



Tumor grade related expression of neuroglobin is negatively regulated by PPAR γ and confers antioxidant activity in glioma progression



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A B S T R A C T

Neuroglobin (NGB), distributed mainly in central and peripheral nervous systems, is a nerve globin with neuroprotective effects against oxidative stress resulting from hypoxia and ischemia. Recent studies have indicated that the expression of NGB is related to neurodegenerative disorders and cancers, but the molecular mechanisms for its transcriptional regulation and protection are not well defined. Here, we report that the expression of NGB in glioma is grade related and is negatively regulated by PPAR γ . Specific PPAR γ agonist reduces the expression of NGB, while its inhibitor enhances the expression. Moreover, NGB participates in regulating the phosphorylation of AKT in glioma cells, which may contribute to the glioma progression where accumulating oxidative pressure presents. Overexpression of NGB could protect glioma cells against 4-HNE induced cell death, and partially reverse PPAR γ 's pro-apoptotic and anti-proliferative abilities. These results display an important role of NGB in glioma progression and a mechanism for its transcriptional regulation, and suggest that the treatment on glioma through PPAR γ agonist appears to be triggered by the modulation of NGB.

1. Introduction

Reactive oxygen species (ROS) are continuously generated in cells largely by the mitochondria during energy metabolism process [1]. It is reported that ROS, in particular hydrogen peroxide, have physiological roles as signaling molecules involved in cell growth, differentiation and death [2]. However, when the ROS production overwhelms the clearance conducted by the endogenous antioxidant systems, excessive ROS accumulation may cause progressive oxidative damages to lipids, proteins and DNAs, and may trigger cell death [3,4]. These damages are collectively defined as oxidative stress, which has been implicated in a wide variety of human diseases, such as neurodegenerative disorders, cardiovascular diseases and cancers [5,6]. Whereas, multiple researches have indicated that cancer cells maintain much higher ROS levels than normal cells, which are critical for its high proliferation rate, accelerated metabolism and tumor progression [7–9]. Therefore, it is fundamentally important for cancer cells to balance the ROS levels with further oxidative damage that cells can bear. However, how cancer cells

deal with such draconic oxidative pressure is complex and remains incompletely understood.

Gliomas are the most common primary tumors in central nervous system, which make up about 30% of all brain and central nervous system tumors and 80% of all malignant brain tumors [10]. Gliomas can be graded from I to IV according to the World Health Organization (WHO), or divided into low grade (WHO grade I and II) and high grade (WHO grade III and IV), which are determined by the pathologic evaluation of the tumors [11]. Despite the frequency of gliomas, the molecular mechanism and etiology for this thorny tumor are not well defined, making its treatment a challenge. Multiple molecules have been reported with potentiality in glioma treatment, and peroxisome proliferator-activated receptors (PPARs) family member – PPAR γ – is one of them. PPARs are ligand-activated transcription factors regulating various genes involved in lipid metabolism and energy homeostasis, and its three subtypes, PPAR α , PPAR δ and PPAR γ , have been identified [12,13]. PPAR γ is widely involved in different cancers, and its agonists, including thiazolidinediones (TDZs) and nonthiazolidinediones, are

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emerging to block the motility and invasiveness of glioma cells. Grommes et al. have indicated that micromolar doses of pioglitazone (30 μ M) counteract C6 rat glioma cell invasiveness in vivo [14]. Eyupoglu et al. have found that troglitazone treatment could effectively block glioma progression and brain invasion with an in vitro glioma invasion model [15]. However, the mechanisms evoked by PPAR γ agonists in glioma treatment are not fully understood.

Neuroglobin (NGB) is a recently identified protein, distributed mainly in central and peripheral nervous systems [16]. It's one of the vertebrate globin family members, together with hemoglobin, myoglobin and cytoglobin [17]. The expression of NGB increases in vitro cells subjected to ischemia and hypoxia, and protects cells from these insults [18–20]. Meanwhile, multiple researches have shown that NGB could protect rats and mice against experimental stroke in vivo, while the expression of NGB has been found increased in the peri-infarct region and essentially absent from the infarct core in human stroke patients [21,22]. These works show the protective effects of NGB against oxidative stress results from ischemia and hypoxia. Moreover, increasing evidences have shown the close relationship between NGB and cancers. Fiocchetti et al. reported that NGB could promote estrogen receptor α -positive cancer cells survive under oxidative stress [23], but zhang et al. indicated that NGB functions as a tumor suppressor in Hepatocellular Carcinoma [24]. Therefore, more work are needed to investigate the role NGB plays in cancers. Here, we report that the expression of NGB, modulated by PPAR γ , is required for glioma to resist oxidative pressure during its progression. We find that the expression of NGB is directly correlated with the grade of the glioma, and is inversely correlated with PPAR γ expression. Activation of PPAR γ results in suppressed NGB expression, while inhibition of it enhances the latter. Overexpression of NGB protects against oxidative stress resulting from PPAR γ agonists treatment and 4-HNE insult, indicating the important role of enhanced NGB expression in glioma progression. Our work reveals a mechanism for the regulation of NGB expression, and demonstrates that NGB is an important part of the antioxidant defense required for glioma progression.

