

Retinopathy in a novel model of metabolic syndrome and type 2 diabetes: new insight on the inflammatory paradigm

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ABSTRACT The pathogenesis of diabetic retinopathy (DR) in metabolic syndrome (MetS) and type 2 diabetes (T2D) is not well studied, partly because an appropriate model has not been developed. Recently, we introduced a novel model of spontaneous T2D and MetS that replicates the relevant features of the human disease. In the current study, we investigated the retinal vascular changes in these animals. Experimental DR in streptozotocin (STZ)-injected rodents is described as an inflammatory disease, in which intercellular adhesion molecule 1 (ICAM-1) plays a key role. In comparison, advanced diabetes (HbA1c>10%) in the Nile grass rat (NGR) was associated with lower ICAM-1 protein expression when compared with that in normal or moderately diabetic animals. Vascular cell adhesion molecule 1 (VCAM-1) expression, however, was unaffected by the disease state. As opposed to the STZ-induced model of DR, in diabetic NGRs, most leukocytes accumulated in the retinal arteries. Consistent with the ICAM-1 reduction, leukocyte accumulation was significantly reduced in advanced disease. Similarly, leukocyte adhesions were significantly lower, with elevated plasma triglycerides (>200 mg/dl), and cholesterol (>240 mg/dl). However, these adhesions were significantly higher in animals with higher plasma insulin (>5 μ IU/ml) and leptin (>20 ng/ml), suggesting a role for these hormones in diabetic retinal leukostasis. Diabetic NGRs showed substantial retinal endothelial injury, primarily in the microvessels, including vascular tortuosity, obliterated acellular capillaries, and pericyte

ghosts. The NGR provides a convenient and realistic model for investigation of retinal changes in MetS/T2D with convincing advantages over the commonly used STZ-induced T1D.— Noda, K., Nakao, S., Zandi, S., Sun, D., Hayes, K. C., Hafezi-Moghadam, A. Retinopathy in a novel model of metabolic syndrome and type 2 diabetes: new insight on the inflammatory paradigm. *FASEB J.* 28, 2038–2046 (2014). www.fasebj.org

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CURRENTLY, THERE IS A GROWING epidemic of metabolic syndrome (MetS) and type 2 diabetes (T2D) in industrialized and developing countries. MetS is characterized by the variable coexistence of abdominal obesity, dyslipidemia, hyperinsulinemia, and hypertension (1, 2). When insulin-resistant individuals fail to maintain the hyperinsulinemia needed to keep a normal blood glucose (BG) level, fulminant disease ensues (3). MetS is associated with a markedly increased risk of the development of diabetic complications. A debilitating complication and a leading cause of vision loss is diabetic retinopathy (DR; ref. 4). Currently, there is no satisfactory model for the study of the pathogenesis of DR in MetS/T2D.

Streptozotocin (STZ)-induced hyperglycemia is commonly used in retinal research; however, the great need to reach beyond hyperglycemia has been recognized (5). Existing T2D models include the ob/ob mouse (6), the KK mouse (7), the db/db mouse (8), the Goto-Kakizaki (GK) rat (9), the Wistar fatty rat (10), the Otsuka-Long-Evans-Tokushima fatty (OLETF) rat (11), the Torii nonobese rat (12), and the morbidly obese desert sand rat (13). These models have contributed

Abbreviations: BG, blood glucose; BRB, blood–retina barrier; BW, body weight; ConA, concanavalin A; DR, diabetic retinopathy; EB, Evans blue; FITC, fluorescein isothiocyanate; HbA1c, hemoglobin A1c; ICAM-1, intercellular adhesion molecule 1; MetS, metabolic syndrome; NGR, Nile grass rat; PBS, phosphate-buffered saline; PI, propidium iodide; STZ, streptozotocin; T1D, type 1 diabetes; T2D, type 2 diabetes; TG, triglyceride; VCAM-1, vascular cell adhesion molecule 1

