

Impact of insulin resistance on vasculature and kidney

Ferruh Artunc*, Erwin Schleicher*, Cora Weigert, Andreas Fritsche, Norbert Stefan, Hans-Ulrich Häring.

¹ Department of Internal Medicine IV, Division of Endocrinology, Diabetology, Vascular Disease, Nephrology and Clinical Chemistry, University Hospital Tübingen, Germany

² Institute of Diabetes Research and Metabolic Diseases (IDM) of the Helmholtz Center Munich at the University of Tübingen, Tübingen, Germany

³ German Center for Diabetes Research (DZD), Tübingen, Germany

*shared first authorship

Address for correspondence:

Hans-Ulrich Häring, MD

Department of Internal Medicine

Division of Endocrinology, Diabetology, Angiology, Nephrology and Clinical Chemistry

University Hospital Tübingen

Otfried-Mueller-Str.10

72076 Tübingen, Germany

E-mail: hans-ulrich.haering@med.uni-tuebingen.de

Tel. +49-7071-29 83670

Fax +49-7071-29

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Key points

- Besides the classical insulin target tissues insulin acts on most human organs/cells including arterial vasculature and kidney
- In insulin-resistant states like obesity or type 2 diabetes not only the classical insulin effects are impaired but also the effects on vasculature and kidney
- Insulin stimulates its own delivery to target cells by PI3 kinase-mediated and NO-stimulated feed-forward actions on the vasculature involving increased capillary recruitment and endothelial transcytosis effects which are all impaired in insulin resistance
- Insulin resistance affects kidney function in many aspects such as alterations in hemodynamics, podocyte and tubular function
- Insulin's actions on renal sodium handling is preserved in insulin-resistance contributing to sodium retention and arterial hypertension
- Renal and vascular insulin-resistance can be improved by an integrated approach including life-style and/or pharmacological agents

Abstract

Insulin resistance is a systemic disorder affecting many organs and insulin-regulated pathways. It is characterized by a reduced insulin action in spite of increased insulin concentrations (hyperinsulinemia). The vasculature and the kidney are non-classical insulin target organs with distinct functions, partly differing from the insulin effects on classical target organs. Insulin causes vasodilation by enhancing endothelial NO production via activation of the PI3 kinase pathway. In insulin resistant states this pathway is impaired and the MAPK/ERK pathway, with increased endothelin-1 production, overrides the NO action, resulting in vasoconstriction. While the insulin action on perivascular fat tissue and the subsequent effects on the vascular wall are not fully understood, the hepatokine fetuin-A released by the fatty liver, may provoke pro-inflammatory effects of the perivascular fat. The strong association of salt-sensitive arterial hypertension with insulin resistance indicates an involvement of the kidney in the insulin resistance syndrome. The functional relevance of the insulin receptor bearing renal tubular cells and podocytes has been studied in conditional cell-specific insulin receptor knockout models in mice revealing decisive roles of insulin signaling in podocyte viability and tubular function. Renal sodium transport is preserved in insulin resistance and contributes to salt-sensitivity of blood pressure in hyperinsulinemia. Therapeutically, renal and vascular insulin resistance can be improved by an integrated holistic approach aiming at restoring overall insulin sensitivity and improving insulin signaling.

Introduction

The development of valid determination of plasma insulin concentrations in the 1960s revealed disproportionate hyperinsulinemia in apparently healthy obese individuals, thereafter termed insulin resistance. With the discovery of the insulin receptor (IR) in the early 1970s¹ the mechanisms of reduced insulin action could be studied on a molecular basis in both humans and laboratory animals. Insulin resistance has been shown to occur in the classical insulin-responsive organs liver, skeletal muscle and white adipose tissue ²⁻⁴, particularly in association with obesity and the metabolic syndrome ⁵. This condition is accompanied by impaired glucose tolerance and dyslipidemia with high triglyceride plasma concentrations. Insulin effects other than those affecting metabolism have been described in non-classical insulin-responsive organs such as pancreatic β -cell survival, endothelium-dependent vasodilation in the vasculature and renal sodium transport ⁶. Unexpectedly, insulin has been shown to influence brain function to regulate glucose metabolism and food-seeking behavior as reviewed recently ⁷. In the present review we describe insulin's action on the vasculature and the kidney, including molecular mechanism of insulin signal transduction and its impairment in the insulin resistant states, of the otherwise healthy obese individual with normal kidney function. The development of insulin resistance in chronic kidney disease and uremia is covered elsewhere ⁸.

