Extracellular Hsp90α, which participates in vascular inflammation, is a novel serum predictor of atherosclerosis in type 2 diabetes

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ABSTRACT

Introduction Atherosclerosis is the main pathological change in diabetic angiopathy, and vascular inflammation plays an important role in early atherosclerosis. Extracellular heat shock protein 90 (eHsp90) is secreted into the serum and is involved in various physiological and pathological processes. However, the specific mechanism of eHsp90 in early atherosclerosis remains unclear. This study explored the relationship between Hsp90 and diabetic lower extremity arterial disease and investigated the expression of eHsp90 in vascular endothelial cells under environmental stimulation and the function and mechanism of eHsp90α in involved in diabetic atherosclerosis.

Research design and methods One hundred and three selected patients were divided into three groups: the diabetes mellitus group (n=27), the diabetic lower extremity arterial disease group (n=46), and the diabetic critical limb ischemia group (n=30). The relationships among serum Hsp90, oxidative stress indexes, and patient outcomes and the correlations among the indexes were analyzed. H&E staining and immunohistochemistry were used to observe the vasculature of amputated feet from patients with diabetic foot. An oxidative stress endothelial injury model was established under high glucose in vitro to explore the role of eHsp90 release in atherosclerosis progression.

Results The level of serum Hsp90 was upregulated with aggravation of diabetic vascular disease. Hsp90α was correlated with malondialdehyde to some extent and was an independent risk factor in the progression of diabetic vascular disease, with predictive ability. The expression area of Hsp90α was consistent with the area of inflammatory infiltration in the vessel lumen. Vascular endothelial cells were found to increase eHsp90α secretion under stress. Then inhibition of eHsp90α can reduce the degree of cellular inflammation and damage. Endothelial cell-conditioned medium and recombinant human Hsp90α increased monocyte migration via the low-density lipoprotein receptor-related protein 1 (LRP1) receptor to promote disease progression.

Conclusions eHsp90α plays a critical role in the early inflammatory injury stage of atherosclerosis.

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INTRODUCTION

Diabetes mellitus (DM) has become a global public health problem. Approximately 4.2 million people die each year from DM or its complications.1 Lower extremity arterial disease (LEAD) is the most common initial manifestation of cardiovascular disease in individuals with type 2 diabetes mellitus (T2DM).2 Up to 80% of people with T2DM die from cardiovascular complications, and the disease reduces the average life expectancy by
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approximately 10 years. Therefore, early prevention and diagnosis of LEAD is particularly important in patients with diabetes.

The initial pathogenesis of atherosclerosis (AS) includes the formation of advanced glycosylation end products, excessive production of reactive oxygen species (ROS), and other processes. The different pathogenic mechanisms induced by hyperglycemia all reflect a single upstream process, namely overproduction of ROS. Ross proposed that AS is a pathological process characterized by a chronic inflammatory response. When the levels of metabolic abnormalities are increased, superoxide can also activate and participate in the pathogenesis of various complications. However, there is no marker reflecting the current states of the vascular endothelium and the inflammatory response that can be applied for early clinical detection, which makes early diagnosis difficult.

Heat shock protein 90 (Hsp90) is an important molecular chaperone that is widely involved in immune regulation. Lei et al. proposed that Hsp90 is involved in the pathogenesis of diabetic angiopathy. In recent years, Hsp90 inhibitors have been found to have protective effects against diabetic angiopathy. For example, Lazaro et al. reported that application of an Hsp90 inhibitor (Alvespimycin (17-DMAG)) effectively reduced renal injury in a diabetic mouse model. Kim et al. found that inhibiting Hsp90 can reduce the formation of AS by inhibiting the migration and proliferation of vascular smooth muscle cells (VSMCs). Madrigal-Matute et al. found that Hsp90 is highly expressed in the thin fibrous cap area of human carotid atherosclerotic plaques. All the above findings suggest that Hsp90 may be a new target for AS therapy. Extracellular Hsp90 (eHsp90) has been reported to be involved in a variety of pathological processes and plays a different biological role than intracellular Hsp90. Our team found that eHsp90 can be released outside the cell through exosomes under stress conditions, which in turn affects cytoskeletal proteins to promote the migration of epithelial cells, participating in damage repair. Although Hsp90 has been indicated to be correlated with diabetic vascular disease, previous studies have focused more on its role as a molecular chaperone. Whether eHsp90 is involved in vascular inflammation or fibrosis is still unclear.

