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Regulation of Microvascular Flow and Metabolism: An Overview

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Abstract

Skeletal muscle is an important site for insulin to regulate blood glucose levels. It is estimated that skeletal muscle is responsible for ~80% of insulin-mediated glucose disposal in the post-prandial period. The classical action of insulin to increase muscle glucose uptake involves insulin binding to insulin receptors on myocytes to stimulate glucose transporter 4 (GLUT 4) translocation to the cell surface membrane, enhancing glucose uptake. However, an additional role of insulin that is often under-appreciated is its action to increase muscle perfusion thereby improving insulin and glucose delivery to myocytes. Either of these responses (myocyte and/or vascular) may be impaired in insulin resistance, and both impairments are apparent in type 2 diabetes, resulting in diminished glucose disposal by muscle. The aim of this review is to report on the growing body of literature suggesting that insulin-mediated control of skeletal muscle perfusion is an important regulator of muscle glucose uptake and that impairment of microvascular insulin action has important physiological consequences early in the pathogenesis of insulin resistance. This work was discussed at the 2015 Australian Physiological Society Symposium “Physiological mechanisms controlling microvascular flow and muscle metabolism”.

Key words: Insulin action, blood flow, insulin resistance

Introduction

Skeletal muscle makes up a high proportion of body mass (40% of total body weight) due to its important mechanical role in the body. The contractile function of skeletal muscle necessitates increased metabolic activity, the requirements of which are met by alterations in blood flow to facilitate nutrient delivery and waste removal via an intricate vascular network.

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Due to its considerable mass, high metabolic rate, and capacity to store glycogen, skeletal muscle is a major site for glucose disposal and thus an important tissue for whole body glucose homeostasis.

Since the early 20th century it has been known that skeletal muscle blood flow increases dramatically during increased metabolic demand. It is becoming clear that increasing total blood flow to muscle does not necessarily improve nutrient exchange by myocytes (1, 2). These consistent findings strongly suggest that total muscle blood flow may not be tightly coupled with metabolism. Rather, the vascular system in muscle has a further level of complexity which involves modulation of blood flow at the microvascular level to control myocyte nutrient exchange. This review highlights the growing body of literature suggesting that insulin-mediated control of the microvasculature in skeletal muscle is an important physiological regulator of muscle glucose uptake, and that loss of microvascular insulin action precedes and potentially drives the development of myocyte insulin resistance.

Insulin and total muscle blood flow

Studies from the 1930s demonstrated that insulin could cause vasodilation and modify blood flow (3). However, these studies were conducted with large doses of insulin that resulted in hypoglycaemia. Thus, whether the vascular activity was due to insulin *per se*, or due to an associated hypoglycaemia-induced epinephrine release was unclear. In the 1990s Baron and colleagues performed studies in humans using the euglycaemic hyperinsulinaemic clamp technique and confirmed that insulin was indeed vasoactive and could dilate the vascular network in skeletal muscle (4, 5). They demonstrated that insulin increases total leg blood flow in a dose-dependent fashion, and that this response was paralleled by increases in muscle glucose uptake. Thus, it was Baron and colleagues who championed the notion that insulin-mediated increases in total muscle blood flow augments the metabolic actions of insulin by enhancing the delivery of glucose and insulin to the myocyte (4, 5). A number of independent investigators have produced similar results showing that insulin stimulates total muscle blood flow in both humans (6-11) and experimental animals (12-14).

At that time, however, the effect of insulin to stimulate muscle blood flow was viewed as highly controversial. Many studies that reported increases in total blood flow used supra-physiological doses of insulin or required extended exposure times (several hours) before seeing any effect on muscle blood flow (8, 15, 16). These findings raised concerns about the

physiological relevance of insulin-stimulated blood flow, particularly given that insulin levels typically remain elevated for only 90-120 min following a meal in healthy people (17, 18). Yki-Jarvinen and colleagues challenged this theory by demonstrating that insulin-stimulated glucose uptake in skeletal muscle occurred before any augmentation of total muscle blood flow (19-21). These findings raised a number of questions regarding the impact of increased blood flow on insulin-mediated muscle metabolism (19-21). This temporal discordance argument against an important vascular role of insulin is invalid due to the assumption that insulin's vascular action is restricted only to increases in total muscle blood flow. We, together with our collaborators (1, 2), hypothesise that modulation of blood flow within the microvascular network is a more physiologically relevant mode of nutrient and hormone delivery to the myocyte than total limb blood flow.

