

## TRANSLATIONAL PHYSIOLOGY |

# Heat therapy promotes the expression of angiogenic regulators in human skeletal muscle

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**Kuhlenhoelter AM, Kim K, Neff D, Nie Y, Blaize AN, Wong BJ, Kuang S, Stout J, Song Q, Gavin TP, Roseguini BT.** Heat therapy promotes the expression of angiogenic regulators in human skeletal muscle. *Am J Physiol Regul Integr Comp Physiol* 311: R377–R391, 2016. First published June 29, 2016; doi:10.1152/ajpregu.00134.2016.—Heat therapy has been shown to promote capillary growth in skeletal muscle and in the heart in several animal models, but the effects of this therapy on angiogenic signaling in humans are unknown. We evaluated the acute effect of lower body heating (LBH) and unilateral thigh heating (TH) on the expression of angiogenic regulators and heat shock proteins (HSPs) in healthy young individuals. Exposure to LBH ( $n = 18$ ) increased core temperature ( $T_c$ ) from  $36.9 \pm 0.1$  to  $37.4 \pm 0.1^\circ\text{C}$  ( $P < 0.01$ ) and average leg skin temperature ( $T_{leg}$ ) from  $33.1 \pm 0.1$  to  $39.6 \pm 0.1^\circ\text{C}$  ( $P < 0.01$ ), but did not alter the levels of circulating angiogenic cytokines and bone marrow-derived proangiogenic cells (CD34<sup>+</sup>CD133<sup>+</sup>). In skeletal muscle, the change in mRNA expression from baseline of vascular endothelial growth factor (VEGF), angiopoietin 2 (ANGPT2), chemokines CCL2 and CX3CL1, platelet factor-4 (PF4), and several members of the HSP family was higher 30 min after the intervention in the individuals exposed to LBH ( $n = 11$ ) compared with the control group ( $n = 12$ ). LBH also reduced the expression of transcription factor FOXO1 ( $P = 0.03$ ). Exposure to TH ( $n = 14$ ) increased  $T_{leg}$  from  $32.8 \pm 0.2$  to  $40.3 \pm 0.1^\circ\text{C}$  ( $P < 0.05$ ) but  $T_c$  remained unaltered ( $36.8 \pm 0.1^\circ\text{C}$  at baseline and  $36.9 \pm 0.1^\circ\text{C}$  at 90 min). This intervention upregulated the expression of VEGF, ANGPT1, ANGPT2, CCL2, and HSPs in skeletal muscle but did not affect the levels of CX3CL1, FOXO-1, and PF4. These findings suggest that both LBH and TH increase the expression of factors associated with capillary growth in human skeletal muscle.

angiogenesis; skeletal muscle; heat therapy; blood flow

SKELETAL MUSCLE CAPILLARY rarefaction is a common feature and a significant contributor to exercise intolerance in several chronic disease states, including peripheral artery disease (PAD) (62), chronic heart failure (CHF) (14), and chronic obstructive pulmonary disease (COPD) (65). Emerging evidence also indicates that a progressive reduction in skeletal muscle microvessel density plays an important role in the development and progression of metabolic syndrome (17).

Promoting capillary growth is, therefore, a major therapeutic goal to restore skeletal muscle function and exercise capacity in these populations. Surprisingly, few therapeutic strategies are known to effectively activate angiogenic signaling and increase skeletal muscle capillarization in humans. Exercise training is undoubtedly one of the most potent angiogenic therapies (18), but few patients engage and adhere to structured exercise programs (47, 61). Further, this option is challenging, or not amenable, for patients with severe disease and restricted locomotion. Gene and cell therapy are promising alternatives, but the results of most therapeutic angiogenesis trials to date have been largely disappointing (2). An urgent need remains for novel, more accessible strategies that stimulate angiogenesis in human skeletal muscle.

Heat therapy is a promising strategy that has recently been successfully used to treat patients with CHF (31, 49), severe COPD (76), and PAD (66, 74). Mounting evidence indicates that repeated exposure to whole body heat stress reduces the clinical symptoms and improves exercise tolerance in patients with these conditions (31, 49, 66, 76). The exact mechanisms underlying these documented clinical benefits are unclear, but several pieces of evidence indicate that heat therapy might work, in part, by promoting angiogenesis. First, whole body heat stress induced by far-infrared dry sauna or a heating blanket promotes the expression of the proangiogenic mediator vascular endothelial growth factor (VEGF) and increases capillary density in the myocardium in healthy (21) and hypertensive (28) rats, as well as in a model of myocardial infarction (69). Second, repeated treatment with far infrared dry sauna increases skeletal muscle capillary density in a mouse model of PAD (1, 40) and diabetes (26) and in rats treated with glucocorticoids (43). This angiogenic response in the heart and in skeletal muscle is closely coupled with increased expression of heat shock proteins (HSPs), molecular chaperones that are known to modulate the angiogenic process (40, 67, 71). Third, endothelial cells exposed to mild heat stress have an enhanced capacity for vascular tube formation, which is indicative of angiogenesis activity (60). Fourth, sauna therapy induces an increase in the number of circulating CD34<sup>+</sup> progenitor cells (49, 66), which promote vascular growth and repair (36). Despite this compelling evidence derived primarily from cell culture and animal studies, the impact of systemic and local

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Table 1. *Subject characteristics*

|                                     | Protocol 1  | Protocol 2     |             | Protocol 3  |
|-------------------------------------|-------------|----------------|-------------|-------------|
|                                     |             | Control (33°C) | LBH (48°C)  |             |
| Number of subjects<br>(male/female) | 18 (12/6)   | 12 (8/4)       | 11 (6/5)    | 14 (12/2)   |
| Age, yr                             | 22 ± 1.1    | 21 ± 0.6       | 24 ± 1.7    | 21 ± 0.8    |
| Body weight, kg                     | 70.3 ± 2.9  | 72.4 ± 4.6     | 70.7 ± 4.3  | 73.4 ± 3.0  |
| Height, cm                          | 172.2 ± 1.5 | 171.0 ± 2.5    | 173.5 ± 4.0 | 175.0 ± 2.4 |
| BMI, kg/m <sup>2</sup>              | 23.7 ± 0.9  | 24.6 ± 1.2     | 23.3 ± 0.9  | 23.9 ± 0.8  |

Values are expressed as mean ± SE. LBH, lower body heating; BMI, body mass index.

heat stress on angiogenic signaling in humans remains to be determined.

Skeletal muscle angiogenesis stems from the cumulative effects of transient changes in the abundance of proangiogenic mediators coupled with the inhibition of angiostatic factors (34, 51). Defining the time course and magnitude of changes in the transcriptional levels of these factors in response to a single stimulus is, therefore, critical toward understanding how the remodeling process is initiated and coordinated (15, 57). The goal of the present study was to determine the effects of a single 90-min session of heat therapy on systemic and skeletal muscle levels of key angiogenic regulators in humans. Water-circulating trousers were used to create two distinct experimental paradigms: 1) lower body heating (LBH), which induced an

increase in body core temperature and significant changes in systemic hemodynamics, and 2) unilateral thigh heating (TH), in which the heating stimulus was confined to one thigh, and body core temperature remained unaltered. These distinct experimental approaches allowed us to examine the potential contribution of systemic responses to heat stress to local changes in the expression of angiogenic genes in skeletal muscle. Systemic heat stress has been shown to induce physiological responses that can impact skeletal muscle angiogenesis, including activation of the sympathetic nervous system (7) and increased levels of bone marrow-derived proangiogenic cells (CD34<sup>+</sup>CD133<sup>+</sup>) (49, 66). Therefore, we hypothesized that LBH would increase the circulating levels of proangiogenic cytokines and cells and promote an increase in the mRNA level of angiogenic regulators and HSPs in skeletal muscle. Because a systemic angiogenic response would not occur during TH, we hypothesized the changes in the expression of proangiogenic factors in skeletal muscle following this protocol would be smaller compared with LBH.

## METHODS

### Participants

Fifty-five young, normally active individuals (Table 1) were recruited to participate in three separate studies (*protocol 1*,  $n = 18$ ; *protocol 2*,  $n = 23$ ; and *protocol 3*,  $n = 14$ ). Participants were asked to fill out a medical history questionnaire. Individuals were excluded

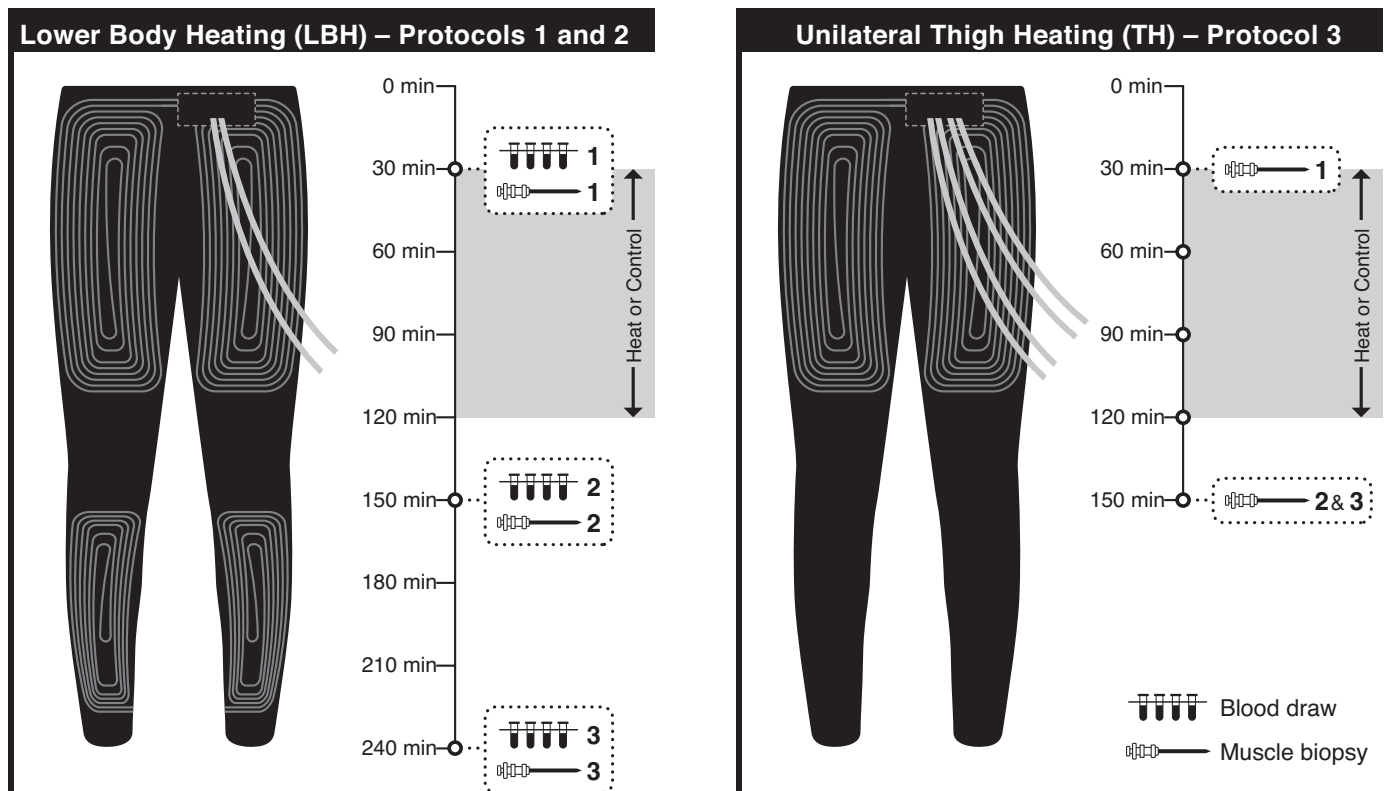


Fig. 1. Illustration of the water-circulating garments used for heat treatment and timelines for the experimental protocols. *Left*: lower body heating (LBH) was achieved by circulating hot water through tube-lined pants that covered the legs, thighs, and buttocks. This approach was used on *protocols 1* and *2*. Blood samples (*protocol 1*) and muscle biopsies (*protocol 2*) were taken prior to and 30 and 120 min following exposure to LBH or a control intervention. *Right*: a custom garment with tubing around the thigh was used for local heat treatment. This garment has a separate tubing circuit for each limb, which allows for one thigh to be exposed to heat treatment while the contralateral thigh serves as a control. This approach was used in *protocol 3*. Muscle biopsies were taken prior to and 30 min after the interventions.

