



Published in final edited form as:

Bone. 2017 June ; 99: 26–38. doi:10.1016/j.bone.2017.02.014.

TNF α Contributes to Diabetes Impaired Angiogenesis in Fracture Healing

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Abstract

Diabetes increases the likelihood of fracture, interferes with fracture healing and impairs angiogenesis. The latter may be significant due to the critical nature of angiogenesis in fracture healing. Although it is known that diabetes interferes with angiogenesis the mechanisms remain poorly defined. We examined fracture healing in normoglycemic and streptozotocin-induced diabetic mice and quantified the degree of angiogenesis with antibodies to three different vascular markers, CD34, CD31 and Factor VIII. The role of diabetes-enhanced inflammation was investigated by treatment of the TNF α -specific inhibitor, pegsunercept starting 10 days after induction of fractures. Diabetes decreased both angiogenesis and VEGFA expression. The reduced angiogenesis and VEGFA expression in diabetic fractures was rescued by specific inhibition of TNF *in vivo*. In addition, the TNF inhibitor rescued the negative effect of diabetes on endothelial cell proliferation and endothelial cell apoptosis. The effect of TNF α *in vitro* was enhanced by high glucose and an advanced glycation endproduct to impair microvascular cell proliferation and

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Disclosures

All authors state that they have no conflicts of interest.

Authors's roles: Study design: LG, TE and DG. Study conduct and data interpretation: all authors. Drafted and revised manuscript: JL, KK, DG. Approved final version of manuscript: all authors. DG is responsible for the integrity of the manuscript.

stimulate apoptosis. The effect of TNF, high glucose and an AGE was mediated by the transcription factor FOXO1, which increased expression of p21 and caspase-3. These studies indicate that inflammation plays a major role in diabetes-impaired angiogenesis in endochondral bone formation through its effect on microvascular endothelial cells and FOXO1.

Keywords

blood vessel; cytokine; endothelial; forkhead; hyperglycemia; inflammation; vascularization; VEGF

Introduction

Diabetes impairs bone formation and delays fracture healing, presenting a challenge in the management of diabetic fractures. Diabetics also have more complications during fracture healing [1–3]. Fracture healing requires angiogenesis and an adequate blood supply [4, 5]. Interference with angiogenesis impairs fracture healing [6, 7] while treatment with angiogenic factors such as VEGFA and FGF-2 improves bone formation during repair [8, 9]. Diabetes leads to reduced angiogenesis that may contribute to the pathologic outcomes in diabetic complications [10]. Endothelial progenitor cells are significantly reduced in type 1 and 2 diabetic patients and in animal models [11]. Injection of CD34⁺ endothelial progenitors in diabetic mice augments vascularization and improves wound healing [12] and the application of angiogenic factors promotes vascularization and diabetic fracture healing in a rat model [13]. Although it is known that diabetes reduces angiogenesis in fracture repair the mechanisms for this decrease have not been established [13].

Diabetes has been shown to increase inflammation in fracture calluses that promotes early cartilage resorption and reduced mesenchymal stem cell numbers [14, 15]. The early inflammatory response is beneficial by inducing recruitment of mesenchymal stem cells and leukocytes, which produce factors to stimulate tissue repair and angiogenesis [16]. However, prolonged inflammation leads to deficient bone formation and impaired fracture healing [17–19]. In particular, tumor necrosis factor-alpha (TNF α) levels remain elevated in diabetic fractures, resulting in early apoptosis of chondrocytes and mesenchymal stem cells [15, 20]. Thus, elevated levels of TNF α later in fracture repair may contribute to poor fracture healing in diabetics.

In the current study, we demonstrate that diabetes hampers angiogenesis in areas of endochondral ossification during fracture healing and reduces VEGFA expression. Specific inhibition of TNF α with pegsunercept after the early phase restored blood vessel formation in diabetic fractures. Mechanistically, the TNF α -dependent changes were due to reduced expression of VEGFA and reduced proliferation and increased apoptosis of endothelial cells. The latter were mediated *in vitro* by the transcription factor FOXO1. Our findings indicate that the vascular deficit associated with fracture healing is due in part to diabetes-enhanced TNF α and that the control of inflammation during fracture repair may offer a pragmatic approach to augment diabetic fracture healing.

Materials and Methods

Animals

Diabetes was induced in 8-week-old male CD-1 male mice (Charles River Laboratories, Wilmington, MA) with a daily intraperitoneal injection of streptozotocin (STZ, 40mg/kg, Sigma-Aldrich, St. Louis, MO) for 5 days. Control mice were treated with vehicle alone (10mM citrate buffer). Mice were considered hyperglycemic when blood glucose levels were greater than 12.48mmol/l. STZ-induced diabetic mice received insulin treatment through insertion of slow release insulin implants as described previously [21] or i.p injection of pegsunercept, a TNF specific inhibitor as described below. A simple transverse closed fracture of the tibia (insulin studies) or femur (TNF inhibitor studies) was performed as previously described [21]. The articular surface of the tibia or femur was exposed and a 27-gauge spinal needle was inserted for fixation. After closure of the incision a fracture was created by blunt trauma. Any fractures not consistent with standardized placement criteria (mid-diaphyseal) or grossly comminuted were excluded. Animals were subsequently euthanized at the indicated time points. Bone was harvested with most of the muscle and soft connective tissue removed. Mice were hyperglycemic for at least 3 weeks prior to fracture. In some experiments, animals were treated with TNF- α inhibitor pegsunercept (4mg/kg, Amgen, Thousand Oaks, CA) by intraperitoneal injection every 3 days starting at 10 days post-fracture [15]. At the onset of experiments normoglycemic groups had mean glucose values of 6–8mmol/l and diabetic groups had mean values of 25–28mmol/l, which was not significantly affected by treatment with pegsunercept. Diabetic mice treated with insulin had mean glucose levels of 6mmol/l. Animals were euthanized at 10, 16 and 22 days after fracture. All procedures were approved by the Institutional Animal Care and Use Committee (IACUC).

Immunofluorescence/Immunohistochemistry

Fracture samples were fixed and decalcified as previously described [14]. Transverse cross-sections were cut at 5 μ m closest to the fracture callus center. Sections were deparaffinized and subjected to antigen retrieval in 10mM citric acid (pH 6.0) by pressure heating. ESM table 1 provides the list of antibodies and reagents utilized from this study. Tyramide amplification and 3,3'-Diaminobenzidine or Alexa-546 were used to localize the signals. Areas of endochondral ossification or mature bone were examined at 100 \times or 400 \times original magnification and 5–8 images per specimen were captured using a Nikon Eclipse 90i microscope. Image analysis was performed using NIS-Elements software (Nikon) under blinded conditions. Blood vessels were described as small or moderate vessels depending on the number of endothelial cells associated with the vessel. Blood vessels with 2–4 endothelial cells lining the vessel were categorized as small blood vessels, while vessels with 5 or more cells associated with it were considered moderate vessels. VEGFA immunopositive hypertrophic chondrocytes were counted in cartilage areas.

Cell culture, transfection, and qPCR

In vitro experiments were carried out with human microvascular endothelial cells (HMVEC) from Cell Systems (Kirkland, WA) and maintained in EGM-2 MV growth medium (Lonza, Walkersville, MD) with 5% FBS. HMVECs were transfected with 10nM FOXO1 siRNA

using GenMute siRNA transfection reagent according to the manufacturer's instructions. Total RNA was isolated using Quick-RNA MicroPrep kit (Zymo Research, Irvine, CA), and real-time PCR was performed on a StepOnePlus real-time PCR system (ABI). Results were normalized to gene levels of ribosomal protein L32.

In vitro experiments—HMVECs were grown to 70% confluence and then starved in EGM-2 MV medium with 0.1% FBS and without growth supplements for 24 hours. Cells were then cultured in the same media without or with TNF α (2 or 10 ng/mL) for 2 days, high glucose supplementation (17 mM) for 5 days, or advanced glycation endproduct (200 μ g/mL) (AGE; carboxymethyllysine modified albumin) for 3 days prepared as previously described [22]. Apoptosis was measured with annexin-V kit and caspase-3 antibody. Serum-stimulated proliferation with 5% FBS was measured with p21 antibody and BrdU assay kit (Cell Signaling, Danvers, MA) according to manufacturer's instructions. Quantification of BrdU was determined using a microplate reader (Infinite M200 PRO, Tecan, Morrisville, NC) set to read absorbance at 450 nm. Images were captured with a Nikon Eclipse 90i microscope and mean fluorescence intensity (MFI) measured using NIS Elements software. HMVECs were grown to 70% confluence with 5% FBS EGM2-MV and then incubated in media supplemented with TNF (2 ng/mL), high glucose (17 mM) or an AGE, CML-albumin (200 μ g/mL) for 3 days prepared as previously described [22]. Cells were then transferred to 96 well plates coated with Matrigel Basement Membrane Matrix (LDEV-free, Corning) in triplicate and cultured in EBM media (Lonza). Formation of tubes by human microvascular endothelial cells was examined 18 hrs later using HMVECs cultured in Matrigel as described [23]. The tube number and tube length per well were counted using NIS Elements software (Nikon).

Data analysis

All data are expressed as mean \pm standard error. Statistical analyses were determined using SPSS software (SPSS, Chicago, IL). Statistical significance was defined as $p < 0.05$. One-way ANOVA with Tukey post-hoc test was performed to determine statistical significance at a single time point, while two-way ANOVA with Tukey post-hoc was used to analyze data across time points.