Insulin and microvascular blood flow

As highlighted above, most studies have focused on insulin's action on total muscle blood flow. However, the effect of insulin to alter the distribution of microvascular blood flow within skeletal muscle may be more physiologically relevant. Total blood flow in skeletal muscle is controlled by constriction/dilation of the 1st-3rd order arterioles. The control of capillary (microvascular) perfusion in skeletal muscle is controlled by the 3rd-5th order arterioles (see Figure 1). At any given point in time, not all capillaries are perfused (22, 23) and this is due vasomotion (24, 25). Vasomotion is a process where blood vessels contract and dilate at regular time intervals to modulate blood flow through different capillary modules to maintain efficient nutrient supply. Therefore, taking a muscle biopsy to count capillaries is not sufficient to understand the extent of microvascular blood flow *in vivo*.

In order to determine the microvascular actions of insulin in skeletal muscle, we developed two techniques for measuring blood flow at this level of the vascular tree *in vivo*. The first technique we developed relies on metabolism of exogenously infused 1-methylxanthine (1-MX) to 1-methylurate (1-MU) by microvascular xanthine oxidase (13, 14, 26-30). 1-MX and 1-MU can be quantified in plasma using high performance liquid chromatography (31). Xanthine oxidase is located primarily on capillary endothelial cells, and not large arteries and veins, or in myocytes (32, 33). The disappearance of 1-MX across the hindleg (A-V difference x femoral artery blood flow) can therefore be used as a biochemical marker for the extent of microvascular blood flow in muscle.

The second technique was an adaptation of an ultrasound imaging technique (contrast-enhanced ultrasound, CEU) to skeletal muscle (14, 34-41). This technique involves infusion of albumin or phospholipid microbubbles (perfluorocarbon gas filled) into the systemic circulation. The microbubbles are similar in size and rheology to red blood cells, and importantly they remain intravascular (42) making them an excellent perfusion tracer. The microbubbles oscillate in size when exposed to ultrasound, and can be destroyed with high energy pulses of ultrasound. The oscillation and destruction of microbubbles results in the generation of a signal that can be measured as acoustic intensity. Following ultrasonic destruction of microbubbles, the rate of reappearance of microbubbles is a reflection of blood velocity, while the microvascular blood volume can be measured by the plateau of tissue opacification. The contribution of fast filling arteries, arterioles, veins and venules can be background subtracted due to the high flow velocity and thus the residual signal reflects only the microcirculation. Replenishment curves are fitted to $y = A(1 - e^{-\beta t})$, where y is acoustic intensity, t is the time from the destructive ultrasound pulse, A is plateau video intensity (microvascular blood volume), and β is the rate constant, which provides a measure of flow velocity in the microvasculature.

Using both 1-MX and CEU techniques, we were the first to demonstrate that insulin increases microvascular blood flow in skeletal muscle in a similar fashion to muscle contraction (36, 43) (see Figure 1). Importantly, this microvascular action of insulin contributes to ~50% of insulin-stimulated glucose disposal, a finding observed in both humans and experimental animals (13, 14, 35-40). In addition, the microvascular response to insulin can occur independent of its effects on total muscle blood flow, and that even low physiological doses of insulin produced marked increases in microvascular blood flow (14, 30, 36, 39), (44). Interestingly, this insulin-stimulated increase in microvascular blood flow occurs early (by 7 min) in rats and precedes stimulation of the insulin signalling cascade in the myocyte or any significant increase in muscle glucose uptake (39).

Given that insulin is released into the circulation in response to a meal, it is plausible that the vascular actions of insulin occur first to facilitate increased delivery of glucose and insulin itself to the myocyte for glucose disposal. Indeed, we have shown that insulin-mediated microvascular responses occur whether insulin is infused intravenously (euglycaemic hyperinsulinaemic clamp) (35, 45) or secreted from the pancreas following the ingestion of a

mixed meal (37). When taken together, these data highlight the sensitive and rapid nature of insulin's microvascular actions in skeletal muscle and emphasise the integral role that the microvasculature has in normal insulin-mediated muscle glucose uptake.