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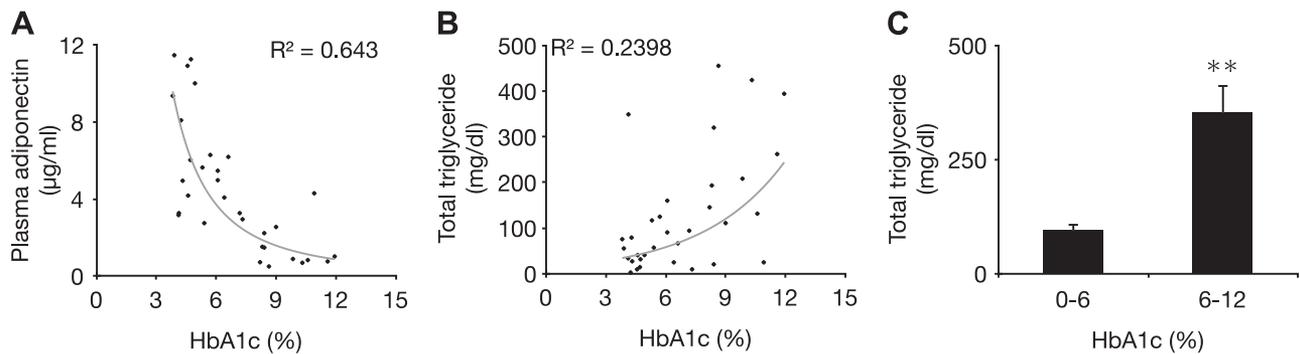


Figure 1. Metabolic characteristics of the NGR. *A*) Adiponectin is a key indicator of the metabolic state. Negative correlation between plasma adiponectin and plasma HbA1c in NGRs shows the close resemblance of this model with the human T2D and MetS. *B*) Distribution of plasma TG levels in relation to plasma HbA1c as an indicator of disease state in NGRs. *C*) Total plasma TGs in normal (<6%) and diabetic (>6% HbA1C) NGRs. ** $P < 0.01$.

Statistical analysis

All values are expressed as means \pm SEM. Student's *t* test was used for statistical analysis. Differences between the experimental groups were considered statistically significant at $P < 0.05$.

RESULTS

Metabolic levels in the examined animals

As an important hormonal correlate of the metabolic state, non-food-deprived levels of plasma adiponectin were measured in the animals that were subsequently used for retinal analysis. Plasma adiponectin correlated inversely with HbA1c (Fig. 1A). By contrast, plasma TGs correlated positively with HbA1c (Fig. 1B), in line with our original observation in a larger population (20, 26). Plasma TGs in the normal nondiabetic controls (HbA1c<6%) were within normal range and significantly lower than in the diabetic (HbA1c<6%) animals,

confirming the fidelity of this aspect of the model to the human MetS/T2D (Fig. 1C).

Lower ICAM-1 expression in advanced disease

ICAM-1 is up-regulated in STZ-induced diabetes and thus is thought to be mechanistically involved in DR pathogenesis (17). Staining for ICAM-1 in normal and diabetic NGRs revealed that the molecule was present in the retinal vessels of the animals (Fig. 2A). Since little is known about the dynamics of ICAM-1 expression in MetS/T2D, ICAM-1 in retinas of the NGRs was measured and found comparable in the normal (HbA1c <6%) and diabetic (HbA1c 6–10%) animals. In the animals with late-stage disease (HbA1c >10%), ICAM-1 was substantially reduced (Fig. 2B). Indeed beyond an HbA1c of 6%, ICAM-1 correlated negatively with HbA1c (Fig. 2C). When ranked in subgroups by HbA1c, there was no difference between the normal (HbA1c <6%) and the diabetic (HbA1c 6–10%) animals, whereas the advanced diabetic rats (HbA1c >10%) showed significantly lower ICAM-1 levels (Fig. 2D).

