


# Hypobaria-Induced Oxidative Stress Facilitates Homocysteine Transsulfuration and Promotes Glutathione Oxidation in Rats with Mild Traumatic Brain Injury

Journal of Central Nervous System Disease  
Volume 13: 1–11  
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DOI: 10.1177/1179573520988193



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## ABSTRACT

**BACKGROUND:** United States service members injured in combat theatre are often aeromedically evacuated within a few days to regional military hospitals. Animal and epidemiological research indicates that early exposure to flight hypobaria may worsen brain and other injuries. The mechanisms by which secondary exposure to hypobaria worsen trauma outcomes are not well elucidated. This study tested the hypothesis that hypobaria-induced oxidative stress and associated changes in homocysteine levels play a role in traumatic brain injury (TBI) pathological progression caused by hypobaria.

**METHODS:** Male Sprague Dawley rats were exposed to a 6 h hypobaria 24 h after mild TBI by the controlled cortical impact. Plasma and brain tissues were assessed for homocysteine levels, oxidative stress markers or glutathione metabolism, and behavioral deficits post-injury in the absence and presence of hypobaria exposure.

**RESULTS:** We found that hypobaria after TBI increased oxidative stress markers, altered homocysteine metabolism, and promoted glutathione oxidation. Increased glutathione metabolism was driven by differential upregulation of glutathione metabolizing genes. These changes correlated with increased anxiety-like behavior.

**CONCLUSION:** These data provide evidence that hypobaria exposure after TBI increases oxidative stress and alters homocysteine elimination likely through enhanced glutathione metabolism. This pathway may represent a compensatory mechanism to attenuate free radical formation. Thus, hypobaria-induced enhancement of glutathione metabolism represents a potential therapeutic target for TBI management.

**KEYWORDS:** Hypobaria, homocysteine, oxidative stress, glutathione, traumatic brain injury

**RECEIVED:** July 28, 2020. **ACCEPTED:** December 18, 2020.

**TYPE:** Original Research Article

**FUNDING:** The authors disclosed receipt of the following financial support for the research, authorship, and/or publication of this article: This work was supported by the Department of Defense, U.S Air Force, Grant # FA8650-17-2-6H10

**DECLARATION OF CONFLICTING INTERESTS:** The authors declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

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## Introduction

Traumatic brain injury (TBI) is a public health problem with more than 175 000 victims among United States service members during the Iraq and Afghanistan wars and this number continues to grow as some of them remain in combat zones.<sup>1</sup> Wounded service members, including approximately 10% who suffered from psychiatric disorders such as TBI, were aeromedically evacuated to hospitals in select countries for advanced care critical for survival between 2001 and 2013.<sup>2</sup> During these up to 9 h flights, the cabin pressure is usually maintained at the equivalent to 8000 ft altitude. The mild hypoxia that accompanies such hypobaric conditions is thought to be one of several stressors that could cause secondary injury to TBI patients during these flights.<sup>3,4</sup>

Over the past decade, there has been significant interest in understanding the impact of hypobaria on TBI pathological progression using various injury models and hypobaria conditions. Mice who underwent 5 h hypobaria (8000 ft) after a

closed head injury 3 h earlier presented worse outcomes in cerebral inflammation and neuronal death than those exposed to hypobaria a day later.<sup>5</sup> In a recent report, rats subjected to blast overpressure and exposed 48 h later to hypobaria for 12 h presented with exacerbation of multi-organ injuries, including cerebral lesions.<sup>6</sup> Work from our laboratory showed that 6 h exposure to 8000 ft hypobaria worsened axonal injury in rats subjected to under-vehicle blast.<sup>7</sup> Furthermore, hypobaria-exposure aggravated inflammation, neuronal death, and cognitive deficits in rats subjected to TBI by lateral fluid percussion and exposed to hypobaria up to 7 days post-injury.<sup>8</sup> Despite current advances in understanding the impact of hypobaria-exposure on TBI pathophysiology, hypobaria-associated biochemical changes and their effects on injury pathological outcomes remain unclear.

Elevated levels of the non-proteinogenic amino acid homocysteine (HCY), known as hyperhomocysteinemia (HHCY), is a stress biomarker and stress inducer with substantial



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neurotoxic properties.<sup>9-13</sup> Homocysteine metabolism is tightly regulated to ensure it is continuously remethylated to form methionine, its precursor, or transsulfurated to contribute to glutathione formation, an essential endogenous antioxidant.<sup>14,15</sup> There is very limited data on homocysteine metabolism and pathological impact in hypobaria. A previous study by Sharma and colleagues reported that prolonged stay at high altitude (7500 ft) for 3 months, showing an increase in homocysteine levels.<sup>16</sup> We recently reported in our model of methionine-induced moderate HHCY worsens TBI-induced lesion volume, BBB disruption, inflammation, anxiety-like behavior, and working memory in a rat model of mild TBI by controlled cortical impact.<sup>17</sup> In a prior retrospective study involving 150 participants, Rahmani et al<sup>18</sup> found a significant correlation between plasma HCY levels, TBI severity, and associated mortality. In the current study, we tested the hypothesis that hypobaria-induced oxidative stress and associated changes in homocysteine levels play a role in traumatic brain injury (TBI) pathological progression caused by hypobaria.

Mild traumatic brain injury (mTBI) model used in this study is unlike moderate and severe TBI, defined in rodents by no breaking of the dura, subtle pathological changes around the impact region and fast recovery from behavioral deficits.<sup>17,19</sup>

## Materials and Methods

### *Animals and housing*

Male Sprague Dawley rats (300-350 g; Envigo, CA) were used. They were housed in pairs in a temperature-controlled environment, maintained at  $23 \pm 2^\circ\text{C}$ , with a 12 h light/dark cycle, with continuous access to food and water *ad libitum*. All animal procedures were approved by The University of Maryland, Baltimore, Animal Use and Care Committee (IACUC # 1118014) and, the USAF Animal Use Programs Office of Research Oversight & Compliance, and the USAMRMC Animal Care and Use Review Office (ACURO). Animals were randomly assigned to experimental groups composed of 4 to 6 rats/group for biochemical assays and histological analyses, and 8 rats/group for behavioral studies.

### *Induction of controlled cortical impact injury*

The Controlled Cortical Impact (CCI) method was used to induce mTBI.<sup>17,19</sup> Animals were maintained under general anesthesia with 2.5% isoflurane throughout surgery. Rat body temperature was maintained at  $37^\circ\text{C}$  using a heating pad coupled with a rectal probe. Following craniotomy, CCI was performed over the left parietal cortex. Using a stereotaxically positioned 4 mm diameter piston tip, impact-injury was performed with an electromagnetically controlled impactor (Leica Biosystems, IL) at a piston velocity of 5 m/sec and depth penetration of 1.5 mm. The bone flap was replaced, the skin closed, and the anesthesia terminated. Sham-operated rats were also

anesthetized, followed by craniotomy and replacement of the bone flap in the absence of cortical impact.