Results

Diabetes decreases angiogenesis in areas of endochondral bone formation

To examine the effect of diabetes on angiogenesis, immunostaining was carried out in mouse fracture calluses 10 and 16 days post-fracture, during which robust angiogenesis takes place in association with cartilage removal and new bone formation. Antibody to CD34 was used to identify single endothelial progenitor cells as well as newly formed blood vessels. Approximately 15% of the total cell population in normoglycemic mice was CD34⁺ at 10 and 16 days after fracture (Fig. 1C). The number of CD34⁺ single cells was decreased in diabetic mice by 30% ($p < 0.05$) (Fig. 1C). Insulin treatment reversed this reduction, indicating that it was not an artifact of streptozotocin treatment, but rather a direct result of diabetes ($p < 0.05$) (Fig. 1C). The number of small and moderate vessels increased between days 10 and 16 by approximately 50% concurrent with the transition from cartilage to bone

(Fig 1 A, B, D and E)($p<0.05$). Diabetes reduced the number of small and moderate sized CD34⁺ vessels by 40–50%, which was rescued by insulin ($p<0.05$) (Fig 1 A, B, D and E). Blood vessels were also identified by antibody to Factor VIII. The number of vessels was increased by 65%–80% in normoglycemic mice between day 10 and day 16 ($p<0.05$) (Fig 1F–G). Diabetes reduced the number of Factor VIII⁺ small and moderate sized blood vessels at 10 and 16 days by 44–50%, which was rescued by insulin ($p<0.05$) (Fig 1F–G). The number of Factor VIII⁺ vessels on day 22 were also reduced in the diabetic group, a time point when primary bone formation has largely been completed ($p<0.05$) (Fig. 1H–I).

Diabetes induced reduction in angiogenesis is mediated by elevated TNF α

The hypothesis that elevated TNF α in diabetic animals affected angiogenesis was tested with a TNF α -specific inhibitor, pegsunercept starting at 10 days post fracture to not interfere with early events in repair. Diabetic animals had reduced numbers of blood vessels which was largely restored by inhibition of TNF (Supplemental Fig 1). The number of individual CD34⁺ endothelial cells and the number of small and moderate sized blood vessels was reduced by 35–50% in diabetic mice ($p<0.05$), which was largely reversed by treatment of diabetic mice with pegsunercept ($p<0.05$) (Fig. 2A–C). Pegsunercept had little effect on angiogenesis in normal mice (Fig. 2A–C). The same pattern was observed when CD31 or Factor VIII was used as a marker of endothelial cells (Fig. 2E–G). Pegsunercept treatment rescued the reduction in CD31⁺ single cells and blood vessels (Fig. 2E–G) and Factor VIII⁺ vessels (Fig. 2H–I) caused by diabetes ($p<0.05$) but had little effect on angiogenesis in normoglycemic controls.

TNF Suppresses VEGFA Expression in Vivo and Inhibits Tube Formation in Vitro

Expression of VEGFA is an important early step in angiogenesis and one of the principal drivers of angiogenesis in bone. We assessed VEGFA expression in areas of bone formation during fracture healing. Hypertrophic chondrocytes were shown to be high expressers of VEGFA (Supplemental Fig 2). Diabetes significantly reduced the number of hypertrophic chondrocytes that expressed VEGFA by ~50% compared to normoglycemic fractures ($p<0.05$) (Fig 3A). Treatment of diabetic animals with the TNF α inhibitor pegsunercept rescued the negative effect of diabetes on VEGFA expression ($p<0.05$) (Fig 3A).

Tube formation reflects terminal aspects of blood vessel formation and is a critical step in angiogenesis. We examined the impact of three factors that are elevated in diabetes, TNF α , an advanced glycation endproduct and the effect of high glucose (Fig 3B and C). Moderate doses of TNF α (2ng/ml), AGE (200 μ g/ml) or high glucose (17mM) inhibited tube formation measured as either the number of tubes formed or the total tube length. The combination of all three was significantly greater than TNF, AGE or high glucose alone.

TNF α reduces proliferation of endothelial cells

Double immunofluorescence was carried out with antibodies to CD34 and Ki67 to examine the effect of diabetes on endothelial cell proliferation *in vivo*. The total number of proliferating Ki-67⁺ cells was reduced 45–50% by diabetes ($p<0.05$) (Fig. 4A) and the number of proliferating single endothelial cells (Ki67⁺/CD34⁺) or proliferating endothelial

cells associated with blood vessels was reduced by 40% in diabetic mice ($p < 0.05$) (Fig. 4B). Insulin reversed the impaired angiogenesis in the diabetic group ($p < 0.05$) (Fig. 4A–D).

Diabetes-reduced proliferation for all cell types was reversed by treatment with either insulin or pegsunercept ($p < 0.05$) (Fig. 4A and 5A). Proliferating single Ki67⁺/CD34⁺ endothelial cells or those associated with blood vessels were decreased by 40 – 54% in the diabetic group and reversed by inhibiting TNF ($p < 0.05$) (Fig. 5B–D). Pegsunercept had little effect in the total number of proliferating cells (Fig 5A) or in the number of proliferating endothelial cells in normoglycemic mice (Fig. 5B & E). Endothelial cell proliferation was reduced by diabetes and largely restored by inhibition of TNF when CD31 was used as a marker of endothelial cells (Fig. 5E–G) and little effect was seen in vessels of normoglycemic mice.

TNF α , AGE and HG inhibit endothelial cell proliferation that is FOXO1 dependent

In vitro experiments were carried out with primary cultures of microvascular endothelial cells to investigate the impact of TNF α and TNF α in combination with high glucose and advanced glycation end-products (AGE) based on evidence that each is elevated by diabetes [24]. TNF α , AGE and high glucose each reduced microvascular endothelial cell DNA synthesis ($p < 0.05$) (Fig. 6A). Low dose TNF α with high glucose and AGE had a similar effect as high dose TNF α (Fig. 6B). We determined whether the transcription factor FOXO1 mediated the effect of TNF α , AGE and high glucose on vascular endothelial cell proliferation. Both the effect of low dose TNF α +HG+AGE and high dose TNF α was mediated by FOXO1 as shown by substantial reversal with FOXO1 silencing ($p < 0.05$) (Fig. 6B). The knockdown efficiency of siFOXO1 was approximately 70% (Supplemental Fig 3).

To better understand the effect of FOXO1 in inhibiting DNA synthesis, p21, which inhibits cell cycle progression [25] was measured. TNF α increased p21 mRNA 2-fold and p21 protein 3.3-fold ($p < 0.05$) (Fig. 9A and C). The addition of high glucose and AGE elevated the expression of p21 7.6-fold ($p < 0.05$) (Fig. 9C). High dose TNF α and low dose TNF α +HG+AGE stimulated p21 was largely reversed by FOXO1 knockdown ($p < 0.05$) (Fig. 9C).

TNF α increases apoptosis of endothelial cells in diabetic fractures

Another potential mechanism for reduced angiogenesis is increased endothelial cell apoptosis. The overall level of apoptosis reflected by the total number of TUNEL⁺ cells was increased ~2 fold by diabetes ($p < 0.05$) (Fig. 6A) and restored by insulin treatment. The number of TUNEL⁺/CD34⁺ single cells and TUNEL⁺/CD34⁺ cells associated with blood vessels was elevated almost 3-fold in diabetic mice ($p < 0.05$), which was substantially improved with insulin treatment ($p < 0.05$) (Fig. 6B–D).

Pegsunercept treatment reversed the level of apoptosis for all cell types in diabetic animals but had little effect on the normoglycemic group ($p < 0.05$) (Fig. 7A). Diabetes doubled the number of TUNEL⁺/CD34⁺ positive apoptotic single endothelial cells or TUNEL⁺/CD34⁺ endothelial cells associated with blood vessels ($p < 0.05$), which was returned to almost normal levels with pegsunercept treatment ($p < 0.05$) (Fig. 7B–D). Pegsunercept had little effect on apoptosis in normal mice (Fig. 7A–D).

The effect of TNF α on microvascular endothelial cell apoptosis and proliferation is enhanced by high glucose and an advanced glycation end product

In vitro experiments were carried out with primary cultures of microvascular endothelial cells to determine the impact of TNF α with high glucose and advanced glycation end-products (AGE) that are also elevated by diabetes. TNF α , AGE and high glucose each reduced HMVEC DNA synthesis ($p < 0.05$) (Fig. 8A). Low dose TNF α with high glucose and AGE had a similar effect as high dose TNF α (Fig. 8C). We determined whether the transcription factor FOXO1 mediated the effect of TNF α , AGE and high glucose on vascular endothelial cell proliferation. Both the effect of low dose TNF α +HG+AGE and high dose TNF α was mediated by FOXO1 as shown by substantial reversal with FOXO1 silencing ($p < 0.05$) (Fig. 8C). The knockdown efficiency of siFOXO1 was approximately 70% (Fig 8B and D). Endothelial cell apoptosis was also increased by TNF α , AGE and high glucose ($p < 0.05$) (Fig. 8E) and substantially reversed by FOXO1 siRNA compared to scrambled siRNA ($p < 0.05$) (Fig. 8F).