Dysregulation of insulin signaling in peripheral tissues

Insulin is the central hormone in the control of glucose and lipid metabolism. It mediates its biological effects via binding to IR consisting of two extracellular insulin binding α -subunits and two intracellular tyrosine kinase β subunits. Alternate splicing of the single insulin receptor gene results in the two IR isoforms A and B ^{9,10}. Insulin may also bind to the highly homologous IGF-1 receptor or insulin/IGF-1 receptor heterodimers albeit with reduced affinity ¹¹. The insulin-mediated receptor activation leads to activation and autophosphorylation of the receptor tyrosine kinase ¹² and initiates different cascades of phosphorylation events. The two major arms of the activated insulin receptor signal

transduction are the insulin receptor substrate (IRS) – PI3-kinase (PI3K) – Akt/PKB pathway and the Grb2 – SOS – Ras – MAPK pathway (Figure 1A). IRS isoforms are intracellular adaptor proteins that are recruited to and phosphorylated on tyrosine residues by the activated insulin receptor initiating the recruitment and activation of PI3K ^{13,14}. Activated PI3K phosphorylates phosphatidylinositol (4,5)-bisphosphate (PIP2) to phosphatidylinositol (3,4,5)-triphosphate (PIP3), which as a membrane-anchored lipid second messenger activates 3-phosphoinositide-dependent protein kinase 1 (PDK1) and recruits Akt/PKB. The serine/threonine kinase Akt/PKB is a central node in regulating the biological effects of insulin, since it phosphorylates and activates essential downstream kinases (glycogen synthase kinase 3 (GSK3), mTORC1, ribosomal protein S6 kinase), transcriptional regulators such as Forkhead box O family members (FoxO), sterol regulatory element binding protein (SREBP), peroxisome proliferator-activated receptor γ coactivator 1 (PGC1 α), and the GTPase-activating protein Akt substrate 160 KDa (AS160) ¹⁵. Subsequently, a network of metabolic regulators is controlled on the level of enzymatic activity or gene expression leading to enhanced glucose uptake, glucose storage, lipid synthesis, protein synthesis and suppression of gluconeogenesis and lipolysis. PDK1 also activates atypical protein kinases C PKC ζ and PKC λ/ι contributing to insulin-stimulated glucose uptake and lipid synthesis ¹⁶. The insulin-induced Grb2-SOS-Ras pathway leads to activation of MAPK ERK1/2 and is a major control mechanism of cell proliferation and differentiation. Other non-metabolic effects of insulin mediated via the PI3K/Akt/PKB pathway include inhibition of apoptosis and promotion of cell survival ¹⁷.

Insulin signal transduction must be tightly controlled to avoid severe metabolic and proliferative perturbations. The negative regulators are often activated by insulin as feedback mechanism to inhibit the signaling pathway on the critical nodes insulin receptor/IRS or Akt/PKB. Dysregulation of these regulators by chronic hyperactivation contributes to the development of insulin resistance (figure 1B). Negative regulators are phosphotyrosine and phosphoserine/threonine protein phosphatases (PTP1B, PP2A,B,C) ^{18,19}, lipid phosphatases controlling PIP3 levels (PTEN, SHIP) ^{20,21}, and adaptor proteins of the

insulin receptor and IRS (Grb, SOCS) ^{22,23}. Another well-studied inhibitory mechanism of the insulin signaling pathway is the serine/threonine phosphorylation of the insulin receptor and IRS via the insulin-mediated activation of serine/threonine kinases, predominantly by c-Jun amino-terminal kinase (JNK), I κ B kinase (IKK), PKC, S6K1, and ERK (Fig. 1B) ²⁴⁻²⁸.

In the insulin resistant state, the cellular response to insulin is reduced and the activation of the insulin signaling pathway requires higher concentrations of insulin. Important contributors to insulin resistance are hyperinsulinemia, hyperglycemia, inflammation, excess of lipids, mitochondrial dysfunction, and ER stress ²⁹. Serine/threonine kinases are activated by the elevated plasma levels of insulin, glucose or free fatty acids, by lipids or intermediates of lipid metabolism (ceramides, diacylglycerol), by increased concentrations of cytokines, ER stress and increased levels of reactive oxygen species. Hyperactivation of protein phosphatases ³⁰, increased expression of adaptor proteins, and enhanced O-N-acetylglucosamine-modification of insulin signaling molecules ³¹ are also related to the development of insulin resistance. As a consequence, the modified proteins show an impaired interaction with their signaling partners and altered affinities, or enhanced protein degradation.