Mechanism of insulin-mediated microvascular recruitment

Local infusion of insulin into the forearm of healthy subjects, at a dose that does not have any significant spill-over to the systemic circulation, can stimulate microvascular blood flow and is paralleled by increased muscle glucose uptake (41). In contrast, central (intracerebroventricular) administration of insulin in healthy rats does not produce any metabolic or haemodynamic effects in skeletal muscle (46). Together, these studies suggest that insulin's effects on microvascular blood flow in muscle are mediated by local, rather than central actions of insulin.

Nitric oxide (NO) has been strongly implicated in insulin-mediated vasodilation (47, 48). Quon and colleagues have elucidated the insulin signalling cascade in cultured vascular endothelial cells which involves the activation of the insulin receptor/IRS-1/PI3K/Akt/eNOS pathway leading to NO production (49-51). The resulting NO then acts to relax adjacent vascular smooth muscle cells thus causing vasodilation. Insulin's ability to stimulate microvascular blood flow in muscle is inhibited by systemic infusion of the nitric oxide synthase inhibitor, N ω -nitro-L-arginine-methyl ester (L-NAME) (39, 48). Importantly, insulin-mediated microvascular recruitment can also be blocked when L-NAME is infused locally, implicating local NOS activation in insulin's vascular actions (52).

Endothelin-1 (ET-1) has also been implicated in the regulation of insulin-mediated microvascular blood flow. In addition to NO, insulin can also stimulate ET-1 release via a MAPK-dependent pathway (53). Systemic infusion of ET-1 into rats can block insulin-mediated microvascular blood flow (54). The vasodilator effects of insulin on large artery dilation are augmented by ET-1 receptor blockade (55, 56). Thus, insulin causes generation of both NO and ET-1, and it is the balance between these two vasoactive agents that contributes to the regulation of microvascular blood flow in skeletal muscle (55, 57).

Epoxyeicosatrienoic acids (EETs) are signalling molecules formed from arachidonic acid. New evidence is emerging to suggest that EETs are involved in insulin-mediated microvascular recruitment in skeletal muscle, particularly when insulin-mediated NO dilation is blunted (58). There is strong evidence that these compounds are one of the endothelial-

derived hyperpolarising factors causing vasodilation of skeletal muscle microvasculature (58). A role for EET production in insulin sensitisation is suggested from a study from Xu and colleagues (59) showing that CYP2J3 gene delivery *in vivo* increased EET generation, reduced blood pressure, and reversed insulin resistance, albeit by an unknown mechanism.

Impaired microvascular blood flow and insulin resistance

Insulin resistance is defined as a reduced biological response to insulin (60, 61). Given the evidence detailed above regarding insulin's vascular actions, it is logical that the development of insulin resistance also includes a haemodynamic mechanism.

Laakso and colleagues have demonstrated that obese, insulin resistant people display reduced total muscle insulin-mediated vasodilation (4). Others have reported a relationship between insulin resistance and endothelial dysfunction (62-64). While these studies have restricted measurements to large blood vessel responses to insulin, microvascular responses to insulin also appear to be reduced during obesity and insulin resistance. De Jongh and colleagues assessed microvascular function in skin of healthy lean and obese individuals in response to insulin (65). The obese, insulin resistant cohort showed reduced insulin-mediated microvascular blood flow in skin and this correlated with whole body insulin resistance (65).

Vasoconstrictors such as α -methylserotonin (26) and ET-1 (54), inhibit both insulin-stimulated microvascular blood flow and glucose uptake *in vivo*. The loss of vascular insulin action is also apparent during acute infusion of factors known to be elevated in various insulin resistant states, such as TNF α (29) and elevated free fatty acids (elevated by infusion of Intralipid and heparin) (66). In addition, chronic animal models of insulin resistance, including the high fat-fed (27), Zucker obese (28), and Zucker Diabetic fatty (34) rats display reduced microvascular and metabolic responses to insulin. Activity restricted insulin resistant primates also display impaired muscle microvascular responses to marked insulin secretion elicited by an intravenous glucose tolerance test (67). Insulin infusion in humans (35) or the ingestion of a mixed meal (37) similarly act to increase microvascular blood flow and these microvascular responses are blunted in obese insulin resistant subjects (35, 37).

