodes of 8-hour hypoxia and similar passage-age normoxic cells, it was observed that three genes exhibited significant down-regulation, as indicated in Table 2. Conversely, eleven genes displayed significant up-regulation, as demonstrated in Table 3.

The alterations in gene expression in PANC-1 cells, which were subjected to 40 episodes of 8-hour hypoxia, were assessed using PCR. After conducting a comparison between the H40-PANC-1 pancreatic cancer cells and the H0-PANC-1 cells, it was observed that five genes exhibited down-regulation, while three genes showed up-regulation. This information is

presented in Table 4 and Table 5, respectively.

**Wound healing results**

The utilization of the in vitro wound healing assay to monitor the migratory behavior of PANC-1 cells at the forefront of the scratch is exemplified in Figure 1. The photographs taken at the onset and conclusion of a 48-hour incubation period exhibited notable cellular migration towards the scratch. The photographs captured at the initial time point of 0 hours can be juxtaposed with the concluding photographs taken at 48 hours to ascertain the extent of closure for each individual scratch. The analysis of the images revealed that the control PANC-1 cells (located on the left side) and H40 PANC-1 cells (located on the right side) exhibited a slower migration rate in comparison to the H20 PANC-1 cells, where the scratch was nearly closed after 48 hours.

**Table 2.** Genes Profoundly down-regulated in H20-PANC-1 compared to H0-PANC-1 cells.

Gene Symbol	Gene Description	Fold Regulation	Gene Function (Qiagen,USA)
EGLN1	Egl-9 family hypoxia inducible factor 1	-6.0349	Angiogenesis, regulation of cell proliferation
VDAC1	Voltage dependent anion channel 1	-5.0159	Metabolism, regulation of apoptosis
NFKB 1	Nuclear factor kappa B subunit 1	-2.1628	Regulation of apoptosis, regulation of cell proliferation

**DISCUSSION**

PANC-1 pancreatic cancer cells were subjected to recurring episodes of hypoxia lasting 8 hours, repeated three times per week, over a period of up to six months. Gene expression profiling was conducted at two-time points during the study: midway through the experiment and at its conclusion. The findings suggest that hypoxia is a significant factor in the emergence of a chemo-resistant phenotype in the PANC-1 pancreatic cancer cell line. It is evident that there are notable disparities in gene expression profiling between cells that were subjected to 20 episodes of hypoxia. We have achieved the enhancement of chemo-resistance to Doxorubicin through the induction of hypoxia in PANC-1 cells for a total of 20 episodes,

**Table 3.** Genes profoundly up regulated in H20-PANC1 compared to H0-PANC1 cells.

Gene Symbol	Gene Description	Fold Regulation	Gene Function (Qiagen,USA)
BTG1	B cell translocation gene 1 anti-proliferative	9.8383	Regulation of apoptosis, regulation of cell proliferation, angiogenesis.
HIF1AN	Hypoxia inducible factor 1 alpha subunit inhibitor	6.9431	HIF 1 & co-transcription factors
TP53	Tumor protein P53	6.6093	Regulation of apoptosis
HMOX1	Heme oxygenase (decycling) 1	4.529	Angiogenesis, apoptosis, metastasis
GPI	Glucose - 6- phosphate isomerase	3.9178	regulation of cell proliferation, regulation of apoptosis , metabolism
PKM	Pyruvate kinase, muscle	3.4092	Metabolism, apoptosis
SLC16A3	Solute carrier family 16 member 3	2.8492	Transporter, metabolism
ADORA2B	Adenosine A2b receptor	2.3495	Angiogenesis
TFRC	Transferrin receptor	2.1186	Regulation of cell proliferation
BHLHE40	Basic helix-loop-helix family member e40	2.0144	Cell differentiation, proliferation, apoptosis, and metabolism.
NCOA1	Nuclear receptor coactivator 1	2.0017	receptors, regulation of cell proliferation