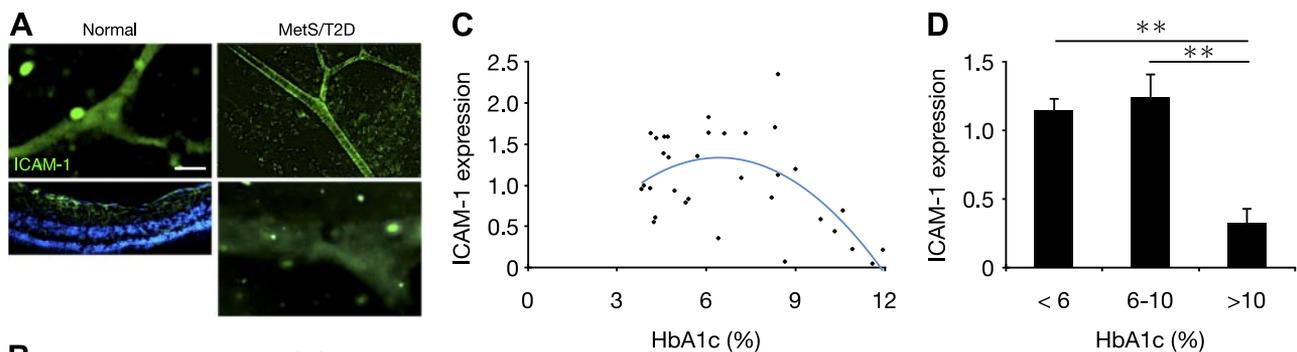


Figure 2. Dynamic of ICAM-1 expression in retinas of NGR. *A*) Immunohistochemistry for ICAM-1 in retinal vessels of normal (<6% HbA1c, left) and diabetic NGR (6–10% HbA1c, top right), and late stage disease (>10% HbA1c, bottom right micrograph) Scale bar = 50 μ m. *B*) Representative Western blot bands for ICAM-1 and β -tubulin in normal (<6%), diabetic (6–10%), and late-stage (>10% HbA1c) diabetic animals. *C*) Distribution of retinal ICAM-1 expression in relation to plasma HbA1c. *D*) Quantitative analysis of Western blots of retinal ICAM-1 in normal (<6%), diabetic (6–10%), and late-stage (>10% HbA1c) diabetic animals. Standardization was determined as the ratio of the ICAM-1 band densities through the respective β -tubulin internal control bands. ** $P < 0.01$.

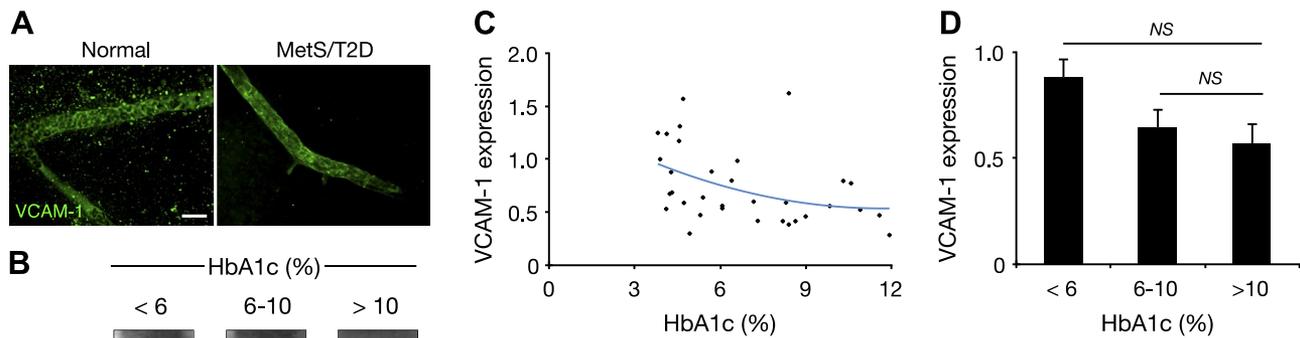


Figure 3. Unchanged VCAM-1 in retinas of NGRs. *A*) Immunohistochemistry for VCAM-1 in retinal vessels of normal (<6% HbA1c) and diabetic NGRs (6–10% HbA1c). Scale bar = 30 μ m. *B*) Representative Western blot bands for VCAM-1 and β -tubulin in normal (<6%), diabetic (6–10%), and late-stage diabetic (>10% HbA1c) animals. *C*) Distribution of retinal VCAM-1 expression in relation to plasma HbA1c. *D*) Quantitative analysis of retinal VCAM-1 in normal (<6%), diabetic (6–10%), and late-stage diabetic (>10% HbA1c) animals. NS, not significant.

Unchanged VCAM-1 with disease progression

As a comparison with another important endothelial injury marker, we examined VCAM-1 expression in the retinas of the same NGRs. Retinal vessels of the normal and MetS/T2D animals showed positive staining for VCAM-1 (Fig. 3A). When measured in Western blots, VCAM-1 was at comparable levels in the normal (HbA1c <6%), diabetic (HbA1c 6–10%), and late-stage diabetic (HbA1c >10%) animals (Fig. 3B). However, as opposed to ICAM-1, no correlation was observed between VCAM-1 and HbA1c levels (Fig. 3C). That rats with advanced diabetes (HbA1c >10%) showed no significant difference in VCAM-1 expression with normal and diabetic animals suggests that reduced endothelial density *per se* is not the cause of lower adhesion molecule expression (Fig. 3D).