### *Rat exposure to hypobaria*

At twenty-four hours post-surgery, rats were housed in cages with *ad libitum* food and water access and placed in a hypobaria chamber. A vacuum pump connected to the chamber was gradually adjusted to depressurize the chamber to 568 mmHg, which is the ambient pressure at 8000 ft., altitude. As previously reported, the chamber was kept normoxic (21% O<sub>2</sub>) with supplemental O<sub>2</sub>, were at a near-normal inspired oxygen level of 30%, the SpO<sub>2</sub> was 97.0% at a hypobaria level 8000 ft. Rats were maintained under these conditions for 6 h, followed by depressurization every 15 min to sea level ambient pressure (757 mm Hg).<sup>7</sup> As control, other rats were placed in the chamber under normobaric conditions, equivalent to 80 ft., altitude, 21% O<sub>2</sub> for the same duration. Some sham or CCI rats (6/group) received intraperitoneal injections of 2-dihydroethidium (2-DHE) in 50% DMSO (6 mg/kg) to track superoxide formation<sup>20</sup> during hypobaria or normobaria-exposure.

### *Behavioral testing*

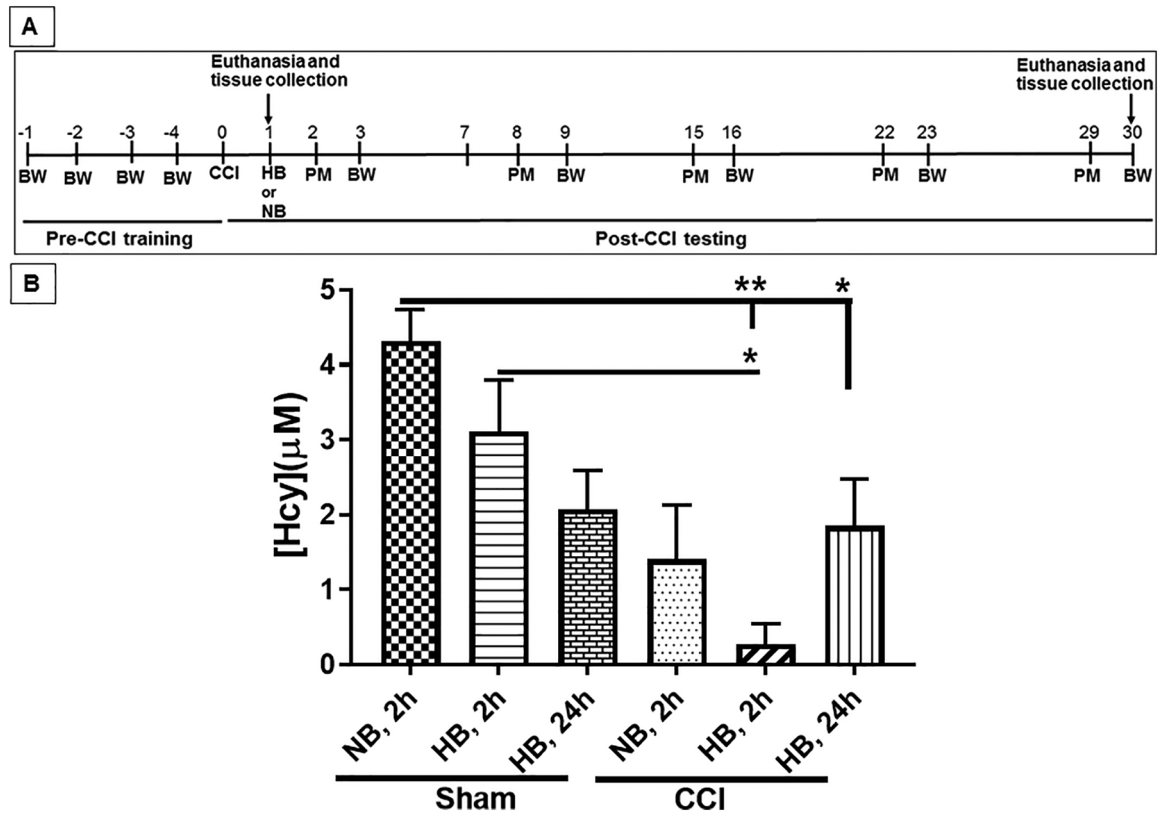
All behavioral tests were performed at time points described in the experimental time (Figure 1A)

*Beam walk test.* This test was performed to assess fine motor movements, as previously described.<sup>21,22</sup> Briefly, animals were trained to walk on a 1.5 mm wooden beam for 3 consecutive days, and baseline data determining the number of foot faults over a total of fifty steps were collected on day 4.<sup>22</sup> Test to assess the effect of hypobaria-exposure on fine motor movement in sham and CCI rats overtime was performed on days 3, 9, 16, and 30 post-surgery.

*Elevated Plus maze test.* The Elevated Plus maze was used to assess rats for anxiety-like behavior on days 2, 8, 15, and 29 post-surgery using a Plus-sign shape device as previously described.<sup>23,24</sup> Briefly, rats were individually placed in the central area of the device and allowed to explore the maze for 10 min. Animal movements were analyzed by an overhead camera and data collected and analyzed by Any-Maze software (SD instrument, CA) to provide several behavioral parameters including the time spent in open arms. Anxiety-like behavior was interpreted as inversely proportional to the time an animal spent in the open arms.

### *Tissue collection and processing*

Brain and blood were collected at 2 h, 24 h, or 30-days post-surgery. Rats were anesthetized and blood collected and processed to obtain plasma for biochemical assays as previously described.<sup>17,24</sup> Rats were thereafter euthanized and brain



**Figure 1.** Hypobarbia exposure enhances plasma homocysteine elimination: (A) a detailed timeline of behavioral tests and endpoints of experimental procedures starting on day 4 before surgery (study day -4). BW, beam walk test; CCI, controlled cortical impact; HB, hypobarbia exposure; NB, normobarbia exposure; PM, plus-maze test and (B) histogram of homocysteine quantitation showing that hypobarbia enhanced its elimination in sham rats. Homocysteine elimination was potentiated, leading to scanty or undetectable levels among hypobarbic CCI-rats ( $n=4/\text{group}$ ;  $*P < .05$ ,  $**P < .01$ ).

collected and processed for slicing in 40  $\mu\text{m}$  thick coronal sections stored at  $-20^\circ\text{C}$  for histology as described.<sup>17,24</sup>

#### Plasma homocysteine measurement

Homocysteine levels were measured in rat's plasma collected 2 h or 24 h post-hypobarbia or normobarbia exposure ( $n=4/\text{group}$ ). The assay was performed using an enzymatic assay kit (BioVision Inc., CA) following the manufacturer's instructions.