Regulation of the insulin signaling cascade and its dysregulation leading to insulin resistance are well-described in the classical insulin target tissues skeletal muscle, liver and adipose tissue, which play an important role in the regulation of glucose and lipid metabolism. However, the insulin receptor is ubiquitously expressed and functional insulin signaling is found in other peripheral, non-classical insulin target tissues, including the vasculature ³² and the kidney ³³. Binding of insulin to its receptor is found in the cells of the glomerulum, in podocytes ³⁴, in mesangial cells ³⁵, endothelial and epithelial cells ³⁶, and the tubulus ³⁷. This serves diverse functions ranging from glucose uptake as energy fuel to regulation of glomerular function, gluconeogenesis and tubular transport. In this review, we describe the metabolic and non-metabolic effects of the insulin signaling cascade in the vasculature and the kidney, as well as causes and consequences of insulin resistance in these tissues.

Organ cross-talk in the insulin-resistant state

Insulin resistance is a hallmark of obesity, metabolic syndrome and type 2 diabetes mellitus. It indicates a condition in which target cells show an attenuated response to insulin which is compensated by increased insulin secretion. Insulin resistance affects many organs and cellular insulin-regulated pathways, however, the extent of insulin resistance can vary considerably. Insulin resistance affects also organ crosstalk by altered metabolic or hormonal signals promoting organ dysfunction and disease (Fig. 2). In the kidney, obesity and insulin-resistance is an important risk factor for a decline of glomerular filtration rate (GFR) as well as the onset and progression of chronic kidney disease (CKD) ³⁸⁻⁴². Particularly visceral obesity, e.g. as estimated by increased waist circumference, was found to be a stronger predictor of end-stage renal disease than the elevated body mass index (BMI) ⁴³. Increased visceral obesity may cause CKD by promoting metabolic diseases. However, studies indicate that even in the absence of the well-known risk factors hypertension or diabetes, obesity per se may be harmful to the kidney by causing hyperfiltration ⁴⁴⁻⁴⁶. In this respect it was shown that insulin resistance, which often accompanies visceral obesity, is a strong marker of incident CKD, independent of other risk factors, including age and fasting plasma glucose ⁴⁷. Thus, insulin resistance itself and/or factors promoting insulin resistance may early on play a role in the development of CKD ⁴⁸.

But what are the major organs and pathophysiological mechanisms that are involved in both, the development of insulin resistance and impaired kidney function? Defects in skeletal muscle, white adipose tissue and liver resulting in impairment of insulin signaling have been shown to induce hyperglycemia, predominantly by impairing glucose disposal and increasing hepatic glucose production ⁴⁹. Recent evidence suggests that impaired brain insulin signaling ⁷ and an altered gut microbiome ⁵⁰ are involved in this process. Because the onset and progression of CKD may also be independent of glycemia, other mechanisms like humoral signals of metabolic tissues may be relevant for the early stages of impaired renal function. In this respect the expansion of visceral adipose tissue, which is being infiltrated by immune cells, is involved in this process ⁵¹. Pro-inflammatory cytokine

signaling not only induces insulin resistance, but also impairs kidney function ⁵². Furthermore, the infiltration of expanded visceral adipose tissue by immune cells and the pro-inflammatory cytokine signaling in adipose tissue decreases the secretion of the anti-inflammatory adipokine adiponectin. This protein is considered important for sustaining whole body insulin sensitivity and is thought to have beneficial effects on the kidney ⁵³. Furthermore, the expansion of adipose tissue results in increased production of the adipokine leptin. This adipokine is thought to induce proteinuria and type IV collagen expression in the kidney by increased production of TGF- β 1 and, thereby, to contribute to the development of glomerulosclerosis ^{54,55}. In addition, increased visceral obesity may be linked with the onset and progression of CKD via hyperinsulinemia, inappropriate activation of the renin-angiotensin-system and oxidative stress in the kidney. The resulting pathology includes impaired pressure/natriuresis relationship, increased salt sensitivity of blood pressure, aldosterone excess, glomerular hypertension, endothelial dysfunction and vasoconstriction as well as matrix expansion ^{42,56}.