Table 2. List of genes included in the custom PCR array used in protocol 2

| Gene Symbol | Official Full Name  |
|-------------|---|
| VEGFA       | Vascular endothelial growth factor A                                  |
| NOS3        | Nitric oxide synthase 3   |
| PPARGC1A    | Peroxisome proliferator-activated receptor gamma, coactivator 1 alpha |
| CCL2        | Chemokine (C-C motif) ligand 2  |
| MMP9        | Matrix metalloproteinase 9  |
| MMP2        | Matrix metalloproteinase 2  |
| TIMP1       | TIMP metalloproteinase inhibitor 1                                    |
| ANGPT2      | Angiopoietin 2  |
| ANGPT1      | Angiopoietin 1  |
| TEK         | TEK tyrosine kinase, endothelial                                      |
| THBS1       | Thrombospondin 1  |
| PF4         | Platelet factor 4   |
| CXCL12      | Chemokine (C-X-C motif) ligand 12                                     |
| NCL         | Nucleolin   |
| FOXO1       | Forkhead box O1   |
| FOXO3       | Forkhead box O3   |
| IL6         | Interleukin 6   |
| IL8         | Interleukin 8   |
| TNF         | Tumor necrosis factor   |
| CX3CL1      | Chemokine (C-X3-C motif) ligand 1                                     |
| HSP90AA1    | Heat shock protein 90-kDa alpha class A member 1                      |
| HSP90AB1    | Heat shock protein 90-kDa alpha class B member 1                      |
| HSPA1A      | Heat shock 70-kDa protein 1A  |
| HSPA1B      | Heat shock 70-kDa protein 1B  |
| HSPA8       | Heat shock 70-kDa protein 8   |
| HSPD1       | Heat shock 60-kDa protein 1   |
| HSPB1       | Heat shock 27-kDa protein 1   |
| GAPDH       | Glyceraldehyde-3-phosphate dehydrogenase                              |
| PPC         | Positive PCR control  |
| RTC         | Reverse transcription control   |
| HGDC        | Human genomic DNA contamination                                       |

if they were obese (BMI > 30 kg/m<sup>2</sup>), used tobacco products, were diabetic, were taking any medication other than birth control, participated in any kind of supervised physical activity or exercised for more than 3 days/wk. Female participants were tested during the early follicular phase of their menstrual cycle (*days 1–7*) or during the placebo phase if they were taking oral contraceptives. The Institutional Review Board at Purdue University approved all experimental procedures, and verbal and written consent were obtained from all participants.

#### Instrumentation and Experimental Protocols

Three separate protocols were conducted to determine the angiogenic responses to LBH (*protocols 1 and 2*) and TH (*protocol 3*). The experimental design with time points for blood draws and muscle biopsies for each protocol is depicted in Fig. 1. All experimental sessions were completed in the morning in a temperature-controlled room (average temperature of 24.2 ± 0.2°C). Participants were asked to avoid caffeine and alcoholic beverages for 12 h and intense exercise for at least 24 h prior to the experimental sessions. Participants were also asked to abstain from food for at least 1 h prior to each laboratory visit.

The purpose of *protocol 1* was to determine the effect of LBH on the circulating levels of angiogenic, inflammatory, and vasoactive mediators, as well as the number of circulating proangiogenic cells. Participants completed two experimental sessions, at least 72 h apart, in a randomized cross-over design. Subjects were given a wireless telemetry pill (HQ, Palmetto, FL) for core temperature monitoring during the experiments and were instructed to ingest it the night before the experiment (~7–9 h before the experimental sessions) (82). Upon arrival at the laboratory, subject body weight and height were recorded, and four thermocouples (MLT422; ADInstruments, Colo-

rado Springs, CO) were taped to the calf and thigh for measurement of mean leg skin temperature. The medial aspect of the calf on both legs was gently shaved, and laser-Doppler flow probes (VP12; Moor Instruments, Axminster, UK) were placed inside local heaters (VHP1; Moor Instruments) and were attached to the skin for the measurement of skin red blood cell flux (moorVMS-LDF2; Moor Instruments). Heart rate (HR) was monitored via a three-lead electrocardiogram (FE132; Bio Amp, ADInstruments). Systolic and diastolic blood pressures (BP) were measured from the left arm using an automated device (Tango+, Suntech Medical, Morrisville, NC). An intravenous catheter was placed in an antecubital vein of the right arm for blood sampling. Participants were asked to put on water-circulating trousers on top of shorts or underwear (Med-Eng, Ottawa, Canada). This garment is made of a tight-fitting elastic fabric, with an extensive network of medical-grade polyvinyl chloride tubing sewn onto the fabric and was designed to cover the calves, thighs, and buttocks (Fig. 1, *left*). In addition, to minimize heat loss during the interventions, participants also wore polyvinyl chloride pants and had their legs covered with a thermal foil blanket. After instrumentation, participants rested quietly for 30 min in the supine position, and a baseline blood sample was taken. Next, the water-circulating garment was connected to a heated bath circulator (Sahara S21, Thermo Scientific). In the LBH session, water at 48°C was perfused through the garment for 90 min with a goal to increase leg skin temperature to ~39.5–40°C (24). In the control intervention, water at 33°C was circulated through the garment to maintain leg skin temperature at baseline levels. At the end of the intervention period, the garment was disconnected from the water circulator, and the participants remained supine for another 2 h. Blood samples were taken 30 and 120 min following the completion of the trials (Fig. 1, *left*). In the last 20 min of the protocol, the temperature of the local skin heaters placed on the calf was increased to 43°C at a rate of 0.1°C/s to promote maximal increases in skin blood flow and allow for the calculation of maximal cutaneous vascular conductance (CVC<sub>max</sub>) (11). Leg skin temperature, cutaneous red blood cell flux, and HR were recorded continuously, while systolic and diastolic BP and core body temperature were recorded every 5 min for the entire duration of the protocol.

In *protocol 2*, enrolled participants were randomly allocated to either a group exposed to LBH (*n* = 11) or to a control group (*n* = 12). The purpose of this protocol was to determine the effect of LBH on the mRNA expression of angiogenic mediators in skeletal muscle. Participants were instrumented as described above in *protocol 1* (Fig. 1, *left*). After 30 min of rest in the supine position, the thighs were exposed, and a biopsy was taken from the vastus lateralis muscle of a randomly selected thigh using a 5-mm Bergstrom biopsy needle (Pelomi Medical, Albruslund, Denmark), as described previously (19,

Table 3. List of genes included in the custom PCR array used in protocol 3

| Gene Symbol | Official Full Name                               |
|-------------|--|
| VEGFA       | Vascular endothelial growth factor A             |
| CCL2        | Chemokine (C-C motif) ligand 2                   |
| ANGPT2      | Angiopoietin 2                                   |
| ANGPT1      | Angiopoietin 1                                   |
| PF4         | Platelet factor 4                                |
| FOXO1       | Forkhead box O1                                  |
| CX3CL1      | Chemokine (C-X3-C motif) ligand 1                |
| HSP90AA1    | Heat shock protein 90-kDa alpha class A member 1 |
| HSP90AB1    | Heat shock protein 90-kDa alpha class B member 1 |
| HSPA1B      | Heat shock 70-kDa protein 1B                     |
| HSPA8       | Heat shock 70-kDa protein 8                      |
| HSPD1       | Heat shock 60-kDa protein 1                      |
| HSPB1       | Heat shock 27-kDa protein 1                      |
| GAPDH       | Glyceraldehyde-3-phosphate dehydrogenase         |
| RTC         | Reverse transcription control                    |
| HGDC        | Human genomic DNA contamination                  |

20, 64). Next, the water-circulating garment was perfused with either 48°C water (LBH group) or 33°C water (control group) for 90 min. Two additional muscle biopsies were taken 30 and 120 min after the end of the intervention (Fig. 1, *left*). The second biopsy was taken from the opposite leg used for the first biopsy, and the third biopsy was taken from a site at least ~3 cm away from the first biopsy site.

