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# **Hypoxia promotes non-small cell lung cancer cell stemness, migration, and invasion via promoting glycolysis by lactylation of SOX9**

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#### **ABSTRACT**

**Background:** Lung cancer is the deadliest form of malignancy and the most common subtype is non-small cell lung cancer (NSCLC). Hypoxia is a typical feature of solid tumor microenvironment. In the current study, we clarified the effects of hypoxia on stemness and metastasis and the molecular mechanism.

**Methods:** The biological functions were assessed using the sphere formation assay, Transwell assay, and XF96 extracellular flux analyzer. The protein levels were detected by western blot. The lactylation modification was assessed by western blot and immunoprecipitation. The role of SOX9 in vivo was explored using a xenografted tumor model.

**Results:** We observed that hypoxia promoted sphere formation, migration, invasion, glucose consumption, lactate production, glycolysis, and global lactylation. Inhibition of glycolysis suppressed cell stemness, migration, invasion, and lactylation. Moreover, hypoxia increased the levels of SOX9 and lactylation of SOX9, whereas inhibition of glycolysis reversed the increase. Additionally, knockdown of SOX9 abrogated the promotion of cell stemness, migration, and invasion. In tumor-bearing mice, overexpression of SOX9 promoted tumor growth, and inhibition of glycolysis suppressed tumor growth.

**Conclusion:** Hypoxia induced the lactylation of SOX9 to promote stemness, migration, and invasion via promoting glycolysis. The findings suggested that targeting hypoxia may be an effective way for NSCLC treatment and reveal a new mechanism of hypoxia in NSCLC.

#### **Introduction**

<span id="page-0-5"></span><span id="page-0-4"></span>Lung cancer is still the first cause of cancer-related death globally. The most common type non-small cell lung cancer (NSCLC) accounts for 85% of lung cancer.<sup>1</sup> The diagnosis of NSCLC is delayed due to the lack of obvious symptoms in the early stage.<sup>[2](#page-8-1)</sup> Over the past years, molecular targeted therapy and immunotherapy have significantly improved NSCLC clinical outcomes. $3,4$  $3,4$  Although the therapy strategies are developing, the 5-year survival rate of NSCLC at advanced stage remains poor.<sup>5</sup> Therefore, novel therapeutic methods of NSCLC are urgently required.

<span id="page-0-9"></span><span id="page-0-8"></span><span id="page-0-7"></span><span id="page-0-6"></span>Growing evidence has confirmed that the tumor microenvironment (TME) promotes cancer development. Soluble factors, enriched in TME, induce abnormal proliferation, metastasis, angiogenesis, and drug resistance of tumor cells.<sup>6</sup> Hypoxia is a typical characteristic of TME in almost all solid tumors.<sup>7</sup> The rapid growth of tumor cells leads to a decrease in oxygen levels in the surrounding area. Chronic hypoxia can induce necrosis of tumor cells around blood vessels. However, once the tumor cells adapt to the hypoxic environment, the <span id="page-0-10"></span>tumor cells will become more aggressive and drug-resistant.<sup>[8](#page-8-7),[9](#page-8-8)</sup> In addition, hypoxia is associated with the phenotypes of cancer stem cells (CSCs) and induces epithelial-mesenchymal transition (EMT) and chemoradiotherapy resistance of CSCs.[10,](#page-8-9)[11](#page-8-10) However, the underlying mechanisms of hypoxia are complex and remain largely unclear.

<span id="page-0-13"></span><span id="page-0-12"></span><span id="page-0-11"></span>SRY-related high mobility group-box 9 (SOX9), a transcription factor, is a member of the SOX family. SOX9 is highly conserved and regulates embryonic development and determines sex.<sup>[12](#page-8-11)</sup> SOX9 is overexpressed in NSCLC and related to poor prognosis. SOX4 promotes cellular prolifera-tion, invasion, and EMT of NSCLC.<sup>[13,](#page-8-12)[14](#page-8-13)</sup> Additionally, the expression levels of SOX9 in TME are regulated by noncoding RNA, RNA methylation, and protein modifications such as phosphorylation and acetylation.<sup>[15](#page-8-14)</sup> However, whether hypoxia could regulate SOX9 in NSCLC is still unknown.

