HIF-1α affects trophoblastic apoptosis involved in the onset of preeclampsia by regulating FOXO3a under hypoxic conditions

ZHAN ZHANG, CHENXI HUANG, PING WANG, JUNJUN GAO, XIN LIU, YINGYING LI, SHUJUN YAN and YING SHI

Departments of 1Clinical Laboratory, and 2Obstetrics and Gynecology,
The Third Affiliated Hospital of Zhengzhou University, Zhengzhou, Henan 450052, P.R. China

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Abstract. Preeclampsia (PE) is a pregnancy-specific syndrome that has severe implications on perinatal mortality and morbidity. Excessive apoptosis of trophoblasts induced by hypoxia may be associated with the development of PE, but the exact pathogenesis is unknown. Forkhead box O transcription factor 3a (FOXO3a) is activated under hypoxic conditions. Furthermore, hypoxia-inducible factor-1α (HIF-1α) is sensitive to variations in partial oxygen pressure. Thus, the aims of the present study were to investigate the expression levels of HIF-1α and FOXO3a in placental samples of early onset severe PE, and their effect on trophoblastic apoptosis under hypoxic conditions. Cobalt chloride was used to establish the hypoxic model. The present study examined the expression levels of HIF-1α and FOXO3a in the placental tissues and HTR8/SVneo cells under hypoxic conditions. It was found that HIF-1α and FOXO3a were highly expressed in placental tissues of patients with PE and in HTR8/SVneo cells under hypoxic conditions. Furthermore, knockdown of FOXO3a using a specific small interfering RNA (siRNA) decreased apoptosis in HTR8/SVneo cells. Moreover, it was found that after knockdown of HIF-1α using siRNA, FOXO3a expression and the apoptotic rate were reduced in HTR8/SVneo cells. Therefore, the present results indicated that the elevated expression of HIF-1α increased trophoblastic apoptosis by regulating FOXO3a, which may be involved in the pathogenesis of PE.

Introduction

Preeclampsia (PE) is a pregnancy-specific complication with a global incidence of 4-5% (1). PE is characterized by the development of hypertension with or without proteinuria, after 20 weeks of gestation (2). While, PE leads to the morbidity and mortality of mothers and perinatal infants, its pathogenesis has not been fully elucidated (1,3). The placenta is crucial for the development of PE (4). It has been shown that placental hypoxia is considered to be the main factor contributing to the pathogenesis of PE, and is associated with excessive apoptosis of trophoblasts, which results in decreased trophoblast invasion and insufficient spiral artery remodeling (5,6).

Hypoxia-inducible factor 1 (HIF-1) is a transcriptional factor that helps maintain oxygen homeostasis and can react quickly to low oxygen tension (7). Furthermore, it is a heterodimer containing two subunits, α and β (8). HIF-1α is an oxygen-regulated subunit that responds to changes in cellular oxygen, while HIF-1β is a constitutively expressed subunit (8). Under normal levels oxygen HIF-1α is degraded, and under hypoxic conditions degradation of HIF-1α is inhibited, thus resulting in rapid accumulation of the protein (9). Moreover, HIF-1α is critical for placental development, and prolonged expression of HIF-1α causes pregnancy-associated disorders in placental trophoblasts (10). In PE, placental hypoperfusion and ischemia can result in a hypoxic microenvironment, which induces the expression of HIF-1α (11). Previous studies have demonstrated that HIF-1α can participate in cell apoptosis by regulating the expression of Forkhead box O transcription factor 3a (FOXO3a) (12). FOXO3a is a member of the FOXO transcription factor family, and four FOXO family members have been identified in humans: FOXO1, FOXO3a, FOXO4 and FOXO6 (13). Together these transcription factors control various biological functions such as cellular metabolism, cell cycle regulation, apoptosis and regulation of stress response (14). Moreover, previous studies have found that hypoxia significantly increases the expression of FOXO3a, and decreases the phosphorylation of Akt and FOXO3a, thus resulting in increased nuclear accumulation (14,15). Hu et al (16) revealed that FOXO3a is a downstream effector of HIF-1α and is activated by hypoxia. Furthermore, it has been shown that knockdown of FOXO3a increases apoptosis of human umbilical vein endothelial cells (HUVECs) cells under hypoxia (16).