In this study, in vivo serological studies and in vitro cell models were used to explore the direct relationship between serum Hsp90 concentrations and subclinical AS in patients with T2DM and the role of eHsp90 in diabetic vascular disease in an oxidative stress injury model. The findings will provide a new perspective for investigation into early therapeutic targets for diabetic vascular disease.

RESEARCH DESIGN AND METHODS

Study subjects

From August 30, 2019 to May 28, 2021, a total of 103 patients at the Endocrinology and Metabolism Department of the Nanfang Hospital of Southern Medical University were randomly enrolled in this study. The patients were divided into three groups: the diabetes mellitus group (DM, 27 patients), the diabetic lower extremity arterial disease group (DLEAD, 46 patients), and the diabetic critical limb ischemia group (DCLI, 30 patients). The subjects were grouped according to their clinical presentations and imaging studies.

The following were the inclusion criteria:

- **DM group:**
  - T2DM was diagnosed according to the WHO 1999 criteria.
  - Screening for LEAD was performed in patients without a diagnosis of LEAD.

- **DLEAD group:** patients were diagnosed with T2DM according to the above DM diagnostic criteria and LEAD was diagnosed according to the International Working Group on the Diabetic Foot (IWGDF) 2019 guidelines on the basis of the following:
  - Symptoms and signs of AS (intermittent claudication, resting pain, decreased or absent pulse in the dorsal foot artery, etc).
  - An ankle-brachial arterial pressure index (ABI) <0.9, a toe-brachial arterial pressure index <0.7, no three-phase foot pulse graph waveform, or a percutaneous oxygen partial pressure <30 mm Hg.

- **DCLI group:** DCLI was diagnosed in patients who met the above diagnostic criteria for DLEAD and who had lower extremity ischemic infection, ulceration and/or deep tissue destruction for more than 4 weeks, and an ABI <0.4.

The patients and/or their families were informed and agreed to participate in the study. Patients were excluded if they met any of the following exclusion criteria before admission:

- History of diabetic ketoacidosis or hyperosmolar status within 30 days.
- History of diabetic coma within 3 months or any hypoglycemic event within 1 month.
- History of cancer, immune system disease, or any hypoglycemia event within 1 month.
- History of other types of DM.
- Age less than 18 years old.

Sample collection

Blood samples (3 mL) were collected under aseptic conditions from the peripheral veins of the subjects. Serum samples were obtained via cold centrifugation at 3000 revolutions per minute (rpm) for 15 min. The clear supernatants were stored at −80°C until they were used for biochemical analysis.
Assessment of Hsp90α and Hsp90β

The levels of serum Hsp90α and Hsp90β were measured using ELISA kit (JingKang Bio, China). All procedures were performed according to the manufacturer’s instructions. Absorbance was measured at a wavelength of 450 nm.

Histopathology and immunohistochemistry

Blood vessels were obtained from amputated tissue discarded during surgery from three patients with DCLI, immersed in paraformaldehyde for at least 24 hours for fixation, paraffin-embedded, and cut into 2 μm sections. The samples were stained with HE for histopathological assessment. Immunohistochemistry for Hsp90α was performed.

Cell culture

Human umbilical vein endothelial cells (HUVECs) and the human monocyte line THP-1 were obtained from the China Center for Type Culture Collection. HUVECs were cultured in DMEM/F12 (Gibco, USA), and THP-1 cells were grown in Roswell Park Memorial Institute (RPMI) 1640 medium supplemented with 10% fetal bovine serum (FBS) (ExCell Bio, China) and 1% penicillin/streptomycin in a 37°C incubator with a humidified atmosphere and 5% CO₂.

Antibodies and reagents

The primary antibodies included Hsp90α (07-2174; Merck, USA), Hsp90β (SPC-177; StressMarq Biosciences, Canada), low-density lipoprotein receptor-related protein 1 (LRP1) (BS9805M; Bioworld Technology, USA), p-Akt (664444g; Proteintech, USA), Akt (#2920S; CST, USA), and β-actin (RM2001; RayBiotech, China).

The primary reagents included recombinant human Hsp90α protein and recombinant human Hsp90β protein (SPR-101C and SPR-102C; StressMarq, Canada) and 17-allylamo-17-demethoxygeldanamycin, (17AAG) (HY10211; MCE, USA). A non-commercial antisericeted Hsp90α monoclonal antibody, 1G6-D7, was kindly donated by the University of Southern California Keck School of Medicine (USA).