<span id="page-0-14"></span>In this study, we aimed to investigate the effects of hypoxia on NSCLC cell stemness and metastasis. We found that hypoxia promoted the stemness, migration, and invasion of NSCLC cells by mediating SOX9 lactylation. The findings identified a new

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Non-small cell lung cancer; hypoxia; metastasis; stemness; SOX9; lactylation

mechanism of hypoxia in NSCLC and suggested that targeting hypoxia might be a novel treatment for NSCLC.

## **Materials and methods**

## *Cell culture*

A549 and H838 cells (ATCC, USA) were cultured in Dulbecco's Modified Eagle Medium (DMEM; Gibco, USA) supplemented with 10% fetal bovine serum (FBS, Gibco, USA) and 1% penicillin/streptomycin in an incubator at  $37^{\circ}$ C with 5% CO<sub>2</sub>.

### *Hypoxia induction*

A549 and H838 cells were seeded into 6-well plates at the concentration of  $2 \times 10^5$  cells/well. The cells were incubated at 37°C with 1%  $O_2$  for 8, 16, and 24 h. The cells in the 0 h group were maintained in normoxia conditions (20%  $O_2$ ).

## *Cell transfection*

A549 and H838 cells were seeded into 6-well plates at the concentration of  $2 \times 10^5$  $2 \times 10^5$  cells/well and incubated for 24 h before transfection. Then, the cells were transfected with short hairpin RNA (Sh)-SOX9 and corresponding negative control (sh-NC) acquired from Genepharma (China) using Lipofectamine 2000 (Invitrogen, USA). 6 hours later, the medium was replaced with fresh medium. After 48 h, the transfected cells were harvested.

#### *2-deoxyglucose (2-DG) treatment*

<span id="page-1-0"></span>A549 and H838 cells were treated with 20 mM glycolysis inhibitor 2-DG (Sigma-Aldrich, USA) for 48 h as previously reported.<sup>[16](#page-8-15)</sup>

## *Sphere formation assay*

CSCs were sorted from A549 and H838 cells using the spherocyte medium. Briefly, A549 and H838 cells were digested and washed with PBS twice. The cells were seeded into 6-well ultralow cluster plates and incubated with DMEM/F12 without serum containing 20 ng/mL epidermal growth factor, 20 ng/ mL b-fibroblast growth factor, 2% B27, and 4 μg/mL insulin (all from Sigma-Aldrich) for 10 days. The spheres were observed under a microscope (Olympus, Japan) and the sphere formation rate was quantified.

#### *Transwell assay*

Cell migration was evaluated using 8 μm 24-well chambers (Corning, USA). The cell suspension containing  $2 \times 10^5$  cells was placed on the top chambers, and the DMEM with 10% FBS (500 μL) was filled with the bottom chambers. Following 24 h of incubation at 37°C with 5%  $CO<sub>2</sub>$ , the migrated cells were fixed with 4% paraformaldehyde for 20 min and stained with 0.1% crystal violet for 20 min. The stained cells were counted under a microscope (Olympus).

Cell invasion was assessed using the same chambers as used in cell migration detection but pre-coated with Matrigel (BD Biosciences, USA). The other operations were the same as that of cell migration mentioned above.

## *Determination of glucose consumption*

Glucose consumption was detected using a glucose uptake assay kit (Sigma-Aldrich) according to the manufacturer's instructions. The cells were lysed using lysis buffer on ice. The samples (50 μL) were incubated with 8 μL assay buffer and 2 μL enzyme mix for 60 min at 37°C. Next, glutathione reductase (20 μL), substrate-DTNB (16 μL), and recycling mix (2 μL) were added in each well. The absorbance was measured at 412 nm.

### *Determination of lactate production*

Lactate production was detected using a lactic acid assay kit (Sigma-Aldrich) following the manufacturer's protocol. Briefly, cells were homogenized in lactate assay buffer and centrifuged at 13,000  $\times$ g for 10 min. The supernatant (50 µL) was incubated with 50 μL reaction mix (containing 46 μL lactate assay buffer, 2 μL lactate enzyme mix, and 2 μL lactate probe) at room temperature for 30 min. The absorbance was measured at 570 nm.