#### Determination of oxidative stress markers in plasma and brain

**Malondialdehyde measurements.** Lipid peroxidation levels were determined by measuring the amount of thiobarbituric acid reactive substance (TBARS) malondialdehyde (MDA) in rat plasma samples ( $n=6/\text{group}$ ). The assay was performed using a kit (Cayman Chemical, MI) following the manufacturer's instructions.

**Superoxide production.** The formation of superoxide anion was monitored in rat brains during hypobarbia using 2-dihydroethidium (Molecular Probes, OR), a fluorescent probe that incorporates between nucleic acid strands of superoxide producing cells.<sup>20</sup> DHE was dissolved in 50% DMSO and administered intraperitoneally at 6 mg/kg immediately before hypobarbia or

normobarbia exposure ( $n=4/\text{group}$ ). Rats were euthanized at two-hours post-exposure and brain removed, processed, sectioned, and preserved as described.<sup>17,24</sup> DHE incorporated cells were microscopically visualized with a 20 $\times$  objective at 594 nm wavelength.

#### Measurements of reduced and oxidized plasma glutathione levels

Plasma samples were collected during euthanasia from sham or brain-injured rats at 2 h, 24 h, or 30 days post-hypobarbia or normobarbia exposure. Total and oxidized glutathione concentration was measured in rat plasma samples ( $n=4/\text{group}$ ) using a colorimetric assay kit (Life Technologies Inc, MD). In brief, the assay used a colorimetric substrate that reacted with the free thiol group on glutathione to generate a highly colored product. Reduced glutathione concentration was the difference between total and oxidized plasma glutathione concentration measured separately, following the manufacturer's instructions.

#### Transcriptional analysis of glutathione metabolizing genes

Total RNA was extracted from snap-frozen samples of the ipsilateral prefrontal cortex, parietal cortex, and hippocampus from sham and CCI-rats euthanized 2 h post-hypobarbia or

**Table 1.** Presentation of forward (F) and reverse (R) primer sequences used for transcriptional analysis of glutathione metabolizing genes GS, GPX-1, and GPX-4. The table also shows the annealing temperature of these primers, their respective target amplicon size, and the mRNA sequence's access code from which each primer set was derived.

MRNA	RAT PRIMER SEQUENCE (5'-3')	ANNEALING TEMP (°F)	AMPLICON (BP)	ACCESS CODE
GS-F	ACA CAT CTT CAT TGC CCT CTC	57	101	NM_012962.1
GS-R	TCT ATG CCC ATC TAC CCT CTT			
GPx-1-F	GTT CCA GTG CGC AGA TAC A	57	111	NM_030826.4
GPx-1-R	CCA GAT ACC AGG AAT GCC TTA G			
GPx-4-F	CCC ACT GTG GAA ATG GAT GA	58.5	105	NM_017165.4
GPx-4-R	ACG CAG CCG TTC TTA TCA A			
GAPDH-F	CCC TTC ATT GAC CTC AAC TAC	55	102	NM_017008.4
GAPDH-R	GAT GAC CAG CTT CCC ATT CT			

normobaric exposure (n = 4/group). RNA extraction was done using the PureLink RNA Mini kit (Ambion/Life Technologies Inc., CA) following the manufacturer's instructions. Extracted RNA concentration was determined using a Spectrophotometer Nanodrop 2000C (Thermo Scientific, CA), and RNA tubes were stored immediately at  $-80^{\circ}\text{C}$  until used.

First, an Access RT-PCR system kit (Promega, WI) was used to amplify 2  $\mu\text{g}$  RNA and determine the optimal annealing temperature of primer sets designed for the quantitative amplification of glutathione metabolizing genes *GS*, *GPx-1*, and *GPx-4*, as well as the housekeeping control gene glyceraldehyde 3-phosphate dehydrogenase (*GAPDH*) (Table 1).

Quantitative reverse transcriptase-polymerase chain reaction (Q-RT-PCR) was performed using the Power SYBR Green RNA-to-CT 1-Step kit (Applied Biosystem, MA). In brief, a complete RT-PCR mix containing 150 nM of primer (IDT, CA) that yielded the best annealing temperature for the respective *GS*, *GPx-1*, or *GPx-4* gene of interest and *GAPDH* control was mixed with approximately 15  $\mu\text{g}$  of individual RNA template and placed in a 96 wells plate. Q-RT-PCR test of samples in the 96 wells plate was run in a QuantStudio-3 machine (Applied Biosystems, CA) for a total of 35 cycles after reverse transcriptase reaction at  $48^{\circ}\text{C}$  for 30 min and enzyme activation at  $95^{\circ}\text{C}$  for 10 min (n = 4/group). Each of the 35 Q-PCR cycles consisted of 15-s denaturation at  $95^{\circ}\text{C}$ , and 1 min annealing/extension at the temperature noted in Table 1. Each sample was run in triplicate for the target genes *GS*, *Gpx-1*, or *GPx-4* and duplicate for the reference gene (*GAPDH*). At the end of the run, the amplification of a single gene product was verified by the presence of a single peak in the melting curve analysis. Normalized cycle threshold ( $\Delta\text{Ct}$ ) values were obtained by subtracting the  $\text{Ct}$  of the reference gene from that of each of the target genes. The double delta  $\text{Ct}$  ( $\Delta\Delta\text{Ct}$ ) for each target gene was determined by deducting the mean  $\Delta\text{Ct}$  of the target gene in the sham normobaric group from the  $\Delta\text{Ct}$  of the respective target gene. The fold-change of each target gene compared to the normobaric sham group was calculated as  $2^{-\Delta\Delta\text{Ct}}$ .