2. Material and methods

2.1. Subjects

The study was approved by the Ethics Committee of Shuangnan Hospital, Chengdu and the Committee of Sichuan Agricultural University. We ensured that all research subjects in the study have gave informed consent. Thirty-nine patients with brain glioma admitted by the Department of Neurosurgery, Shuangnan Hospital of Chengdu from June 2010 to October 2014 were randomly selected. There were 21 male and 18 female patients, aged between 15 and 73 years. All patients received cranial MRI scan, and the glioma was confirmed by pathological examination after surgery. Their gliomas were dissected and divided into two pieces, with one rapidly frozen in liquid nitrogen and preserved at -80°C pending for western blot, qRT-PCR, and NADPH/NADP assay, and the other one fixed in paraformaldehyde (4%) was used for further immuno-histochemical staining. Before the western blot, qRT-PCR, and NADPH/NADP assay were performed, each of the frozen samples was further divided into three pieces pending for these assays.

2.2. Immuno-histochemical staining

Part of the tissue was fixed in 4% paraformaldehyde before paraffin sections were performed. Then, immune-histochemical staining was performed with standard methods and the primary antibodies of NGB (ab37258, Abcam, MA, USA; 1:100) and PPAR γ (ab45036, Abcam, MA, USA; 1:500). Positive signals were visualized using colorimetric detection with diaminobenzidine (DAB), and the hematoxylin indicated the nucleus. Finally, the images were photographed with a microscope

(BX43, OLYMPUS).

2.3. Immunofluorescence staining

Parts of the tissue was fixed in 4% PFA and kept in 30% sucrose at 4°C before frozen sectioning was performed. Standard Immunofluorescence staining was carried out to examine the expression of NGB in different grade of glioma using anti-NGB antibody (ab37258, Abcam, MA, USA; 1:100) as primary antibody, and the Goat anti-Mouse IgG antibody (Alexa Fluor 488, Invitrogen, A-11029, MA, USA, 1:1000) was used as a secondary antibody. DAPI indicates the nucleus and the images were photographed with a fluorescent microscope (BX63, OLYMPUS) with excitation at 488 nm.

2.4. Quantitative realtime PCR

RNA extraction and real-time PCR were performed as previously reported. Briefly, total RNA was extracted from the tissues using Trizol reagent (Invitrogen, Waltham, MA, USA). RNA was subjected to reverse transcription with reverse transcriptase according to the manufacturer's instructions (Fermentas, Waltham, MA, USA). Quantitative real-time PCR was performed using the Bio-Rad CFX96 system, and the relative gene expression was normalized to internal control Actin. Primer sequences for SYBR Green probes of NGB are: Forward: 5'-ggc acc gtc ctg ttt gcc ag-3' and Reverse: 5'-cga gca tca cct tcc tga tg-3'; β -Actin: Forward: 5'-tcc ttc ctg ggc atg gag t-3' and Reverse: 5'-aaa gcc atg cca atc ta tc-3'.

2.5. Western blotting

Standard Western blotting procedures were carried out as previously reported [25] with the following antibodies: anti-4-Hydroxy-2-Nonenal (4-HNE) antibody (ab48506, Abcam, MA, USA; 1:1000) to detect lipid peroxidation, anti-PTEN (ab32199, Abcam, MA, USA; 1:500) and anti-AKT (ab8805, Abcam, MA, USA; 1:1000), anti-NGB antibody (ab37258, Abcam, MA, USA; 1:500), anti-PPAR γ (ab45036, Abcam, MA, USA; 1:1000), anti-Phospho-Akt (Ser473) antibody (mAb #4060, Cell signaling, Danvers, MA; 1:1000), anti- Phospho-Akt (T308) (mAb #13038, Cell signaling, Danvers, MA; 1:1000), anti- β -actin (BA0410, Boster, Wuhan, China; 1:1000), anti-GAPDH (A00227, Boster, Wuhan, China; 1:1000). Briefly, samples from each patient were lysed and solubilized in PBS buffer containing 1% triton X-100, 1% SDS, Protease inhibitor cocktail III (BioVision) and Phosphatase Inhibitor Cocktail IV (BioVision). Ultrasonication was performed, and the total protein was collected after refrigerated centrifugation at 4°C . Protein concentrations were determined by Bradford assay using BSA as a standard. 15–20 μ g of protein from each sample was separated on a 12.5% Tris-HCl denaturing gel by SDS-PAGE, transferred to nitrocellulose membranes for 1 h at 300 V, 300 mA, blocked with 5% non-fat milk, incubated with primary antibodies overnight at 4°C and followed by a series of 3 washes in PBS. Membranes were then incubated with a HRP-conjugated secondary antibody for 1 h at room temperature followed by 5 washes in PBS. HRP detection was performed using the Super Signal West Pico Chemiluminescent Substrate (PIERCE) and membranes were exposed to standard FUJI film using an automated film developer.

2.6. Cell culture

U87 cells were routinely cultured in DMEM (Gibco, Waltham, MA, USA), supplemented with 10% fetal bovine serum (FBS) in an incubator under an atmosphere of 5% CO $_2$ at 37°C . Cells were plated in 6-well cell plates, 24 h after the PPAR γ agonist troglitazone and pioglitazone (Sigma-Aldrich) supplement, or 12 h after the PPAR γ inhibitor GW9662 (Sigma-Aldrich) treatment, cells were collected for genes and proteins expressions detection. To investigate the protection of NGB, 50 μ M 4-

Hydroxy-2-nonenal (4-HNE, Cat#2083, BioVision) or pioglitazone were added 24 h after the NGB transfection in 96-well cell plate, and cell viability was assayed with CCK-8 kit 12 h after the insults.