Caspase-3 is one of the primary effector caspases induced by TNF α . TNF α stimulated a 2.2 fold increase in caspase-3 mRNA and a 3.9-fold increase in activated caspase-3 protein ($p < 0.05$) (Fig. 9B–D). The combination of TNF α , high glucose and AGE led to 6.5-fold increase in activated caspase-3, which was attenuated by FOXO1 silencing ($p < 0.05$) (Fig. 9D). Thus, the addition of an AGE and high glucose to TNF α further stimulates the expression of genes that regulate cell cycle and apoptosis in a FOXO1 dependent manner.

Discussion

Studies presented here demonstrate that diabetes markedly reduces angiogenesis in fracture healing in areas undergoing new bone formation. Angiogenesis was examined using three different markers, CD34, CD31 and Factor VIII to assess the full spectrum of endothelial cells. The anti-angiogenic effect of diabetes coincided with a decrease in VEGFA expression. Both reduced angiogenesis and reduced VEGFA expression were mediated by TNF α as demonstrated by rescue with a TNF-specific inhibitor. Moreover, diabetes enhanced apoptosis and decreased proliferation of endothelial cells *in vivo*. TNF α inhibition restored proliferation and prevented apoptosis, leading to a recovery in the number of blood vessels found in the fracture callus. TNF α reduced the capacity of microvascular endothelial cells to respond to growth factors present in serum *in vitro* and stimulated apoptosis of microvascular endothelial cells. The effect of TNF α was enhanced by other factors elevated in diabetics, high glucose and AGEs in a FOXO1 dependent manner.

In this study, markers specific to CD34, CD31 and Factor VIII were used to assess angiogenesis in fracture healing. CD34 positive cells in the healing callus represent putative endothelial progenitor cells [26]. CD34 is also expressed by endothelial cells in capillaries and small blood vessels [27]. CD31 is expressed in immature microvascular endothelial cells as well as mature endothelial cells in small to large-size vessels [28] and Factor VIII is primarily expressed in mature blood vessels [28]. Diabetes significantly reduced the number of endothelial cells and blood vessels regardless of which marker was examined in areas of endochondral ossification. This is likely to be problematic as angiogenesis is needed for fracture healing and is indispensable for new bone formation [29]. Reduced angiogenesis

can be explained by lower levels of endothelial cell proliferation and increased apoptosis. It may also be due to reduced numbers of circulating CD34+ and CD31+ cells [30]. Insulin treatment restored angiogenesis in diabetic animals, indicating that impaired angiogenesis was a direct result of diabetes and not an artifact of streptozotocin treatment.

TNF α expression is dysregulated and prolonged in diabetic fracture calluses and in areas of endochondral ossification [15, 20]. We previously reported that local levels of TNF α in the fracture callus are increased in diabetic mice compared to normoglycemic mice [15, 20]. Although TNF α is reported to be a pro-angiogenic cytokine that is necessary for vascular infiltration in tumorigenesis [31] and to promote physiologic fracture healing in the early phases [32], our findings indicate that in later stages of diabetic fracture healing TNF α interferes with angiogenesis. In contrast, TNF α inhibition in normal animals in the same timeframe had little effect on angiogenesis.

The proliferative capacity of microvascular endothelial cells is critical for proper angiogenesis [33]. Our results suggest that diabetes reduces the expression of VEGFA and that reduced VEGFA expression is rescued by inhibiting TNF α . The results are striking given a recent report that VEGFA plays a critical role in angiogenesis as well as osteoblast differentiation associated with new bone formation [34]. Moreover, TNF α reduced the proliferative capacity of endothelial cells to respond to growth factors. The latter is mediated through FOXO1 and enhanced by the presence of other factors that are elevated in diabetes, namely, high glucose and advanced glycation end products. Previous studies have shown that endothelial progenitor cells isolated from type II diabetic patients have decreased proliferative capacity *in vitro*, consistent with our *in vivo* data [30]. Elevated TNF α can also affect endothelial cell numbers through increased apoptosis. Inhibition of TNF α rescued the high levels of endothelial cell apoptosis *in vivo*, and TNF α *in vitro* induced endothelial cell apoptosis.

We showed that the effect of TNF α , high glucose and an AGE on microvascular endothelial cells was mediated by FOXO1. All three reduced endothelial proliferation and increased apoptosis. FOXO1 was needed for each of these effects and was shown to enhance expression of p21, which reduces progression through the cell cycle and increase caspase-3, which is a major effector of apoptosis. TNF α and FOXO1 have also been shown to play a role in apoptosis of microvascular endothelial cells in diabetic retinopathy [35]. Our *in vitro* results suggest that the effect of TNF, AGEs and high glucose on endothelial cells is direct although this does not rule out an indirect effect from the impact the diabetes on hypertrophic chondrocytes. This indirect effect may be important since we show here that chondrocytes may be an important source of VEGFA. Thus, FOXO1 mediates at least some of the negative effects of diabetes on angiogenesis during fracture healing and as a result is likely to impair fracture healing under diabetic conditions. FOXO1 may also have a significant effect on chondrocytes in diabetic fractures based on our previous reports that it mediates increased chondrocyte apoptosis and expression of chemokines by chondrocytes [36, 37].

In summary, diabetes has a significant effect on angiogenesis during fracture repair, reducing vessel formation and VEGFA expression, which can be directly linked to the impact of

diabetes-induced TNF α dysregulation. Furthermore, TNF α along with other factors present in diabetes such as high glucose and an AGE impair the ability of microvascular endothelial cells and their progenitors to respond to proliferative factors and promote apoptosis of these cells. Effects of TNF, high glucose and AGE may be mediated by FOXO1 that regulates antiproliferative and pro-angiogenic factors. In contrast, TNF α levels present during normal fracture repair are not problematic and did not limit angiogenesis.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

This work was supported by NIH grants AR060055 and DE019108. We thank Chandani Patel and Srujana Kraleti for their assistance.

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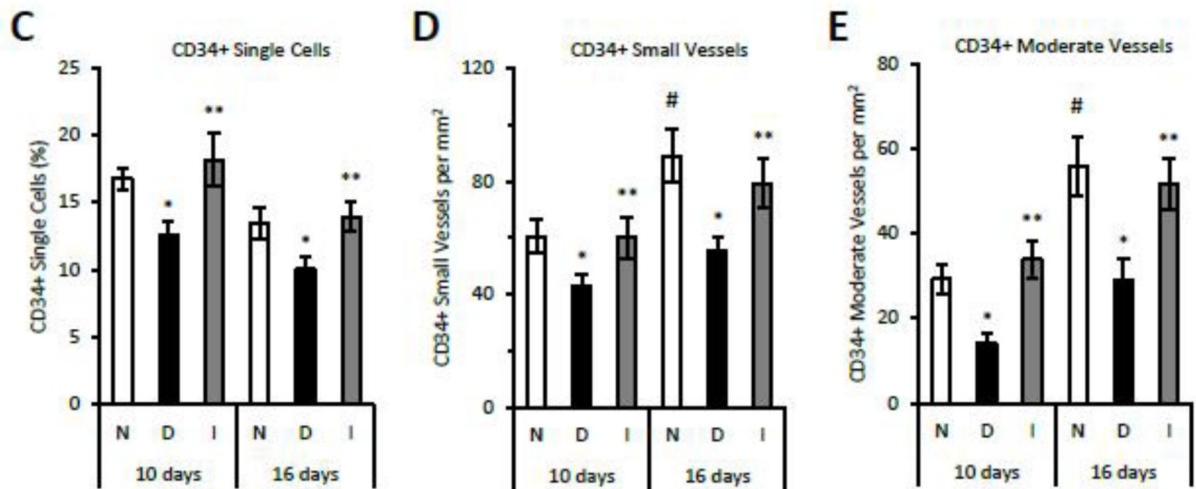
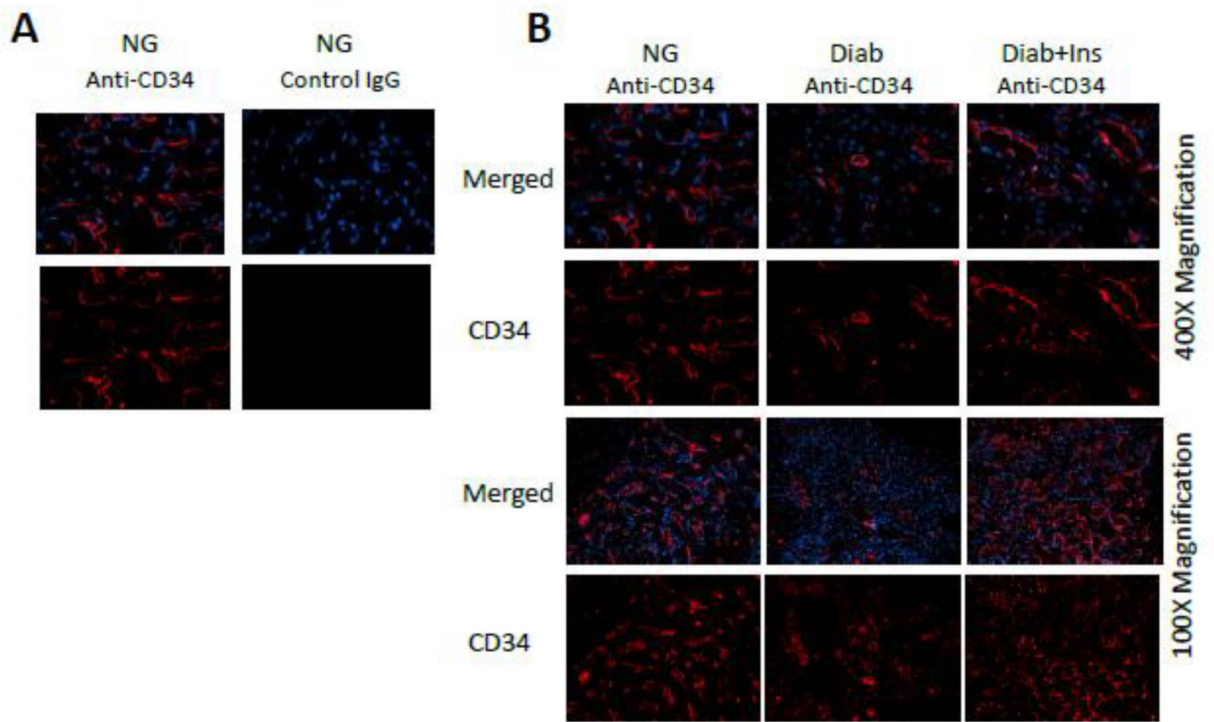
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Highlights