Cell viability assay

Cell viability was evaluated with a Cell Counting Kit-8 (CCK-8) assay. Briefly, cells were seeded into a 96-well plate at a density of 5×10⁴ cells/well and grown to 80% confluence. After treatments, the medium was replaced with fresh medium containing 10% CCK-8 reagent. Absorbance at 450 nm was measured after a 2-hour incubation at 37°C. Triplicate wells were included for each group.

Detection of MDA content and SOD activity

Following treatment, supernatants were collected and the levels of lipid peroxidation were determined with a micro-malondialdehyde (MDA) assay reagent kit and an superoxide dismutase (SOD) assay kit (KGT003 and KGT00150, Keygen Biotech, China) according to the manufacturer’s instructions.

Measurement of cellular and mitochondrial ROS

Intracellular changes in ROS generation were measured with an ROS assay kit (S0033; Beyotime, China). According to the manufacturer’s instructions, cells were incubated with 10 μM 2,7’-dichlorodihydrofluorescein diacetate (DCFH-DA) in a cell incubator with 5% CO₂ at 37°C for 20 min. The fluorescence intensity of the cells was then detected using a flow cytometer.

Mitochondrial ROS activity was measured with MitoSOX Red assay (M36008, Invitrogen). Cells were incubated with 5 mM MitoSOX Red probe for 10 min at 37°C. Red fluorescence was determined using an Olympus FV1000 confocal laser scanning microscope (Tokyo, Japan).

Protein extraction and protein expression analysis

HUVECs were plated in 10 cm dishes. After treatment, conditioned medium (CM) from the serum-free cultures was collected, centrifuged, and filtered through Millipore Amultra-4 (50K) column. The total cell lysates were centrifuged at 13,000×g for 15 min at 4°C, and the total protein concentration was determined using BCA Protein Assay Kit (Keygen, China).

Immunofluorescence

Cells were fixed with 4% paraformaldehyde for 10 min at room temperature. Permeation was performed by incubating the pretreated cells with ice-cold methanol at −20°C for 15 min. The cells were incubated with primary antibodies (1:100) at 4°C overnight. Images were captured using an Olympus FV1000 confocal laser scanning microscope.

mRNA expression analysis

Total cell RNA was isolated with RNAiso (Takara, China), and 1000 ng of total RNA were used for reverse transcription with PrimeScript RT Reagent Kit. Real-time quantitative PCR was performed using SYBR Premix Ex Taq and Premix Taq with the indicated primers (online supplemental table 1).

Measurement of 8-oxoG, ET-1, and MCP-1

Following treatment, cells and supernatants were collected and the levels of DNA damage were measured with an 8-oxoguanine (8-oxoG) ELISA kit (JingKang Bio, China). Endothelial cell injury markers were then detected with a human endothelin 1 (ET-1) ELISA kit (Cusabio, China) and a human monocyte chemotactic protein 1 (MCP-1) ELISA kit (JingKang Bio) according to the manufacturer’s instructions.

Cell migration assay

Twenty-four Transwell inserts with pore sizes of 3 μm (Corning, USA) were employed. A total of 10⁶ THP-1 monocytes in 200 μL of serum-free RPMI-1640 medium were loaded into the upper chamber of each Transwell insert. Then, 600 μL of endothelial cell CM were treated.
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with hydrogen peroxide (H₂O₂) and 17AAG, or RPMI medium was added to the lower chamber. The cells were then allowed to migrate for 8 hours. The cells that migrated across the membrane to the lower chamber were counted under an inverted microscope (Nikon Eclipse TS100, Japan).

Statistical analysis

All experiments were repeated three times. SPSS V.22.0 was used for statistical processing of the data. Differences between groups were analyzed by one-way analysis of variance, Bonferroni test, Kruskal-Wallis test (K-W test), and χ² test according to the characteristics of the data. Spearman correlation analysis was used to analyze the correlations of related indexes. A multivariate logistic regression model was used to identify potential independent risk factors. A receiver operating characteristic (ROC) curve was drawn to evaluate the predictive value. Significance level was at p<0.05.