The present study investigated the expression levels of HIF-1α and FOXO3a in placental tissues of patients with early onset severe PE, and examined its effect on trophoblastic apoptosis under hypoxia.

Key words: preeclampsia, hypoxia, apoptosis, hypoxia-inducible factor-1α, forkhead box O transcription factor 3a
Materials and methods

Case selection. Patients were recruited for the study between May 2017 and December 2018 at The Third Affiliated Hospital of Zhengzhou University. In total, 30 women (mean age, 32.90±5.41 years) with early onset severe PE were chosen as the experimental group and 30 women (mean age, 32.45±4.66 years) with a normal pregnancy constituted the control group. Women who were from the Chinese Han population selected cesarean sections. Inclusion and exclusion criteria for early onset severe PE were strictly based on guidelines of the American College of Gynecologists, Task Force on Hypertension, published in 2013 (17). The exclusion criteria included multi-fetal pregnancies, gestational diabetes mellitus, chronic hypertension, connective tissue diseases and smoking. The study was approved by The Ethics Committee of The Third Affiliated Hospital of Zhengzhou University, and informed consent was obtained from all the patients. Detailed clinical information of patients in the two groups is shown in Table I.

Sample collection. The biopsies were separated from the maternal aspect of the placenta after delivery. Regions with calcification, necrosis and infarction were not collected. Blood in the tissues was removed using sterile filter paper. Specimens were fixed with 10% buffered formalin for 24 h at room temperature and embedded in paraffin at room temperature to be used for immunohistochemistry (IHC). The remaining samples were immediately stored at -80°C for RNA and protein extraction.

IHC staining. Placental tissues were cut into 4 µm sections for IHC analysis. The tissue sections were heated to 60°C for 2 h and deparaffinized using xylene, and sequentially rehydrated using a series of graded ethanol (100, 95, 85 and 75%) for 5 min at room temperature. This was followed by microwave oven heating to a boil in 10 mM citrate buffer (pH 6.0; Invitrogen; Thermo Fisher Scientific, Inc.) for 15 min to achieve antigen retrieval. Tissues were incubated with 3% H2O2 for 15 min at 37°C to suppressed endogenous peroxidase activity. Then, sections were incubated with a rabbit anti-human FOXO3a monoclonal antibody (1:800; cat. no. 12829S; Cell Signaling Technology, Inc.), overnight at 4°C. Negative controls were treated for 2 h with 10 mM PBS following the same method. Then, tissues were incubated with a biotin-conjugated secondary antibody (1:200; cat. no. SP-9001; OriGene Technologies, Inc.) for 1 h at room temperature. The product obtained using a 3,3'-diaminobenzidine tetrahydrochloride substrate kit (ZSGB-BIO) was observed for 2-5 min at room temperature. Counterstaining of the sections were performed using 0.1% hematoxylin for 5 min at room temperature. The staining of the sections were independently evaluated by two pathologists, and was based on the estimated staining intensity scale (18) of 0-3: i) 0, No staining and 0-5%, positive staining; ii) 1, buff staining and 6-25% positive staining; iii) 2, pale brown staining and 26-75% positive staining; and iv) 3, sepia staining and 75-100% positive staining. Light microscopy images were captured at x200 magnification. The immunohistochemical score was the positive percentage multiplied by staining intensity, and was defined as: 0, negative; <4, weakly positive; 4-8, positive; >8, strong positive.

Cell culture and treatment. The HTR8/SVneo cell line (American Type Culture Collection) was incubated with DMEM at high glucose (HyClone; GE Healthcare Life Sciences), supplemented with 10% FBS (Biological Industries), 100 U/ml ampicillin and 100 U/ml streptomycin at 37°C in 5% CO2-humidified incubators. HTR8/SVneo cells were inoculated in 6-well plates (1x104 cells/well). When cells had grown to reach a fusion of 60%, they were treated with 0, 125, 250 and 500 µmol/l cobalt chloride (cat. no. e8661; Sigma-Aldrich; Merck KGaA) for 0, 24, 48 and 72 h at 37°C. After hypoxia treatment, cellular proteins were extracted using RIPA lysis buffer (Beijing Solarbio Science & Technology Co., Ltd.) and protease inhibitor, and the optimal concentration and time were determined using western blot analysis, as described below. In the present study, 250 µmol/l cobalt chloride was selected as the concentration and 48 h as the duration of hypoxic condition for follow-up experiments.