### Lesion volume assessment

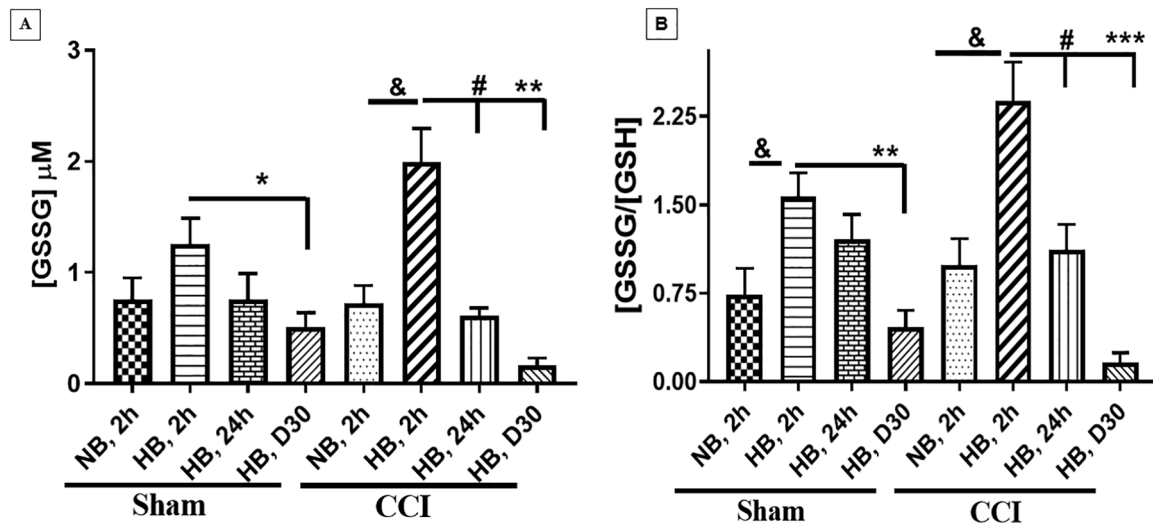
Hematoxylin and eosin (Sigma, MO), stained brain sections were dehydrated, cleared, and mounted for lesion volume estimation. Injury lesion volume was quantified from 8 sections corresponding to every 24 serial expanding from bregma 1.05 to  $-6.30$  based to cover the complete lesion area (n = 5/group). Quantification was done using the Cavalieri estimator method of unbiased stereology using the stereo-investigator software (MBF Biosciences, VT). The contusion region (impact-induced damaged region resulting in tissue loss) and penumbra region (the pathologically affected region around impact-induced lesion that could be salvaged with an effective therapeutic intervention) were outlined in the ipsilateral hemisphere of all sections with a grid spacing of 100  $\mu\text{m}$ , and the injury lesion volume was determined and expressed in  $\text{mm}^3$ .<sup>25,26</sup>

### Quantitation and statistical analysis

Image J software (NIH) was used to determine the percent area covered by DHE incorporated cells. To achieve that, 30 microscopic fields of each serial section from DHE or saline-injected rats were captured with SRS fractionator of the stereo-investigator software (MBF Biosciences, VT). Each image was optically segmented with a threshold greyscale value of 0.3  $\mu\text{m}$ , and the percent DHE-positive area was automatically measured. The mean of all DHE positive microscopic fields was determined.

Statistical analyses were performed using GraphPad InStat-3 software (GraphPad Software, Inc., CA). Data from biochemical and histological tests were analyzed using One-Way ANOVA and repeated ANOVA for behavioral data. Both tests were coupled to the Tukey-Kramer post-test to compare differences among the various groups. Results were expressed as mean  $\pm$  standard error of the mean (SEM). Statistical significance was defined as  $P < .05$ . The individuals performing histologic, biochemical, and behavioral assays were blinded to animal group identifications, using codes revealed after data collection. glyceraldehyde 3-phosphate dehydrogenase





**Figure 2.** Hypobaric exposure enhances glutathione oxidation: (A) quantitative graph of oxidized glutathione levels showing that glutathione oxidation shows a non-significant increase by hypobaric at 2h post-exposure in sham rats ( $n=4$ ). It is significantly intensified by the combination of mTBI and hypobaric exposure ( $n=4$ ,  $\&P < .05$ ,  $**P < .01$ ) and (B) histogram of the ratio of oxidized to reduced glutathione showing a 1.8-fold increase among hypobaric shams relative to normobaric shams ( $n=4$ ;  $\&P < .05$ ) at 2h post-exposure. In contrast, there was more than a two-fold difference between hypobaric-CCI rats compared to normobaric-CCI rats ( $n=4$ ;  $\&P < .05$ ). Glutathione oxidation reduced significantly over time after hypobaric ( $\#P < .05$ ,  $**P < .01$ ,  $***P < .001$ ).

## Results

### *Hypobaric exposure reduces circulating homocysteine levels*

To determine the effect of acute hypobaric-exposure on HCY metabolism, we measured plasma HCY levels at 2h and 24h after 6h of 8000ft hypobaric-exposure using normobaric-exposure as control. We found that mTBI alone significantly reduced plasma HCY levels in normobaric-rats after 2h after exposure,  $1.41 \pm 0.723 \mu\text{M}$ , compared to normobaric-shams,  $4.32 \pm 0.431 \mu\text{M}$  (Figure 1;  $Q=4.31$ ,  $P < .05$ ). At the same 2h time point, HCY levels were further reduced or undetectable in the plasma of TBI rats exposed to hypobaric to  $0.27 \pm 0.28 \mu\text{M}$  and were significantly lower than levels in normobaric ( $3.12 \pm 0.63 \mu\text{M}$ ) and hypobaric ( $4.32 \pm 0.431 \mu\text{M}$ ) shams (Figure 1;  $Q=0.41$ ,  $P < .05$ ;  $Q=5.88$ ,  $P < .01$  respectively). HCY levels in CCI-hypobaric animals were still significantly reduced 24h after hypobaric  $1.85 \pm 0.63 \mu\text{M}$  compared to normobaric shams (Figure 1;  $Q=4.23$ ,  $P < .05$ ). Hypobaric-exposure also caused a non-significant decrease in HCY levels among non-TBI sham rats with levels of  $3.12 \pm 0.63 \mu\text{M}$  at 2h post-exposure and  $2.07 \pm 0.52 \mu\text{M}$  at 24h post-exposure compared to normobaric-shams  $4.32 \pm 0.431 \mu\text{M}$  (Figure 1;  $P > .05$ ).

### *Hypobaric-induced increases oxidized glutathione*

We measured reduced and oxidized glutathione levels in that plasma. This showed that hypobaric-exposure caused a modest increase in oxidized glutathione in sham rats at 2h post-exposure ( $1.25 \pm 0.23 \mu\text{M}$ ) compared to normobaric-shams ( $0.76 \pm 0.19 \mu\text{M}$ ;  $P > .05$ ; Figure 2A). TBI alone had no effect

on glutathione oxidation among normobaric-rats at 2h post-exposure ( $0.723 \pm 0.158 \mu\text{M}$ ) compared to normobaric-shams ( $0.76 \pm 0.19 \mu\text{M}$ ;  $Q=0.56$ ,  $P > .05$ ; Figure 2A). However, the combination of TBI and hypobaric significantly increased plasma oxidized glutathione to  $1.998 \pm 0.296 \mu\text{M}$  compared to CCI-normobaric and CCI-hypobaric 24-h and 30days post-surgery (Figure 2A;  $Q=4.41$ ,  $P < .05$  and  $Q=9.23$ ,  $P < .01$  respectively).