- Diabetes decreased both angiogenesis and VEGFA expression.
- The reduced angiogenesis and VEGFA expression in diabetic fractures was rescued by specific inhibition of TNF *in vivo*.
- TNF inhibition rescued the negative effect of diabetes on reduced endothelial cell proliferation and increased endothelial cell apoptosis.
- The effect of TNF α *in vitro* was enhanced by high glucose and an advanced glycation endproduct to impair microvascular cell proliferation and enhance apoptosis.
- The effect of TNF, high glucose and an AGE was mediated by the transcription factor FOXO1, which increased expression of p21 and caspase-3.



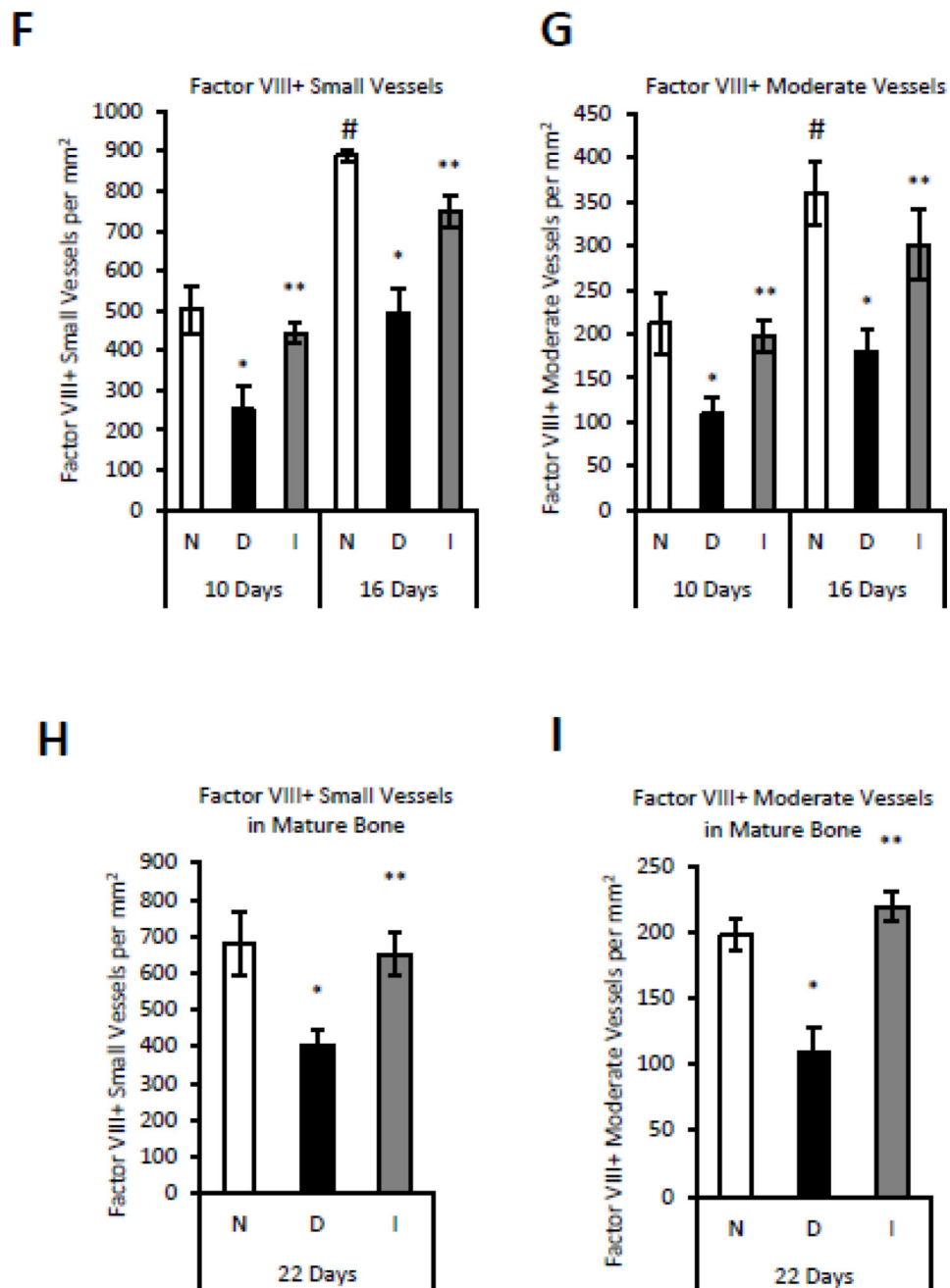
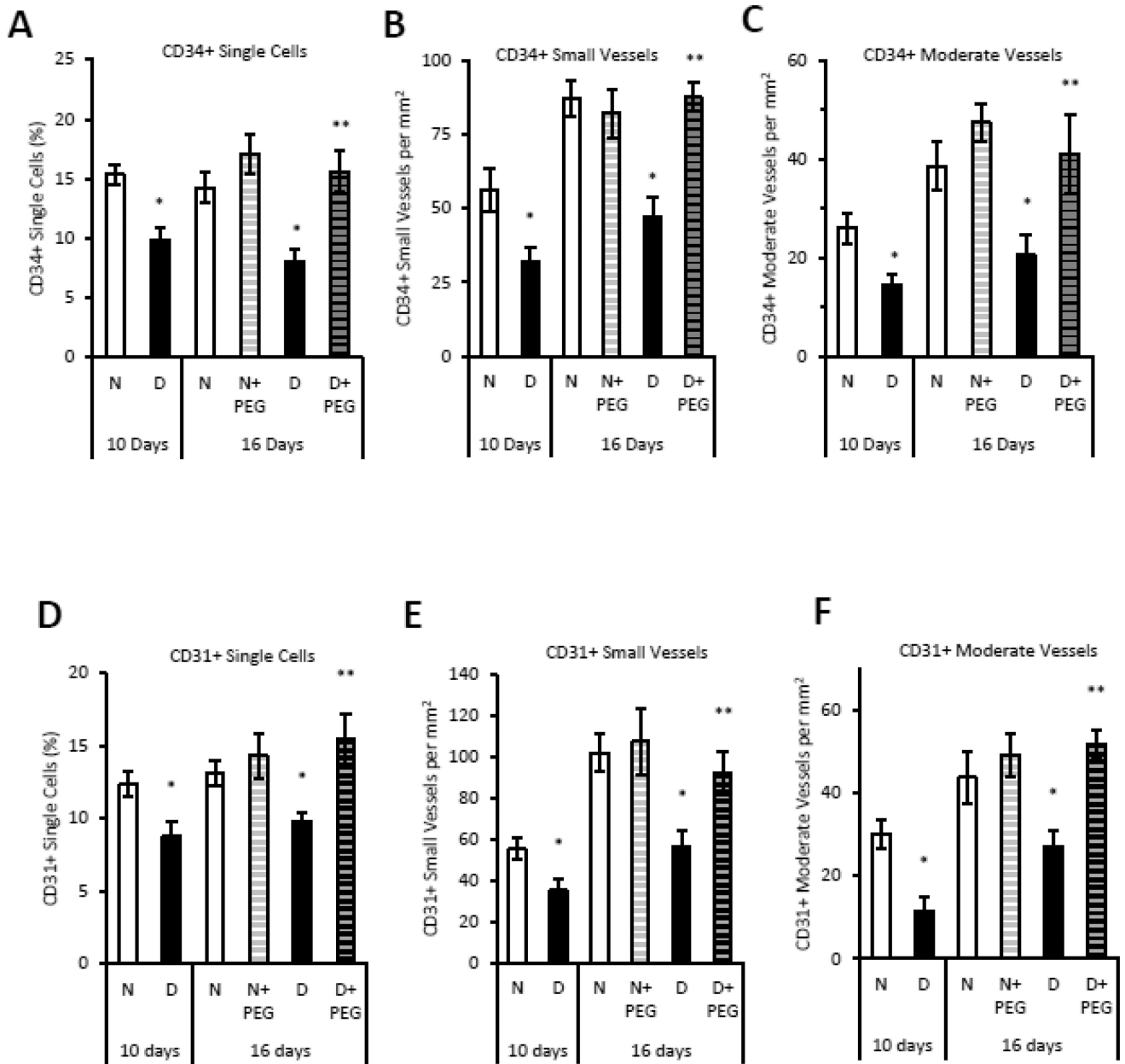


Figure 1. Diabetes decreases CD34+ and Factor VIII+ blood vessels in areas of endochondral bone formation and mature bone during fracture repair

A: Immunofluorescent images following incubation with anti-CD34⁺ IgG or matched control IgG in normoglycemic mice and **B:** immunofluorescent images of CD34⁺ expression in normoglycemic (N), diabetic (D) and insulintreated diabetic (I) mice from areas of bone formation in 16 day fracture specimens (400× or 100× magnification). **C–E:** CD34 immunopositive single cells, small vessels or moderate-sized vessels were analyzed as described in the Methods at 10 and 16 days after fracture. **F–G:** Factor VIII positive blood vessels were detected by immunohistochemistry in areas of bone formation at 10 and 16

days post fracture. **H-I:** Factor VIII positive blood vessels were quantified in areas of mature bone at 22 days after fracture. N=5-6 per group. *p<0.05 between normal vs diabetic groups, **p<0.05 between diabetic and insulin-treated groups, # p<0.05 between corresponding groups on day 10 and 16.