Cell viability assay. HTR8/SVneo cells were inoculated into 96-well plates (5x103 cells/well) and a Cell Counting Kit-8 (CCK-8; Dojindo Molecular Technologies, Inc.) assay was used to determine cell viability, according to the manufacturer's protocol. Briefly, when the cells had grown to reach a fusion of 60%, they were treated with 250 µmol/l cobalt chloride for 0, 24, 48 and 72 h at 37°C. Subsequently, 10 µl CCK-8 solution was added to each well and incubated at 37°C for 1 h. Absorption values were obtained using a microplate reader (Bio-Rad Laboratories, Inc.) at 450 nm.

Reverse transcription-quantitative PCR (RT-qPCR). Total RNA of tissues and HTR8/SVneo cells was extracted using TRIzol® reagent (Invitrogen; Thermo Fisher Scientific, Inc.), and RT into cDNA using a ReverTra Ace RT- qPCR kit (cat. no. 651600; Toyobo Life Sciences) under the following conditions: 37°C for 15 min, 50°C for 5 min and 98°C for 5 min. SYBR-Green Realtime PCR Master mix (cat. no. 722100; Toyobo Life Sciences) was used for specific gene amplification on a StepOnePlus RT PCR system (Applied Biosystems; Thermo Fisher Scientific, Inc.). The following thermocycling conditions were used: Initial denaturation at 95°C for 60 sec, followed by 40 cycles of amplification at 60°C for 15 sec and a final extension step at 72°C for 45 sec. The primers used for HIF-1α were: Forward, 5'-GCCGCTGGAGACACACAA TC-3' and reverse, 5'-TCCATCGGAAGGACTAGGTTG-3'. The primers used for FOXO3a were: Forward, 5'-GGTGGCTTCTCAAGCTCTCC-3' and reverse, 5'-AATGGCGCTT GAGTTCAAAAGG-3'. The primers used for β-actin were: Forward, 5'-GGGAAAATCTGCGTGACATTAAGG-3' and reverse, 5'-CAGGAGGAGGGCTGGAGATGTC-3'. All results were normalized to the expression of β-actin. The 2^ΔΔCq method was used to calculate the relative change of all the target genes (19).

Western blotting. RIPA lysis buffer and protease inhibitor (Beijing Solarbio Science & Technology Co., Ltd.) were used at a ratio of 100:1 to lyse the tissues and cells to collect the supernatant. Protein concentrations were measured using a bicinchoninic acid protein assay kit (Thermo Fisher Scientific, Inc.). Soluble proteins (40 µg) were separated using 8 and 12% SDS-PAGE gel and transferred onto PVDF membranes.
Membranes were then blocked with 5% non-fat milk in TBS-0.1% Tween-20 (TBST) for 2 h at room temperature and incubated with a rabbit anti-human HIF-1α (1:1,000; cat. no. ab51608; Abcam), rabbit monoclonal anti-FOXO3a (1:1,000; cat. no. 12829S; Cell Signaling Technology, Inc.) and rabbit polyclonal anti-β-actin (1:2,000; cat. no. ab8227; Abcam) overnight at 4°C. The membranes were incubated at room temperature for 1 h and washed three times for 10 min with TBST. Subsequently, membranes were incubated with a horse-radish peroxidase-conjugated secondary antibody (1:10,000; cat. no. iH-0012; Beijing dingguo changsheng Biotechnology Co., Ltd.) for 1 h at room temperature. The protein bands were visualized using an enhanced chemiluminescence Plus kit (Beyotime Institute of Biotechnology) and an AI600 imaging system (Fisher Scientific, Inc.). After 48 h of transfection, extraction of total RNA or protein of cells was assessed to confirm efficiency for subsequent experiments.