In *protocol 3*, a custom-made garment was used to heat one thigh, while the opposite thigh served as a control (Fig. 1, *right*). The goal of this experiment was to investigate the effects of local heating, which does not evoke systemic responses, such as changes in core body temperature, on the mRNA expression of angiogenic mediators in the vastus lateralis muscle. Participants were asked to complete two experimental sessions. On the first session, two skin thermocouples were taped to each thigh for measurement of skin temperature, and participants were asked to put on the custom water-circulating garment and polyvinyl chloride pants and had their legs covered with a

thermal foil blanket. A baseline muscle biopsy was taken after 30 min of rest in the supine position from one randomly selected thigh. Next, the garment was connected to the water circulators. In the thigh assigned to receive the heat treatment, hot water (48–52°C) was circulated through the garment for 90 min to increase skin temperature to ~39–40°C as in *protocols 1* and 2. In the opposite thigh, skin temperature was maintained at 33°C for the entire duration of the protocol. Thirty minutes after the end of the treatment, one muscle biopsy was taken from both the control and heated thighs, as described above (Fig. 1, *right*). The biopsies were only taken at 30 min after the intervention in this protocol because most of the changes in mRNA levels occurred at this time point in *protocol 2*. The second session was conducted at least 1 wk after *session 1*. The purpose of this session was to characterize the physiological responses to TH. Subjects were instructed to ingest the wireless telemetry pill (HQ, Palmetto, FL) for core temperature monitoring, as described for *protocol*

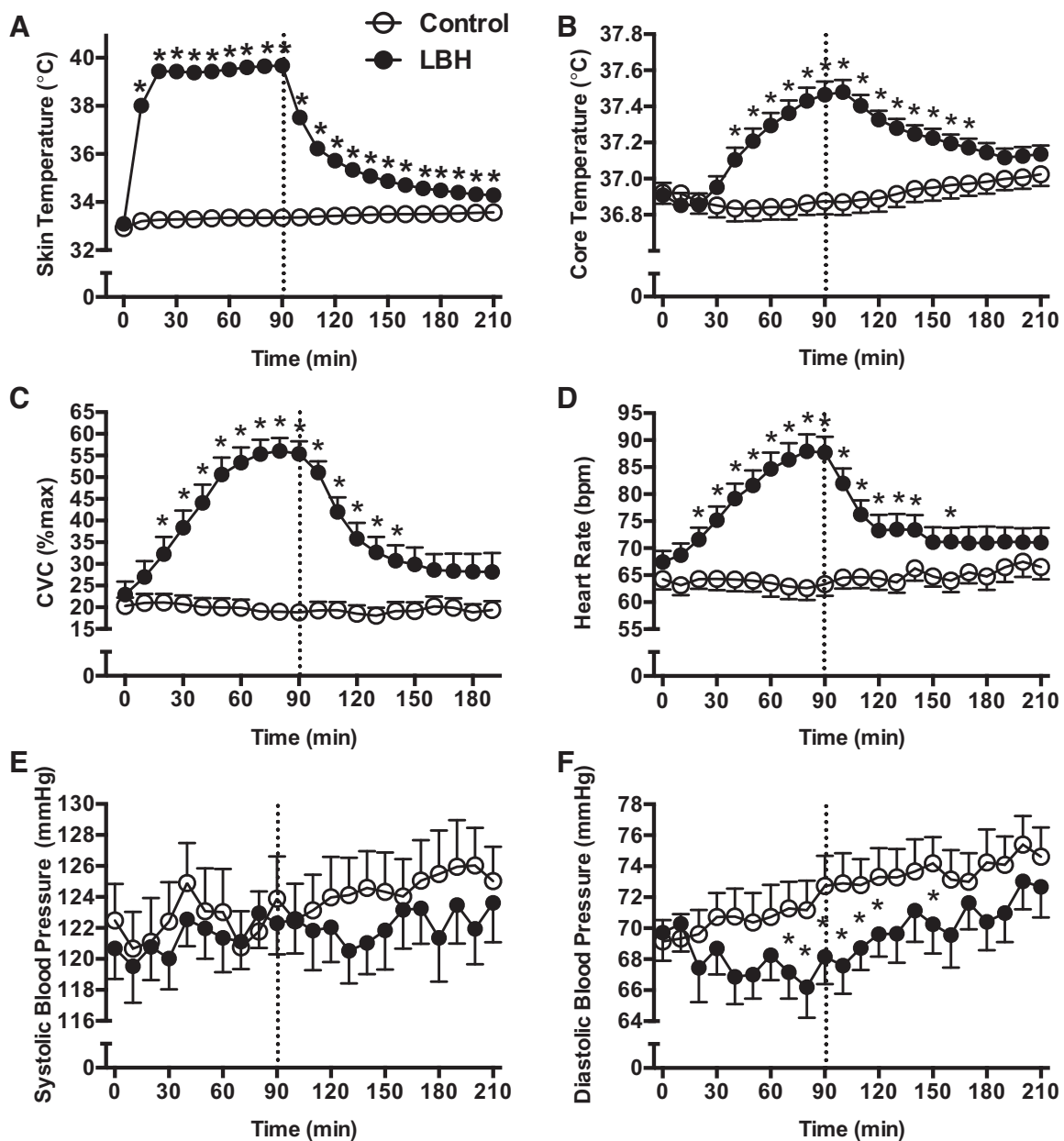


Fig. 2. Skin (A) and core temperature (B), CVC (C), HR (D), and systolic (E) and diastolic (F) blood pressure before, during, and after exposure to LBH (●) or a control intervention (○) for 90 min. Dashed line indicates the end of the intervention period. \* $P < 0.05$  vs. Control.

1. Upon arrival, subject weight and height were recorded, and skin thermocouples were taped to the thigh for the measurement of skin temperature. Participants were then fitted with the custom garment and polyvinyl chloride pants, as detailed above. A blood pressure cuff was wrapped around the left arm for blood pressure monitoring during the protocol using an automated device (Tango+, Suntech Medical, Morrisville, NC). After instrumentation, subjects were allowed to rest quietly for 30 min in the supine position. Next, the groin area was exposed, and the common femoral artery was imaged on both legs using a 10-MHz multifrequency linear probe attached to a high-resolution ultrasound machine (T3000; Terason, Burlington, MA). Diameter and velocity signals were acquired simultaneously in a duplex mode and were corrected with an insonation angle of 60°. Images were recorded for 1 min in each leg using screen capture software (Camtasia Studio, TechSmith). The garment was then connected to the water circulators for application of local heating and control treatments, as described above. Diameter and blood flow recordings were made every 30 min during the intervention, and once 30 min after the end of the heating protocol. Blood pressure and core body temperature were measured every 5 min, while skin temperature was recorded continuously for the entire duration of the protocol.

#### Measurements

**Circulating factors.** Venous blood samples were collected in serum-separating tubes (BD Vacutainer SST plus), allowed to clot for 30 min at room temperature, centrifuged for 15 min at 1,200 rpm, aliquoted, and stored at  $-80^{\circ}\text{C}$  until analysis. A custom Milliplex assay kit (EMD Millipore, Billerica, MA) was used to determine the concentrations of several angiogenic and inflammatory mediators, including granulocyte colony stimulating factor (G-CSF), granulocyte macrophage stimulating factor (GM-CSF), interleukin 6 (IL-6), interleukin 8 (IL-8), chemokine (C-C motif) ligand 2 (CCL2), tumor necrosis factor alpha (TNF- $\alpha$ ), vascular endothelial growth factor (VEGF), fibroblast growth factor 2 (FGF-2), and chemokine (C-X3-C motif) ligand 1 (CX3CL1). The assay was performed following the manufacturer's protocol using the Bio-Plex 200 System (Bio-Rad, Hercules, CA) at the Bio-Plex Core Facility of Indiana University. The serum concentration of endothelin-1 was determined using an ELISA kit (R&D Systems, Minneapolis, MN), according to manufacturer's instructions.

**Circulating proangiogenic cells.** A multiparametric flow cytometry method (45, 46) was used to identify circulating  $\text{CD34}^+\text{CD133}^+$  cells. Briefly, venous blood samples were collected in a cell preparation tube with sodium heparin (BD Vacutainer CPT tubes) and taken to the Flow Cytometry and Cell Separation Facility (Bindley Bioscience Center, Purdue University). Samples were centrifuged within 2 h of collection for 15 min at 1,500 RCF, as suggested by the manufacturer. Mononuclear cells were isolated, washed in PBS with 0.5%

BSA, and stained with antibodies against cell surface antigens, including anti-CD34-PE (BioLegend, San Diego, CA; no. 343506), anti-CD31-FITC (BioLegend, no. 303104), anti-CD45-BV510 (BioLegend, no. 304014), and anti-CD133-APC (Miltenyi, no. 130-090-826). In addition, to exclude dead/apoptotic cells, red blood cells, and monocytes, mononuclear cells were costained with Live/Dead stain-violet (no. L34955; Life Technologies, Carlsbad, CA), anti-CD235a-Pacific Blue (BioLegend, no. 306612), and anti-CD14-BUV395 (BD Biosciences, no. 563561), respectively. The cells were initially stained with Live/Dead stain for 30 min at  $4^{\circ}\text{C}$ , washed once with PBS/BSA, incubated with antibodies for 20 min at  $4^{\circ}\text{C}$ , washed twice in PBS/BSA, and analyzed using a BD FACS Aria III Cell Sorter. At least 1,000,000 events were collected for each sample. The data were analyzed using Flowjo v9.4.3 (Treestar).

**Skeletal muscle gene expression.** Biopsy samples were initially placed in a microcentrifuge tube containing 1 ml of RNeasy lysis buffer (Qiagen, Carlsbad, CA). The sample was kept at  $4^{\circ}\text{C}$  overnight and then stored at  $-80^{\circ}\text{C}$  until analysis. Approximately 30 mg of tissue was used for RNA extraction using the TRIzol Reagent (Invitrogen, Life Technologies), according to manufacturer's instructions. The quality and concentration of total RNA were measured by a spectrophotometer (Nanodrop 3000, ThermoFisher), as described previously (84). cDNA was prepared using the RT<sup>2</sup> first strand kit following the manufacturer's instructions (Qiagen, Valencia, CA). In *protocol 2*, the expression of 27 genes related to angiogenesis and inflammation, as well as members of the HSP family, was determined using a custom RT<sup>2</sup> Profiler PCR array kit (Qiagen) and the Roche LightCycler 480 PCR System (Roche Diagnostics, Indianapolis, IN). The three samples from each subject (baseline, 30 min and 120 min postintervention) were loaded in the same array plate. The list of genes analyzed is shown in Table 2 and includes the housekeeping gene glyceraldehyde-3-phosphate dehydrogenase (GAPDH), as well as a reverse-transcription control (RTC), a positive PCR control (PPC), and a human genomic DNA contamination control. Data were analyzed using the GeneGlobe Data Analyses Center (Qiagen, Valencia, CA). GAPDH was used to normalize cycle threshold (Ct) values. The comparative Ct method was used to calculate the changes in gene expression of each target mRNA relative to the baseline sample (33). A similar strategy was used to analyze the samples obtained in *protocol 3*, with the exception that the custom array had only 13 select genes, in addition to GAPDH, a reverse transcription control, and a human genomic DNA contamination control (Table 3).