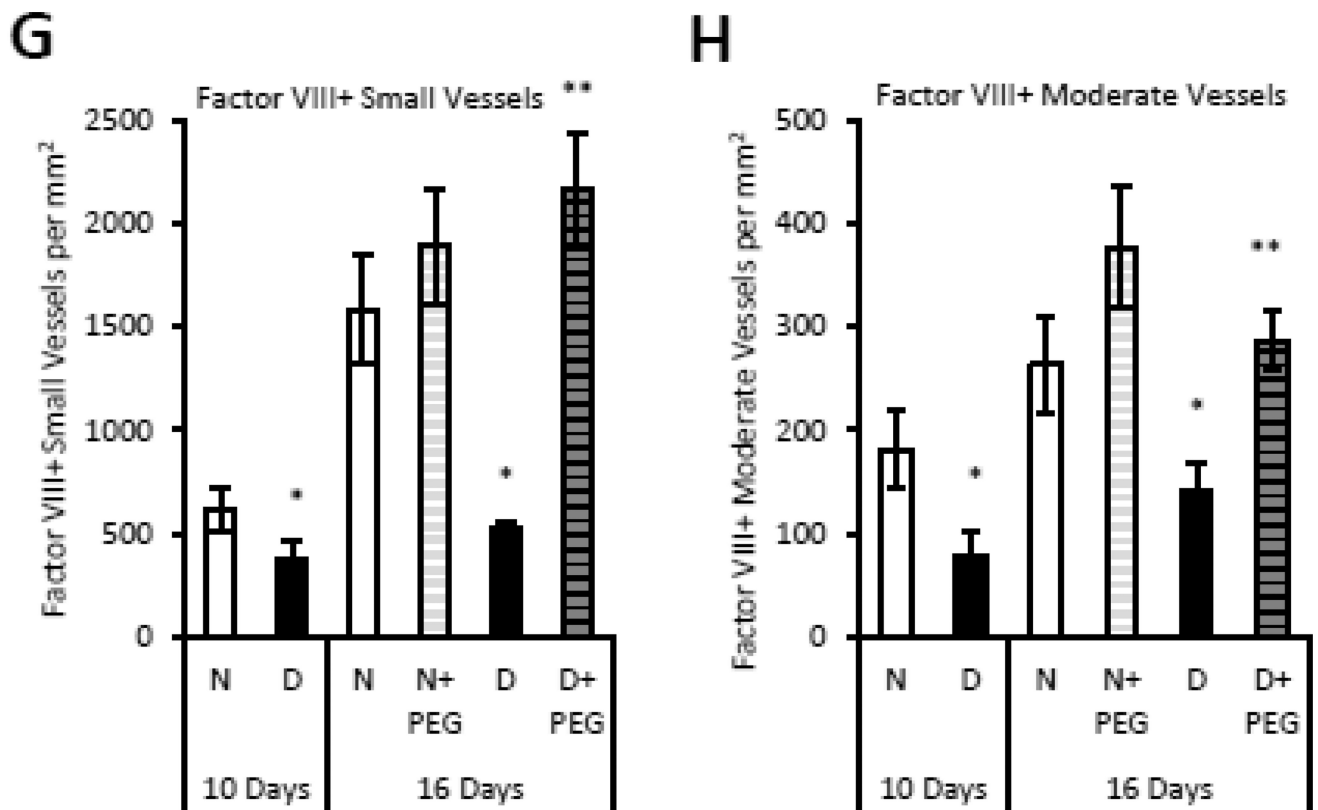


Figure 2. Reduced angiogenesis in diabetic fracture healing is mediated by elevated TNF α . Areas of endochondral ossification from fracture calluses were examined by immunofluorescence and immunohistochemistry in normoglycemic mice (N), diabetic mice (D) or diabetic mice treated 10 days post-fracture with TNF α -specific inhibitor pegsunercept (D+PEG). **A–C:** CD34 immunopositive cells and blood vessels in fracture specimens from normoglycemic, diabetic and diabetic+PEG mice in areas of bone formation. **D–F:** CD31 immunopositive cells and blood vessels were counted in areas of bone formation in healing fractures. **G–H:** Factor VIII immunopositive small and moderate-sized blood vessels were counted in healing fractures. N=5–6 per group. *p<0.05 between normal vs diabetic groups, **p<0.05 between 16-day diabetic and pegsunercept-treated diabetic groups.

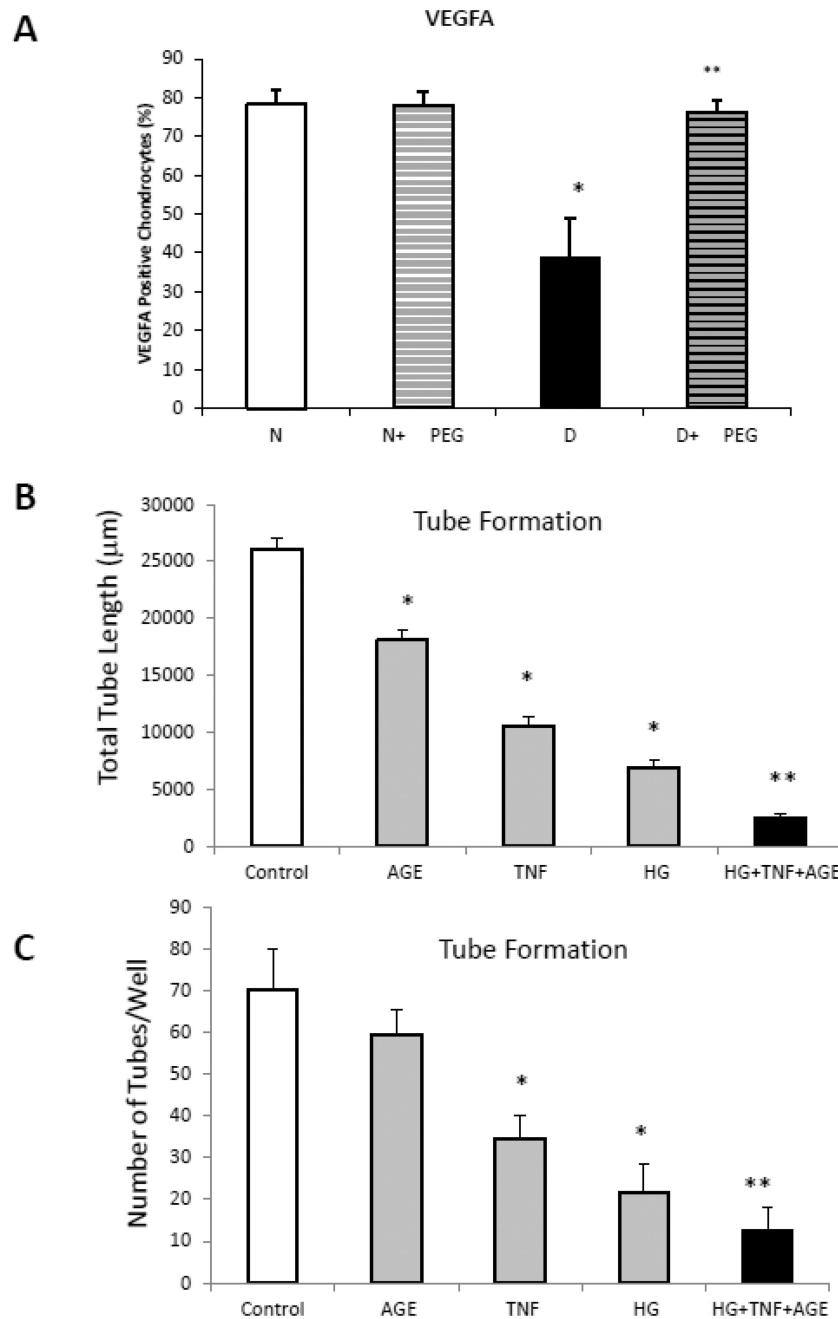


Figure 3. VEGFA expression and tube formation are inhibited by TNF

A. VEGFA expression by hypertrophic chondrocytes was measured by immunofluorescence. **B–C.** Tube form by microvascular endothelial cells was examined in vitro without or with incubation in media supplemented with an AGE (CML-albumin, 200 $\mu\text{g}/\text{ml}$), TNF α (200ng/ml), high glucose (17mM) or a combination of all three. Data are expressed as total tube length per well (B) or number of discrete tubes per well (C). * $P < 0.05$ compared to untreated control; ** $P < 0.05$ compared to individual factors.