The ratio of oxidized to reduced glutathione is an important metric of the metabolic impact of TBI and hypobaric on glutathione metabolism. When we assessed the ratio of oxidized to reduced glutathione, we found that hypobaric-exposure substantially enhanced oxidized glutathione over reduced glutathione in both sham and CCI rats resulting in  $1.57 \pm 0.202$  and  $2.38 \pm 0.323$ -ratios respectively, compared to  $0.741 \pm 0.164 \mu\text{M}$  and  $0.99 \pm 0.22 \mu\text{M}$  for normobaric-sham and CCI rats (Figure 2B,  $P < .05$ ). This ratio decreased significantly overtime at 24h and 30-days (Figure 2B;  $Q=4.61$ ,  $P < .05$  and  $Q=6.79$ ,  $P < .01$  respectively) post-exposure in a time-dependent manner. The elevation of oxidized glutathione implies increased free radical scavenging through glutathione oxidation,<sup>27</sup> which could activate the endogenous glutathione oxidizing metabolic pathways.

### *Hypobaric induces upregulation of glutathione metabolizing genes*

We examined the transcriptional profile GS, GPx-1, and GPx-4 in 3 regions of the ipsilateral hemisphere, the prefrontal cortex, parietal cortex, and hippocampus, at 2h post-exposure, to determine how their expression responds to hypobaric-exposure. All results are presented in Table 2. These

**Table 2.** Presentation of transcriptional analysis results of glutathione metabolizing genes GS, GPX-1, and GPX-4 in the prefrontal cortex, the parietal cortex of shams, and CCI rats exposed to normobaric or hypobaric conditions. These results represent the fold expression change of each of these genes in different experimental groups compared to normobaric-shams used as control.

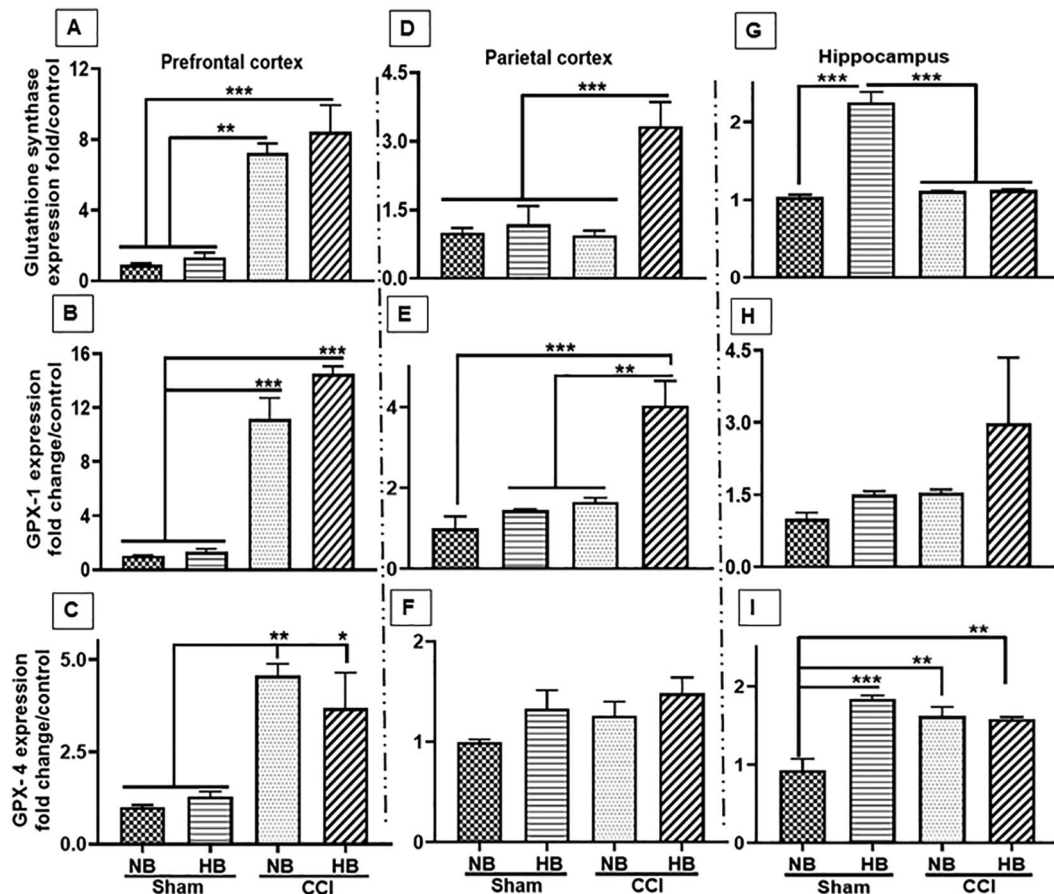
STUDY GROUPS	PREFRONTAL CORTEX				PARIETAL CORTEX				HIPPOCAMPUS					
	GS		GPX-1		GS		GPX-1		GS		GPX-1		GPX-4	
	GS	GPX-1	GS	GPX-1	GS	GPX-1	GS	GPX-1	GS	GPX-1	GS	GPX-1	GS	GPX-4
Sham, NB	1 ± 0.086	1 ± 0.045	1 ± 0.05	1 ± 0.059	1 ± 0.3	1 ± 0.021	1 ± 0.034	1 ± 0.124	1 ± 0.034	1 ± 0.021	1 ± 0.034	1 ± 0.124	1 ± 0.138	1 ± 0.138
Sham, HB	1.33 ± 0.24	1.35 ± 0.19	1.28 ± 0.137	1.2 ± 0.2	1.45 ± 0.03	1.33 ± 0.18	2.25 ± 0.14	1.5 ± 0.05	2.25 ± 0.14	1.33 ± 0.18	2.25 ± 0.14	1.5 ± 0.05	1.84 ± 0.04	1.84 ± 0.04
CCI, NB	7.23 ± 0.53	11.16 ± 0.56	4.5 ± 0.32	0.95 ± 0.05	1.65 ± 0.1	1.26 ± 0.14	1.12 ± 0.005	1.54 ± 0.06	1.12 ± 0.005	1.26 ± 0.14	1.12 ± 0.005	1.54 ± 0.06	1.62 ± 0.12	1.62 ± 0.12
CCI, HB	8.43 ± 1.51	14.5 ± 0.58	3.68 ± 0.95	3.33 ± 0.26	4.04 ± 0.6	1.46 ± 0.15	1.12 ± 0.02	2.99 ± 1.36	1.12 ± 0.02	1.46 ± 0.15	1.12 ± 0.02	2.99 ± 1.36	1.58 ± 0.023	1.58 ± 0.023

results show a heterogeneous expression pattern in different brain regions. In the prefrontal cortex, the expression of all 3 genes was enhanced by the cortical impact-induced brain injury (Table 2; Figure 3A–C;  $Q=4.51$ ,  $P<.05$ ;  $Q=6.90$ ,  $P<.01$ ;  $Q=9.83$ ,  $P<.001$ ) with no significant effect of hypobaric-exposure only on the expression compared to normobaric-shams.