Together, these studies highlight the important link between microvascular and metabolic actions of insulin in muscle, and implicate the loss of microvascular insulin sensitivity in the development and progression of myocyte insulin resistance.

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Microvascular-induced muscle insulin resistance

The mechanisms of skeletal muscle insulin resistance are multifactorial, however most investigators have focused on myocyte insulin action. It is important to note that most animal models used to investigate insulin resistance develop both myocyte and microvascular insulin resistance. However, we and others have evidence demonstrating that the microvasculature can become insulin resistant prior to changes in myocyte insulin sensitivity suggesting the vasculature is an early contributor to development of insulin resistance (68-71).

We have recently characterised two dietary animal models that develop microvascular insulin resistance (68, 69). The first of these models is the moderately raised dietary fat model, in which dietary fat in rats is increased from 5% to 9% wt./wt. (68), rather than the more common 5- to 7-fold increase employed in many studies (27, 72, 73). The second model we have characterised is the high salt-fed rat in which dietary NaCl is increased from 0.3% to 8.0% wt./wt (69). After a four-week dietary intervention, both of these animal models develop whole body, skeletal muscle and microvascular insulin resistance *in vivo* when compared to control rats. Insulin-mediated myocyte glucose uptake is normal in both of these animal models when assessed using the constant-flow pump-perfused hindleg technique. In this *ex vivo* preparation the vasculature is fully intact, but does not dilate in response to insulin, enabling insulin and glucose delivery to the myocyte in the absence of insulin-mediated changes in blood flow (68). The reduction in insulin-stimulated muscle glucose uptake *in vivo* therefore is driven by microvascular, rather than myocyte insulin resistance in these two animal models. These data position the loss of normal microvascular function as an early event in the development of muscle insulin resistance.

The endothelial IRS2 knockout mouse displays microvascular insulin resistance and impaired insulin-mediated muscle glucose uptake *in vivo*, and this is associated with reduced activation of eNOS in the vascular endothelium (71). When muscles from these animals are incubated with insulin *in vitro* (where delivery to the myocyte occurs by diffusion and not via the vasculature), myocyte glucose uptake is comparable to control animals. These data implicate reduced eNOS activation by insulin as a contributor to reduced insulin-mediated microvascular recruitment and muscle glucose uptake *in vivo*.

Recent evidence suggests an important role of muscle capillary density in insulin action. Muscle-specific vascular endothelial growth factor (VEGF) knockout animals have reduced

skeletal muscle capillary density and display whole body and muscle insulin resistance *in vivo* (70). However, when muscle samples are isolated and incubated to assess myocyte insulin action, their degree of myocyte insulin sensitivity was similar to control animals. Similarly, humans with reduced skeletal muscle capillary density are also muscle insulin resistant (74). When capillary density in muscle is enhanced with prazosin, muscle glucose uptake improves without any measurable effects on muscle insulin signalling, suggesting enhanced vascular delivery of insulin and glucose to the myocyte (75).

Thus, changes in diet (e.g. elevated dietary fat or sodium), impaired eNOS activation, or even reduced muscle capillary density can impair insulin-mediated microvascular function, even in the presence of normal myocyte insulin sensitivity.

Transport of insulin from the vasculature into the interstitial space

Another potential rate-limiting step for insulin's metabolic actions in skeletal muscle is the movement of insulin from the vasculature to the interstitial space. It has been consistently shown that the concentration of insulin in the interstitial space in skeletal muscle is ~50% lower than the concentration in plasma (76-78). Given that the time-course for insulin-mediated glucose uptake in skeletal muscle is delayed in insulin resistance and type 2 diabetes, there may be a delay in the transit of insulin from the vasculature to the interstitium in these pathologies (79).

In cell culture experiments, insulin transport into the vascular endothelium is insulin receptor mediated (80, 81). In addition, trans-endothelial transport of insulin is dependent on PI3-K, MAPK and cSrc-family tyrosine kinase signalling pathways and activation of eNOS (82). Importantly, inflammatory cytokines such as TNF α and IL-6, which are elevated during states of insulin resistance, impair insulin uptake into the endothelium (83). Thus, these findings suggest that impaired or delayed insulin delivery to the interstitial space in contact with myocytes is a potential mechanism in the development of insulin resistance. However, whether trans-endothelial transport of insulin is a rate limiting step for glucose uptake in insulin resistant muscle still remains to be confirmed *in vivo* (84).