#### Data and Statistical Analysis

Leg skin temperature, cutaneous red blood cell flux, and HR were recorded at 40 Hz using a data acquisition system (PowerLab and LabChart, ADInstruments, Colorado Springs, CO), and the last 2 min of every 5-min bin was averaged for the entire protocol. Systolic and

Table 4. Serum concentrations of angiogenic and inflammatory factors and the percentage of circulating proangiogenic cells ( $\text{CD34}^+\text{CD133}^+$ ) before and 30 and 120 min following exposure to lower body heating or to a control intervention (*protocol 1*)

|  | Control ( $33^{\circ}\text{C}$ ) |                   |                   | LBH ( $48^{\circ}\text{C}$ ) |                   |                   |
|--|----------------------------------|-------------------|-------------------|------------------------------|-------------------|-------------------|
|  | Pre                              | 30 min post       | 120 min post      | Pre                          | 30 min post       | 120 min post      |
| TNF- $\alpha$ , pg/ml                        | 13.2 $\pm$ 2.9                   | 16.6 $\pm$ 3.8    | 16.2 $\pm$ 4.2    | 15.0 $\pm$ 3.3               | 17.5 $\pm$ 4.5    | 15.2 $\pm$ 3.1    |
| IL-8, pg/ml                                  | 34.7 $\pm$ 9.6                   | 39.6 $\pm$ 10.3   | 35.1 $\pm$ 8.7    | 33.4 $\pm$ 7.7               | 34.9 $\pm$ 8.5    | 38.9 $\pm$ 8.8    |
| VEGF, pg/ml                                  | 656.6 $\pm$ 146.8                | 747.9 $\pm$ 165.8 | 737.1 $\pm$ 149.3 | 648.5 $\pm$ 123.6            | 704.8 $\pm$ 128.5 | 685.6 $\pm$ 135.4 |
| FGF2, pg/ml                                  | 126.0 $\pm$ 39.6                 | 139.9 $\pm$ 37.8  | 137.3 $\pm$ 34.6  | 115.6 $\pm$ 28.8             | 135.2 $\pm$ 35.2  | 136.2 $\pm$ 38.4  |
| G-CSF, pg/ml                                 | 71.5 $\pm$ 22.8                  | 81.6 $\pm$ 25.3   | 83.5 $\pm$ 27.1   | 63.1 $\pm$ 18.6              | 63.9 $\pm$ 16.7   | 70.8 $\pm$ 23.9   |
| CCL2, pg/ml                                  | 666.1 $\pm$ 40.2                 | 507.8 $\pm$ 36.0  | 504.1 $\pm$ 36.9  | 670.0 $\pm$ 48.1             | 511.2 $\pm$ 36.2  | 500.3 $\pm$ 35.9  |
| ET-1, pg/ml                                  | 1.1 $\pm$ 0.06                   | 1.5 $\pm$ 0.09    | 1.6 $\pm$ 0.1     | 1.2 $\pm$ 0.08               | 1.0 $\pm$ 0.08*   | 1.4 $\pm$ 0.08    |
| $\text{CD34}^+\text{CD133}^+$ , % live cells | 0.2 $\pm$ 0.03                   | 0.19 $\pm$ 0.03   | 0.18 $\pm$ 0.03   | 0.21 $\pm$ 0.05              | 0.21 $\pm$ 0.04   | 0.19 $\pm$ 0.03   |

Values are expressed as means  $\pm$  SE. VEGF, vascular endothelial growth factor; FGF2, fibroblast growth factor 2; G-CSF, granulocyte colony-stimulating factor; CCL2, chemokine (C-C motif) ligand 2; ET-1, endothelin-1. \* $P < 0.05$  vs. Control.

diastolic BP was recorded every 5 min, and mean arterial pressure was calculated as diastolic pressure plus one-third pulse pressure. Cutaneous vascular conductance (CVC) was calculated as red blood cell flux divided by mean arterial pressure and expressed as a percentage of  $CVC_{max}$ , as determined by the maximal value obtained during local heating at 43°C for 20 min.

Analysis of diameter and Doppler flow profile in the femoral artery in *protocol 3* was performed using computerized edge-detection and wall-tracking software (FMD Studio, Quipu, Pisa, Italy). The Doppler flow analysis computed time-averaged anterograde ( $V_{ant}$ ) and retrograde ( $V_{ret}$ ) values of velocity ( $V$ ), which were used for the calculation of positive and negative blood flows (BF) using the following equation:  $BF = (V \times 3.14 \times d^2)/400$ , where  $d$  is the vessel diameter. Mean BF was computed as the algebraic sum of anterograde and retrograde BF. Anterograde and retrograde wall shear rates (SR) were calculated using the following equation:  $SR = 4 \times V/d$ . The oscillatory shear index (OSI) was calculated as  $OSI = (SR_{ret})/(SR_{ant} + SR_{ret})$ .

All statistical analyses were conducted using SAS (version 9.4; SAS Institute, Cary, NC), with results expressed as means  $\pm$  SE. A two-way mixed-effect (with random individual effect and repeated measurements) ANOVA was used to compare the physiological responses and the levels of circulating factors and progenitor cells between trials in *protocol 1*, followed by Bonferroni post hoc comparisons when appropriate. In *protocol 2*, subject characteristics were compared between groups using unpaired  $t$ -tests. Because of nonnormal distribution, gene expression responses at each time point (30 min and 120 min) were compared between groups using the Wilcoxon Rank-Sum test. In *protocol 3*, a one-way repeated-measures ANOVA model was used to identify significant changes from baseline in heart rate, systolic and diastolic BP, and skin and core body temperature. A two-way mixed-effect ANOVA was employed to compare the blood flow parameters between the leg exposed to TH and the control leg. Post hoc analysis (Bonferroni) was performed when appropriate. As in *protocol 2*, gene expression responses had a nonnormal distribu-

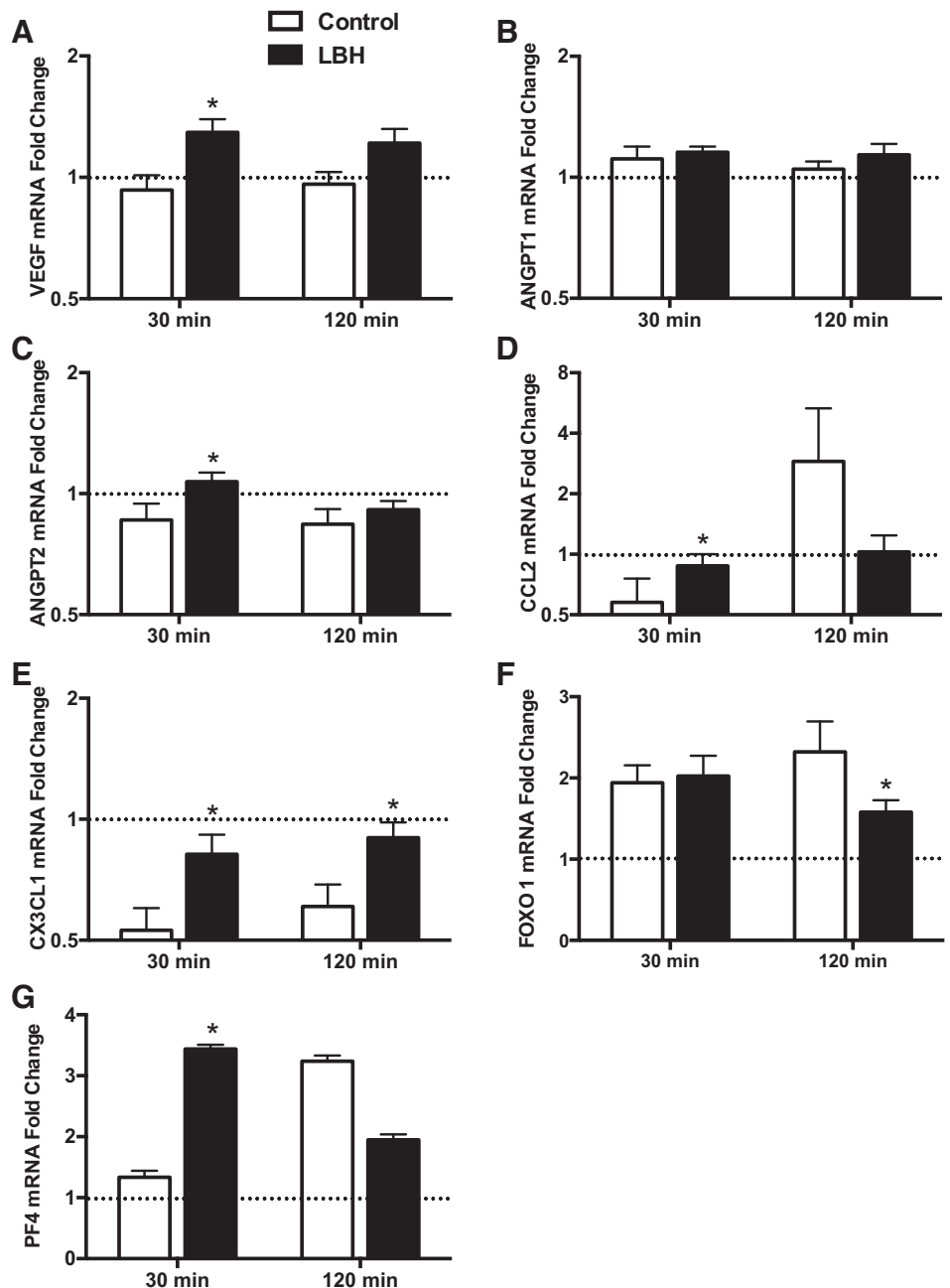


Fig. 3. Fold changes in skeletal muscle mRNA expression relative to the baseline sample of select angiogenic, inflammatory, and angiostatic mediators in the group exposed to LBH (solid bars;  $n = 11$ ) and in the control group (open bars;  $n = 12$ ). Biopsy samples were taken at baseline and 30 and 120 min following the completion of the interventions. The baseline sample was assigned a value of 1 and is represented as a dashed line. \* $P < 0.05$  vs. Control.

tion in *protocol 3*, and differences between thighs were compared using the paired Wilcoxon rank test. To allow for a direct comparison between gene expression responses at 30 min postintervention between *protocols 2* and *3*, each variable was subjected to logarithm transformation. A *t*-test was then used to compare the changes in the expression of each gene between the two distinct interventions. For all analyses,  $P \leq 0.05$  was considered statistically significant.

## RESULTS

### Protocol 1

**Physiological responses to LBH.** Figure 2 depicts the changes in skin and core temperatures, HR, systolic and diastolic blood pressure, and CVC, before, during, and after exposure to LBH or a control intervention. There were no baseline differences between trials for any of the physiological variables. As expected, exposure to LBH for 90 min induced marked increases from baseline in leg skin temperature (from  $33.1 \pm 0.1$  to  $39.6 \pm 0.1^\circ\text{C}$ ,  $P < 0.01$ ), CVC (from  $22.9 \pm 3.0$  to  $56 \pm 3.0\%$  max,  $P < 0.01$ ), and HR (from  $67 \pm 2$  to  $87 \pm 2$  bpm,  $P < 0.01$ ), as well as a modest increase in core temperature (from  $36.9 \pm 0.1$  to  $37.4 \pm 0.1^\circ\text{C}$ ,  $P < 0.01$ ). Systolic blood pressure was not affected by LBH, but diastolic blood pressure was significantly lower during the last 30 min of heating, and early in the recovery period after LBH compared with the control trial ( $P < 0.05$ , Fig. 2F).

**Effect of LBH on circulating cytokines and bone marrow-derived proangiogenic cells.** The serum concentrations of angiogenic and inflammatory mediators and the percentage of proangiogenic cells (CD34<sup>+</sup>CD133<sup>+</sup>) measured prior to and 30 and 120 min following exposure to LBH or the control intervention are shown in Table 4. The concentration of three (IL-6, CX3CL1, and GM-CSF) out of the nine factors measured was at or below detection levels and, therefore, the data were excluded. There were no differences between trials for any of the measured growth factors, cytokines, or the percentage of CD34<sup>+</sup>CD133<sup>+</sup> cells. The concentration of the potent vasoconstrictor ET-1 was significantly lower at 30 min following the intervention in the LBH trial compared with the control condition ( $P < 0.01$ ) (Table 4).