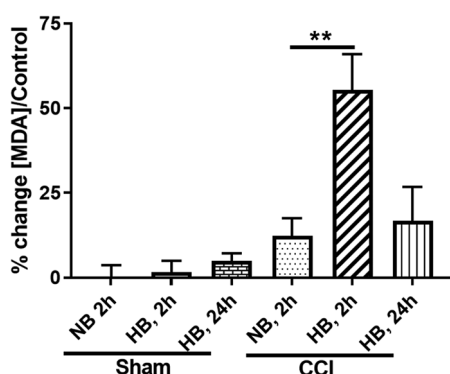
In contrast, the combination of cortical brain injury and hypobaric-exposure upregulated the expression of GS and GPX-1 in the parietal cortex compared to normobaric brain-injured rats as well as hypobaric and normobaric shams (Table 2; Figure 3D, E;  $Q=6.26$ ,  $P<.01$ ). There was no significant change in GPX-4 expression in the parietal cortex of rats in any group (Table 2; Figure 3F;  $P>.05$ ). In the hippocampus, GS expression was significantly upregulated by hypobaric alone in shams compared to other groups (Table 2;  $Q=15.32$ ,  $P<.001$ ). Similar to the parietal cortex, hypobaric or CCI only increased the expression of GPX-1 in the hippocampus and the combination of both resulted in more upregulation, although all are non-significant (Table 2; Figure 3H,  $P>.05$ ). The expression of GPX-4 in the hippocampus was increased by injury or hypobaric alone, and no additional effect of the 2 combined (Table 2,  $Q=7.1$ ,  $P<.01$ ;  $Q=9.96$ ,  $P<.001$ ). In summary, the expression of glutathione metabolizing genes was differentially affected by brain injury, hypobaric, or by the combination of both factors, dependent on the targeted brain region.

#### *Hypobaric conditions increase oxidative stress*

The biochemical markers above indicate that there may be considerable oxidative stress causing a compensatory reduction in HCY and elevation in the processing of glutathione antioxidant systems. Consistent with this conjecture, we found that TBI caused a non-significant  $12.3 \pm 5.23\%$  increase in plasma malondialdehyde levels in normobaric-CCI rats at 2 h post-exposure compared to normobaric-shams (Figure 4;  $P>.05$ ). In addition to the modest increase in TBI, when combined with hypobaric-exposure, there was a significant increase in MDA formation of  $55.38 \pm 10.5\%$  over baseline (Figure 4;  $Q=6.83$ ,  $P<.01$ ). Although there is generally a good correlation between blood and brain biomarkers in TBI studies,<sup>28</sup> these measures of oxidative stress do not give information on where in the body they originate. To verify which brain regions were impacted by hypobaric induced free radical formation during hypobaric exposure, we used 2-DHE for in vivo tracking of superoxide production in brain cells (Figure 5A) and compared conditions by measuring the percentage of pixels in defined brain regions that are 2-DHE positive. There was significant increase in 2-DHE incorporated cells in normobaric-CCI rats ( $2.42 \pm 0.18\%$  area covered) compared to normobaric-shams ( $1.02 \pm 0.16\%$ ; Figure 5B;  $Q=0.46$ ,  $P<.05$ ). Hypobaric-exposure alone did not cause a significant change in superoxide production in



**Figure 3.** Glutathione metabolic genes expression were differentially altered by hypobaric exposure and mTBI. Histograms depicting changes in the transcriptional analysis of GS, GPX-1, and GPX-4 in the prefrontal cortex (A-C), parietal cortex (D-F), and hippocampus (G-I;  $n=4/\text{group}$ ). These graphs indicate that (1) in the prefrontal cortex, the expression of all 3 genes was mainly driven by impact-induced brain injury with no or little effect of hypobaric exposure ( $*P < .05$ ,  $**P < .01$ ,  $***P < .001$ ). (2) In the parietal cortex, the expression of GS and GPX-1 was mostly influenced by the combination of brain injury and hypobaric exposure ( $P < .01$ ,  $P < .001$ ), whereas that of GPX-4 showed no significant change among groups ( $P > .05$ ). (3) In the hippocampus, GS expression was significantly upregulated by hypobaric alone in shams compared to other groups (G;  $***P < .001$ ), whereas that of GPX-4 was significantly affected by injury or hypobaric alone (I;  $***P < .01$ ,  $***P < .001$ ). In comparison, the expression of GPX-1 was not significantly affected by hypobaric or injury alone, the combination of both resulting in greater upregulation.



**Figure 4.** Hypobaric exacerbates lipid peroxidation in rats with TBI. Plasma malondialdehyde levels were a non-significant increase in normobaric-CCI rats 2h post-exposure. Hypobaric exposure caused a significant increase of this molecule in CCI rats plasma simultaneously ( $n=6/\text{group}$ ;  $P < .01$ ). Malondialdehyde levels decreased in these rats 24h after hypobaric exposure.

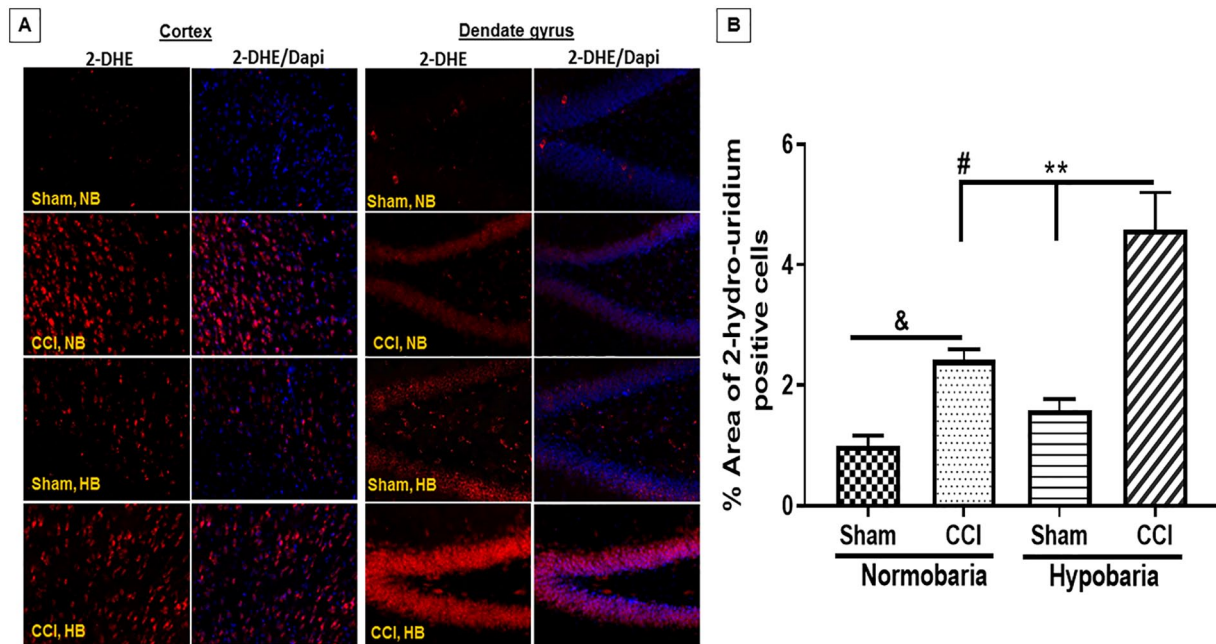
sham rats resulting in  $1.59 \pm 0.18\%$  versus normobaric-shams ( $1.02 \pm 0.16\%$ ; Figure 5B;  $P > .05$ ). However, in the presence of hypobaric with CCI, there was a marked increase in