Surprising inverse relation of leukocyte accumulation with disease progression

DR is an inflammatory disease, and leukocyte accumulation is considered an early mechanistic event (17). When firm leukocyte adhesion was measured in the normal and diabetic/metabolic NGRs, few leukocytes were found to adhere firmly to the retinal vessels. In the diabetic animals, firmly adhering leukocytes were found, but surprisingly, they occurred mostly in the retinal arteries, not in the veins (Fig. 4A). Furthermore, there was a negative correlation between leukocyte accumulation in arteries and the state of disease (Fig. 4B). Indeed, the number of firmly adhering leukocytes in animals with advanced disease (HbA1c >10%) was significantly lower than in the normal control (HbA1c <6%) or diabetic (HbA1c 6–10%) animals (Fig. 4C).

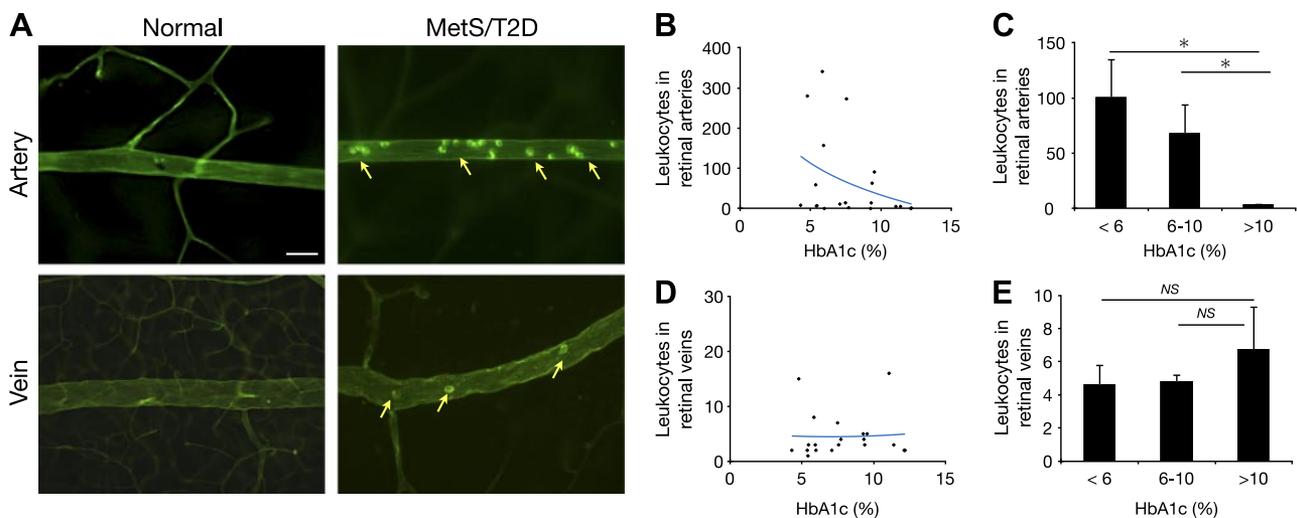


Figure 4. Retinal arterial leukostasis in diabetic animals. Characterization of retinal inflammation in normal and diabetic animals. *A*) Visualization of leukostasis in retinal vessels during diabetes. Micrographs show representative ConA staining in retinas of normal and diabetic animals. Arrows depict firmly adhering leukocytes. Scale bar = 30 μ m. *B*) Distribution of leukocytes in retinal arteries in relation to HbA1c. *C*) Quantitative analysis of arterial leukocytes in normal (<6%), diabetic (6–10%), and late-stage diabetic (>10% HbA1c) animals. *D*) Distribution of the leukocytes in retinal veins in relation to HbA1c. *E*) Quantitative analysis of the venous leukocyte accumulation in normal (<6%), diabetic (6–10%), and late-stage diabetic (>10% HbA1c) animals. * P < 0.05.