Gene expression changes in H40-PANC-1 pancreatic cancer cell line



**Table 4.** Genes profoundly down-regulated in H40-PANC-1 cells.

Gene Symbol	Gene Description	Fold Regulation	Gene Function (Qiagen,USA)
ODC1	Ornithine decarboxylase	-2.5154	Regulation of cell proliferation
HIF3A	Hypoxia inducible factor 3 alpha subunit	-2.4014	HIF1 & Co-transcription factors
COP55	COP9 signalosome subunit 5	-2.198	HIF1 & Co-transcription factors, regulation of cell proliferation
PDK1	Pyruvate dehydrogenase kinase 1	-2.1379	Regulation of cell proliferation
EGR1	Early growth response 1	-2.0106	Angiogenesis, regulation of cell proliferation

**Table 5.** Genes profoundly up-regulated in H40-PANC1 compared to H0-PANC1 cells.

Gene Symbol	Gene Description	Fold Regulation	Gene Function (Qiagen,USA)
MMP9	Matrix metalloproteinase 9	2.3345	Angiogenesis, wound healing, metastasis
EGLN1	EGL-9 family hypoxia inducible factor 1	2.0233	HIF1 & cell proliferation
ADM	Adrenomedullin	2.012	Regulation of apoptosis, regulation of cell proliferation

each lasting 8 hours. This observation was substantiated by a significant elevation in the IC50 values in comparison to the normoxic cells, exhibiting an approximate 2.7-fold increase. Our results are in agreement with<sup>18</sup> who reported that hypoxia induces chemo-resistance. Nevertheless, until this report is written, it is the first study to illustrate generation of chemo-resistance to Doxorubicin in PANC-1 pancreatic cancer cell line through chronic hypoxia.

One of the hypothesized mechanisms by which hypoxia prompts chemo-resistance is by decreasing in the formation of damaging ROS by Doxorubicin, which is decreased because of lacking oxygen<sup>19</sup>. As the MTT cell proliferation assay was done under normoxia, this theory is unlikely to contribute to our results. Another suggested theory relies on the rapid proliferation rate of cancer cells<sup>20</sup>. The theory in question is not applicable to our study, as the rate of cell proliferation in our sample has been found to be equivalent to that of normoxic cells of a similar age. This suggests that the genomic and proteomic characteristics of the cell are responsible for chemo-resistance development.

The primary finding of our study pertains to the up-regulation of the B cell translocation gene 1 anti-proliferative (BTG1) and Tumor protein P53 (TP53) in H20-PANC-1, with fold changes of 9.8383 and 6.6093, respectively, compared to H0-PANC-1. The induction of BTG1 by hypoxia and its observed promotion of apoptosis suggest that it is unlikely to play a role in chemo-resistance induction.

TP53 is one of the most prominent tumor suppressor genes<sup>21</sup>, TP53 is inactivated by mutation in almost 50% of human cancer<sup>22</sup>, including PDAC (50– 75%)<sup>23</sup>. TP53 is known to be induced by hypoxia<sup>24</sup>. The up-regulation of P53 in our results has been confirmed by<sup>25</sup> who found that P53 protein accumulates in cells exposed to hypoxia, this accumulation increases with the duration of the hypoxia incubation along with the decrease in the pO2 level<sup>25</sup>.

Mutant p53 proteins interfere with regulation of cell survival, DNA damage repair and drug resistance<sup>26</sup>. Our results can also concur with the work of<sup>27</sup> who have found that mutant p53

stimulates chemo-resistance of pancreatic adenocarcinoma cells to the standard drug gemcitabine. Heme oxygenase (decycling) 1 (HMOX1) was found to be up-regulated by 4.529 folds in H20 PANC-1 compared to H0 PANC-1, HMOX1 is involved in preventing chemo toxic-induced apoptosis, this concur with the findings of<sup>28</sup> who have indicated that HMOX-1 is involved in conferring the chemo-resistance to Doxorubicin in breast cancer cells by preventing apoptosis and autophagy. Another prominent gene in our study that is associated with apoptosis is Pyruvate kinase, muscle (PKM), which was up-regulated by 3.4092 folds in H20-PANC-1 compared to H0-PANC-1. Increased PKM levels have been associated with resistance to 5-fluorouracil in patients with colorectal cancer<sup>29</sup>. Also, genetic silencing of PKM potentiates the effects of docetaxel and cisplatin in in vitro and lung cancer xenograft models<sup>30</sup>. A recent study observed that a significant proportion (68%) of human pancreatic adenocarcinoma specimens exhibited a robust expression of PKM2. Additionally, nearly all of the analyzed pancreatic cancer cell lines also demonstrated a high level of PKM expression. Furthermore, the detection of PKM was found to have detrimental effects on cellular proliferation and enhanced apoptosis when examined in vitro<sup>31</sup>. Accordingly, we can conclude that HMOX-1 and PKM play an important role in defending PANC-1 cells from hypoxia- and chemotoxic-induced apoptosis.