2.7. NADPH/NADP ratio assay

The NADPH/NADP ratio assay was performed on 39 glioma extracts using the NADP/NADPH assay kit (Cat#K347, BioVision) according to the manufacturer's instructions. Briefly, ~20 mg samples were extracted in 400 μ L of the recommended extraction buffer, and 50 μ L were processed following instructions for each duplicate. OD450 measurements were made on a plate-reader (Thermo, Waltham, MA, USA) at 25 °C, and the data was converted to nmol/sample using a standard curve and values were used for ratio as previously reported [26].

2.8. Cell viability assay

Cell viability was measured by Cell Counting Kit-8 (CCK-8) system (Dojindo, CK04-11, Minato-ku, Tokyo, Japan) according to the manufacturer's instructions. Briefly, CCK-8 solution (10 μ L per 100 μ L of medium in each well) was added, and the plates were then incubated at 37 °C for 1 h. The absorbance of each well was read at 450 nm using a microplate reader (Thermo, Waltham, MA, USA).

2.9. ROS level analysis

U87 Cells were plated in 6-well cell plates, and they were transfected with Vehicle (GFP) and GFP-NGB plasmid using Lipofectamine 2000 (Invitrogen). Twenty-four hours after the transfection, pioglitazone (50 μ M) was supplemented for another 24 h. Then, fresh culture medium with ROS-sensitive fluorescent probes (Ethidium, Invitrogen) was added, and incubated for 30 min at 37 °C. Finally, the cells were collected and intracellular ROS was detected by fluorescence-activated cell sorting FACS of ~10,000 GFP positive cells.

2.10. Statistical analysis

Data represent the means and SEM. One-way ANOVA and post hoc tests were performed for all statistical significance analysis using GraphPad Prism software (GraphPad Software, Inc., La Jolla, CA, USA). * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

3. Results

3.1. Patients information

As shown in Table 1, thirty-nine patients with tumor were enrolled in this study, including 21 males and 18 females. Age are between 15 and 73 years, with the mean age of 42.5. Every patient received a cranial MRI scan before surgery. The results revealed that most of the lesions located in the frontal, parietal and temporal lobe, while some others located in basal ganglia, lateral ventricles and thalamus. All these lesions were confirmed as glioma by pathological examination after operation, and were graded from I to IV. There were 2 cases in grade I, 16 in grade II, 7 in grade III, and 14 in grade IV, and all of these cases were divided into Low grade (grade I and II, 18 cases) and High grade (grade III and IV, 21 cases).

3.2. Tumor grade related expression of NGB is negatively correlated with that of PPAR γ

Literatures have reported the expression of NGB in lots of tumors [27–29]. To investigate the expression of NGB in glioma, 18 low-grade and 21 high grade glioma samples were examined. We have found that the expression of NGB presented in glioma, and the mRNA level of NGB

Table 1
General information and clinical diagnosis of patients with glioma.

No.	Gender	Age	Lesion area	WHO grade	Low or High grade
1	Male	39	Frontal and Temporal Lobe	2	Low
2	Male	27	Temporal Lobe	2	Low
3	Female	43	Lateral Ventricles	2	Low
4	Female	17	Frontal and Temporal Lobe	2	Low
5	Male	22	Parietal Lobe	2	Low
6	Male	38	Temporal Lobe (right)	1	Low
7	Male	49	Frontal and Temporal Lobe	2	Low
8	Female	42	Basal Ganglia, Insula	4	High
9	Male	54	Frontal Lobe	2	Low
10	Male	30	Frontal and Parietal Lobe	2	Low
11	Male	73	Temporal Lobe	2	Low
12	Male	48	Frontal and Temporal Lobe	2	Low
13	Male	53	Basal Ganglia	3	High
14	Female	48	Basal Ganglia	3	High
15	Male	33	Parietal and Temporal Lobe	4	High
16	Female	45	Frontal, Parietal and Temporal Lobe	4	High
17	Female	38	Parietal and Frontal Lobe	2	Low
18	Female	55	Frontal and Temporal Lobe	3	High
19	Male	40	Frontal, Parietal and Temporal Lobe	4	High
20	Female	42	Frontal, Parietal and Temporal Lobe	4	High
21	Male	42	Frontal, Parietal and Temporal Lobe	3	High
22	Male	27	Frontal and Temporal Lobe	4	High
23	Female	64	Frontal Lobe, Lateral Ventricles	4	High
24	Male	39	Parietal Lobe	2	Low
25	Female	56	Basal Ganglia, Insula	4	High
26	Female	41	Frontal, Parietal and Temporal Lobe	4	High
27	Female	72	Lateral Ventricles	2	Low
28	Male	42	Basal Ganglia, Insula	4	High
29	Female	53	Frontal and Temporal Lobe	2	Low
30	Female	28	Frontal, Parietal and Temporal Lobe	1	Low
31	Male	26	Frontal, Parietal, Temporal Lobe and Basal Ganglia	4	High
32	Male	15	Thalamus	3	High
33	Female	31	Frontal, Parietal and Temporal Lobe	4	High
34	Female	66	Frontal, Parietal and Temporal Lobe	3	High
35	Female	45	Frontal and Temporal Lobe	4	High
36	Male	54	Frontal and Temporal Lobe	3	High
37	Male	52	Lateral Ventricles	2	Low
38	Female	45	Frontal and Temporal Lobe	2	Low
39	Male	25	Frontal Lobe, Lateral Ventricles	4	High