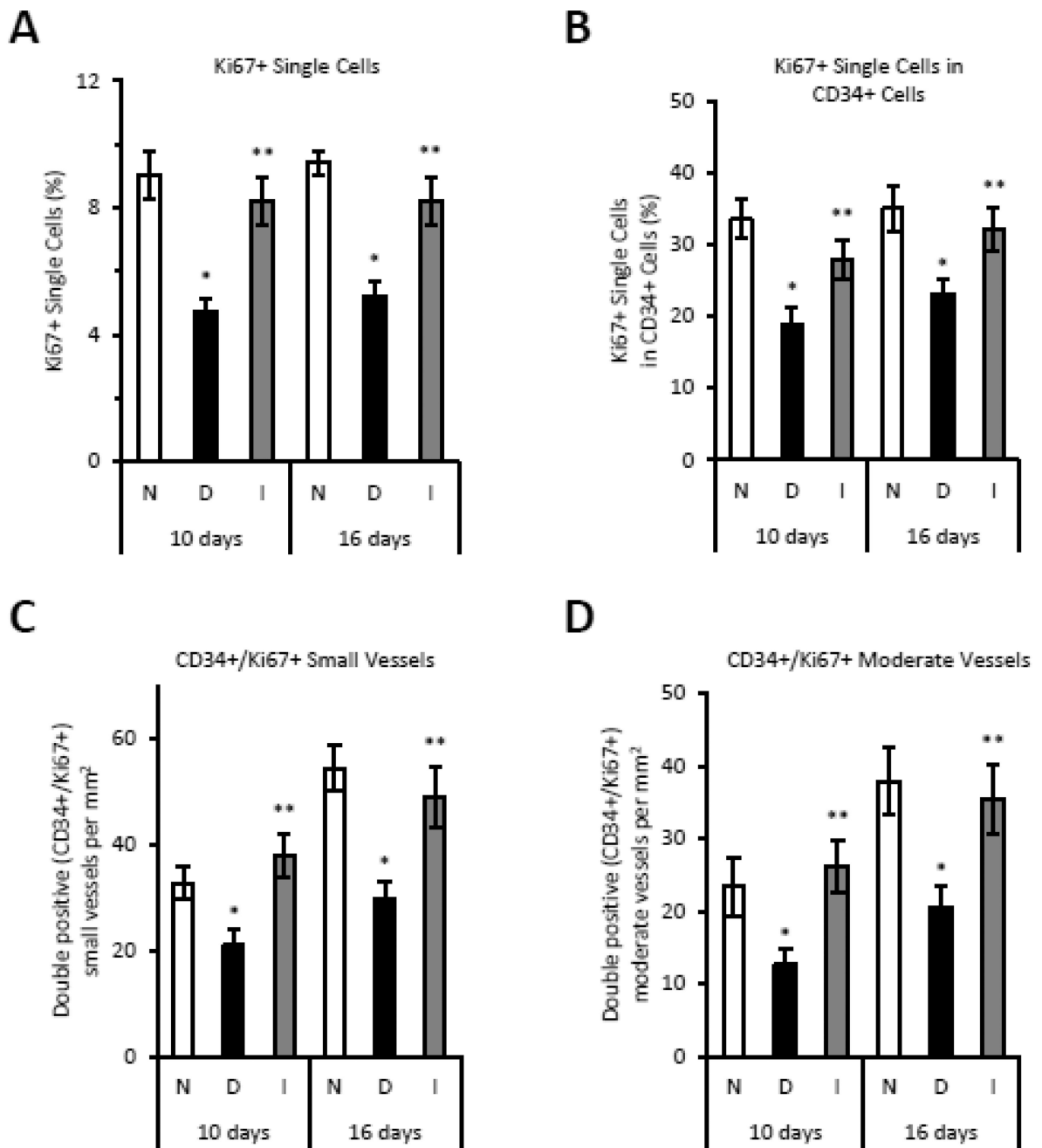


Figure 4. Proliferation of endothelial cells is decreased in diabetes in areas of endochondral bone formation

CD34 and Ki67 positive cells or blood vessels were detected by single or double immunofluorescence at 10 and 16 days after fracture. Groups included were normoglycemic (N), diabetic (D) or diabetic treated with insulin (I). **A:** The total percent Ki67 immunopositive single cells. **B:** Percent CD34 immunopositive single cells that are also Ki67 immunopositive. **C–D:** Percent CD34 immunopositive blood vessels that are also Ki67 immunopositive. * $p < 0.05$ between normal vs diabetic groups, ** $p < 0.05$ between diabetic and insulin-treated groups.

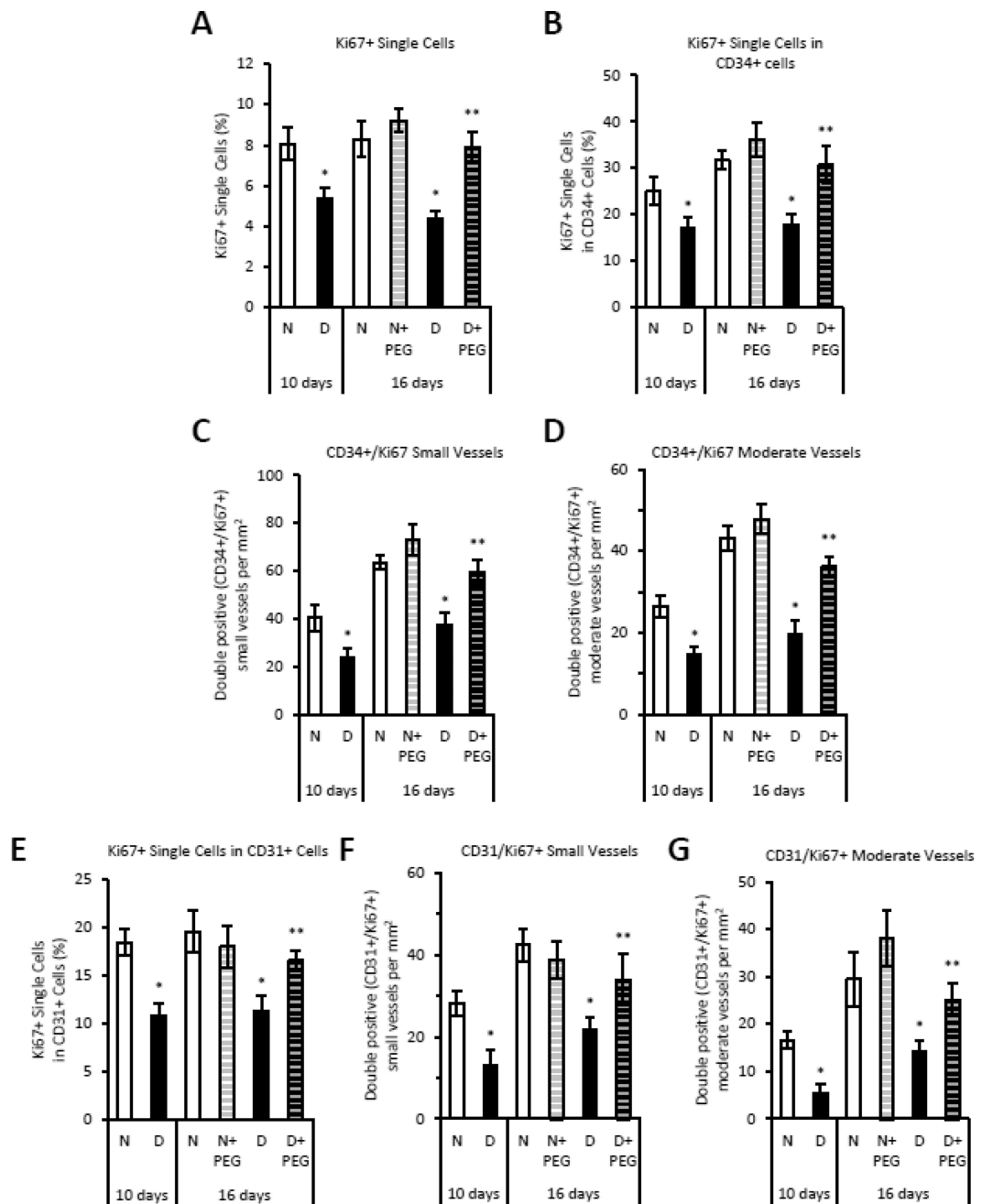


Figure 5. TNF α inhibition reverses the negative effect of diabetes on endothelial cell proliferation

Fracture calluses were examined in areas of endochondral bone formation in sections prepared from normoglycemic mice (N), diabetic mice (D) or diabetic mice treated starting 10 days post fracture with TNF-specific inhibitor pegsunercept (PEG). CD34 and Ki67 single and double immunopositive cells or blood vessels were identified by immunofluorescence. **A:** Total percent Ki67 immunopositive single cells. **B:** Percent CD34 immunopositive single cells that were also Ki67 immunopositive. **C–D:** Percent CD34 immunopositive blood vessels that were also Ki67 immunopositive. **E:** Percent CD31 immunopositive single cells that were also Ki67 immunopositive. **F–G:** Percent CD31

immunopositive blood vessels that were also Ki67 immunopositive. N=5–6 per group.
* $p < 0.05$ between normal vs diabetic groups, ** $p < 0.05$ between 16-day diabetic and
pegsunercpt-treated diabetic groups.