2-DHE positive cells present with 5 times the area compared to sham (TBI/CCI  $4.58 \pm 0.62\%$  of the brain region; Figure 5B;  $Q=0.68$ ,  $P < .01$ ) and normobaric-CCI rats ( $Q=0.46$ ,  $P < .05$ ). Thus, hypobaric exposure exacerbates TBI associated oxidative stress.

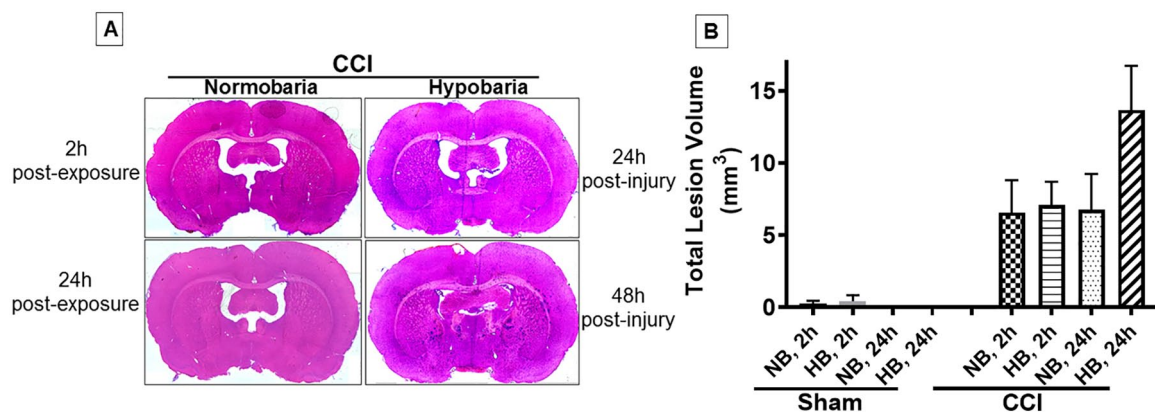
#### TBI lesion volume was not significantly affected by hypobaric exposure

To assess the effect of hypobaric exposure on brain injury lesion, we used the Cavalieri method of unbiased stereology to measure injury-induced lesion volume at 2h and 24h post-exposure. At 2h post-exposure, impact-induced injury lesion volume was approximately the same between normobaric,  $6.56 \pm 2.25 \text{ mm}^3$ , and hypobaric-CCI rats,  $7.09 \pm 1.42 \text{ mm}^3$  (Figure 6A, B;  $P > .05$ ). But at 24h post-exposure, corresponding to 48h post-surgery, brain injury lesion volume presented a non-significant increase following hypobaric exposure,  $13.68 \pm 3.05 \text{ mm}^3$  compared to  $6.78 \pm 2.48 \text{ mm}^3$  among normobaric-rats (Figure 6A, B;  $P > .05$ ). This suggests that hypobaric exposure has no acute negative impact on TBI brain tissue loss.





**Figure 5.** Superoxide formation intensifies in rat with brain injury after hypobaria exposure: (A) representative microscopic images showing 2-DHE incorporation in superoxide producing cells in the injury penumbra region and the hippocampus dentate gyrus. Quantitation of the %area of 2-DHE positive cells and (B) indicates that brain injury alone caused a significantly increased superoxide formation ( $n=4$ ;  $&P < .05$ ) and this was aggravated by hypobaria-exposure ( $n=4$ ;  $\#P < .05$ ,  $**P < .01$ ). The scale bar is 100  $\mu\text{m}$ .



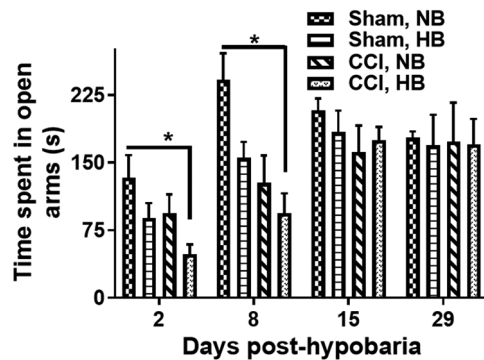
**Figure 6.** Hypobaria exposure does not affect TBI-induced cortical lesion: (A) Representative hematoxylin and eosin-stained coronal brain sections showing cortical impact-induced brain lesions. (B) Lesion volume quantitation indicating that hypobaria did not affect CCI-induced cortical brain lesion 2h post-exposure. However, 24h post-exposure, the lesion was greater but non-significant in hypobaric rat brains ( $n=5$ ), compared to those exposed to normobaria.

### *Hypobaria exposure increases anxiety-like behavior in rats with mild TBI*

Next, we wanted to determine whether hypobaria exposure following mTBI in animals exacerbates behavioral deficits. We assessed the effect of hypobaria-exposure on anxiety-like behavior usually associated with TBI-related psychiatric impairment such as posttraumatic stress disorder.<sup>29,30</sup> Anxiety was assessed with a standard elevated plus-maze in which reduced time spent in the open arms of the maze is interpreted as elevated anxiety-like behavior. As a control, fine motor behaviors classically associated with cerebellar function, a brain region removed from the injury site, were not significantly affected as assessed by the Beam Walk test. On day 2

post-exposure, which is an early time point when changes due to hypobaria are mostly likely detectable, normobaric sham and CCI rats respectively made  $1.28 \pm 0.98$  and  $1.87 \pm 0.62$ -foot faults over 50 steps. In comparison, hypobaric sham and CCI rats made  $1.39 \pm 0.75$  and  $1.77 \pm 1.02$  foot faults ( $P > .05$ , all compared to normobaric shams). However, brain-injured rats exposed to hypobaria spent significantly less time in the open arms of Plus maze on days 2 and 8 post-injury, indicative of increased anxiety compared to normobaria-shams. These animals spent  $48.21 \pm 10.74\text{s}$  and  $94.08 \pm 21.49\text{s}$  in the open arms of the Plus maze on days 2 and 8 respectively versus  $133.04 \pm 25.28\text{s}$  and  $241.82 \pm 29.12\text{s}$  for normobaria-shams (Figure 7;  $Q=5.21$  and  $5.58$  respectively;  $P < .05$ ). To a lesser