were much higher in high-grade gliomas than that in low-grade ones (Fig. 1A). Western blots and immunofluorescence staining revealed the same findings of NGB protein levels in glioma (Fig. 1B, C and E). Positive correlation between NGB expression and the grade of glioma suggests an important role of NGB in glioma progression. PPAR γ , which is widely expressed in multiple cancers, functions as a tumor suppressor [30]. In glioma, western blots and immune-histochemical staining displayed suppressed expression of PPAR γ in high-grade gliomas (Fig. 1B, C and F), and the expressions of NGB and PPAR γ were inversely related (Fig. 1D, E and F). These findings reveal the completely contrary expressions of NGB and PPAR γ related to the grade of glioma.

3.3. PPAR γ modulates the expression of NGB in vitro

PPAR γ functions as a regulator of genes involved in multiple processes. PTEN, the downstream target of PPAR γ , is upregulated after PPAR γ activation, resulting in dephosphorylation of phosphatidylino-

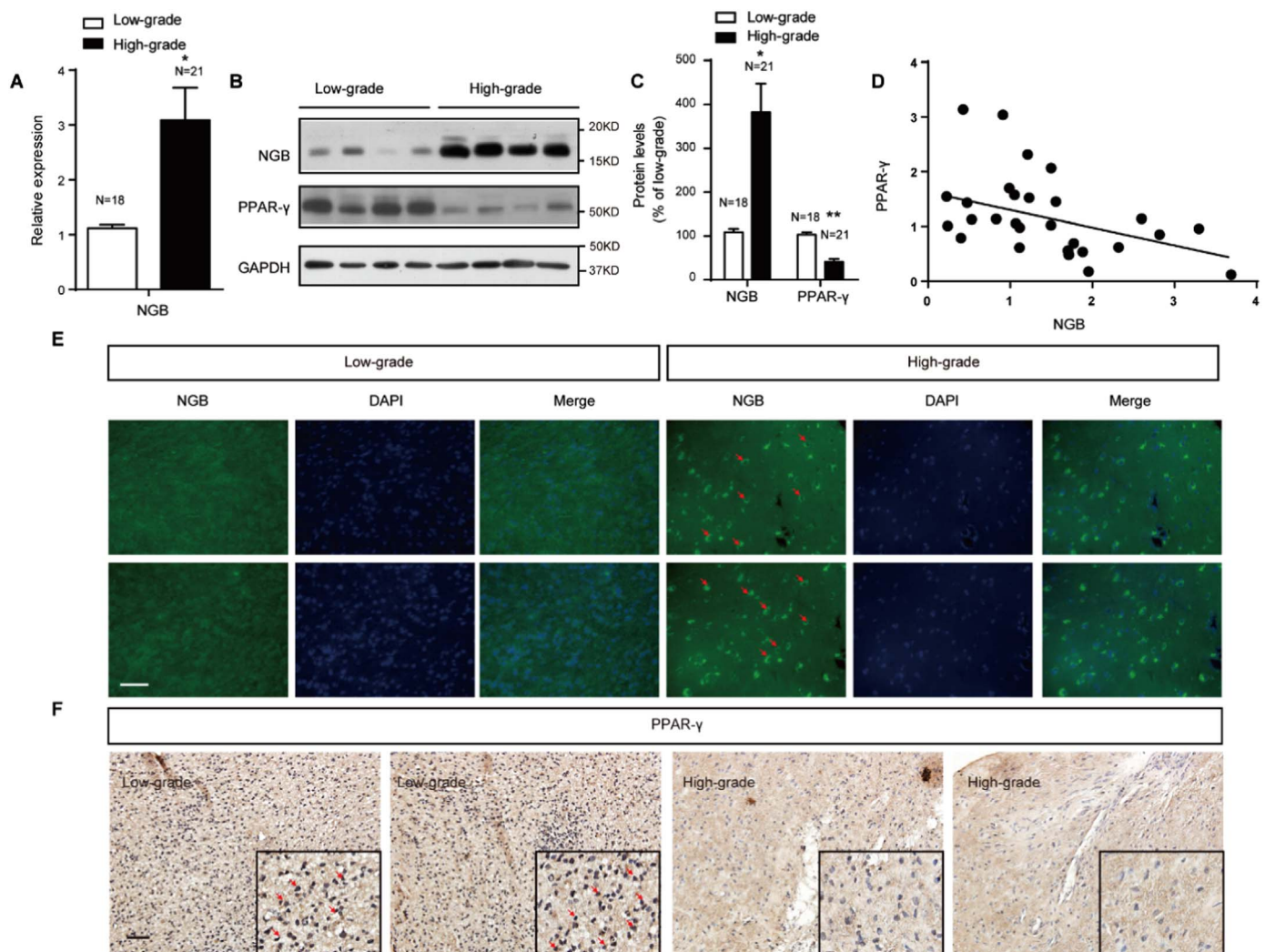


Fig. 1. Tumor grade related expression of NGB is negatively correlated with that of PPAR γ . (A) qRT-PCR shows a robust expression of NGB mRNA in high-grade gliomas (N=21) compared with low-grade gliomas (N=18). Error bars indicate SEM. * $p < 0.05$. (B and C) Representative images of western blots and quantification show higher NGB and less PPAR γ protein levels in high-grade gliomas (N=21) than those in low-grade gliomas (N=18). Error bars indicate SEM. * $p < 0.05$, ** $p < 0.01$. (D) Scatter dots distribution shows the change trend of NGB/ β -actin in different grade gliomas with PPAR γ / β -actin. (E) Immunofluorescence staining with NGB antibody shows an increase in NGB signaling (green, indicated by arrows) in high-grade gliomas; Bar, 50 μ m. (F) Immuno-histochemical staining shows the expression pattern of PPAR γ in different grade of gliomas. Bar, 100 μ m. Dark brown indicates PPAR γ . (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article).