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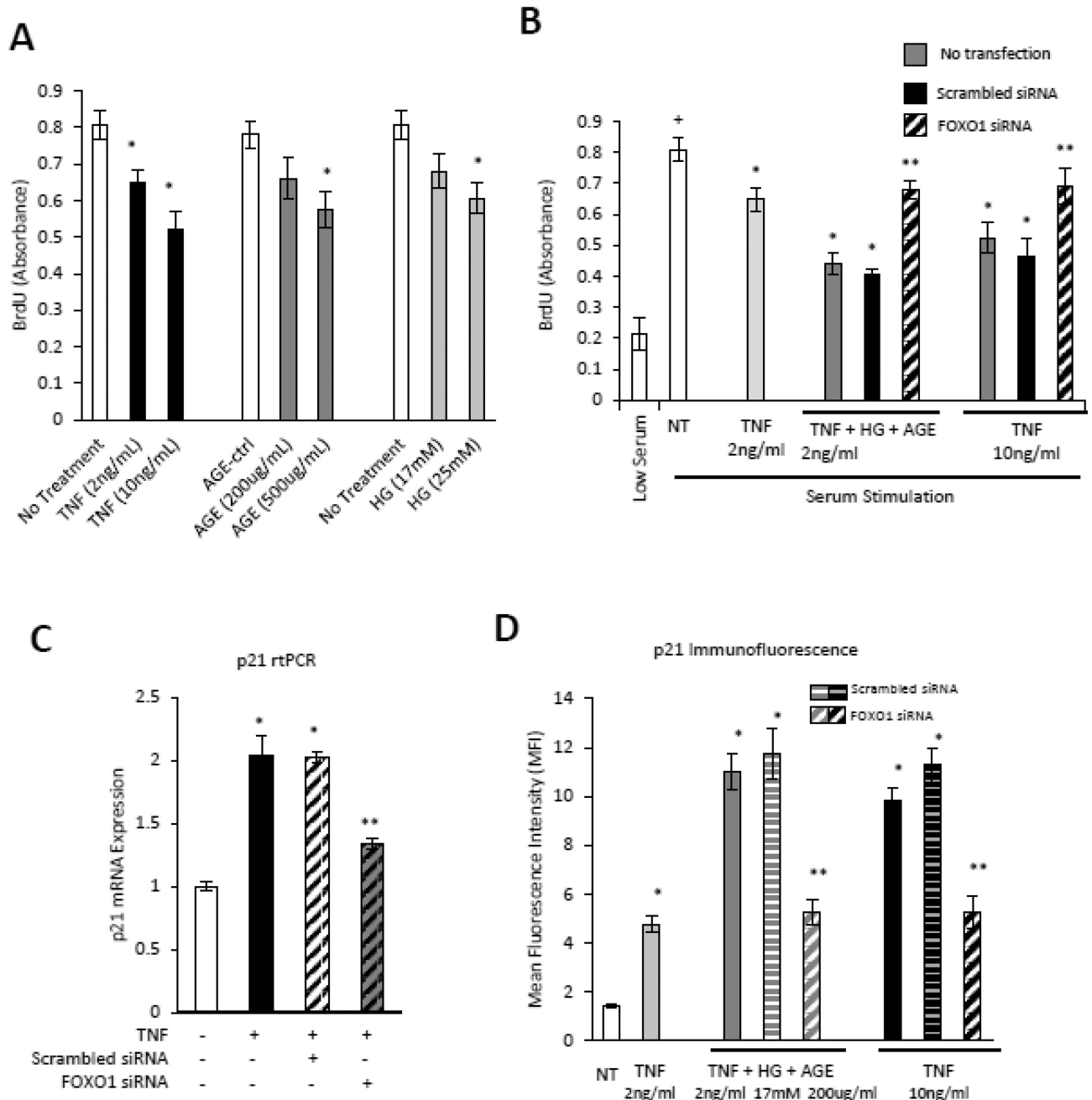


Figure 6. High glucose, TNF α and an AGE inhibit endothelial cell proliferation
 Human microvascular endothelial cells (HMVECs) were incubated with TNF α , high glucose or an AGE, CMLalbumin. **A:** DNA synthesis in serum stimulated HMVECs incubated with TNF α , AGE or high glucose was measured by BrdU assay. **B:** HMVECs were incubated with TNF α , AGE or high glucose alone or combined TNF (2ng/ml)+AGE (200ug/ml)+high glucose (17mM) with or without transfection with scrambled siRNA or FOXO1 siRNA and then stimulated with serum (10% FBS) as indicated. **C:** p21 mRNA levels were measured by PCR in TNF stimulated HMVEC following transfection with scrambled or FOXO1 siRNA. **D:** HMVEC were incubated with TNF α or TNF α +high

glucose+AGE with or without transfection with scrambled or FOXO1 siRNA. Mean fluorescence intensity (MFI) of p21 was measured by quantitative immunofluorescence. + indicates P<0.05 cells incubated in low-serum, * indicates P<0.05 compared to matched control, ** indicates P<0.05 compared to matched scrambled siRNA control.

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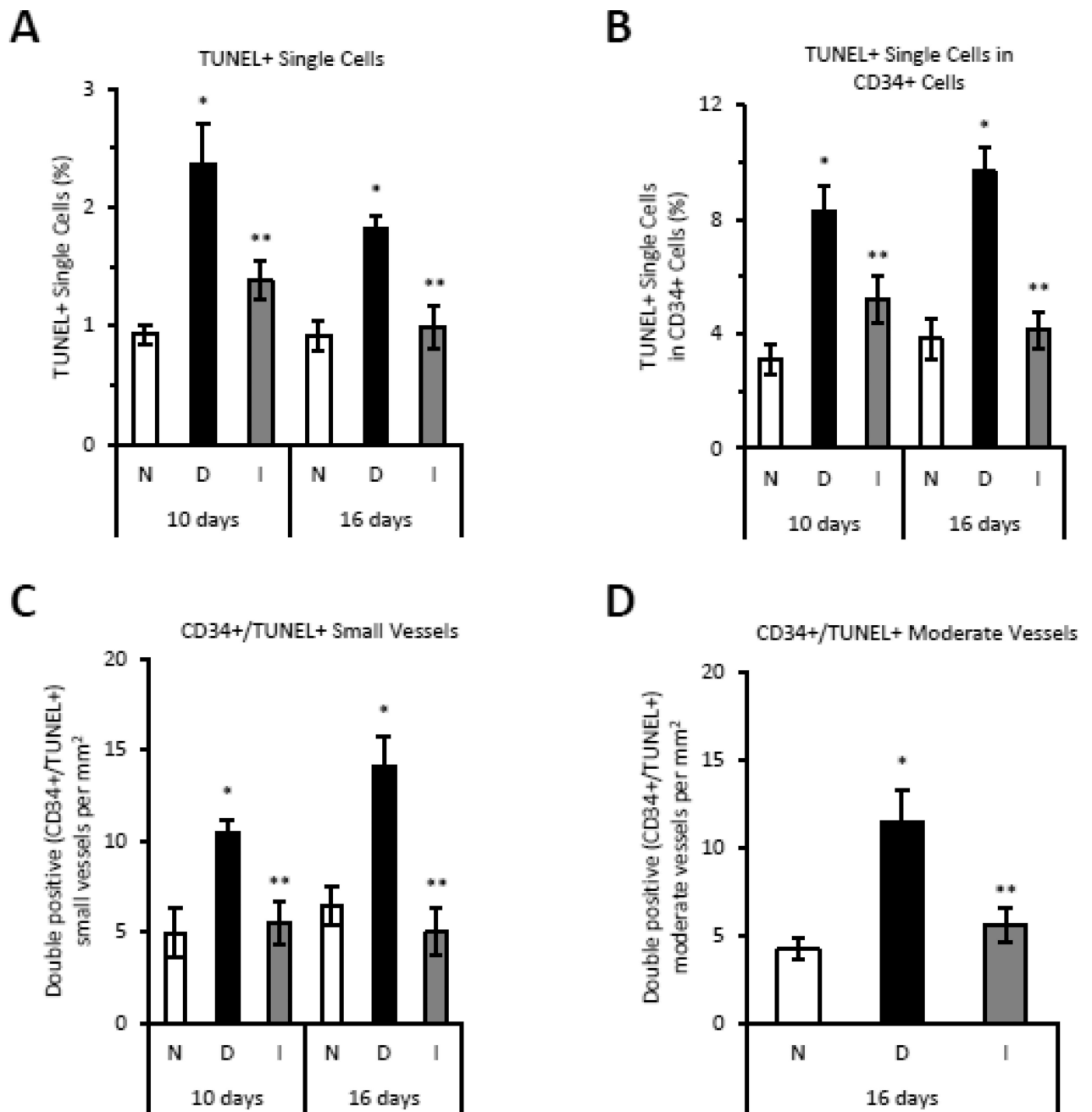


Figure 7. Diabetes enhances endothelial cell apoptosis during fracture healing that is reversed by insulin treatment

Sections from fracture calluses were examined in normoglycemic (N), diabetic (D) or diabetic plus insulin treated (I) mice in areas of bone formation. CD34 positive cells were identified by immunofluorescence and apoptosis was quantified by TUNEL assay. **A:** Total percent single cells that were TUNEL positive. **B:** The percent CD34 immunopositive single cells that were also TUNEL positive. **C–D:** The percent CD34 immunopositive blood vessels that were also TUNEL positive. N=5–6 per group. * $p < 0.05$ between normal vs

diabetic groups, ** $p < 0.05$ between 16-day diabetic and pegsunercept-treated diabetic groups.