**Figure 7.** Traumatic brain injury-induced anxiety-like behavior is aggravated by hypobaria exposure. Graph depicting the time spent by rats in the open arms of the elevated plus maze. Both brain injury and hypobaria alone caused a non-significant in anxiety-like behavior. The combination of mTBI and hypobaria significantly increased anxiety-like behavior on days 2 and 8 post-surgery compared to normobaric-shams ( $n=8/\text{group}$ ;  $*P < .05$ ). Two and 4 weeks post-HB, this behavioral deficit was resolved, and rats in all groups spent closely the same time in the plus-maze open arms ( $P > .05$ ).

extent, hypobaria-sham and normobaria-CCI rats also exhibited some deficits in anxiety-like behavior on days 2 and 8 post-surgery (Figure 7;  $Q < 4.96$  thresholds for significance;  $P > .05$ ), compared to normobaria-shams. After 8 days, rats in all groups recovered from this deficit and exhibited Plus maze performance close to normobaric-shams.

## Discussions

Reports from several investigators suggest that flying at high altitude is associated with increased physiological stress,<sup>31,32</sup> thus representing a risk for TBI patients who require aeromedical evacuation for advanced care. Results from the present study indicate that exposure of rats with mTBI, which represents up to 80% of all TBI cases,<sup>33,34</sup> to hypobaria induces an acute increase of neural oxidative stress, which subsequently triggers an enhanced response of the antioxidant defense system that uses HCY and glutathione in a compensatory attempt to attenuate free radical formation.

Levels of oxidative stress markers malondialdehyde and superoxide increased in rats subjected to mTBI 2 h post-hypobaria (Figures 4 and 5). This observation suggested that oxidative damage during aeromedical evacuation produced changes that could represent targets for therapeutic interventions. This is a key concept, as unmitigated free radical formation mediates the induction of many TBI pathological markers including, inflammation and neuronal degeneration that are exacerbated by hypobaria-exposure.<sup>5-8,35,36</sup> Therefore, our findings suggest that hypobaria-induced oxidative stress is a significant mediator or indicator of TBI pathological progression during and after hypobaria exposure.

Hypobaria-induced oxidative stress was coupled with very low to undetectable levels of plasma homocysteine (Figure 1). Homocysteine is a non-proteinogenic amino acid

with substantial cytotoxic potential when accumulated.<sup>37-39</sup> Therefore, its metabolism is tightly regulated to ensure it is continuously eliminated.<sup>15,40</sup> This occurs via its remethylation to methionine, its precursor, or by its transsulfuration to cystathionine, contributing to the downstream formation of glutathione, a very potent antioxidant.<sup>14,15</sup> Therefore, we postulated that the drastic reduction of homocysteine levels under oxidative stress at 2-h post-hypobaria was indicative of its flush through the transsulfuration pathway. This hypothesis was potentiated by a significant increase of glutathione oxidation resulting in a two-fold increase of oxidized glutathione versus reduced glutathione at 2 h post-hypobaria exposure (Figure 2A, B). The return of oxidized glutathione to baseline levels at day 30 post-exposure suggests an attenuation of oxidative stress. This observation correlates with the significant improvement in anxiety-like behavior to that of normobaric-sham rats. The therapeutic potential of homocysteine transsulfuration to promote glutathione formation has been underscored in several neurological disorders. For instance, the supplementation of N-acetylcysteine to promote glutathione biosynthesis through the transsulfuration pathway was neuroprotective in models of Huntington's disease,<sup>41,42</sup> Alzheimer's disease,<sup>43,44</sup> and TBI.<sup>45,46</sup> Moreover, this therapeutic approach is in a trial for pediatric TBI management.<sup>47</sup>

Glutathione has free radical scavenging and anti-inflammatory properties.<sup>48,49</sup> Its metabolism is regulated by genes, including *GS*, which controls its biosynthesis through the transsulfuration pathway and *GPx*, which catalyzes glutathione oxidation to attenuate free radical formation.<sup>14,15</sup> *GPx* has 8 isozymes, among which isozymes 1 and 4 are abundant in the brain.<sup>50-53</sup> We analyzed the transcriptional profile of these genes in the prefrontal cortex and the hippocampus that are very vulnerable to oxidative stress<sup>54-57</sup> and the parietal cortex around the cortical injury penumbra region. The upregulation of *GS* and *GPx* in the prefrontal cortex (Figure 3A-C) attests to its vulnerability to oxidative stress, relative to its distant location to the injury region in the parietal cortex. In the parietal cortical, the combination of CCI and hypobaria resulted in an increase in *GS* and *GPx* that was not present with CCI or hypobaria alone (Figure 3D-F). This suggests that the CCI penumbra region may be uniquely stressed by a synergy between the injury and hypobaria. To our knowledge, this study is the first to describe changes in glutathione and associated gene expression in a model of TBI and Hypobaria exposure.

Although the mechanism for the distinctive expression patterns in different brain regions is not fully elucidated, increased *GPx* expression in the cerebral cortex of aged rats was associated with higher astrocyte density.<sup>58</sup> Furthermore, *GPx* activity was derived primarily from astrocytes relative to neurons in chicks' forebrain.<sup>59</sup> Also, increased *GPx* expression or increased glutathione oxidation was reported in several oxidative stress associated neurological disorders.<sup>60,61</sup>

In summary, this study provides evidence that hypobaria exacerbates oxidative stress in rats with mTBI. This was associated with an enhanced flush of homocysteine through the trans-sulfuration pathway to promote glutathione biosynthesis, which is necessary to attenuate free radical formation. Increased glutathione metabolism was putatively driven by the upregulation of glutathione metabolizing genes, suggesting that enhanced glutathione metabolism represents a therapeutic target for TBI management. Further studies would be needed to understand how this antioxidant redox system would respond to interventions during and after hypobaria-exposure following moderate to severe TBI.

### Acknowledgements

We acknowledge Ms. Fengying Li and Ms. Lyric Ramsue for their technical assistance

### Author Contributions

Experimental design: FT, GF, CM; Laboratory experiments and data collection: FT, MG; Data analysis: FT, AP, GF, MG; Manuscript writing: FT, GF; Manuscript editing: AP, MG, FT.

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