RESULTS

Characteristics of the study subjects

This is a hospital-based, cross-sectional study. The baseline demographic and clinical information of the 103 patients (58 male, 56.3%; 45 female, 43.7%) included in this study is displayed in Table 1. All patients were considered to have T2DM as defined by the WHO 1999 criteria. Univariate analysis showed significant differences in age, course, hypertension, hyperlipidemia, family history, C reactive protein (CRP), cholesterol (CHOL), low-density lipoprotein (LDL), white cell count (WCC), and hemoglobin among the three groups (p<0.05).

| Parameter                  | Patients | F(χ²)  | P value |
|----------------------------|----------|--------|---------|
| n (%)                      | DM       | DLEAD  | DCLI    |
| Male/female (n)            | 27/13    | 46/18  | 16/14   |
| Age (years)                | 44.41±11.40 | 57.07±10.19 | 64.53±10.46 |
| BMI (kg/m²)                | 22.13±2.93 | 23.54±3.60 | 23.19±2.74 |
| Course (%)                 |          |        |         |
| <5 years                   | 23 (85.2)| 15 (32.6)| 8 (26.7)|
| 5–10 years                 | 2 (7.4)  | 11 (23.9)| 4 (13.3)|
| ≥10 years                  | 2 (7.4)  | 18 (39.1)| 17 (56.7)|
| Smoking (%)                | 7 (25.9)| 15 (32.6)| 6 (20.0)|
| Current                    | 5        | 14     | 5       |
| Former                     | 2        | 1      | 1       |
| Never                      | 20       | 31     | 24      |
| Hypertension (%)           | 7 (25.9)| 19 (41.3)| 18 (60.0)|
| Hyperlipidemia (%)         | 6 (22.2)| 23 (50.0)| 13 (43.3)|
| Family history (%)         | 3 (11.1)| 19 (41.3)| 5 (16.7)|
| 2hPBG (mM)                 | 10.51±4.41| 11.67±3.91| 9.75±5.12|
| GLU (mM)                   | 7.65 (5.23–14.46)| 7.11 (4.92–10.13)| 9.41 (6.39–14.97)|
| HbA1c (mmol/mol)           | 9.70 (7.21–12.41)| 9.25 (8.20–11.20)| 7.80 (7.10–9.50)|
| CRP (mg/L)                 | 1.12 (0.57–5.66)| 2.40 (0.85–7.54)| 21.35 (2.63–62.07)|
| TG (mM)                    | 1.40 (1.06–2.53)| 1.57 (0.94–2.44)| 1.61 (1.28–2.77)|
| CHOL (mM)                  | 4.87 (3.91–5.55)| 4.22 (3.22–5.24)| 3.77 (2.93–4.58)|
| HDL-C (mM)                 | 1.09 (0.82–1.30)| 0.94 (0.76–1.17)| 0.84 (0.72–0.97)|
| LDL-C (mM)                 | 3.09±0.88| 2.67±0.94| 2.43±1.05|
| WCC (×10⁹/L)               | 6.66 (5.31–9.02)| 7.61 (6.50–9.50)| 8.74 (7.21–11.71)|
| Hb (g/L)                   | 138.6 (128.5–145.5)| 127.5 (113.0–143.0)| 106.0 (85.0–118.5)|

Table 1 Baseline characteristics of the study subjects

Data are mean±SD or median (IQR).

BMI, body mass index; CHOL, cholesterol; CRP, C reactive protein; DCLI, diabetic critical limb ischemia; DLEAD, diabetic lower extremity arterial disease; DM, diabetes mellitus; GLU, glucose; Hb, hemoglobin; HbA1c, hemoglobin A1c; HDL-C, high-density lipoprotein cholesterol; 2hPBG, 2-hour postprandial blood glucose; LDL-C, low-density lipoprotein cholesterol; TG, triglyceride; WCC, white cell count.
Comparison and correlation analyses of serum levels of MDA, Hsp90α, Hsp90β, WCC, and CRP in the study subjects

The levels of MDA, Hsp90α, and Hsp90β in the three groups are shown in figure 1A–C. In pairwise comparisons, the levels of the above indicators were significantly higher in patients with DLEAD or DCLI than in patients with DM. The results showed that patients with T2DM were characterized by a state of continuous and chronic oxidative stress. The relationships of Hsp90α and Hsp90β with WCC, CRP, and MDA levels in the study subjects are shown in figure 1D–I. Pearson’s correlation analysis showed that there were significant correlations between Hsp90α and MDA levels (r=0.233, p=0.018).

Multivariate logistic regression model of risks associated with diabetic vascular disease

The results of the collinearity analysis of independent variables showed that triglyceride and CHOL were collinear (online supplemental table 2). The clinical variables with statistical significance that were identified in the univariate logistic regression analysis (online supplemental tables 3 and 4) were included in the multivariate logistic regression analysis (figure 2A). As depicted, age, Hsp90α, course, and family history were independent risk factors for DLEAD. Moreover, age, Hsp90α, Hsp90β, and course were independent risk factors for DCLI.