sitol 3,4,5-triphosphate (PIP $_3$) and AKT [31]. We have noticed decreased PTEN protein levels in high-grade gliomas, where the PPAR γ was suppressed, compared with low-grade gliomas (Fig. 2A and B). Consistent with this, the phosphorylation of AKT (S473 and T308) exhibited a robust increase in high-grade gliomas (Fig. 2A and B), contributing to the proliferation and survival of glioma cells [32]. To investigate whether PPAR γ modulates the expression of NGB, in vitro experiments were performed. We found that the mRNA and protein levels of NGB were suppressed after the treatment of typically PPAR γ agonist, troglitazone and pioglitazone, accompanied by increased PTEN expression and decreased phosphorylation of AKT (Fig. 2C, D and E). Moreover, the expression of NGB exhibited robust increase under the insult of PPAR γ inhibitor GW9662 (Fig. 2F, G and H). Previous reports have shown that PPAR γ could negatively regulate gene expression after its activation [33,34], and our results display the same regulation effect of NGB expression through PPAR γ activation.

3.4. NGB regulates the phosphorylation of AKT in glioma cells

Recent studies have shown that AKT/PKB is able to regulate cell survival by monitoring the activities of transcriptional factors, such as

FOXO family, CREB and YAP, which are responsible for pro- and anti-apoptotic genes [35]. To figure out whether the increased NGB expression is due to enhanced phosphorylation of AKT, we stimulated AKT pathway through IGF-1 treatment, which resulted in an enhanced phosphorylation of AKT, but neither the mRNA nor protein levels of NGB increased under this stimulation in U87 cells (Fig. 3A, B and C). Furthermore, we transfected HA-AKT plasmid into U87 cells, but no changes of NGB protein levels were observed (Fig. 3D and E). Researchers have reported that NGB could attenuate tau hyper-phosphorylation and promote neurite outgrowth through the regulation of AKT phosphorylation [36,37]. We have noticed an increased phosphorylation of AKT in U87 cells, which were transfected with NGB plasmids, suggesting that the activation of AKT pathway in glioma may partly result from robust enhanced expression of NGB (Fig. 3F and G, Fig. 1B). Meanwhile, the expression of NGB could partly reverse the inhibition of AKT activity induced by pioglitazone treatment (Fig. 3F and G). All these indicate that NGB could regulate the phosphorylation of AKT in glioma cells, thus the enhanced expression of NGB may contribute to the activation of AKT pathway in glioma and promote glioma progression.