Small interfering RNA (siRNA) transfection. Specific siRNAs for HIF-1α (siHIF-1α; 5’-CAAATCAAGAAGTTGCAATT A-3’) and FOXO3a (siFOXO3a; 5’-GCACAGAGTGGATG AAGT-3’) were purchased from Guangzhou riboBio co., ltd. Then, 250 µl RNase-free water was used to the synthesize siRNAs until they reached a storage concentration of 20 µM. HTR8/SVneo cells were cultured in 6-well plates (1x10^4/well). Cells with 60% confluence were replaced with fresh culture serum-free medium 2 h prior to transient transfection. Using a final concentration of 50 nM siHIF-1α, siFOXO3a and negative control (siNC), 5 µl siRNAs were transfected into cells using 5 µl Lipofectamine 3000 reagent (Invitrogen; Thermo Fisher Scientific, Inc.) and incubated at 37°C for 24 h. Cells were digested using 1X TrypLE Express (Gibco; Thermo Fisher Scientific, Inc.) and resuspended in a 1X binding buffer (BD Biosciences). Then, the cells were double-stained with 20 µg/ml Annexin V-FITC and PI (BD Biosciences) for 15 min at room temperature. Cell apoptosis was determined using an Epics XL. MCL flow cytometer (Beckman Coulter, Inc.) and analyzed using FlowJo version 10.6 software (FlowJo LLC) (21).

Statistical analysis. All statistical analyses were performed using the SPSS 21.0 (IBM Corp) and GraphPad Prism 6.0 software (GraphPad Software, Inc.) (22). Data are presented as the mean ± SD, or the medians and interquartile ranges. Data between two of groups with normal distribution were compared using independent samples t-test, while data between two groups with a skewed distribution were compared using a Mann Whitney U test. The immunohistochemical staining intensity of FOXO3a expression levels were determined using a χ² test. A Kruskal Wallis test with a Bonferroni’s correction post hoc test was used to analyze multiple groups. P<0.05 was considered to indicate a statistically significant difference.
Results

HIF-1α and FOXO3a expression in the placental tissues of patients with PE is higher compared with healthy pregnant women. HIF-1α and FOXO3a at mRNA level in placental tissues were detected using RT-qPCR. It was found that the mRNA expression levels of HIF-1α and FOXO3a in the PE group were significantly increased compared with the healthy control group (Fig. 1A). HIF-1α and FOXO3a protein levels were analyzed using western blotting, and it was demonstrated that the protein expression levels of both HIF-1α and FOXO3a were increased in the PE group (Fig. 1B and C). Subsequently, the localization of FOXO3a expression in the placenta was assessed using IHC staining. The results suggested that the trophoblast expression of FOXO3a was located in the cytoplasm and nucleus (Fig. 1D). Moreover, the staining intensity of FOXO3a was higher in the PE placental tissues compared with healthy controls ($\chi^2=22.31; P<0.001$; Table II).

Cobalt chloride-induced hypoxia leads to increased protein expression levels of HIF-1α and FOXO3a in HTR8/SVneo cells. HTR8/SVneo cells were cultured in 0, 125, 250 and 500 µmol/l cobalt chloride. Western blot analysis results identified that the expression levels of HIF-1α and FOXO3a were significantly elevated after treatment with cobalt chloride (Fig. 2A-C). However, it was found that the expression of HIF-1α was decreased in 500 µmol/l cobalt chloride. Moreover, the protein expression of both HIF-1α and FOXO3a gradually increased with longer treatment periods with 250 µmol/l cobalt chloride. (Fig. 2D-F). The present results also indicated that cell viability was significantly decreased with increasing treatment duration, especially when treatment time was $>24$ h (Fig. 2G). Therefore, based on the cell viability at ~50%, 250 µmol/l cobalt chloride for 48 h was used to induce hypoxic conditions.

Hypoxia induces nuclear translocation of HIF-1α and FOXO3a, and increases the rate of apoptosis in HTR8/SVneo cells. In order to observe intracellular localization of HIF-1α and FOXO3a, HTR8/SVneo cells were cultured under normal and hypoxic conditions. Using cell immunofluorescence analysis, it was demonstrated that the expression of HIF-1α and FOXO3a was at low levels in the cytoplasm under normal conditions. Furthermore, nuclear translocation of these factors was significantly increased during hypoxia (Fig. 3A and B). Using flow cytometry analysis, it was identified that the rate

### Table II. Immunostaining of Forkhead box O transcription factor 3a in control and early onset preeclampsia groups.

| Immunohistochemical staining | Group | None | Weak | Moderate | Strong | $\chi^2$ | P-value |
|-----------------------------|-------|------|------|----------|--------|---------|---------|
| FOXO3a                      | Control (n=30) | 7    | 19   | 3        | 1      | 22.31   | <0.001  |
|                             | PE (n=30)    | 2    | 5    | 13       | 10     |         |         |

P<0.001 vs. control. FOXO3a, Forkhead box O transcription factor 3a; PE, preeclampsia.