### *Determination of ATP level*

Cellular ATP level was measured using an ATP assay kit (Beyotime, China) in line with the manufacturer's protocol. Briefly, A549 and H838 cells were lysed and the supernatant was collected after centrifuging at 12,000 ×g for 5 min. ATP detection working solution (100 μL) was mixed with 20 μL supernatant, and the results were detected using a luminometer.

## *Extracellular acidification rate (ECAR) and oxygen consumption rate (OCR) analysis*

A549 and H838 cells at the density of  $1.2 \times 10^4$  cells/well were seeded into XF96 cell culture plates and incubated until 90% confluence. For ECAR and OCR analysis, the culture medium was replaced by the assay medium without bicarbonate provided by Seahorse Bioscience (USA). Then, the ECAR and OCR were assessed using the XF96 extracellular flux analyzer (Seahorse Bioscience). For ECAR analysis, glucose, oligomycin A, and 2-DG were automatically added in turn, and the results were reported as mpH/min. For OCR analysis, oligomycin A, carbonyl cyanide 4-(trifluoromethoxy)phenylhydrazone (FCCP), and antimycin A and rotenone (rote/AA) were automatically added in turn, and the results were reported as pmol/min.

#### *Western blot analysis*

Total proteins were isolated from cells in RIPA lysis buffer. Following detecting the protein concentration using a BCA kit (Pierce, USA), 30 μg proteins were separated by 10% sodium dodecyl sulfate – polyacrylamide gels (SDS-PAGE). Then, the

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Figure 1. Hypoxia promotes cell stemness, migration, and invasion of NSCLC. (a) Sphere formation of CSCs sorted from A549 and H838 cells. (b) The protein levels of the markers of cell stemness including NANOG and CD133. (c) The represent images of cell migration under a microscope. The migration rate was quantified. (d) The represent images of cell invasion under a microscope. The invasion rate was quantified. \**p*<.05, \*\**p*<.01, and \*\*\**p*<.001 vs. 0 h group.

proteins were transferred onto PVDF membranes. The membranes were incubated with specific primary antibodies at 4°C overnight followed by incubating with the secondary antibody. Protein bands were visualized with ECL western blot substrate (Pierce). The antibodies were listed as follows: anti-NANOG (ab109250, 1/3000, Abcam, USA), anti-CD133 (ab222782, 1/ 2000, Abcam), anti-GAPDH (ab181602, 1/10000, Abcam), anti-SOX1 (ab109290, 1/1000, Abcam), anti-SOX2 (ab97959, 1/1000, Abcam), anti-SOX3 (ab183606, 1/2000, Abcam), anti-SOX9 (ab185230, 1/5000, Abcam), anti-SOX12 (23939–1-AP, 1/500; Proteintech, USA). The secondary antibody was goat anti-rabbit IgG conjugated HRP (ab6721, 1/3000, Abcam). Anti-Pan-Histone lysine lactylation (Kla; PTM1401, 1/1000, PTM Biolabs, China) was used to detect total lactylation protein levels.

#### *Immunoprecipitation (IP)*

A549 and H838 cells were lysed using the pre-cold IP lysis buffer. After centrifugation, the supernatant was collected to incubate with anti-SOX1 (ab109290, 1/50, Abcam), anti-SOX2 (ab133337, 1/10, Abcam), anti-SOX3 (ab183606, 1/100, Abcam), anti-SOX9 (ab185230, 1/60, Abcam), anti-SOX12 (23939–1-AP, 1/20, Proteintech) at 4°C overnight. Meanwhile, the protein A/ G-agarose beads were suspended with lysis buffer containing protease inhibitor. These suspensions were added to the samples in the next day to incubate for 4 h at 4°C.