### Protocol 2

**Effect of LBH on the mRNA levels of angiogenic, inflammatory, and angiostatic mediators in skeletal muscle.** Changes in gene expression from baseline for select angiogenic, inflammatory, and angiostatic genes are shown in Fig. 3 and Table 5. The expression of three genes (IL-8, TNF, and MMP9) was near or below detection limits (i.e., Ct values  $\geq 35$  or not detectable), and the data were excluded. The mRNA expression of proangiogenic factors VEGF and angiopoietin 2 (ANGPT2) was increased at 30 min in the group exposed to LBH compared with the control group ( $P < 0.05$ ) (Fig. 3). ANGPT1 expression was not significantly different between groups. The mRNA expression of chemokine CCL2 was higher in the LBH group at 30 min following the intervention ( $P = 0.02$ ), while the levels of chemokine CX3CL1 were higher at both 30 and 120 min following LBH compared with the control group ( $P < 0.05$ ). The expression of the angiostatic transcription factor FOXO1 was lower in the LBH group at 120 min following the end of the intervention period ( $P = 0.03$ ). Conversely, the mRNA expression of the angiostatic factor

platelet factor 4 (PF4) was increased at 30 min in the LBH group relative to the control group ( $P = 0.03$ ). The fold changes in mRNA expression of NOS3, PPARGC1A, MMP2, TIMP1, TEK, THBS1, CXCL12, NCL, FOXO3, and IL-6 were not different between groups (Table 5).

**Effect of LBH on the mRNA levels of heat shock proteins.** The relative mRNA expression of members of several HSP families is shown in Fig. 4. In agreement with what was observed for the angiogenic/inflammatory genes, increased expression of these factors in the LBH group was particularly evident at 30 min following the intervention. The mRNA levels of HSP90AA1 and HSP90AB1, members of the HSP90 family, were higher in the LBH group compared with control group at 30 min ( $P < 0.01$ ; Fig. 4), but there were no differences between groups at 120 min. The expression of HSPA1A, also termed HSP72 and a member of the HSP70 family, was slightly higher in the LBH group at 30 min (fold change, Control:  $0.8 \pm 0.04$  and LBH:  $1.06 \pm 0.09$ ), but this difference did not reach statistical significance ( $P = 0.08$ ; Table 5). The expression of HSPA1B and HSPA8, both members of the HSP70 family, was higher in the group exposed to LBH than in the control group at 30 min ( $P < 0.05$ ; Fig. 4). Similarly, the mRNA levels of mitochondria-encoded HSPD1, a member of the chaperonin family, was higher at 30 min in the group exposed to heat treatment than in the group exposed to the sham intervention ( $P = 0.018$ ; Fig. 4).

### Protocol 3

**Physiological responses to TH.** Skin temperature and body core temperature responses to TH are shown on Fig. 5. By study design, skin temperature in the thigh exposed to local heating increased to  $\sim 40^\circ\text{C}$  during the treatment, while in the control thigh, the temperature was maintained at  $\sim 33^\circ\text{C}$ . Core temperature remained constant throughout the entire protocol. Systolic and diastolic BP and HR were also not significantly altered by TH treatment (data not shown). Compared with the control thigh, mean BF and anterograde SR in the femoral artery increased during TH, while retrograde SR and OSI were markedly reduced during the treatment (Table 6).

Table 5. Fold changes in skeletal muscle mRNA expression relative to the baseline sample of select angiogenic, inflammatory, and angiostatic mediators in the group exposed to LBH and in the control group (*protocol 2*)

|          | Control (33°C)<br>(n = 12) |             | LBH (48°C)<br>(n = 11) |             |
|----------|----------------------------|-------------|------------------------|-------------|
|          | 30 min                     | 120 min     | 30 min                 | 120 min     |
| NOS3     | 1.07 ± 0.11                | 1.38 ± 0.25 | 1.16 ± 0.06            | 1.50 ± 0.14 |
| PPARGC1A | 1.18 ± 0.08                | 1.27 ± 0.16 | 1.32 ± 0.06            | 1.21 ± 0.03 |
| MMP2     | 0.99 ± 0.07                | 1.17 ± 0.12 | 0.98 ± 0.07            | 1.07 ± 0.08 |
| TIMP1    | 0.89 ± 0.09                | 1.18 ± 0.22 | 1.12 ± 0.11            | 0.97 ± 0.07 |
| TEK      | 1.09 ± 0.11                | 1.30 ± 0.16 | 1.17 ± 0.05            | 1.23 ± 0.06 |
| CXCL12   | 0.98 ± 0.06                | 0.97 ± 0.06 | 0.98 ± 0.03            | 0.93 ± 0.03 |
| NCL      | 1.05 ± 0.05                | 1.03 ± 0.04 | 1.09 ± 0.04            | 0.96 ± 0.03 |
| FOXO3    | 1.31 ± 0.06                | 1.34 ± 0.07 | 1.43 ± 0.11            | 1.28 ± 0.11 |
| IL6      | 0.73 ± 0.14                | 0.66 ± 0.14 | 0.65 ± 0.08            | 0.62 ± 0.06 |
| HSPA1A   | 0.80 ± 0.05                | 0.73 ± 0.08 | 1.06 ± 0.10            | 0.93 ± 0.08 |

Biopsy samples were taken at baseline and 30 and 120 min following the completion of the interventions.

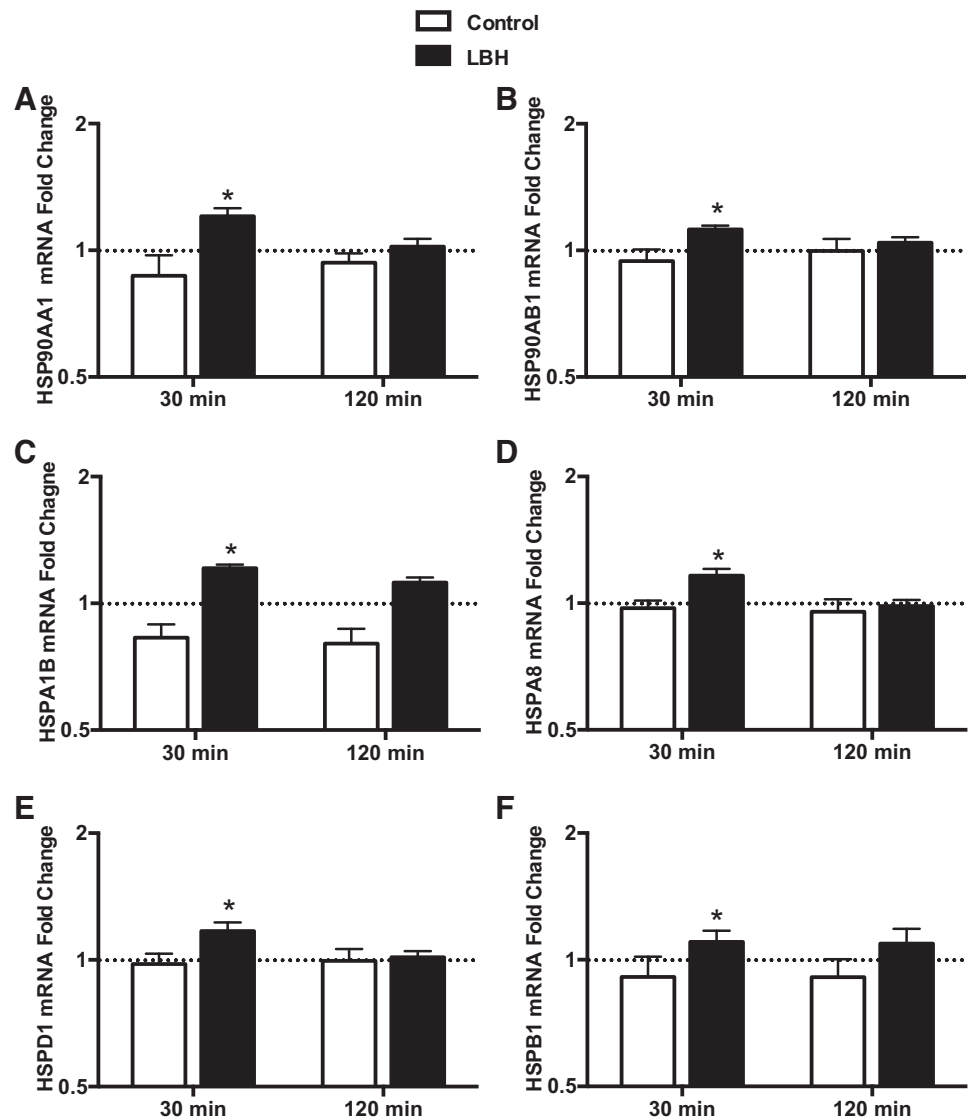


Fig. 4. Fold changes in skeletal muscle mRNA expression relative to the baseline sample of select members of the HSP family in the group exposed to LBH (closed bars;  $n = 11$ ) and in the control group (open bars;  $n = 12$ ). Biopsy samples were taken at baseline and 30 and 120 min following the completion of the interventions. The baseline sample was assigned a value of 1 and is represented as a dashed line. \* $P < 0.05$  vs. Control.

*Effect of TH on the mRNA levels of angiogenic, inflammatory, and angiostatic mediators in skeletal muscle.* Changes in gene expression from baseline for select angiogenic genes following local heat treatment are shown in Fig. 6. In congruence with the findings from protocol 2, TH led to increased mRNA expression of VEGF, ANGPT2, and CCL2 relative to the control thigh ( $P < 0.05$ ). In contrast to LBH, localized heating increased the levels of ANGPT1 ( $P < 0.05$ ) but did not affect the expression of CX3CL1 ( $P = 0.16$ ), FOXO1 ( $P = 0.85$ ), and PF4 ( $P = 0.37$ ).

*Effect of TH on the mRNA levels of heat shock proteins.* In close agreement with the changes induced by LBH, TH increased the expression of members of the HSP90 family (HSP90AA1 and HSP90AB1), HSP70 family (HSPA1B and HSPA8), and the chaperonin family (HSPD1) (Fig. 7).

*Comparison of skeletal muscle gene expression responses between LBH and TH.* The comparison between the LBH and TH protocols for the changes in the expression of select angiogenic, inflammatory, and angiostatic factors and heat shock proteins 30 min after the interventions is shown in Table 7. Contrary to our initial hypothesis, the magnitude of change

in the expression of these factors was not different between the two interventions.