![Figure 1](image-url)
of apoptosis of HTR8/SVneo cells was significantly enhanced during hypoxia (Fig. 3C and D). Therefore, the present results suggested that hypoxia may cause nuclear translocation of HIF-1α and FOXO3a in HTR8/SVneo cells, and augments apoptosis of trophoblasts.

Knockdown of FOXO3a suppresses hypoxia-induced apoptosis in HTR8/SVneo cells. In order to verify the knockdown efficiency of FOXO3a, the mRNA expression of FOXO3a after transfection with a siRNA was measured. Based on the results, siFOXO3a2 was used in follow-up experiments (Fig. 4A) to knockdown FOXO3a in HTR8/SVneo cells. Subsequently, western blot analysis results identified a decrease in the expression of FOXO3a after transfection with siRNA (Fig. 4B and C). However, flow cytometry analysis demonstrated that knockdown of FOXO3a suppressed trophoblastic apoptosis (Fig. 4D and E). Collectively, the present results suggested that knockdown of FOXO3a could suppress hypoxia-induced apoptosis of HTR8/SVneo cells.

Under hypoxia, HIF-1α affects the apoptosis of trophoblast cells by regulating the expression of FOXO3a in HTR8/SVneo cells. In order to measure the knockdown efficiency of HIF-1α, the mRNA expression of HIF-1α was detected after transfection with siRNA. Based on the results, siHIF-1α1 was used in follow-up experiments (Fig. 5A) to knockdown HIF-1α in HTR8/SVneo cells. It was found that the protein expression levels of HIF-1α and FOXO3a were significantly decreased following the transfection with siHIF-1α1 (Fig. 5B-D). Flow cytometry analysis indicated that knockdown of HIF-1α suppressed trophoblastic apoptosis (Fig. 5E and F). Furthermore, cell immunofluorescence results demonstrated that knockdown of HIF-1α decreased the expression and nuclear translocation of FOXO3a under hypoxic conditions (Fig. 5G and H). Therefore, the present results suggested that knockdown of HIF-1α repressed the expression of FOXO3a and the apoptosis of trophoblasts. Thus, it is speculated that HIF-1α affects apoptosis of trophoblasts by regulating the expression of FOXO3a under hypoxia in HTR8/SVneo cells.

Discussion

Previous findings have shown that oxygen tension may regulate cytotoxic trophoblast proliferation and differentiation, which affects the development of the placenta (23). At 8-10 weeks of gestation, the placenta undergoes a physiological hypoxic phase in which trophoblasts continue proliferating and are poorly differentiated (24,25). Afterwards, along with an increase in oxygen pressure, trophoblast cells begin to differentiate normally and invasion increases (25). When spiral artery remodeling is complete, adequate maternal blood supply can be provided to the placenta (26). However, long-term hypoxia causes an increase in the apoptosis of trophoblasts and a shallow cell invasion of the uterus, which can result in the development of PE (27). HIF-1α is a master regulator of oxygen homeostasis and can regulate diverse cellular functions (28). It has been
Figure 3. HIF-1α and FOXO3a nuclear translocation and increased apoptotic rate are associated with hypoxia. Cell immunofluorescence analysis of the expression levels of (A) HIF-1α and (B) FOXO3a (magnification, x400). These proteins are at low levels in the cytoplasm under normal conditions, while nuclear translocation of HIF-1α and FOXO3a increased significantly during hypoxia. (C) Flow cytometry analysis found that the apoptotic rate of HTR8/SVneo cells was significantly increased under hypoxia. (D) Quantification of the flow cytometry results. Data are presented as the medians and interquartile ranges of four independent experiments. *P<0.05. FOXO3a, Forkhead box O transcription factor 3a; HIF-1α, hypoxia-inducible factor-1α.