## *Quantitative real-time polymerase chain reaction (qRT-PCR)*

Total RNA was extracted by the TRIzol reagent (Invitrogen) and 1 μg RNA was used to perform reverse transcription to obtain cDNAs using the SuperScript™ IV first-strand synthesis system (ThermoFisher Scientific, USA). qPCR was carried out using the SYBR Green PCR Mastermix (Solarbio life science, China) and the cDNA as templates. The relative expression of SOX9 was calculated using the 2<sup>-ΔΔCT</sup> method. The specific primers of SOX9 were: F 5ʹ-CAAGAAGGACCACCCGGATT -3ʹ and R 5ʹ-AAGATGGCGTTGGGGGAGAT-3ʹ. The specific primers of the loading control ACTB were: F 5ʹ-CCTCTCCCAAGTCCACACAG-3ʹ and R 5ʹ- GGGCACG AAGGCTCATCATT-3ʹ.

#### *Animal study*

The nude mice were randomly allocated into four groups (6 mice per group): vector, vector + 2-DG, SOX9, and SOX9 + 2-DG. A549 cells were transfected with SOX9 overexpressing vectors and empty vectors and adjusted cell density at  $5 \times 10^{7}$  $5 \times 10^{7}$  $5 \times 10^{7}$ cells/mL. The mice were subcutaneously injected with 100 μL transfected cells to establish a tumor-bearing mouse model. When the tumors reached about  $50 \text{ mm}^3$ , mice were

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Figure 2. Hypoxia promotes glycolysis in NSCLC cells. (a) The glucose consumption, (b) lactate production, and (c) ATP level in A549 and H838 cells. (d) The glycolytic capability of NSCLC cells was evaluated using ECAR curve. (e) The respiration was evaluated by OCR curve. (f) The lactylation proteins in A549 and H838 cells after normoxia (n) and hypoxia (h) treatment. \**p*<.01 vs. normoxia group.

<span id="page-3-1"></span>intraperitoneal injected with 500 mg/kg/day 2-DG, $^{17}$  once every 2 days during the experiment. Tumor volume was measured using a vernier caliper every week and quantified using the formula: Volume  $(mm^3)$  $(mm^3)$  $(mm^3)$  =(length×width<sup>[2](#page-8-1)</sup>/2. After the fourth measurement of tumor volume, the mice were sacrificed. The tumors were isolated from all mice.

#### *Immunohistochemistry (IHC) assay*

Tumor tissue paraffin sections (4 μm) were incubated with anti-SOX9 (ab185230, 1/2000, Abcam) at 4°C overnight followed by incubating with the secondary antibody (ab6721, 1/ 1000, Abcam) at room temperature for 0.5 h. Then, the sections were stained with DAB solution for 3 min at room temperature. After washing using moving water and sealing, the images were visualized under a microscope.

## *Statistical analysis*

Data analysis was performed using GraphPad Prism 8.0 software. The results are shown as mean  $\pm$  SD. The comparisons between two groups were assessed using Student's t-test and the comparisons among multiple groups were assessed using one-way analysis of variance. *p* < .05 indicated statistically significant.

#### **Results**

## *Hypoxia promotes cell stemness, migration, and invasion of NSCLC*

To explore the biological functions of hypoxia, we evaluated sphere formation of CSCs and metastasis of NSCLC cells. Compared with normoxia conditions, sphere formation rate was increased by hypoxia in a time-dependent way [\(Figure 1A](#page-2-0)). The protein levels of NANOG and CD133 were upregulated in a time-dependent manner ([Figure 1B\)](#page-2-0). Additionally, the capability of cellular migration and invasion of NSCLC cells was facilitated by hypoxia in a time-dependent manner ([Figure 1C,D](#page-2-0)). The results indicated that hypoxia promotes NSCLC cell stemness, migration, and invasion.