Diagnostic efficacy of Hsp90α and Hsp90β for DLEAD

ROC curve analysis was conducted to assess the diagnostic efficacy of Hsp90α and Hsp90β for DLEAD (figure 2B). Hsp90α showed better diagnostic performance in the DLEAD group than in the DM group. The area under the curve (AUC) of Hsp90α was 0.701 (61.84% sensitivity and 77.78% specificity) and the cut-off was 263.88 (Youden index: 0.396), while the AUC of Hsp90β was 0.660 (31.58% sensitivity and 96.29% specificity) and the cut-off was 325.88 (Youden index: 0.279).

Histopathological sections

Both inflammatory cell infiltration and elevated Hsp90α expression were present. Tissue sections prepared from human atherosclerotic lesions were subjected to HE staining (figure 2C,E) and immunohistochemistry (figure 2D,F). Inflammatory cells infiltrated the intima heavily in patients with DCLI. Hsp90α was abundantly expressed in the endothelium and smooth muscle. In addition, the expression area of Hsp90α was consistent with the area of inflammatory infiltration.

Vascular endothelial cell model of oxidative injury in vitro

The results of H2O2 concentration screening (figure 3A) indicated that after treatment with 300 μM H2O2 for 24 hours, the cell survival rate was nearly 80%. HUVECs exposed to H2O2 at 200 μM and 300 μM exhibited significantly different MDA contents (figure 3B) and SOD activities (figure 3C) than control cells. Cellular ROS (figure 3D) and mitochondrial ROS detection (figure 3E) indicated that the ROS levels of HUVECs treated with 300 μM H2O2 for 24 hours were increased. At the same time, the expression of the DNA damage marker 8-oxoG and endothelial cell damage factor ET-1 and MCP-1 was increased (figure 3F–I). These results suggest that 300 μM H2O2 treatment for 24 hours can successfully establish an oxidative stress injury model in endothelial cells.

Increased eHsp90 secretion in the vascular endothelial injury model

HUVECs were treated with H2O2 at a range of doses. eHsp90α and eHsp90β secretion increased, but the total amount of intracellular Hsp90 did not change significantly (figure 3I). After H2O2 treatment, the total amount of eHsp90α and eHsp90β from HUVECs was the highest at 24 hours (figure 3J). Treatment with H2O2 and 17AAG alone or in combination significantly increased the eHsp90 of endothelial cells. In addition, H2O2 had little effect on the expression of Hsp90AA1 and Hsp90AB1 (figure 3K,L). The immunofluorescence results (figure 3M) showed that Hsp90α localization shifted with treatment; specifically, the fluorescence at the cell edges increased, suggesting that the expression of Hsp90α in the membrane and periphery may have increased.

eHsp90α enhances the inflammatory injury state of endothelial cells

To explore the specific biological role of eHsp90 in the inflammatory injury model, endothelial cells were treated with human recombinant (hr)Hsp90 protein to assess the expression of related functional indicators. The results showed that eHsp90α increased the expression of ET-1 and MCP-1 in endothelial cell culture medium (figure 4A,B). However, the effect of eHsp90β on endothelial cell dysfunction was not significant. Under oxidative stress, 17AAG and 1G6-D7 were added to inhibit eHsp90α and the expression of ET-1 decreased, suggesting that endothelial cell functional damage was alleviated (figure 4C).

eHsp90 recruits monocytes through LRP1 receptor activation of p-Akt kinase

To study the effect of chronic oxidative stress on the development and progression of AS, we next examined the effect of endothelial cell CM on monocytes. Compared with the control medium, H2O2-treated CM induced THP-1 migration. CM from endothelial cells treated with 17AAG alone or H2O2 in combination exhibited a significantly decreased ability to induce THP-1 migration (figure 4D). We observed that hrHsp90α exposure increased the number of THP-1 cells that migrated to the lower chamber. The chemotactic effect of hrHsp90β on THP-1 monocytes was not obvious. We used 1G6-D7 to identify the specificity of Hsp90α for chemotactic THP-1 monocytes. The results showed that the chemotactic ability of monocytes was weakened by the addition of 17AAG or 1G6-D7 (figure 4E).