Furthermore, we have recently proposed a novel concept to explain how nonalcoholic fatty liver disease (NAFLD) affects metabolism. When lipids accumulate in the liver, it is not only that hepatic glucose production increases and that dyslipidemia develops, but proteins with signaling properties in other tissues (hepatokines) are also being differently regulated ⁵⁷⁻⁵⁹. Among these, fetuin-A is the best studied hepatokine. Its expression is increased in NAFLD ⁶⁰⁻⁶³ and fetuin-A is well known to inhibit insulin signaling ⁶⁴. We could show that fetuin-A induces cytokine expression in monocytes and adipose tissue ⁶⁵ and we and others found fetuin-A to predict incident diabetes ^{66,67} as well as cardiovascular disease ^{68,69}. Recently in animals as well as *in vitro* it was shown that fetuin-A serves as an adaptor protein for saturated fatty acids, allowing them to activate toll-like receptor 4. Thereby, fetuin-A induces inflammatory signaling and insulin resistance ⁷⁰, which are important factors driving the development of type 2 diabetes and CVD. We could show that these animal data and *in vitro* findings can be translated to humans ^{71,72}.

Fetuin-A may also be involved in the pathogenesis of CKD. In advanced CKD and in end stage renal disease fetuin-A may inhibit ectopic calcification, thus protecting the kidney ⁷³. However, in early stages of CKD, when ectopic calcification is not yet relevant, the pro-inflammatory effects of fetuin-A may prevail to a large extent. Of note, elevated fetuin-A levels had been found in women with normal glucose tolerance but with albuminuria; this relationship was independent of well-known predictors of albuminuria ⁷⁴.

Effects of insulin on the vasculature and vascular insulin resistance

Insulin acts on the vasculature and causes endothelium-dependent vasodilation. This was first suggested by hypotensive episodes following s.c. or i.v. administration of insulin ⁷⁵ although this effect was difficult to distinguish from hypoglycemia-induced counter-regulation. In pioneering experiments in the early 1990's Baron et al. ⁷⁶ found an insulin-mediated increase of flow into the skeletal muscle of the leg during a hyperinsulinemic-euglycemic clamp. Compared to lean individuals, the dose-response curve of insulin was gradually right-shifted and flatter in insulin-resistant obese humans and type 2 diabetic patients ^{77,78}. Since both, the dose-response characteristic and the time course for glucose disposal paralleled the insulin-mediated vasodilatory effects, the authors speculated that insulin's effect on the blood flow may contribute to, or even limit, insulin's effects on glucose uptake of the skeletal muscle. Detailed analysis revealed that insulin-mediated vasodilation in skeletal muscle occurs in two phases ⁷⁹: After a few minutes through dilation of the terminal arterioles the number of perfused capillaries is increased without changes in blood flow (Fig. 3A). Within 30 minutes insulin induces the relaxation of larger resistance arterioles, thus increasing the overall limb blood flow with a maximum flow after about 2 hours. Thus in vivo the response to insulin is an integration of both capillary recruitment and increased total blood flow and both insulin actions are reduced and retarded in insulin resistant states like obesity and type 2 diabetes ⁸⁰. In a recent clinical study insulin applied during a euglycemic-hyperinsulinemic insulin clamp improved endothelial function at each of three

levels of the arterial tree in healthy controls, but not in obese insulin resistant subjects, emphasizing the close link between metabolic and vascular insulin resistance in humans ⁸⁰. Previous studies examining the mechanism behind the hemodynamic effects of insulin showed that application of the NO synthase inhibitor L-N-monomethyl-L-arginine abrogated the insulin-induced vasodilation in skeletal muscle, suggesting a major role for insulin-stimulated NO generation ⁸¹. The early finding that the insulin receptor is expressed on vascular endothelium at all levels of the arterial tree further supported an insulin-stimulated, NO-mediated vasodilation ⁸². Independent from endothelium insulin may also cause vasodilation by acting directly on vascular smooth muscle cells (VSMC) ⁸³.

Insulin-mediated endothelial transcytosis

Although insulin promotes its own delivery to the skeletal muscle microvasculature, it still has to cross the tight endothelial barrier to access the interstitium for action on the target cells. Several key experiments in animals showing that the metabolic insulin action closely correlates with the interstitial concentration of insulin, more so than with the plasma levels, suggest that the trans-endothelial transport is the rate-limiting step in insulin availability at target tissues. Previous studies provided evidence for a specific, saturable trans-endothelial insulin transport ⁸⁴. A recent profound study showed that in microvascular capillaries insulin is transported through the endothelium via clathrin-dependent entry and vesicle-mediated exocytosis across endothelial cells, thus by-passing the lysosomal degrading pathway (Figure 3B) ⁸⁵. This trans-endothelial insulin transport is promoted by insulin-induced endothelial NO production (Fig. 3B), indicating that insulin enhances its own transport ⁸⁶. These findings explain why whole-body insulin-induced glucose uptake is maximal after more than one hour, while insulin-induced glucose uptake in isolated adipocytes or skeletal muscle cells occurs within minutes. They also explain the reduced and retarded insulin-stimulated glucose uptake in insulin resistant subjects, where the endothelial insulin effects are impaired (endothelial dysfunction). In summary, insulin stimulates its own delivery to the target tissue via capillary recruitment, increase of

blood flow and, finally its own transcytosis. All of these processes are impaired in insulin resistant states (Fig. 3A,B).