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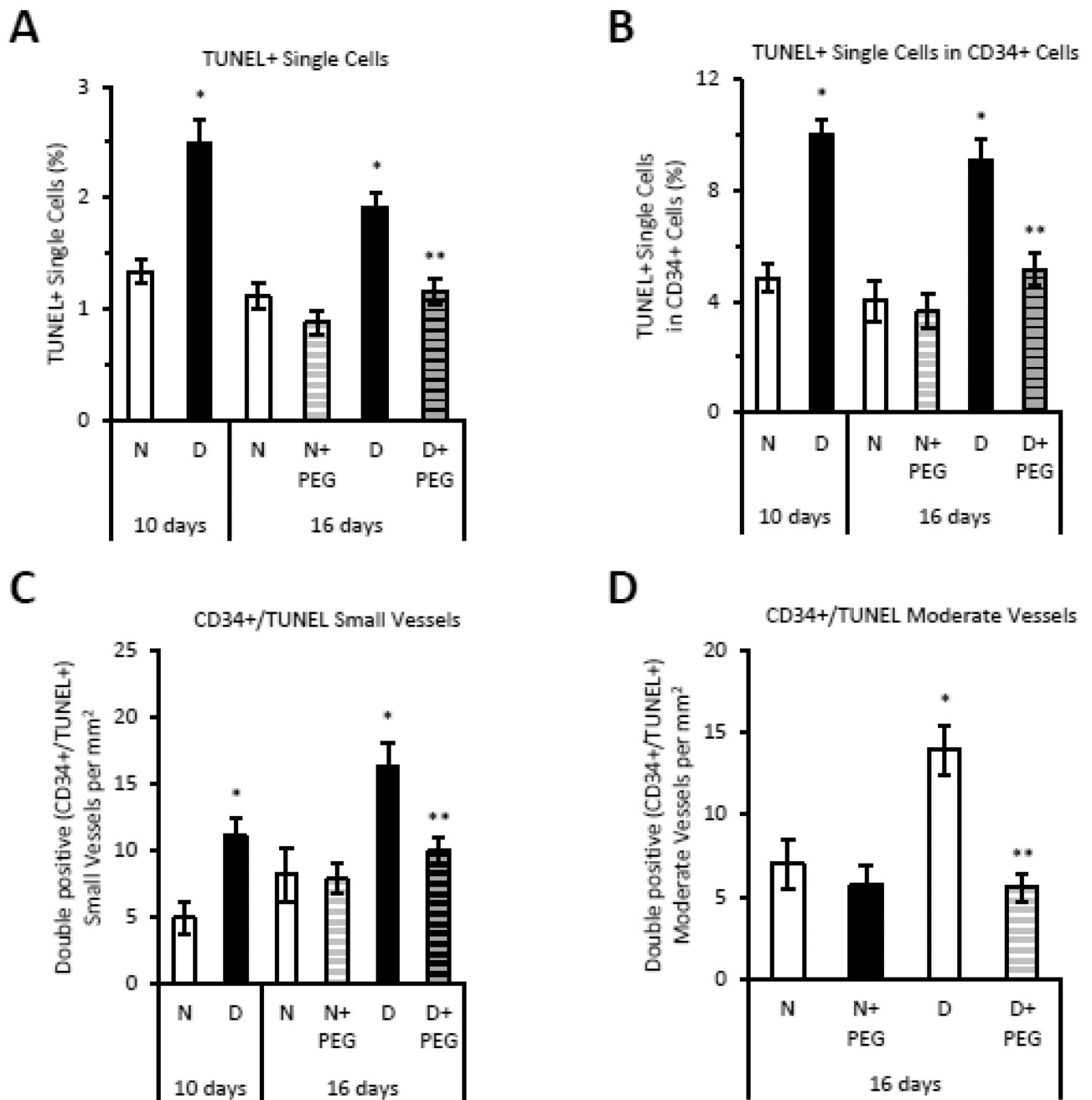


Figure 8. TNF Accounts for Diabetes-enhanced endothelial cell apoptosis during fracture healing Sections from fracture calluses in normoglycemic (N), diabetic (D) or diabetic mice treated with TNF α -specific inhibitor pegsunercept (PEG) starting 10 days post fracture. Sections were examined in areas of endochondral bone formation in the healing fracture callus. CD34 positive cells were identified by immunofluorescence and apoptosis was quantified by TUNEL assay. **A:** TUNEL positive single cells were measured. **B:** The number of TUNEL positive single cells in the CD34 positive single cell population. **C–D:** TUNEL and CD34 double positive blood vessels. N=5–6 per group. * $p < 0.05$ between normal vs diabetic groups, ** $p < 0.05$ between 16-day diabetic and pegsunercept-treated diabetic groups.

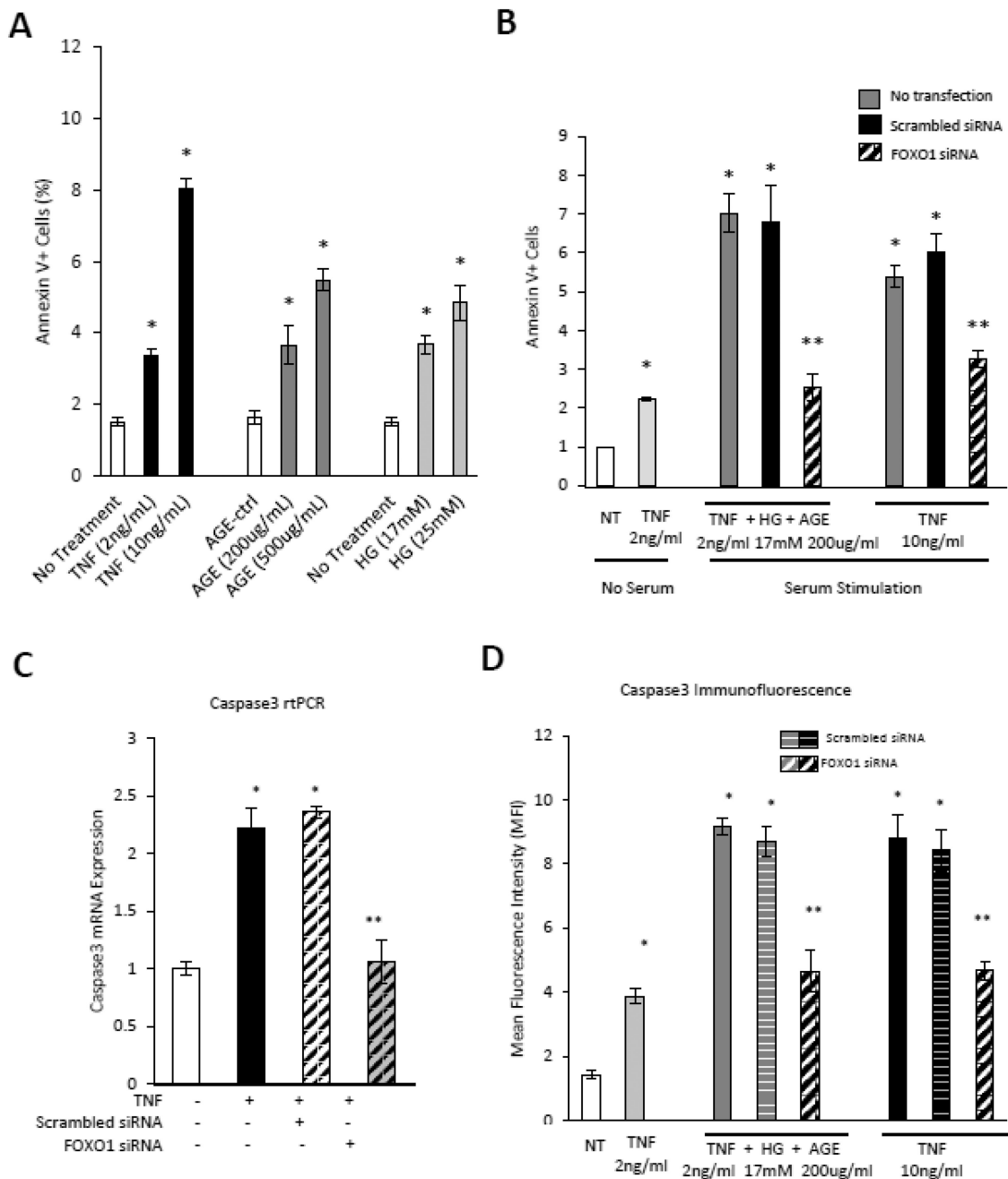


Figure 9. TNF α , high glucose and an AGE regulate the expression of p21 and caspase-3 via FOXO1

Human microvascular endothelial cells (HMVECs) were transfected with scrambled or FOXO1 siRNA and incubated with TNF α , high glucose or an AGE, CML-albumin as indicated. **A–B**: mRNA expression of p21 (A) and caspase3 (B) were evaluated by real-time PCR in HMVECs incubated with TNF α , an AGE, CML-albumin or unmodified albumin (AGE-ctrl), or high glucose. **C–D**: HMVEC were incubated with TNF, high glucose or AGE as indicated in cells without or with transfection with FOXO1 siRNA or scrambled siRNA. Protein levels of p21 and caspase3 were measured by quantitative immunofluorescence and

data expressed as mean fluorescence intensity (MFI). * $p < 0.05$ vs. unstimulated cells, ** $p < 0.05$ vs. scrambled siRNA controls.

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