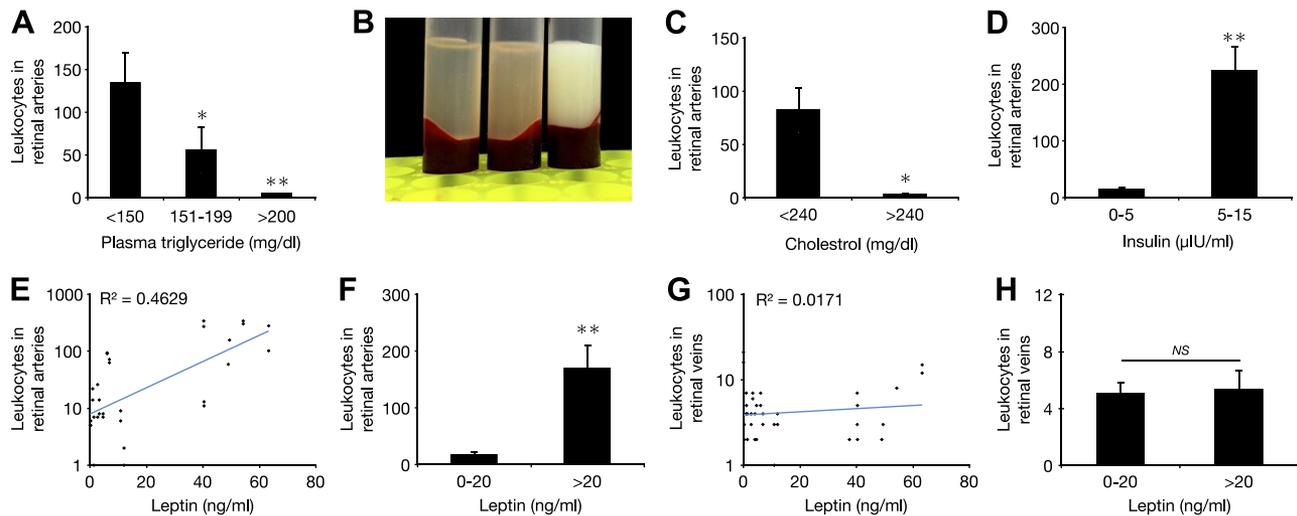


Figure 5. Relation of metabolic parameters and cellular inflammation. *A*) Retinal arterial leukocyte accumulation in relation to plasma TGs. Compared to normal (<150 mg/dl), arterial leukocyte accumulation was significantly lower in animals with marginally increased plasma TG (151–199 mg/dl) and further reduced in animals with markedly increased plasma TG (>200 mg/dl). *B*) Representative plasmas, from left to right: normal (transparent) and moderately and strongly hypertriglyceridemic animals (white opaque). *C*) Retinal leukocyte accumulation in animals with normal (<240 mg/dl) *vs.* animals with elevated plasma cholesterol levels (>240 mg/dl). *D*) Number of retinal arterial leukocytes in relation to fasting plasma insulin level. *E*) Distribution of leukocytes in retinal arteries in relation to plasma leptin. *F*) Number of retinal arterial leukocytes in relation to plasma leptin. *G*) Distribution of the leukocytes in retinal veins in relation to plasma leptin. *H*) Number of retinal venous leukocytes in relation to plasma leptin. * $P < 0.05$; ** $P < 0.01$.

In comparison, in the retinal veins, there was no correlation between leukocyte accumulation and HbA1c (Fig. 4D) (*i.e.*, between the normal, diabetic, or advanced diabetic animals; Fig. 4E).

Characterization of metabolic parameters in retinal inflammation

The role of lipids and adipokines, in DR is not well understood. We measured plasma TGs and quantified leukocyte accumulation. The number of firmly adhering leukocytes was significantly lower in the animals with slightly (151–199 mg/dl) or greatly elevated plasma TGs (>200 mg/dl) compared with that in the normal animals (Fig. 5A). Examples of low, medium, and high plasma TG levels are illustrated by the opacity and color of the plasma from freshly obtained blood samples (Fig. 5B). Similarly, hypercholesterolemia (>240 mg/dl) resulted in significantly less leukocyte adhesion compared to that in the nondiabetic rats with normal cholesterol levels (Fig. 5C). Leukocyte adhesion was significantly increased by hyperinsulinemia (>5 μ IU/ml), compared to that in the normoinsulinemic animals (<5 μ IU/ml) (Fig. 5D). Leptin values correlated positively with retinal arterial leukocyte adhesions (Fig. 5E). Leukocyte adhesions were significantly higher in the animals with higher leptin values compared to that in the normal animals (Fig. 5F). In contrast, plasma leptin did not correlate with retinal venous leukocyte adhesions (Fig. 5G, H).