Next, we treated THP-1 cells with hrHsp90α and Western blotting showed that the LRP1 protein content was significantly elevated in the treated cells. To confirm
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Figure 1  Comparison and correlation analyses of serum levels of MDA, Hsp90α, Hsp90β, WCC, and CRP in the study subjects. Serum MDA (A), Hsp90α (B), and Hsp90β (C) levels were measured in the different groups. *P<0.05, **P<0.01, ***P<0.001, ****P<0.0001 versus the DM or the DLEAD group. Correlation between Hsp90α (D–F) and Hsp90β (G–I) and WCC, CRP, and MDA levels in serum samples. Pearson's coefficient tests were conducted to calculate statistical significance. CRP, C reactive protein; DCLI, diabetic critical limb ischemia; DLEAD, diabetic lower extremity arterial disease; DM, diabetes mellitus; Hsp90, heat shock protein 90; MDA, malondialdehyde; WCC, white cell count.
Figure 2  Multivariate logistic regression model and diagnostic efficacy of Hsp90α and Hsp90β for DLEAD. Histopathological changes in blood vessels. Multivariate logistic regression analysis (A): a forest plot obtained from a multiple logistic regression model as OR with 95% CI. Comparing patients with DLEAD and DM, age, Hsp90α, course, and family history were independent risk factors for the development of DLEAD. Comparing patients with DCLI and DM, age, Hsp90α, Hsp90β, and course were independent risk factors for DCLI. Receiver operating characteristic analysis of the diagnostic efficacy of Hsp90α and Hsp90β for DM and DLEAD (B). Diagnostic ability to distinguish patients with DLEAD from patients with DM. HE staining (C and E) and immunohistochemistry (D and F) (40×, 100×, 200×) of Hsp90α in the blood vessels of patients with DCLI undergoing partial amputation (n=3). CHOL, cholesterol; DCLI, diabetic critical limb ischemia; DLEAD, diabetic lower extremity arterial disease; DM, diabetes mellitus; Hb, hemoglobin; Hsp90, heat shock protein 90; LDL, low-density lipoprotein; MDA, malondialdehyde; WCC, white cell count.
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Figure 3  Increased eHsp90 secretion in a vascular endothelial cell model of oxidative injury in vitro. A CCK-8 assay (A) was used to detect cell viability in HUVECs treated with different concentrations of H$_2$O$_2$ for 2, 4, 6, 8, and 24 hours (n=3). The levels of MDA (B) and SOD (C) were measured with assay kits (n=3). HUVECs were treated with H$_2$O$_2$ (100 µM, 200 µM, and 300 µM) for 24 hours, and then the intracellular levels of ROS (D), mitochondrial ROS (E), and 8-oxoG (F) and the extracellular levels of ET-1 (G) and MCP-1 (H) were detected (n=3). (I) Western blot analysis concentrated medium and total cell extracts from HUVECs cultured with the indicated concentration of H$_2$O$_2$. ImageJ software was used to determine the relative intensities of extracellular and intracellular Hsp90α and Hsp90β bands (n=3). (J) Extracellular and intracellular Hsp90α and Hsp90β expressions were measured after H$_2$O$_2$ stimulation for different durations (n=3). (K) Western blot analysis was performed to assess the expression of related proteins in HUVECs pretreated with 17AAG and then exposed to H$_2$O$_2$ (n=3). (L) qPCR was employed to observe the respective Hsp90α and Hsp90β mRNA levels after HUVECs were stimulated with H$_2$O$_2$ and/or 17AAG for 24 hours (n=3). (M) Representative confocal images of Hsp90α subunit localization in endothelial cells after 24 hours of H$_2$O$_2$ stimulation are shown (n=3). All data represent mean±SD of three biological replicates. Differences between groups were analyzed by one-way ANOVA and the Dunnett test, according to the data features. *P<0.05,**P<0.01,***P<0.001 ****P<0.0001 versus control. 17AAG, 17-allylamino-17-demethoxygeldanamycin; ANOVA, analysis of variance; CCK-8, Cell Counting Kit-8; Ctrl, control; DAPI, 4',6-diamidino-2-phenylindole; eHsp90, extracellular heat shock protein 90; ET-1, endothelin 1; H$_2$O$_2$, hydrogen peroxide; Hsp90, heat shock protein 90; HUVECs, human umbilical vein endothelial cells; MCP-1, monocyte chemotactic protein 1; MDA, malondialdehyde; mRNA, messenger RNA; 8-oxoG, 8-oxoguanine; qPCR, quantitative real-time PCR; ROS, reactive oxygen species; SOD, superoxide dismutase.
**Figure 4** eHsp90α enhanced the inflammatory injury state of endothelial cells and recruited monocytes through LRP1 receptors. Extracellular ET-1 (A) and MCP-1 (B) expressions were measured after hrHsp90α stimulation at different concentrations and durations (n=3). The expression of ET-1 (C) in endothelial cell-CM treated with H2O2 after 17AAG or 1G6-D7 pretreatment (n=3). Representative images with quantification of the results for a THP-1 cell Transwell migration assay in the presence of H2O2-treated and/or 17AAG-treated endothelial cell CM (D) and different reagents (E) (n=5). LRP1 and p-Akt expressions (F) were measured after hrHsp90α stimulation for different durations (n=3). All data represent mean±SD of three biological replicates. Differences between groups were analyzed by one-way ANOVA and the Dunnett test, according to the data features. *P<0.05, **P<0.01, ***P<0.001, ****P<0.0001 versus control. 17AAG, 17-allylamino-17-demethoxygeldanamycin; ANOVA, analysis of variance; CM, conditioned medium; Ctrl, control; eHsp90, extracellular Hsp90; ET-1, endothelin 1; H2O2, hydrogen peroxide; hrHsp90, human recombinant Hsp90; Hsp90, heat shock protein 90; LRP1, low-density lipoprotein receptor-related protein 1; MCP-1, monocyte chemotactic protein 1.
the involvement of the eHsp90α-Akt pathway, we determined that stimulation with eHsp90α induced time-dependent activation of p-Akt (figure 4F).