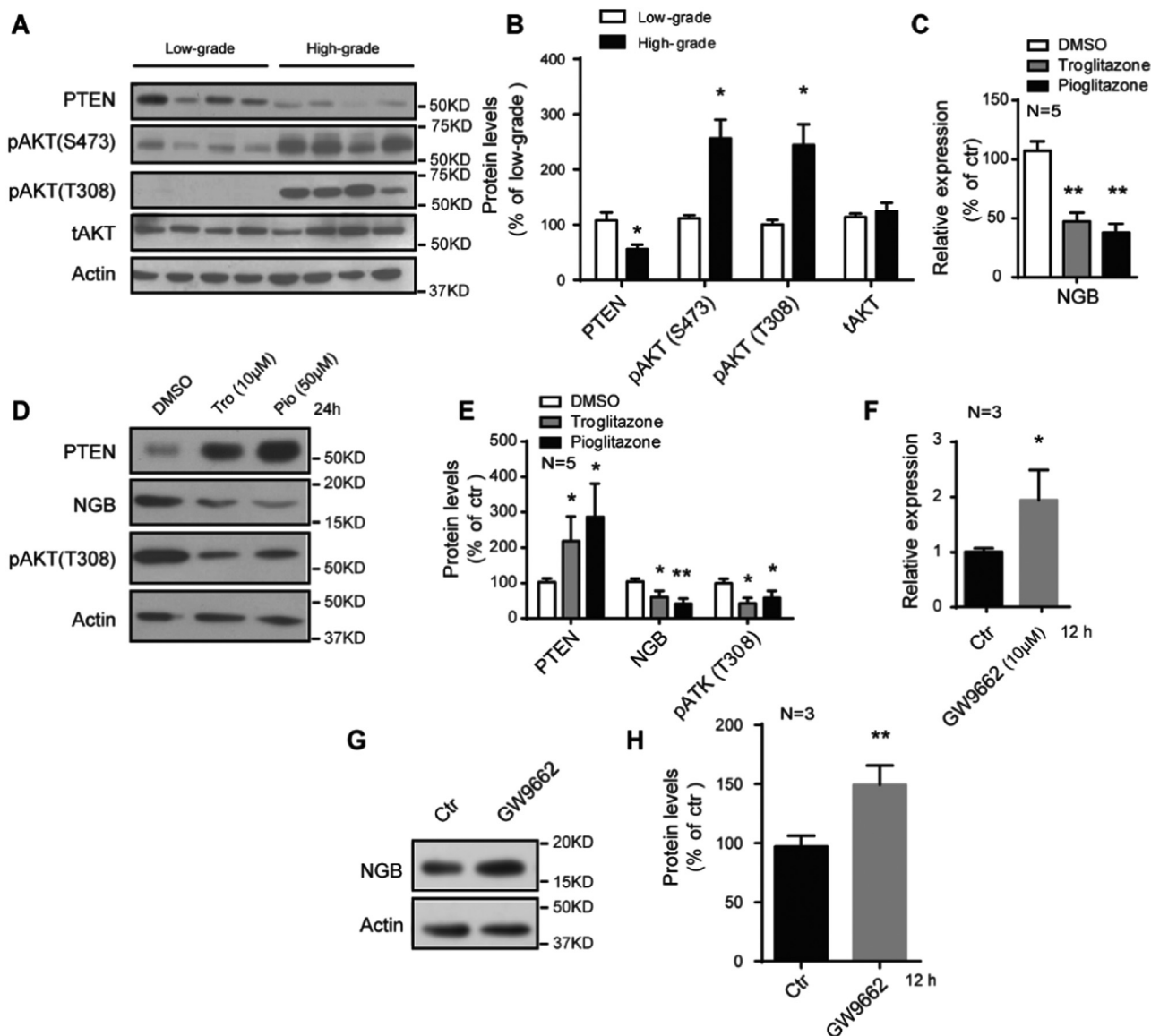


Fig. 2. PPAR γ negatively regulates the expression of NGB. (A and B) Representative images of western blots and quantification show decreased PTEN and increased phosphorylation of AKT in high-grade gliomas (N = 21) than those in low-grade gliomas (N = 18). Error bars indicate SEM. *p < 0.05, **p < 0.01. (C) qRT-PCR shows a suppressed expression of NGB mRNA after troglitazone and pioglitazone treatment in U87 cells. Error bars indicate SEM. *p < 0.05, N = 5. (D and E) Representative images of western blots and quantification show repressed NGB protein levels, accompanied by increased PTEN and decreased phosphorylation of AKT after troglitazone and pioglitazone treatment in U87 cells. Error bars indicate SEM. *p < 0.05, **p < 0.01, N = 5. (F) qRT-PCR shows an enhanced expression of NGB mRNA after GW9662 treatment in U87 cells. Error bars indicate SEM. *p < 0.05, N = 3. (G and H) Representative images of western blots and quantification show increased NGB protein levels after GW9662 treatment in U87 cells. Error bars indicate SEM. **p < 0.01, N = 3.

3.5. NGB protects glioma cells from oxidative stress and functions against PPAR γ 's pro-apoptotic and anti-proliferative abilities

High metabolic rate of cancer cells results in generating more ROS, which have been validated in multiple tumors [9,38]. We have found a suppression in the ratio of NADPH/NADP in high-grade gliomas, indicating oxidative stress resulting from redox imbalance (Fig. 4A). 4-Hydroxy-2-nonenal (4-HNE) is an end product formed by the reaction of ROS with polyunsaturated fatty acids during oxidative stress, the accumulation of which will cause apoptosis [39,40]. Larger amounts of 4-HNE accumulation were detected in high-grade gliomas than the low-grade ones (Fig. 4B and C), but they didn't trigger cell death. This phenotype may be attributed to the protection from the enhanced NGB expression, because we have noticed that the NGB overexpression could protect U87 cells from 4-HNE induced cell death (Fig. 4F). Previous works have shown that PPAR γ agonist could induce glioma cells death and repress the tumor progression [14,15]. Our work indicated that

NGB overexpression could resist this process through mitigating the ROS levels in U87 cells treated with pioglitazone (Fig. 3D and E). These findings display the protection effect of NGB involved in oxidative stress and its antagonistic effect against PPAR γ 's pro-apoptotic and anti-proliferative abilities.

4. Discussion

ROS, a double-edged sword for cancer cells, are critical for high metabolic rate maintenance, survival, and cellular migration, but toxic to cell integrity. Therefore, antioxidant mechanisms responded for cellular redox balance are much more robust in cancer cells rather than those in normal cells. Glutathione (GSH), distributed among all living organisms, exhibits detoxification effects of xenobiotics and some endogenous oxidative compounds either spontaneously or enzymatically with Glutathione peroxidase (GSH-Px) and Glutathione S-transferase (GSTs) [41]. Elevated GSH levels are observed in various types of