### *Hypoxia promotes glycolysis in NSCLC cells*

Subsequently, we explored the effects of hypoxia on glycolysis. A549 and H838 cells treated with normoxia and hypoxia for 48 h were used. The results showed that hypoxia promoted

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Figure 3. 2-DG reverses the cell stemness, migration, invasion, and glycolysis induced by hypoxia. After hypoxia and 2-DG treatment, (A, B) sphere formation of CSCs sorted from A549 and H838 cells. (C) The protein levels of NANOG and CD133. (D) The represent images of cell migration under a microscope. The migration rate was quantified. (E) The represent images of cell invasion under a microscope. The invasion rate was quantified. (F) The total lactylation proteins in A549 and H838 cells. (G) The glucose consumption in A549 and H838 cells. (G) The lactate production in A549 and H838 cells. (H) The ATP production in A549 and H838 cells. (J) The glycolytic capability of NSCLC cells was evaluated using ECAR curve. (K) The respiration was evaluated by OCR curve. \**p*<.01 and \*\**p*<.001 vs. hypoxia group.

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**Figure 4.** Hypoxia induces lactylation of SOX9. (a) The protein levels of SOX1, SOX2, SOX3, SOX9, and SOX12 in A549 and H838 cells under normoxia and hypoxia conditions. (b) The lactylation levels of SOX1, SOX2, SOX3, SOX9, and SOX12 in A549 and H838 cells under normoxia and hypoxia conditions. (c) The protein levels and lactylation levels of SOX9 in NSCLC cells treated with hypoxia and 2-DG.

glucose consumption, lactate production, and ATP level [\(Figure 2A–C\)](#page-3-0). The results of ECAR curve showed that the glycolysis capability was higher in the hypoxia group than that in the normoxia group ([Figure 2D](#page-3-0)). As shown in [Figure 2E](#page-3-0), hypoxia reduced OCR, compared with normoxia, suggesting hypoxia inhibited aerobic respiration. Additionally, the total lactylation proteins were increased after hypoxia treatment ([Figure 2F](#page-3-0)). The results indicated that hypoxia promotes NSCLC cell glycolysis.

## *2-DG reverses the cell stemness, migration, invasion, and glycolysis induced by hypoxia*

To investigate the effects of glycolysis on the biological functions of NSCLC cells, 2-DG was treated to suppress glycolysis. 2-DG decreased sphere formation of hypoxiainduced cells ([Figure 3A,B](#page-4-0)). Moreover, 2-DG downregulated NANOG and CD133 protein levels, compared with hypoxia treatment ([Figure 3C](#page-4-0)). Cell migration and invasion of hypoxia-induced cells were suppressed by 2-DG ([Figure 3D,E](#page-4-0)). As compared with hypoxia, 2-DG reduced the total lactylation of cells [\(Figure 3F](#page-4-0)). Besides, 2-DG abrogated the increase of glucose consumption, lactate production, and ATP level induced by hypoxia ([Figure 3G–I](#page-4-0)). The glycolysis capability was suppressed and anaerobic respiration was promoted by 2-DG treatment in hypoxia-induced cells [\(Figure 3J,K](#page-4-0)). The results showed that suppression of glycolysis inhibited cell stemness, migration, and invasion of NSCLC.

### *Hypoxia induces lactylation of SOX9*

The molecular mechanism of hypoxia was further investigated. The results of western blot showed that hypoxia elevated the protein levels of SOX1, SOX2, SOX3, SOX9, and SOX12, compared with normoxia ([Figure 4A\)](#page-5-0). However, hypoxia only increased the lactylation levels of SOX9, but did not affect the lactylation levels of SOX1, SOX2, SOX3, and SOX12 ([Figure 4B\)](#page-5-0). Moreover, 2-DG downregulated the protein levels and lactylation levels of SOX9 ([Figure 4C](#page-5-0)). The results illustrated that hypoxia promoted SOX9 lactylation modification by promoting glycolysis.

## *Knockdown of SOX9 abrogates cell stemness, migration, and invasion*

To explore the role of SOX9, A549 and H838 cells were transfected with sh-SOX9. As compared with sh-NC, the mRNA and protein levels of SOX9 were downregulated after transfection with sh-SOX9 [\(Figure 5A,B](#page-6-0)). In hypoxiatreated cells, knockdown of SOX9 inhibited sphere formation and downregulated the levels of NANOG and CD133 [\(Figure 5C–E\)](#page-6-0). The migration and invasion were both inhibited by SOX9 knockdown in hypoxia-induced cells [\(Figure 5F–I\)](#page-6-0). The results illustrated that hypoxia promotes cell stemness, migration, and invasion of NSCLC via mediating SOX4 levels.