## DISCUSSION

In the present study, we employed two experimental paradigms to determine the effect of a single session of heat therapy on the expression of angiogenic mediators and HSPs in humans. Mild systemic heat stress was induced by circulating 48°C water through a garment designed to cover the legs, thighs, and buttocks. This strategy increased core body temperature by ~0.6°C and reduced diastolic BP and the serum concentration of the potent vasoconstrictor ET-1. Despite inducing these systemic effects, LBH did not affect the circulating levels of several cytokines or bone marrow-derived proangiogenic cells (CD34<sup>+</sup>CD133<sup>+</sup>). Conversely, in skeletal muscle, LBH evoked a transient increase in the mRNA expression of several important angiogenic factors, including VEGF and HSPs. To explore the mechanisms underlying this response, we designed a custom water-circulating garment that allowed for localized heat application to one thigh, while the contralateral



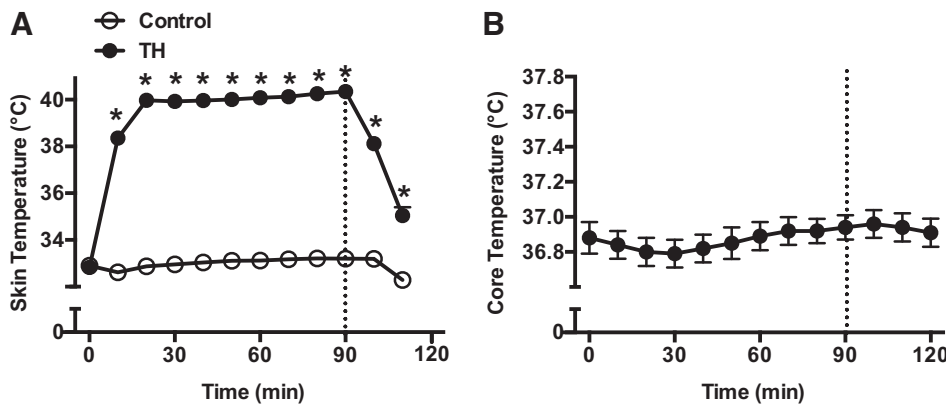


Fig. 5. Skin temperature (A) and core temperature (B) before, during, and after exposure to thigh heating (TH) for 90 min. One thigh was heated (●), while the opposite thigh served a control (○). Dashed line indicates the end of the intervention period. \* $P < 0.05$  vs. Control.

thigh served as a control (Fig. 1, right). In this model, body core temperature and systemic hemodynamics were not significantly altered, while femoral blood flow in the heated thigh increased markedly compared with the control thigh. In close agreement with the responses induced by LBH, local heating also upregulated the expression of angiogenic factors and HSPs in skeletal muscle. These findings suggest that the acute angiogenic response to LBH stems largely from local mechanisms and is not influenced by the systemic responses induced by this treatment. Overall, this study reveals that heat therapy is a simple, noninvasive strategy to activate local angiogenic signaling and possibly promote vascular growth in skeletal muscle.

#### Effects of Heat Therapy on Circulating Angiogenic Factors and Proangiogenic Cells

The reported increase in circulating levels of bone marrow-derived CD34<sup>+</sup> cells in patients with PAD (66) and CHF (49) after treatment with sauna therapy prompted us to investigate the acute effects of LBH on the numbers of these cells, as well as on the concentrations of angiogenic cytokines that participate in the recruitment and mobilization of progenitor cells. The angiogenic potential of CD34<sup>+</sup> cells has been well documented in numerous clinical and preclinical studies (36) and appears to be related to the ability of these cells to produce and secrete angiogenic cytokines (37). Using a multiparametric flow cytometry protocol, we identified and quantified the number of cells that coexpress CD34 and the primitive stem cell marker CD133 (83). Contrary to our hypothesis and the

mentioned sauna studies (49, 66), a single session of LBH did not affect the levels of circulating cytokines and proangiogenic cells (Table 2). Combined, these findings indicate that acute LBH does not evoke a systemic angiogenic response in healthy young participants. Because sauna therapy typically induces larger changes in core temperature ( $\sim 1$ – $1.2^{\circ}\text{C}$ ) (73) than what we observed in the present study ( $\sim 0.6^{\circ}\text{C}$ ), one potential explanation for the lack of effect of LBH on the systemic levels of these proangiogenic mediators is that the heat stress level was not sufficient to trigger the production and/or release of cytokines and promote the recruitment of progenitor cells. Another possibility is that 2 h was not long enough after LBH to induce changes in the levels of these circulating proangiogenic factors and cells. In addition, it is conceivable that multiple or longer sessions of LBH are necessary to effectively activate these angiogenic mediators.

#### Skeletal Muscle Angiogenic Signaling

Several animal studies demonstrated unequivocally that heat therapy promotes angiogenesis in the heart (21, 28, 69) and in skeletal muscle (1, 40, 43), but the molecular mechanisms underlying this adaptation remain poorly defined. One attractive candidate is VEGF, a potent angiogenic inducer that acts by promoting proliferation and enhancing migration and invasion of endothelial cells (5). Increased capillarization in the myocardium following whole body heat stress in rats is, indeed, associated with a marked upregulation of VEGF expression (21, 28, 69). In the present study, an increase in VEGF mRNA expression in skeletal muscle was observed following

Table 6. Flow profile and shear rate responses in the femoral artery to TH application in protocol 3

|                                      | Baseline    |             | 30 min      |                           | 60 min                   |                           | 90 min                   |                           | 30 min post              |                          |
|--------------------------------------|-------------|-------------|-------------|---------------------------|--------------------------|---------------------------|--------------------------|---------------------------|--------------------------|--------------------------|
|                                      | C           | H           | C           | H                         | C                        | H                         | C                        | H                         | C                        | H                        |
| d (cm)                               | 0.66 ± 0.02 | 0.68 ± 0.02 | 0.69 ± 0.02 | 0.70 ± 0.02               | 0.68 ± 0.02              | 0.71 ± 0.02               | 0.69 ± 0.02              | 0.71 ± 0.02               | 0.68 ± 0.02              | 0.72 ± 0.03              |
| V <sub>mean</sub> , cm/s             | 8.6 ± 0.9   | 7.9 ± 0.9   | 11.2 ± 1.6  | 15.7 ± 1.8 <sup>Φ*</sup>  | 12.7 ± 1.3 <sup>Φ</sup>  | 17.8 ± 1.6 <sup>Φ*</sup>  | 14.5 ± 1.7 <sup>Φ</sup>  | 17.3 ± 1.5 <sup>Φ*</sup>  | 12.8 ± 1.6 <sup>Φ</sup>  | 13.1 ± 1.4 <sup>Φ</sup>  |
| V <sub>ant</sub> , cm/s              | 14.9 ± 1.0  | 13.9 ± 0.8  | 17.1 ± 1.4  | 19.9 ± 1.7 <sup>Φ*</sup>  | 17.7 ± 1.3               | 21.3 ± 1.6 <sup>Φ*</sup>  | 19.2 ± 1.5 <sup>Φ</sup>  | 21.2 ± 1.3 <sup>Φ</sup>   | 17.5 ± 1.4               | 17.9 ± 1.2 <sup>Φ</sup>  |
| V <sub>ret</sub> (cm/s)              | 6.3 ± 0.7   | 6.07 ± 0.6  | 6.0 ± 0.7   | 4.3 ± 0.3 <sup>Φ*</sup>   | 4.9 ± 0.5 <sup>Φ</sup>   | 3.5 ± 0.4 <sup>Φ*</sup>   | 4.7 ± 0.5 <sup>Φ</sup>   | 3.9 ± 0.5 <sup>Φ</sup>    | 4.7 ± 0.68 <sup>Φ</sup>  | 4.8 ± 0.7                |
| BF, ml/min                           | 179 ± 21    | 174 ± 24    | 251 ± 39    | 375 ± 58*                 | 283 ± 39                 | 434 ± 57*                 | 321 ± 35                 | 413 ± 48*                 | 286 ± 40                 | 322 ± 40                 |
| SR <sub>mean</sub> , s <sup>-1</sup> | 52.5 ± 5.4  | 47.0 ± 5.3  | 65.6 ± 9.5  | 90.1 ± 10.0 <sup>Φ*</sup> | 75.2 ± 7.4 <sup>Φ</sup>  | 100.4 ± 8.4 <sup>Φ*</sup> | 85.8 ± 11.0 <sup>Φ</sup> | 98.7 ± 8.6 <sup>Φ</sup>   | 75.1 ± 9.1 <sup>Φ</sup>  | 74.8 ± 8.5 <sup>Φ</sup>  |
| SR <sub>ant</sub> , s <sup>-1</sup>  | 90.6 ± 6.1  | 82.9 ± 5.1  | 100.5 ± 8.7 | 114 ± 9.2 <sup>Φ*</sup>   | 104.6 ± 7.2              | 120 ± 7.5 <sup>Φ*</sup>   | 113.1 ± 10 <sup>Φ</sup>  | 121.1 ± 7.7 <sup>Φ</sup>  | 102.8 ± 8.6              | 101.5 ± 7.5 <sup>Φ</sup> |
| SR <sub>ret</sub> , s <sup>-1</sup>  | 38.2 ± 3.8  | 36.0 ± 3.5  | 34.9 ± 4.2  | 24.8 ± 2.1 <sup>Φ*</sup>  | 29.3 ± 3.2 <sup>Φ</sup>  | 19.8 ± 2.7 <sup>Φ*</sup>  | 27.3 ± 2.7 <sup>Φ</sup>  | 22.5 ± 3 <sup>Φ</sup>     | 27.6 ± 3.8 <sup>Φ</sup>  | 26.7 ± 3.8 <sup>Φ</sup>  |
| OSI                                  | 0.29 ± 0.02 | 0.30 ± 0.02 | 0.26 ± 0.02 | 0.19 ± 0.02 <sup>Φ*</sup> | 0.22 ± 0.02 <sup>Φ</sup> | 0.14 ± 0.02 <sup>Φ*</sup> | 0.20 ± 0.02 <sup>Φ</sup> | 0.16 ± 0.02 <sup>Φ*</sup> | 0.21 ± 0.03 <sup>Φ</sup> | 0.21 ± 0.03 <sup>Φ</sup> |

Values are expressed as means ± SE. C, control thigh; H, heated thigh; d, diameter; V<sub>mean</sub>, time-averaged mean velocity; V<sub>ant</sub>, time-averaged antegrade velocity; V<sub>ret</sub>, time-averaged retrograde velocity; SR<sub>mean</sub>, mean shear rate; SR<sub>ant</sub>, antegrade shear rate; SR<sub>ret</sub>, retrograde shear rate; OSI, oscillatory shear index.  $\Phi P < 0.05$  vs. baseline. \* $P < 0.05$  vs. Control.