Increased BRB permeability in MetS/T2D NGRs

To evaluate the BRB function in the diabetic animals, we used the EB technique (28). In the MetS/T2D

animals, we observed significantly higher leakage into the retinal tissue, compared to that in the normal animals (Fig. 6). In T1D, the BRB breakdown in diabetes is thought to be a mechanistic consequence of leukocyte adhesion (28). The current results, however, indicate that other factors, aside from leukocyte accumulation in MetS/T2D, may contribute to the BRB breakdown.

Increased endothelial damage with disease progression

Next, we investigated cellular injury in retinas of the NGRs. In the normal animals, no endothelial propidium iodide (PI) staining was observed (Fig. 7A). In contrast, the diabetic animals showed significantly more PI-positive cells that could be endothelial cells or pericytes (Fig. 7B, C). Notably, most of the PI-positive cells were observed in the microvessels, when compared with the number in the larger vessels (Fig. 7D).

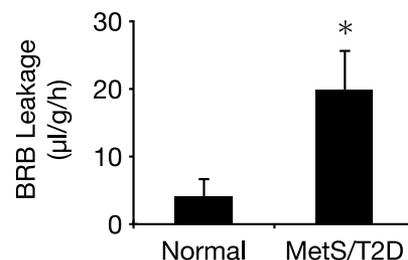


Figure 6. Retinal vascular leakage in normal and diabetic animals. BRB was evaluated in normal (<6% HbA1c) and diabetic (6–10% HbA1c) animals with the EB technique ($n=3$ /group). * $P < 0.05$.

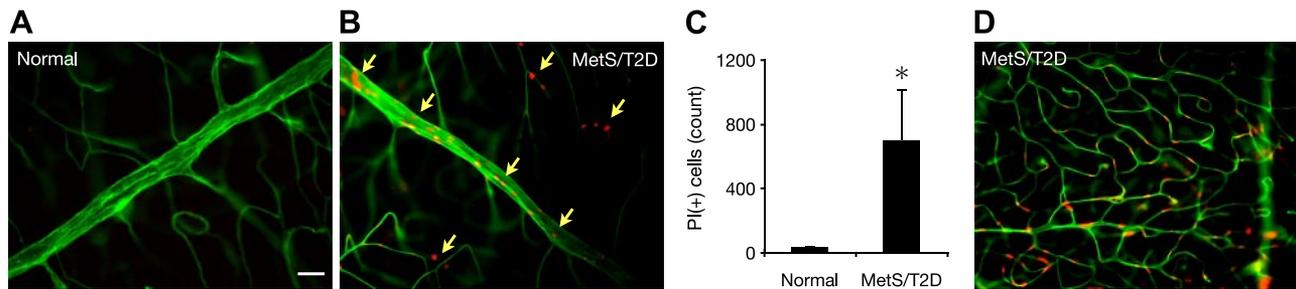


Figure 7. Cellular injury in diabetic NGRs. Retinal endothelial cell damage was detected with PI (red) staining in normal and diabetic NGRs. Vasculature was stained with FITC-ConA (green). *A*) Normal NGR lacking PI-positive staining, indicative of healthy vasculature (6-mo-old male, BW 104.6 g, BG 74 mg/dl, HbA1c 4.8%). Scale bar = 30 μ m. *B*) A representative diabetic NGR (6-mo-old female, BW 90.5 g, BG 473 mg/dl, HbA1c 8.4%) showing PI-positive staining, indicative of cell death. *C*) Quantification of PI-positive cells in the retinas of normal and diabetic NGRs ($n=4$ /group) * $P < 0.05$. *D*) PI-positive endothelial cells or pericytes in the retinal microvessels of a diabetic NGR, indicating early cellular damage.

Structural abnormalities in MetS/T2D NGRs

To further investigate microvascular changes, we performed trypsin digestion of the retinas of the normal and diabetic animals. Retinas of the nondiabetic NGRs showed patent capillaries with endothelial cells and the surrounding pericytes (Fig. 8A and ref. 29). In contrast, MetS/T2D NGR showed numerous obliterated, acellular capillaries, a classic sign of DR (Fig. 8B). In addition, MetS/T2D NGR retinas

showed vascular anomalies that fit the description of pericyte ghosts (Fig. 8C and ref. 30) and other irregularities of the microvascular wall (Fig. 8D). In retinal flat mounts of the FITC-ConA-perfused NGRs, we found widened arterial regions in the diabetic NGRs that were not found in the normal ones (Fig. 8E) or in the STZ-induced diabetic animals (data not shown). Incidentally, these vascular regions coincided with increased leukocyte accumulations (Fig. 8E). As a structural abnormality, irregular venous walls