## *2-DG rescued the promotion of tumor growth induced by SOX9*

Finally, we established the tumor-bearing mouse model to explore the role of 2-DG and SOX9 *in vivo*. The results showed that 2-DG reduced tumor size and weight, whereas SOX9 increased tumor size and weight, compared with the vector group. 2-DG reversed the effects induced by SOX9 [\(Figure 6A–C](#page-7-0)). The lactylation level was decreased by 2-DG and was elevated by SOX9. 2-DG rescued SOX9-mediated lactylation level [\(Figure 6D\)](#page-7-0). The protein levels of SOX9 were downregulated by 2-DG and upregulated by SOX9 overexpression. Moreover, the levels were reversed by 2-DG, compared with the SOX9 group ([Figure 6E\)](#page-7-0). Taken together, SOX9 accelerated tumor growth in glycolysis glycolysis-dependent manner.

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**Figure 5.** Knockdown of SOX9 abrogates cell stemness, migration, and invasion of NSCLC cells. (a) Transfection efficiency of SOX9 following transfection with sh-NC and sh-SOX9 was measured using qRT-PCR. (b) The efficiency SOX9 following transfection was measured using western blot. (c, d) sphere formation of CSCs sorted from A549 and H838 cells. (e) The protein levels of NANOG and CD133. (f, g) the represent images of cell migration under a microscope. The migration rate was quantified. (h, i) the represent images of cell invasion under a microscope. The invasion rate was quantified. \**p*<.001 vs. sh-NC group in (a). \*\**p*<.001 vs. hypoxia +sh-NC group in (d-i).

## **Discussion**

<span id="page-6-3"></span><span id="page-6-2"></span><span id="page-6-1"></span>Advanced NSCLC is accompanied by distal metastasis of the tumor, resulting in cancer-related death. Thus, inhibition of tumor metastasis can improve clinical outcomes of NSCLC. Hypoxia is the main characteristic of TME in solid cancers. It regulates tumor growth, angiogenesis, metabolism, immune response, and drug resistance.<sup>18</sup> On one hand, hypoxia inhibits tumor cell growth due to lack of oxygen; however, on the other hand, hypoxia promotes cell invasion and then accelerates malignant advancement.<sup>[19](#page-9-2)</sup> CSCs regulate self-renewing and differentiation, and hypoxia promotes self-renewal of CSCs.<sup>20</sup>

Herein, we performed sphere formation, western blot, and Transwell assay to analyze NSCLC cellular processes, and found that hypoxia promoted cell stemness, migration, and invasion in a time-dependent manner, suggesting that inhibition of hypoxia could decelerate the progression of NSCLC. Thus, targeting hypoxia may be a novel treatment for NSCLC.

<span id="page-6-4"></span>Glycolysis is a key pathway to provide energy for tumor cells and maintain tumor growth. Otto Warburg observed that cancer cells metabolize glucose through the glycolytic pathway and then convert it to lactic acid, terming Warburg effect.<sup>[21](#page-9-4)</sup> The Warburg effect makes cancer cells have more powerful

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**Figure 6.** 2-DG rescued the promotion of tumor growth induced by SOX9. (a) The represent images of tumors from mice of each group. (b) Tumor volume was measured weekly. (c) Tumor weight. (d) The lactylation levels in tumor tissues. (e) IHC assessed SOX9 levels in tumors. \**p*<.001 and \*\**p*<.01 vs. The vector group. \*\*\**p*<.01 vs. The SOX9 group.

<span id="page-7-2"></span><span id="page-7-1"></span>antioxidant capacity and promotes their metastasis.<sup>[22](#page-9-5)</sup> Hypoxic TME is an important factor to enhance glycolysis in cancer cells. Under hypoxic conditions, tumor cells maintain ATP levels through glycolysis to ensure their survival and growth. $^{23}$  In the current study, we used commercial kits to detect glucose, lactate, and ATP, and used seahorse assay to assess ECAR and OCR. We found that hypoxia promoted glucose consumption, lactate production, ATP level, and gly-colysis, consistent with previous studies.<sup>[24,](#page-9-7)25</sup> Moreover, inhibition of glycolysis by 2-DG abrogated the promotion of cell stemness, migration, and invasion induced by hypoxia, and also suppressed tumor growth *in vivo*. The data suggested that hypoxia facilitated NSCLC cell stemness and metastasis by enhancing glycolysis.