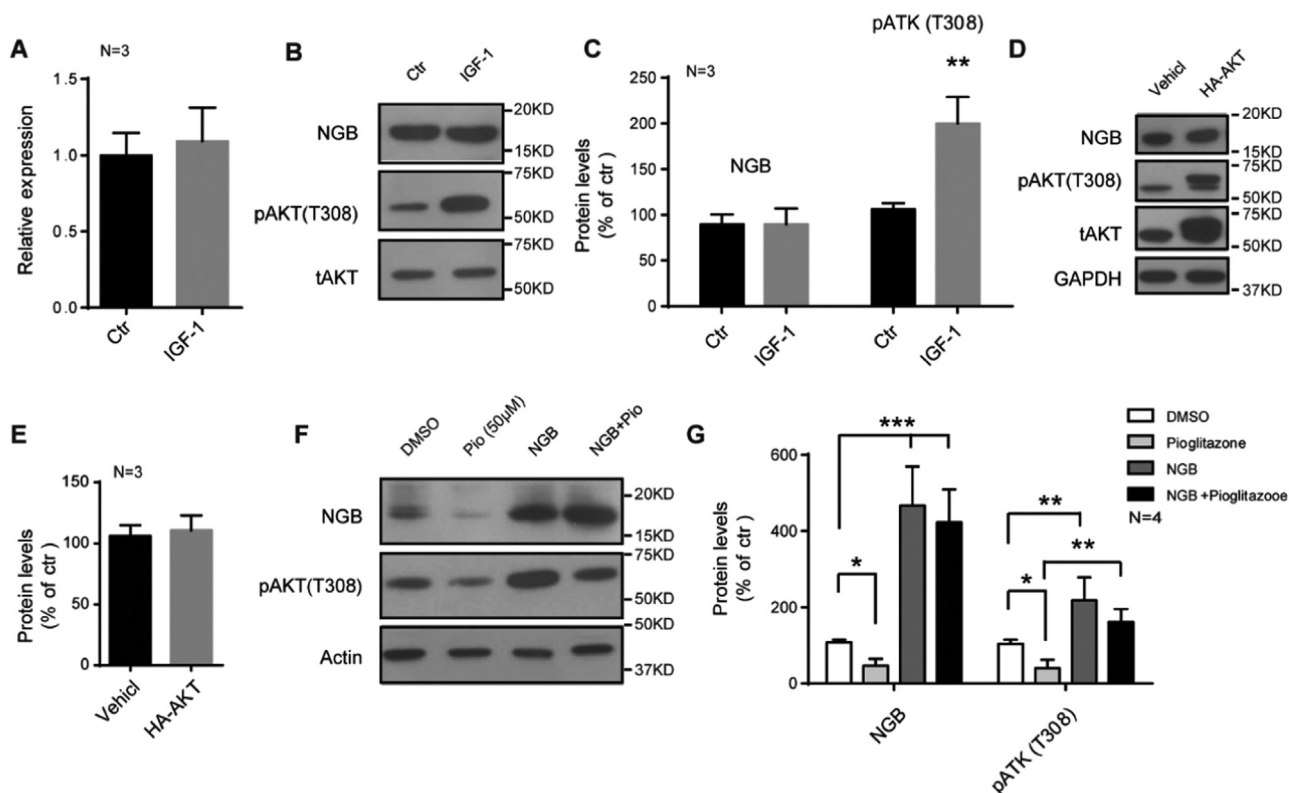


Fig. 3. NGB regulates the phosphorylation of AKT in glioma cells. (A) qRT-PCR shows no changes of NGB mRNA levels after IGF-1 (100 ng/ml) treatment for 6 h in U87 cells. Error bars indicate SEM. N=3. (B and C) Representative images of western blots and quantification show no changes of NGB protein levels after IGF-1 (100 ng/ml) treatment for 6 h in U87 cells. Error bars indicate SEM, **p < 0.01. N=3. (D and E) Representative images of western blots and quantification show no changes of NGB protein levels after the overexpression of HA-AKT in U87 cells. Error bars indicate SEM. N=3. (F and G) Representative images of western blots and quantification show an increased phosphorylation of AKT in NGB transfected U87 cells, and the expression of NGB could reverse the inhibition of AKT activity induced by pioglitazone treatment. Error bars indicate SEM. *p < 0.05, **p < 0.01, ***p < 0.01. N=4.

tumors, and this makes the neoplastic tissues more resistant to chemotherapy [42,43]. Superoxide dismutases (SODs), important antioxidant enzymes responsible for the elimination of superoxide radical, are highly expressed in ovarian cancer [44], colorectal cancer [45] and lung cancer [46]. NGB is a hypoxic and ischemic induced protein, which may function as antioxidant to scavenge reactive oxygen (ROS) and nitrogen species (RNS) [21,47]. We have shown that the expression of NGB in glioma is grade-related, reflecting its important biological role in glioma progression. Accumulating oxidative pressure and lipid oxidation increase the risk of apoptosis, while the enhanced NGB expression may protect against these insults in glioma progression. We have found the imbalance of redox signals, accompanied with increase 4-HNE accumulation, develops more severe in high-grade gliomas, where apoptosis has not been triggered. Meanwhile, in vitro study indicates that NGB overexpression protects glioma cells from 4-HNE induced cell death. Furthermore, we have noticed that the expression of NGB may participate in regulating AKT phosphorylation in glioma cells, suggesting that the protection effects of NGB may function through either itself or AKT related pathways.