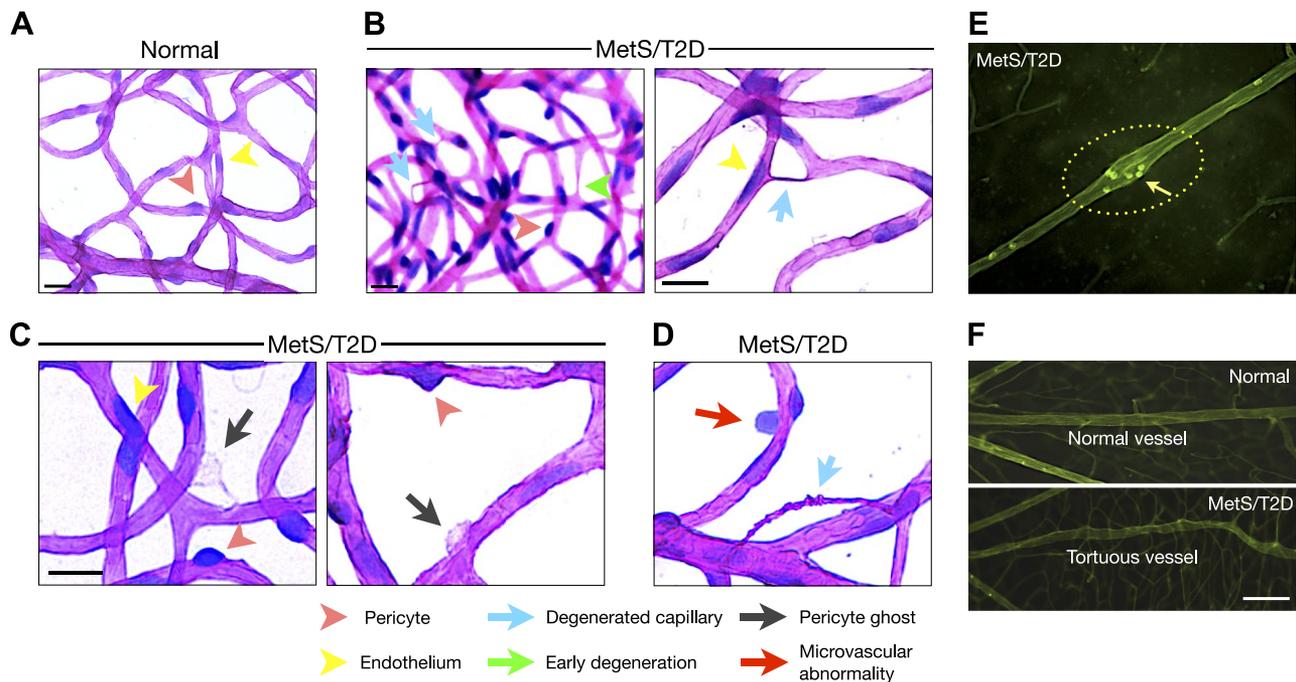


Figure 8. Vascular pathology in diabetic NGRs. Diabetes-induced degenerative vascular changes were visualized in flat mounts of trypsin-digested retinas. *A*) PAS- and hematoxylin-stained flat mounts of trypsin-digested retinas. Normal, nondiabetic animals showed patent retinal capillaries, composed of endothelial cells and surrounded by pericytes. Scale bar = 50 μ m. *B*) T2D/MeS animals showed acellular capillaries, a hallmark of DR. Green arrow, a capillary in an early stage of degeneration; turquoise arrow, obliterated capillaries that are no longer patent. Scale bar = 50 μ m. *C*, *D*) Pericyte ghosts (*C*) and microvascular wall defects (*D*) in retinas of T2D/MeS animals. Scale bar = 50 μ m. *E*) Retinal flat mounts of FITC-ConA-perfused animals, showing widened sections of retinal arterioles in diabetic animals, indicative of early structural damages. Arrow, leukocyte accumulation in the arterial abnormality. *F*) Venous irregularities in diabetic NGRs. In contrast to normal animals (top panel), diabetic NGRs showed tortuous vessels (bottom panel), which has also been reported in human DR (29). Scale bar = 100 μ m.

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