In addition, there is another gene that is associated with apoptosis and notably down-regulated by -5.0159 folds in H20 compared with H0 is Voltage dependent anion channel 1 (VDAC1), VDAC1 is anchor for pro- and anti-apoptotic proteins, respectively, of the hexokinase (HK)<sup>32</sup> and Bcl-2 families of proteins, which contribute to the balance between survival and cell death<sup>33</sup> found that hypoxia induces the appearance of a C-terminal truncated form of VDAC1, so our results is contrast with his finding as VDAC-1 was down-regulated. It has been confirmed that mitochondrial hexokinase I and II (HKI/II) function as anti-apoptotic proteins when bound to VDAC1, but their detachment enabled activation of apoptosis<sup>34</sup>. While VDAC1 was down-regulated by hypoxia, so most probably VDAC1 is not a player in chemo-resistance induction. Another prominent gene in our study is the Egl-9 family hypoxia inducible factor



1 (EGLN1), which is the most significantly down-regulated gene in H20-PANC-1 compared to H0-PANC-1 with -6.0349 folds, Prolyl hydroxylase 2 (PHD2) is an enzyme encoded by the EGLN1 gene and is considered the key oxygen sensor-regulating hypoxia-inducible factor (HIF)<sup>35</sup>. Another important gene that acts as cellular oxygen sensors, directly regulating the transcriptional complex HIF-1 activity that mediates adaptive responses to reduced oxygen level is hypoxia-inducible factor 1 alpha subunit inhibitor (HIF1AN)<sup>36</sup>. The HIF1AN mRNA has been one of the most obviously up-regulated genes in H20-PANC-1 compared to H0-PANC-1 with 6.9431 folds. FIH-1 which HIF1AN encodes. PHDs and FIH-1 are parts of the same superfamily of dioxygenases and both need the same set of cofactors, oxygen, 2-oxoglutarate and ferrous iron, but striking differences exist between these enzymes. FIH-1 possesses a higher affinity to the molecular oxygen in comparison to the PHDs, which recommends that in mild hypoxia FIH-1 remains active while the PHDs are inactive<sup>37</sup>. Our results in agreement with<sup>38</sup> who reported that FIH-1 is capable to control the HIF-1 even under severe hypoxic conditions, when PHD enzymes fail to do so.

One of the surprising results of our study was the up-regulation of Matrix metalloproteinase 9 (MMP9), in H40 PANC-1 compared to H0-PANC-1 by 2.3345 folds while it was barely changed at the level of H20-PANC-1. These results against the expectation as the main function of MMP9 involved in metastasis and wound healing<sup>39,40</sup>. It was expected MMP9 to be up regulated in H20 as the wound closed at faster rate compared to H0 and to be down regulated in H40 when the wound migrated at slower rate compared to H0. These results agree with the huge body of evidence reporting that HMOX-1 is frequently up regulated in tumor tissues and promotes tumor growth and metastasis<sup>41-44</sup> have showed that HMOX1 as well promotes proliferation in pancreatic cancer cell lines. HMOX1 is induced by hypoxia<sup>44,45</sup>. GPI is an autocrine motility factor (AMF), and is also critical for migration, invasion and metastasis of tumor cells<sup>46</sup>. As mentioned in the literature review that the conversion of epithelial cells to more mesenchymal-like cells enabled cell migration, invasion, and metastasis. It has been demonstrated that GPI/AMF suppressed epithelial marker expression and improved mesenchymal marker expression<sup>47</sup>. GPI/AMF is up-regulated by hypoxia and is controlled by HIF-1<sup>48</sup>. As a result, we can suggest that the up regulation of HMOX1 and GPI in our results is consistent with the fact that hypoxia renders cancer cells more progressive and induces metastasis. Among our results Adenosine A2b receptor (ADORA2B) gene is up-regulated by 2.3495 folds in H20-PANC-1 compared to H0-PANC-1. Adenosine itself is a unanimously expressed purine nucleoside that is well-known to have the ability to bridge the gap between metabolic supply and demand<sup>49</sup>. It acts through four G coupled receptors: A1, A2A, A2B and A3 with A2B having the highest threshold for activation<sup>50</sup>. We have thoroughly discussed the role of ADORA2B in literature review under angiogenesis. ADORA2B is known to be induced in stressful conditions including hypoxia<sup>51</sup>.