<span id="page-7-5"></span><span id="page-7-4"></span><span id="page-7-3"></span>Lactic acid is a metabolite of glycolysis that acts as a substrate to support cell growth and regulate biological functions. Histone lysine lactylation (Kla) is a newly identified epigenetic modification, providing potential targets for cancer therapy.<sup>26</sup> Lactylation is involved in numerous diseases, such as inflammation, fibrosis, and malignancy.<sup>27</sup> Hypoxia induces glycolysis to produce lactic acid, stimulating histone lactylation. $^{28}$  In this study, we explored that hypoxia increased the global lactylation levels in NSCLC cells through western blot, consistent with the previous study.<sup>28</sup> Moreover, inhibition of glycolysis reduced lactylation, suggesting that hypoxia promoted lactylation by promoting glycolysis.

<span id="page-7-7"></span><span id="page-7-6"></span>At present, accumulating evidence has shown that multiple proteins have Kla sites except for histones. They widely exist in the nucleus, mitochondria, and cytoplasm.<sup>29</sup> Previous studies have revealed that hypoxia regulates cancer progression by <span id="page-7-10"></span><span id="page-7-9"></span><span id="page-7-8"></span>lactylation of several proteins. For example, Qiao et al.<sup>30</sup> have identified that hypoxia facilitates glycolysis and stemness of esophageal cancer cells by SHMT2 lactylation. Miao et al.<sup>31</sup> have reported that hypoxia lactylates β-catenin promotes colorectal cancer cell proliferation and stemness. In this study, we focused on the proteins of the SOX family, which are involved in tumor initiation and progression. $32$  In view of the role of glycolysis and lactylation, we speculated that SOX proteins could be undergone by lactylation modification. IP and western blot were performed to detect the protein and lactylation levels of the members in the SOX family. The results showed that hypoxia promoted lactylation of SOX9, and did not affect the lactylation of the other members of the SOX family. Moreover, inhibition of glycolysis downregulated the protein levels of SOX9 and the lactylation of SOX9. SOX9 is closely related to the progression of tumors. Importantly, SOX9 is involved in regulation of tumor cellular processes including glycolysis. SOX9 facilitates glycolysis in multiple cancers, such as gastric cancer,  $33$  colorectal cancer,  $34$  and NSCLC.<sup>35</sup> Therefore, we evaluated the role of SOX9 in NSCLC. Rescue experiments indicated that knockdown of SOX9 inhibited cell stemness, migration, and invasion in hypoxia-treated NSCLC cells. In tumor-bearing mice, overexpression of SOX9 accelerated tumor growth, accompanied by an increased lactylation level, and 2-DG rescued the promotion. The findings suggested that hypoxia promoted lactylation of SOX9, and knockdown of which inhibited the progression of NSCLC.

<span id="page-7-11"></span>There are several limitations in this study. We only investigated the effects of hypoxia and SOX9 on stemness, migration, and invasion; however, their effects on other phenotypes, such as apoptosis, angiogenesis, immune escape, and drug resistance, remain unknown. Moreover, the underlying mechanism of hypoxia may be complex, including but not limited to SOX9 lactylation. We will further study this in our future work.

In conclusion, hypoxia facilitated the lactylation modification of SOX9 to promote cell stemness, migration, and invasion of NSCLC by enhancing glycolysis. Moreover, overexpression of SOX9 accelerated tumor growth through inducing glycolysis. The results suggested that inhibition of hypoxia is a novel treatment for NSCLC. How to translate the conclusions into clinical application still needs a lot of research.

#### **Disclosure statement**

No potential conflict of interest was reported by the author(s).

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## **Data availability statement**

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request.

### **Ethical approval**

No ethics approval was required for this study as it involved no human participants or animals.

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