Conclusions

A large body of work over the past two decades has demonstrated that the microvascular effects of insulin are an important component of its capacity to increase glucose disposal in skeletal muscle. Even in the presence of normal myocyte insulin sensitivity, any disruption of

this microvascular action of insulin significantly reduces glucose disposal by skeletal muscle. Thus, microvascular-derived insulin resistance is likely to be one of the first pathogenic changes in the development of insulin resistance, and is therefore an important target for early detection to prevent progression to type 2 diabetes and to inform patient care. An important question that remains is whether correcting this microvascular defect (i.e. restoring insulin-mediated microvascular responses) can effectively treat insulin resistance and type 2 diabetes. This is an area we and others are actively investigating.

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FIGURE LEGEND

Figure 1: Under basal conditions $\sim 1/3^{\text{rd}}$ of capillaries are perfused in skeletal muscle. During contraction arterioles (1° through to 5°) dilate and capillaries fully recruit in an intensity-dependent fashion. In the presence of insulin, approximately $2/3^{\text{rd}}$ of capillaries are perfused, changes in bulk flow may also occur during hyperinsulinemia. During states of insulin resistance, insulin-mediated capillary perfusion of skeletal muscle is markedly impaired.

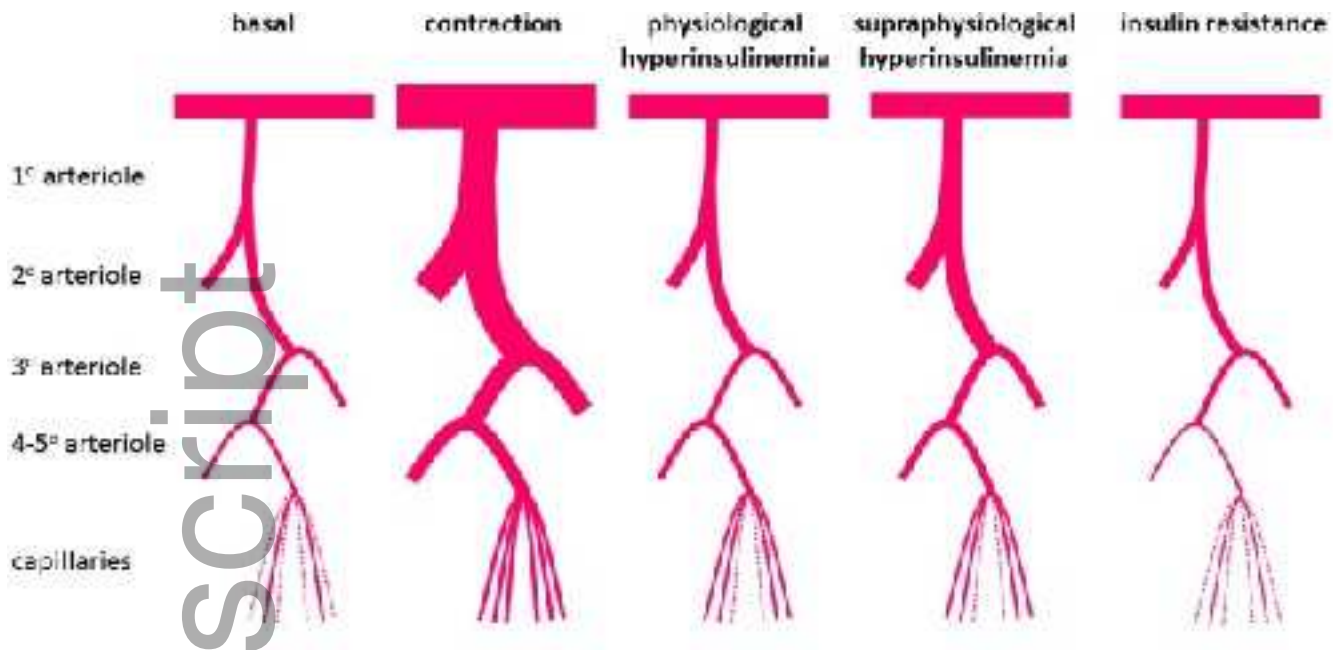


Figure 1:

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