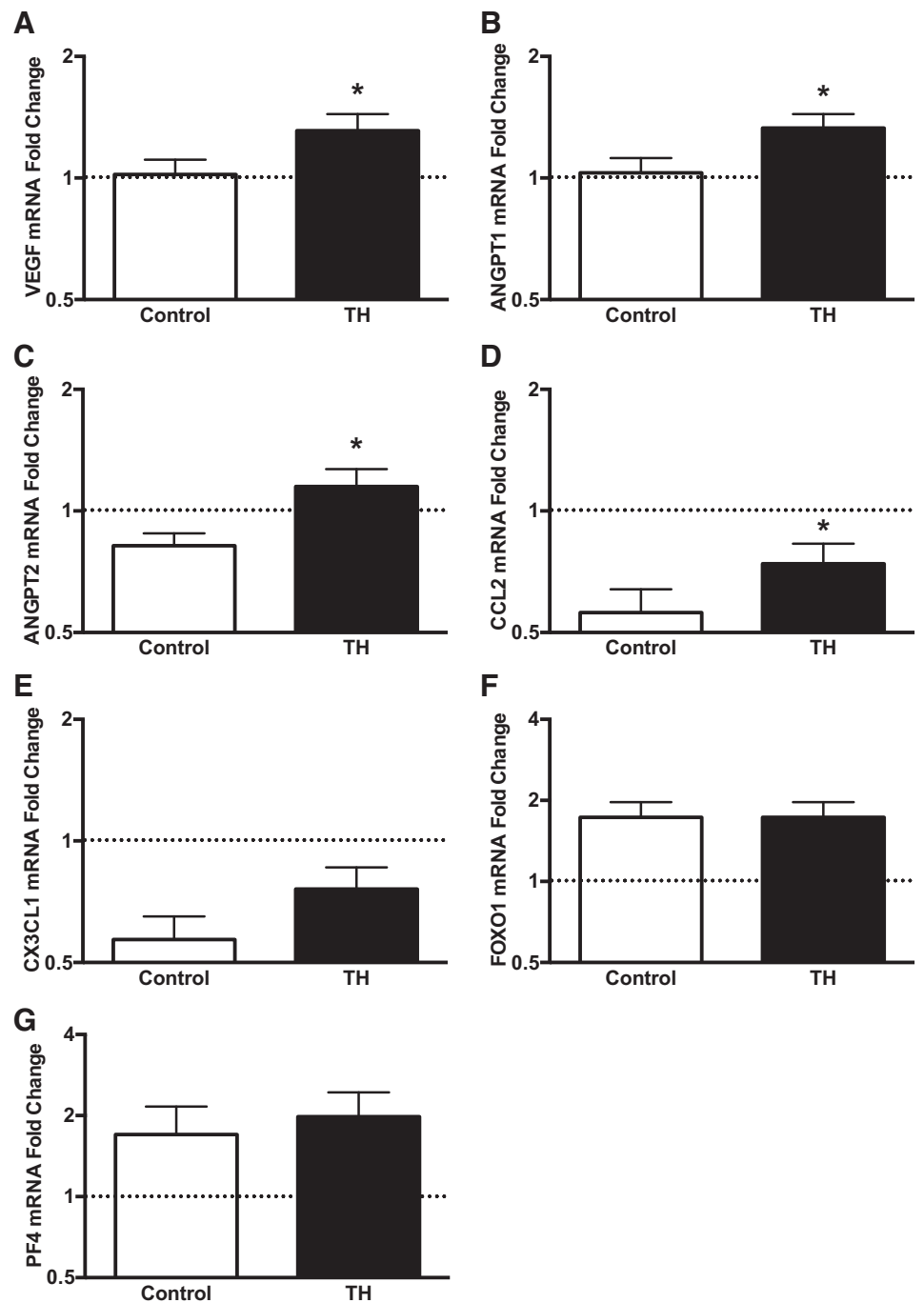


Fig. 6. Fold changes in skeletal muscle mRNA expression relative to the baseline sample of select angiogenic, inflammatory, and angiostatic mediators following exposure to TH. Biopsy samples were taken at baseline and 30 min following the completion of the intervention. The baseline sample was assigned a value of 1 and is represented as a dashed line. \* $P < 0.05$  vs. Control.

both LBH and TH, demonstrating that this central proangiogenic factor might also mediate the angiogenic response to heat therapy in skeletal muscle. VEGF regulates basal and exercise training-induced increases in skeletal muscle capillarization (80), and muscle-specific deletion of this molecule leads to capillary rarefaction (3, 52) and nearly abolishes exercise-induced capillary growth (53).

In addition to VEGF, heat therapy activates other key players involved in the angiogenic cascade. ANGPT1 and ANGPT2 are important proangiogenic factors that act as ligands for Tie-2 receptors in endothelial cells (16). Although the importance of angiopoietin signaling in the vasculature is well defined,

the cellular origins and functional roles of these two factors in skeletal muscle have only recently received attention. ANGPT1 is the principal angiopoietin produced by skeletal myoblasts and myotubes and is an important regulator of myogenesis and angiogenesis (38, 42). Conversely, ANGPT2 has no effect on myoblast proliferation and migration and exerts a relatively weak and context-dependent effect on angiogenesis (41). Mofarrah et al. (42) recently showed that overexpression of ANGPT1 following muscle injury in mice promoted capillary growth and accelerated the recovery of contractile performance. In this scenario, it is noteworthy that TH induced the expression of ANGPT1 in skeletal muscle (Fig. 6). This novel finding might help explain why

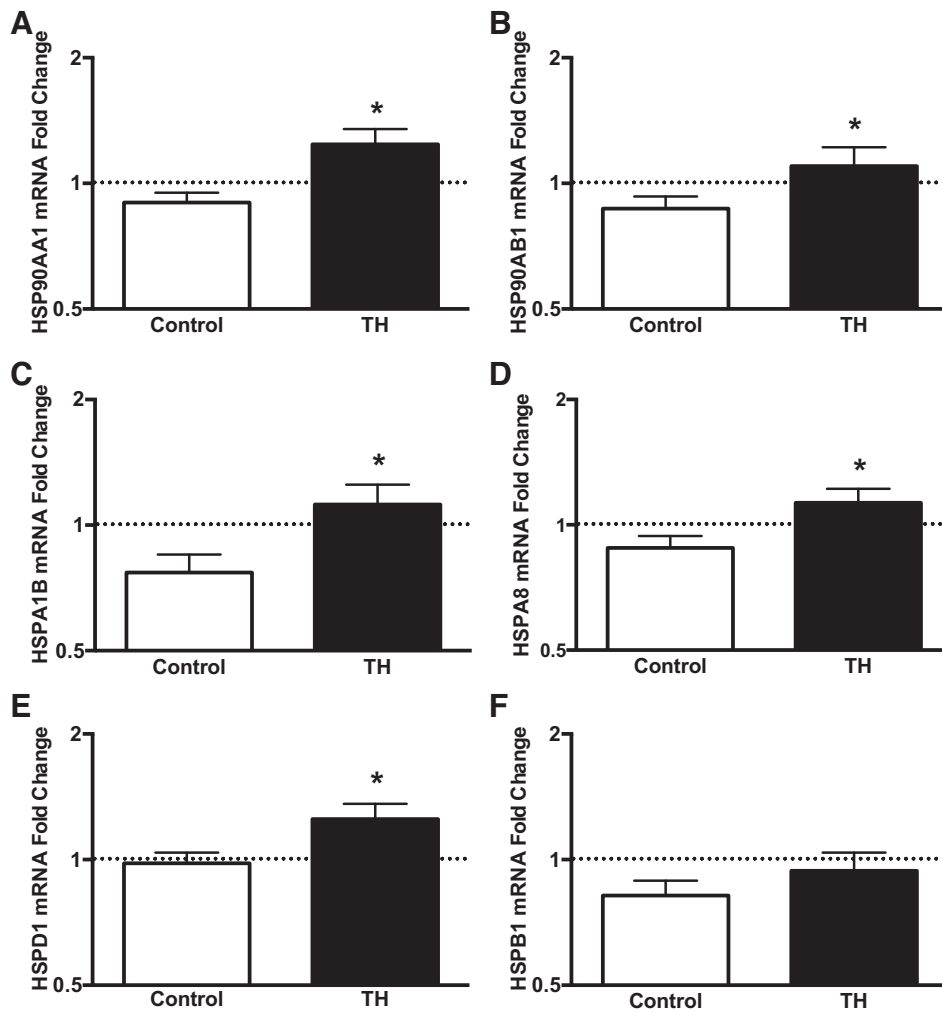


Fig. 7. Fold changes in skeletal muscle mRNA expression relative to the baseline sample of select members of the heat shock protein (HSP) family following exposure to TH. Biopsy samples were taken at baseline and 30 min following the completion of the intervention. The baseline sample was assigned a value of 1 and is represented as a dashed line. \* $P < 0.05$  vs. Control.

repeated heat therapy application facilitates regeneration of injured skeletal muscle (50, 72).

Individuals exposed to LBH had higher mRNA levels of CCL2 and CX3CL1 relative to the control group (Fig. 3). CCL2 expression was also higher in the heated thigh compared with the control thigh on *protocol 3* (Fig. 6). This is an important finding because these myokines/chemokines are involved in the attraction of monocytes/macrophages, which are critical for skeletal muscle remodeling (8). Impaired CCL2 signaling via deletion of its receptor, CCR2, delays angiogenesis and reduces VEGF levels following skeletal muscle injury in mice (48). Recently, Strömberg et al. (70) demonstrated that CX3CL1 stimulation of primary human myoblasts and myotubes promoted marked increases in the expression of proangiogenic factors and chemotactic mediators. These observations, coupled with the notion that the expression of both factors increases markedly in skeletal muscle following an acute bout of exercise in humans (6), indicate that these chemoattractants promote a microenvironment that facilitates angiogenesis and muscle repair (70).

The expression levels of the transcription factor FOXO1 were reduced 2 h after the intervention in the group treated with LBH. FoxO1 regulates the transcription of several angiostatic factors and acts to restrain angiogenesis in skel-

etal muscle (63, 68). A reduction in the expression of this factor might, therefore, allow for the initiation of the angiogenic cascade. Altogether, these findings indicate that both LBH and TH promote the expression of some pivotal factors that mediate capillary growth in skeletal muscle. It is conceivable that these acute responses might translate over time to increased capillary supply in skeletal muscle after repeated treatment. Indeed, evidence derived from exercise training studies indicates that skeletal muscle angiogenesis is preceded by repeated transient mRNA bursts of growth factors and proinflammatory mediators (23, 34). Nonetheless, it is important to highlight that the changes in the expression of some factors in the present study were transient and relatively small. Whether protein levels of these proangiogenic factors are also altered by acute heat therapy remains to be determined. Additional studies are needed to explore the long-term effects of repeated heat therapy on skeletal muscle capillarization in healthy and diseased populations.