DISCUSSION

In our study, we clarified the relationships of oxidative stress with serum Hsp90 levels in patients with diabetes. We propose that serum Hsp90α has reference value for early diagnosis of DLEAD, and our findings indicate that low-dose and long-term ROS exposure can increase eHsp90 secretion from endothelial cells. eHsp90α acts as a chemokine to induce THP-1 cell migration, which may result from a compensatory response of early inflammation and oxidative stress in AS, initiating the atherosclerotic process (online supplemental figure 1).

The incidence and mortality rates of atherosclerotic cardiovascular disease are significantly higher in patients with diabetes than in those without.14 Of patients with T2DM, 50% have complications at the time of diagnosis.15 However, fewer than 10% of patients have typical clinical symptoms and the lack of effective treatments affects patients’ prognoses.16 Recent studies have reported several potential serum markers for screening of diabetic vascular complications, such as serum fibroblast growth factor 23,17 serum sclerostin,18 and circulating Hsp27.19 In addition, heat shock proteins (HSPs) can be transported to the plasma membrane and released extracellularly, resulting in detectable levels of HSP in the blood. This finding has prompted clinical studies exploring the potential use of HSP and anti-HSP antibodies as serum biomarkers for DM complications.20 Available data suggest that circulating Hsp27,21 Hsp60,22 Hsp70,23 and anti-HSP levels may be used as biomarkers for diabetic vascular disease. In type 1 diabetes mellitus, elevated serum Hsp90 levels have been found to be related to cellular autoimmunity in children.24 New findings confirmed the role of Hsp90 as a putative autoantigen triggering inflammation within human carotid atherosclerotic plaques.25 However, there have been no studies on the relationships between eHsp90 and T2DM macrovascular complications. Most of the previous studies focused on the risk factors for LEAD, while there was no clear study on the process from diabetes to vascular disease complications. Therefore, we sought to investigate whether there is an indicator that can be included in primary prevention strategies that has reference value for early prediction. Numerous studies have reported that patients with DM have chronic inflammation, but now many inflammation indicators are closely related to acute infection, such as CRP and procalcitonin (PCT).26, thus, these indicators may not truly reflect chronic inflammation. Serological studies have found that the degree of oxidative stress in patients with diabetes is cumulative and that circulating Hsp90α may reflect a patient’s oxidative stress state. Predictive analysis indicated that serum Hsp90α shows better performance in the initial diagnosis of LEAD in patients with early DM than in patients with diabetes without atherosclerotic changes. However, the number of clinical samples included in this study was still insufficient and no healthy control group was found, so the results could only be limited to patients with diabetes and could not be further extended. The medical treatment of patients was not included in the analysis, so the influence of drugs on relevant indicators could not be excluded. In subsequent studies, we will further improve relevant information to determine whether Hsp90α can be used as a direct risk factor.