Figure 4. Knockdown of FOXO3a suppresses hypoxia-induced apoptosis of HTR8/SVneo cells. (A) mRNA expression of FOXO3a after transfection with the siRNA was detected using reverse transcription-quantitative PCR. Data are presented as the medians and IQRs of three independent experiments. (B) After transfection with siRNA, western blotting results demonstrated that the expression of FOXO3a was significantly decreased. (C) Data are presented as medians and IQRs of three independent experiments. (D) Flow cytometry analysis found that knockdown of FOXO3a suppressed trophoblastic apoptosis. (E) Data are presented as medians and IQRs of four independent experiments. *P<0.05, **P<0.01, ***P<0.001. FOXO3a, Forkhead box O transcription factor 3a; siRNA, small interfering RNA; IQRs, interquartile ranges; NC, negative control.
revealed that the expression of HIF-1α increases in placental tissues in PE (29). Previous findings have also shown that under hypoxic conditions, the increase in HIF-1α expression initiates hypoxia-mediated apoptosis by affecting the expression of downstream molecules (30). This is consistent with the present results, in which the expression of HIF-1α was found to be elevated. The FOXO transcription factor family are thought to be tumor suppressors that mainly regulate the cell cycle, apoptosis, DNA-damage repair and response to oxidative stress (31,32). FOXO3a is a member of the FOXO family, and its expression is enhanced under hypoxia (33). The present results suggested that the expression of FOXO3a was significantly enhanced in placental tissues in PE. Moreover, previous studies have found that HIF-1α can affect the apoptosis of cardiomyocytes and neurons by regulating the expression of FOXO3a (33,34). Therefore, it is hypothesized that HIF-1α may influence the involvement of hypoxia-induced apoptosis in the development of PE by regulating the expression of FOXO3a.

The present study used cobalt chloride to construct a hypoxic model. Furthermore, cobalt chloride-induced hypoxia is able to cause a high expression level of HIF-1α (35). It has also been shown that the expression of HIF-1α is increased in HTR8/SVneo cells under hypoxia and in placental tissues in PE (36). Therefore, the present study investigated the effect of HIF-1α on downstream gene expression and cell functions under hypoxic conditions.

FOXO3a is upregulated under hypoxic conditions, and translocation of FOXO3a from the cytoplasm to the nucleus is induced by serum starvation and hypoxia (33). Hypoxia-activated FOXO3a can also promote the apoptosis of cardiomyocytes (37). A previous study showed that the siRNA knockdown of FOXO3a reduces the protein expression levels of FOXO3a and Bim, as well as inhibiting the apoptosis of...
HUVECs (16). In the present study, it was also found that hypoxia could increase the expression levels of FOXO3a and the rate of apoptosis in HTR8/SVneo cells, whereas inhibiting the expression levels of FOXO3a with siRNA reduced the rate of hypoxia-induced trophoblast apoptosis. Under hypoxic conditions, the HIF-1α subunit becomes stable and interacts with coactivators, such as p300/CREB binding protein, to promote its nuclear transcriptional activity (38). Moreover, the present results suggested that the nuclear expression of HIF-1α was elevated under hypoxia. In addition, suppression of HIF-1α using siRNA is able to decrease the expression of FOXO3a, hypoxia-induced reactive oxygen species accumulation and apoptosis of HUVECs (16). Based on the present result that knockdown of FOXO3a attenuated the apoptosis rate in HTR8/SVneo cells, it is speculated that the hypoxia-induced increase in the expression of HIF-1α and the decrease in trophoblast apoptosis caused by HIF-1α knockdown may be regulated by the expression of FOXO3a.

In conclusion, under hypoxia, elevated expression of HIF-1α leads to an increase in trophoblastic apoptosis via the regulation of FOXO3a, which may be involved in the decrease filtration ability observed in the pathogenesis of PE.

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Availability of data and materials
All data generated or analyzed during this study are included in this published article.

Authors' contributions
CH and ZZ conceived and designed the study, CH, PW, JG and XS performed the experiments and analyzed the data, XL, YL and SY performed sample and data acquisition, CH and ZZ wrote the manuscript. All authors read and approved the final manuscript.

Ethics approval and consent to participate
The study protocol was approved by the Ethics Review Committee of the Third Affiliated Hospital of Zhengzhou University and informed consent was obtained from all the patients (ID no. 2015023).

Patient consent for publication
All patients within the present study provided consent for the publication of their data.

Competing interests
The authors declare that they have no competing interests.

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