One important observation of the current study was that the expression of some genes were altered in the control group on *protocol 2* and in the thigh exposed to the control intervention on *protocol 3*. This response was particularly evident for CCL2 and CX3CL1, which were consistently downregulated follow-

Table 7. Comparison of fold changes in skeletal muscle mRNA expression relative to the baseline sample of select angiogenic, inflammatory and angiostatic mediators and heat shock proteins between the LBH and TH protocols

|          | Protocol 2 (LBH) |              | Protocol 3 (TH) | P     |
|----------|------------------|--------------|-----------------|-------|
|          | C                | LBH          | H-C             |       |
| VEGFA    | -0.07 ± 0.08     | 0.29 ± 0.10  | 0.30 ± 0.10     | 0.479 |
| CCL2     | -0.42 ± 0.18     | -0.13 ± 0.13 | 0.20 ± 0.10     | 0.362 |
| ANGPT2   | -0.14 ± 0.08     | 0.07 ± 0.06  | 0.33 ± 0.11     | 0.685 |
| ANGPT1   | 0.11 ± 0.08      | 0.15 ± 0.04  | 0.30 ± 0.12     | 0.118 |
| PF4      | 0.34 ± 0.53      | 2.44 ± 1.59  | 0.74 ± 0.55     | 0.286 |
| FOXO1    | 0.94 ± 0.21      | 1.02 ± 0.25  | 0.01 ± 0.11     | 0.700 |
| CX3CL1   | -0.47 ± 0.07     | -0.18 ± 0.10 | 0.19 ± 0.09     | 0.462 |
| HSP90AA1 | -0.13 ± 0.10     | 0.21 ± 0.05  | 0.33 ± 0.08     | 0.383 |
| HSP90AB1 | -0.06 ± 0.06     | 0.12 ± 0.02  | 0.23 ± 0.08     | 0.901 |
| HSPA1B   | -0.17 ± 0.06     | 0.21 ± 0.11  | 0.35 ± 0.10     | 0.985 |
| HSPA8    | -0.03 ± 0.04     | 0.16 ± 0.04  | 0.25 ± 0.08     | 0.564 |
| HSPD1    | -0.02 ± 0.06     | 0.17 ± 0.06  | 0.27 ± 0.10     | 0.826 |
| HSPB1    | -0.09 ± 0.11     | 0.08 ± 0.07  | 0.12 ± 0.07     | 0.555 |

Values are expressed as means ± SE. C, control; LBH, lower body heating; TH, unilateral thigh heating; H-C, difference in mRNA fold change between the heated thigh and the control thigh; P, P value for the comparisons between LBH and TH.

ing the control intervention on both protocols (Figs. 3 and 6). This response might be potentially explained by circadian oscillations in gene expression (56), changes in metabolic status (79), or forced physical inactivity for a long period of time. In agreement with our findings, prior studies have demonstrated that the mRNA expression of CX3CL1 tends to decrease slightly over time in resting/nonexercising skeletal muscle (6, 70). Intriguingly, however, the expression of CCL2 has been reported to increase, albeit not significantly, in resting muscle 2–4 h following a baseline biopsy (6, 70, 78). These changes in the expression of key myokines illustrate the critical importance of incorporating a control group in long experiments to account for time-dependent variations in skeletal muscle gene expression in humans.

#### Heat Shock Proteins and Angiogenesis

Most of the salutary effects of heat stress on skeletal muscle are thought to be mediated, in part, by HSPs (25). This highly conserved family of proteins function as molecular chaperones, and HSPs are involved in multiple cellular activities, including angiogenesis signaling (58, 67, 71). For instance, members of the HSP70 and HSP90 families have been reported to modulate angiogenesis in skeletal muscle (58, 67). Pharmacological blockade of HSP70s impairs endothelial cell migration and tube formation in vitro and abrogates capillary growth in a model of peripheral arterial insufficiency (67). Similarly, administration of HSP90 inhibitor 17-DMAG suppressed the angiogenic response to repeated dry infrared sauna therapy in mice with hindlimb ischemia (40). The HSP90 inhibitor 17-AAG has also been shown to impair migration and formation of capillary-like tubes in cultured human umbilical vein endothelial cells (HUVEC) (71). Given the importance of these factors for angiogenesis, it is surprising that few studies have determined the impact of passive heat stress on the expression of HSPs in skeletal muscle in humans. Morton et al. (44) showed that leg immersion in water at 45°C for 1 h had no

effect on muscle protein content of HSPs in healthy young volunteers (44). Vardiman et al. (77) also found that a single session of local leg heating with fluidotherapy did not alter the content of HSP70 and HSP27p in skeletal muscle. In contrast, Touchberry et al. (75) reported that local heating with diathermy and hot packs increase the skeletal muscle content of HSP70 and HSP27p in female but not male subjects. Our study focused solely on the immediate transcriptional responses of several members of distinct HSP families. We report that both LBH and TH promote a rapid and very consistent upregulation of the mRNA expression of HSPs in human skeletal muscle, including members of the HSP90 and HSP70 families (Figs. 4 and 7). As these proteins have been shown to modulate angiogenesis in skeletal muscle, it is conceivable that the angiogenic response to heat therapy is regulated and potentially dependent on heat-induced activation of HSPs.

#### Potential Mechanisms

One hypothesis of the present study was that the skeletal muscle angiogenic response to LBH would be greater than that induced by TH, because in addition to activating local proangiogenic signals, systemic heat stress promotes physiological responses that can influence angiogenesis. For example, as discussed above, whole body heat stress has been shown to trigger the release and recruitment of bone marrow-derived proangiogenic cells (49, 66). Further, whole body heat stress evokes a marked increase in sympathoadrenal activity and an associated elevation in circulating levels of catecholamines (27). Catecholamines have been previously shown to contribute to skeletal muscle angiogenesis in a model of peripheral vascular insufficiency (7). Contrary to our initial predictions, exposure to LBH did not alter the levels of cytokines and proangiogenic cells and, most importantly, the increase in mRNA expression of angiogenic factors in skeletal muscle was remarkably similar between LBH and TH (Table 7). These findings suggest that the angiogenic response to both strategies derive from the activation of local signals capable of initiating the angiogenic cascade. One such mechanism would be the increase in skeletal muscle blood flow and shear stress that occurs during heat therapy application. Increased shear stress is known to induce the expression of several proangiogenic genes and provoke remodeling of the skeletal muscle microvasculature (4). The observed progressive increase in blood flow and shear rates in the femoral artery during TH (Table 6) is consistent with other recent reports (9, 10). Of note, the magnitude of change in thigh blood flow is remarkably similar during isolated leg and moderate whole body heating, which indicates that this hyperemic response is controlled by local mechanisms (9, 10). Until recently, the prevailing view was that these increases in bulk limb blood flow in response to heat stress were solely the result of changes in skin blood flow (see Ref. 12 for review). However, following the seminal report of Keller et al. (29), numerous studies have provided evidence that skeletal muscle blood flow also increases during local hyperthermia (10, 24, 55). For example, Heinonen et al. (24) showed using positron emission tomography that leg heating increases blood flow in the gastrocnemius muscle by ~1 ml·min<sup>-1</sup>·100 g<sup>-1</sup> (24). Although this change is considered small compared with the hyperemic responses induced by interventions such as ex-

ercise (12), it is fair to speculate that sustained increases in local flow during 90 min of heat therapy application and in the recovery period might induce a local change in the expression of angiogenic factors.

On the other hand, emerging evidence indicates that heat stress can promote angiogenesis independent of changes in blood flow. Rattan et al. (60) first showed that preexposure of HUVEC and human dermal microvascular endothelial cells to mild heat stress improved their ability to form capillary-like tubes. More recently, Li et al. (32) demonstrated that mild heat stress increased the expression of VEGF and angiopoietins and enhanced the formation of microvessel-like structures in a coculture system of outgrowth endothelial cells and osteoblasts. These authors speculate that this potent angiogenic response is related to the aforementioned heat-induced expression of HSPs (59), which are known to independently regulate angiogenesis in cultured cells (71) and in animal models (40, 67).

### Clinical Implications

Heat therapy can be applied using several different modalities, including dry sauna (39), immersion in warm water (54), perfusion of hot water through a tube-lined garment (12), and other methods that are commonly used in rehabilitation settings, such as diathermy (75) and fluidotherapy (77). The use of liquid-circulating garments is attractive because this approach is simple, inexpensive, portable, and amenable for home-based application, which makes this option particularly suitable for patients with limited mobility. Although we choose to apply the treatment to the lower limbs, it is important to highlight that these garments can be customized to cover just about every region of the body. For instance, tube-lined suits that cover the whole body allow for a higher rate of heating and more significant changes in body core temperature than observed in the current study (12). This modality has been extensively employed to investigate thermoregulatory responses in older individuals (22), as well as in patients with hypertension (30), Type 2 diabetes (81), and chronic heart failure (13). It is conceivable that the higher magnitude of heat stress attained by this method could induce responses that are different than the ones evoked by targeting just the lower limbs.

On the other hand, local heating methods, such as the TH protocol used herein, afford the possibility of inducing substantial increases in tissue temperature, while maintaining body temperature stable. These strategies might be more appealing for patients that do not tolerate or have contraindications to whole body heat stress. Regardless of whether heat treatment is applied locally or systemically, the findings from the present study indicate that this therapeutic approach may serve as a practical tool to stimulate angiogenic signaling and promote vascular growth in skeletal muscle.

### Limitations

One potential limitation of our experimental design was that subjects were not allowed to drink water during the experiment, and it is possible that sweat-induced dehydration during the LBH protocol might have contributed to the observed changes in skeletal muscle gene expression. However, this confounding effect seems unlikely given that changes in the expression of angiogenic factors and heat shock proteins were remarkably similar between the LBH and TH protocols (Table

7), even though there is negligible dehydration during localized limb heating. In addition, Logan-Sprenger et al. (35) recently reported that mild dehydration induced by overnight fluid restriction had no significant impact on the expression of HSP72 in skeletal muscle in humans. Taken together, these observations indicate that mild dehydration does not seem to impact the expression of angiogenic factors and heat shock proteins in skeletal muscle. It remains to be determined whether more severe levels of dehydration can affect the expression of these factors in humans.

### Perspectives and Significance

Our results show that heat therapy, applied either to both legs or locally to one thigh of healthy young individuals, promotes the expression of key angiogenic mediators in skeletal muscle. These findings set the stage for future studies to test the hypothesis that repeated exposure to this therapy can lead to increased capillarization in healthy and diseased skeletal muscle. If this hypothesis is confirmed, heat therapy may become a practical, noninvasive therapeutic tool to reverse the rarefaction of the skeletal muscle capillary network that is commonly observed in individuals with chronic diseases such as COPD, PAD, and CHF.

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### DISCLOSURES

No conflicts of interest, financial or otherwise, are declared by the authors.

### AUTHOR CONTRIBUTIONS

A.M.K., K.K., D.N., Y.N., A.N.B., T.P.G., and B.T.R. performed experiments; A.M.K., K.K., D.N., Y.N., A.N.B., S.K., Q.S., and B.T.R. analyzed data; A.M.K., K.K., D.N., A.N.B., Q.S., T.P.G., and B.T.R. interpreted results of experiments; A.M.K., K.K., D.N., Y.N., A.N.B., B.J.W., S.K., J.S., Q.S., T.P.G., and B.T.R. approved final version of manuscript; K.K. and B.T.R. prepared figures; K.K., B.J.W., S.K., J.S., T.P.G., and B.T.R. edited and revised manuscript; B.J.W. and B.T.R. conception and design of research; B.T.R. drafted manuscript.

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