It is generally believed that abnormal glucose metabolism damages the arterial endothelial barrier while increasing platelet and inflammatory cell aggregation, which increases local oxidative stress levels. Destruction of the barrier function of vascular endothelial cells is a crucial initial step in this process.27 Oxidative stress is a potential mechanism for the development of endothelial cell dysfunction that is common to all risk factors.28 Previous research has shown that Hsp90 promotes the progression of AS through several mechanisms. oxidized low density lipoprotein (oxLDL) enhanced endothelial nitric oxide synthases (eNOS) association with Hsp90 and impaired endothelium-dependent vasodilation.29 However, statins repress human Dendritic Cells (DC) maturation induced by oxLDL, and repress an atherogenic HSP profile and promote the induction of T regulatory cells.30 Hsp60 has also been found to induce inflammatory cell activation.31 Lentivirus-mediated short hairpin RNA (shRNA) knockdown of Hsp60 and Hsp90 in DCs attenuated the effects of human group X-secreted phospholipase A2 (LDLx) on dendritic cell proliferation.32 However, existing studies have focused on the role of Hsp90 in cells and have not clearly distinguished between Hsp90α and Hsp90β and the possible mechanism of action of eHsp90. Next, we verified cytologically that oxidative stress upregulates the expression of eHsp90 in HUVECs and that this process is not accompanied by an increase in Hsp90 gene expression or a decrease in intracellular protein expression. These findings are in contrast to those of a study by Profumo et al,33 in which oxidative stress upregulates Hsp90 expression on the surfaces of endothelial cells and reduces Hsp90 secretion. There may be a link between the two findings. The expression of eHsp90 may be related to the duration of oxidative stress in cells. After a short (2 hours or 4 hours) treatment, no change in the amount of eHsp90 was observed in the cell culture medium. Previous studies have rarely used indicators to simultaneously measure oxidative stress at both the population and cell levels or used such indicators as references. In our study, we used MDA to measure the oxidative stress state of the population and found that MDA levels were lower in cells after H2O2 treatment than in patient serum. These findings suggest that endothelial cells under low-concentration H2O2 treatment exhibit increased expression of eHsp90, which can stimulate oxidative stress in humans.

AS is a lipid-driven inflammatory disease of the arterial intima. The balance of proinflammatory and
anti-inflammatory mechanisms determines the final clinical outcome, which is characterized by gradual accumulation of lipids and inflammatory cells.34 Two chemotactic processes are involved in AS: monocyte chemotaxis and smooth muscle fibroblast chemotaxis. Circulating leukocytes adhere to and migrate through the endothelial wall to the vascular smooth muscle layer of the intimal membrane.35 The migration of monocytes is an early disease indicator. Ambade et al36 found that the extracellular vesicles (EVs) of mice with alcoholic liver disease can induce macrophage activation through Hsp90. In addition, colocalization of inflammatory cells and Hsp90α was found in patient tissue samples, so we hypothesized that there may be a connection between Hsp90α and inflammatory damage. The results showed that eHsp90α could promote endothelial cell dysfunction and the release of inflammatory cytokines. CM and eHsp90α could promote monocyte migration and initiate the inflammatory process of AS. When we used recombinant Hsp90α to treat monocytes, we observed increased expression of LRP1 in a short period of time, which activated p-Akt, thereby promoting monocyte migration. Our study provides in vitro evidence of the key pathological role of eHsp90α in the control of diabetic vascular complications. This study did not go deep into the cell signaling mechanisms. In the future, we will study the pathway mechanism by knocking down or overexpressing genes, using pathway inhibitors and other means.

For the treatment of AS, there is no specific drug at present and it mainly focuses on the comprehensive management of patients, lowering blood glucose and blood lipids, etc. In the future, targeted therapies targeting different cell types in vivo, such as monocyte-macrophage lines37 and regulating VSMC phenotypic switching,38 could be considered. Another study found that the Hsp90 inhibitor 17-DMAG attenuates oxidative stress in experimental AS.39 This suggests that inhibition of Hsp90 alleviates the progression of AS. Based on our results, new drugs targeting eHsp90 for its proinflammatory properties could be developed in the future.

The study focused on serum Hsp90α, a substance strongly associated with oxidative stress in diabetic AS. This research enriches the understanding of the mechanism and biological function of eHsp90α. Through a clinical study, we combined clinical findings with novel functions of eHsp90α protein. However, as a single-center clinical study, serological results are unrepresentative. There may be admission rate bias, no response bias, etc. Due to selection bias, the research results could not reflect the real connection between a certain factor and the disease. The exploratory study of eHsp90α lacks specific functional area validation. In summary, the current study shows that eHsp90α participates in the inflammatory process of AS and is expected to become a promising new target for disease prediction and for treatment to slow the progression of AS.

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