Stimulation of ADORA2B receptors in cancer cell lines up-regulates the production of angiogenic factors, indicating that tumor A2B receptors may promote revascularization<sup>52-54</sup>. It makes sense for ADORA2B to be up-regulated; not only to empower adenosine, but also to facilitate immune system evasion<sup>55</sup>. Activation of host ADORA2B receptors have been

viewed as an inhibitor of solid tumors rejection and plays an essential role in promoting metastasis<sup>56</sup>.

One of the key metabolic features of cancer cells is rapid utilization of glucose to fuel glycolysis. The metabolic reactions of glycolysis are catalyzed by numerous enzymes, one of the enzymes is PKM which is implicated in cancer<sup>57</sup>. In our study we have found that PKM is up regulated in H20 PANC-1 compared to H0 PANC-1 by 3.4 folds. The rationale behind the up-regulation of PKM is to meet the increased of glucose demand due to hypoxia-induced glycolysis. PKM dimer can translocate to the nucleus, where it will interact with HIF-1a and regulate expression of numerous proglycolytic enzymes<sup>58</sup>. PKM plays a crucial role in aerobic glycolysis<sup>59</sup>. Thus PKM drives the aerobic glycolysis in generating the energy needed for cellular processes

in tumors, a phenomenon termed "the Warburg effect"<sup>60</sup>. Our results agree with<sup>61</sup>, who found that the mRNA levels of PKM was elevated with hypoxic treatment in numerous mouse and human hepatoma cell lines, although HIF-1 mRNA levels were not significantly elevated. In addition, our results found that Solute carrier family 16 member 3 (SLC16A3 also called MCT4), is up-regulated by 2.8 folds in H20-PANC-1. This up-regulation has been reported by<sup>62</sup> who found that SLC16A3 was up-regulated by hypoxia in bladder cancer. The up regulation of MCT proteins is noticed in tumor tissue and supports an apparent need for lactate shuttling to either fuel tumor growth or permit survival under stress conditions<sup>63</sup>. Our findings agree with<sup>64</sup> who found that SLC16A3 in general is elevated in Pancreatic ductal adenocarcinoma (PDAC), but the overexpression of Solute Carrier Family 16 Member 1 (SLC16A1) in PDAC was little versus other cancers<sup>64</sup>. Notable differences in genes were measured in cells exposed to 40 episodes of hypoxia such as Ornithine decarboxylase (ODC) gene. ODC gene is one of the most down-regulated gene in H40 cells compared to H0 cells by -2.5 folds. ODC1 is the first rate-limiting enzyme of polyamine biosynthesis, is found to be associated with cell growth, proliferation and transformation<sup>65</sup>.

## CONCLUSION

We also suggest that this reversal could be owed to the fact that other effectors take over when a stressful situation goes on beyond a specific point. Unfortunately, no other studies have exposed PANC-1 cells for a similar period for comparison purposes. However, the precise determination of a mechanism behind chemo-resistance is beyond the scope of this report. Chemo-resistance was used as an end-point for transformation in light of absence of proliferation rate and morphological changes. Further research is necessary to investigate protein levels in order to uncover the currently elusive mechanism of hypoxia-induced chemotherapy resistance and identify a target for reversing this significant barrier in cancer treatment.

## CONFLICTS OF INTEREST

The authors declare that they have no conflicts of interest.



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