PPAR gamma has been implicated in multiple senescence and senescence-related diseases such as inflammation [48], obesity [49], diabetes [50], and various cancers [30]. The applications implicated with PPARγ in cancer treatment are widely studied, but the results are paradoxical and the mechanisms underline are not well defined [51]. In glioma, PPARγ activation induced by its agonists effectively blocks the glioma progression [14], and some clinical studies show that PPARγ agonists exhibit synergistic anti-tumor effect in patients with high-grade glioma [52–54]. Proliferative advantage over normal tissue represents important hallmarks of cancers cells. One possible mechanism by which PPARγ agonists could inhibit glioma cell proliferation is the induction of cell-cycle arrest in G0/G1 phase and reduction of the

proportion of cells entering S phase [55,56]. Another widely described response of glioma cells to PPARγ agonists is the induction of apoptosis, which may be mediated by Bax-dependent mechanisms, as Bax up-regulation could be caused by PPARγ activation [14,57]. Redox signals are critical to cancer cells, and PPARγ is broadly involved in these processes. Several antioxidant genes (Catalase, MnSOD, GPX, HO-1 etc) are reported to be upregulated after PPARγ activation in normal cells [58], but there exhibits a different situation in glioma cells which were treated with agonists and failed to exhibit an increase in catalase expression and/or activity [25]. Therefore, more work need to be done to understand the exact role of PPARγ in redox balance in glioma. In this report, we have shown that the expression of PPARγ decreased along with the grade of glioma, suggesting its antagonism effect on glioma. PPARγ activation results in excessive ROS accumulation in glioma cells, which finally triggered cell death. Furthermore, our data indicate that the expression of NGB is negatively regulated by PPARγ, which was different from other antioxidant genes. And, NGB over-expression attenuates the toxicity of PPARγ agonists to glioma cells. In general, the expression of PPARγ is repressed during the glioma progression, resulting in enhanced expression of NGB, which further protects glioma cells against accumulating oxidative pressure. These results provide a new perspective to understand the functions of PPARγ and NGB in glioma progression, and identify NGB as a potential target for glioma treatment.

Competing interests

The authors declare that there are no competing interests to disclose.

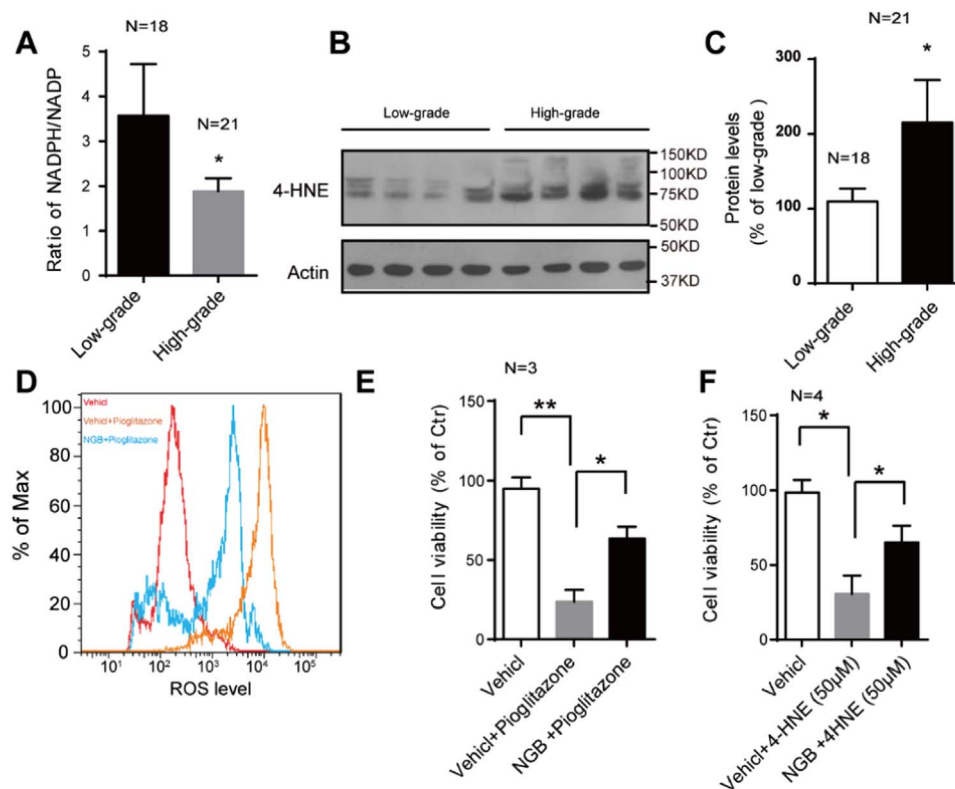


Fig. 4. NGB protects glioma cells from oxidative stress. (A) Quantification shows the ratio of NADPH/NADP in low-grade gliomas (N = 18) and the high-grade ones (N = 21). Decreased ratio is indicative of oxidative stress state. Error bars indicate SEM, *p < 0.05. (B and C) Representative images of western blots and quantification show increased 4-HNE accumulation in high-grade gliomas (N = 21) than that in low-grade ones (n = 18). Error bars indicate SEM. *p < 0.05, **p < 0.01. (D) Pioglitazone treatment causes an increase in ROS levels in U87 cells, while NGB overexpression mitigates this ROS accumulation. (E) Quantification shows overexpression of NGB functions against PPAR γ 's pro-apoptotic and anti-proliferative abilities in pioglitazone treated U87 cells. Error bars indicate SEM. *p < 0.05, **p < 0.01. N = 3. (F) Quantification shows overexpression of NGB could protect U87 cells against 4-HNE induced U87 cell death. Error bars indicate SEM. *p < 0.05. N = 4.

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