Insulin and endothelial NO signaling

Insulin's hemodynamic effects have been identified in different organs, indicating an ubiquitous NO-mediated vasodilatory action. How does insulin promote endothelial NO production? After binding to endothelial IR, insulin activates PI-3K, PDK1 and Akt/PKB (Fig. 1A) leading to phosphorylation of endothelial NO-Synthase (eNOS) at Ser¹¹⁷⁷, enhanced eNOS activity and increased NO production ⁸⁷ (Fig. 4A). As was shown for classical insulin target cells, insulin does not only activate the PI3K pathway, but insulin also stimulates the MAPK/ERK pathway, as outlined in Figure 1A. In endothelial cells activation of this pathway induces the expression and secretion of endothelin 1, stimulating both, vasoconstriction, and VSMC proliferation. In the insulin resistant state, when the PI3K pathway is impaired, and the subsequent NO production is reduced, the MAPK/ERK pathway is not affected, or even enhanced (see below), resulting in an imbalance of both pathways towards vasoconstriction, VSMC proliferation and eventually hypertension ⁸⁸ (Fig.4B).

Recently the importance of APPL1 (adaptor protein, phosphotyrosine interacting with PH domain and leucine zipper 1) has been reported for the regulation of the vascular tone in endothelial cells ⁸⁹. The scaffold protein APPL1 is a major component in the signal transduction of adiponectin, a known insulin sensitizer in classical insulin target tissues ⁹⁰. Transgenic overexpression of APPL1 prevented the age- and obesity-induced impairment of insulin-mediated vasodilation and reversed obesity-induced augmentation of insulin-evoked ET-dependent vasoconstriction. By contrast, deletion of APPL1 caused a selective impairment in the PI3-K/Akt/eNOS pathway and augmented ERK1/2 signaling cascade in vascular endothelium, leading to decreased NO availability and enhanced ET-1 production, respectively. Thus the balance between vasodilatory and vasoconstrictory actions of insulin is shifted towards endothelial dysfunction (Fig.4B). In addition to its function as scaffold protein, APPL1 blocks the binding of the Akt/PKB inhibitor TRB3, thus, supporting the NO production at two intervention

points of the PI3K pathway (Fig. 4A). Activated Akt/PKB phosphorylates Raf-1 at the inhibitory Ser²⁵⁹, thereby, reducing the MAPK /ERK cascade. Hence, adiponectin controls insulin-mediated vasoreactivity via APPL1 and AMPK α 2⁹¹. Besides AMPK activators, glucagon-like peptide 1 induces PKA-mediated NO production (Fig. 4) which in turn stimulates muscle microvascular recruitment⁹². Accordingly, clinical trials revealed that GLP-1 receptor agonist treatment consistently reduced diastolic and systolic blood pressure up to 5 mm Hg⁹³.

Lessons from mouse models

To prove the causal involvement of the insulin signal transduction in insulin-mediated vasodilation a vascular endothelial cell IR knockout mouse was generated⁹⁴. Unexpectedly, in these mice the arterial blood pressure was not increased, but tended to be lower. They had normal glucose homeostasis. Since both, eNOS and ET-1 mRNA expression were reduced by approximately 50%, the reduction of both, the pro-vasodilatory and the pro-vasoconstrictive effect may have outweighed each other, resulting in no net effect. Of note, mice lacking eNOS displayed hypertension, insulin-resistance, fasting hyperinsulinemia, hyperlipidemia, and a 40% lower insulin-stimulated glucose uptake than control mice⁹⁵.

To further reveal the role of the insulin signaling cascade in regulating the vasculature tone mice with deleted IRS-1 were generated that showed elevated plasma triglyceride levels and an increased blood pressure⁹⁶. A relevant role of IRS-1 for the regulation of human blood pressure was recently demonstrated. Carriers of the IRS-1 Gly972Arg mutation had higher blood pressure and lower plasma nitrate/nitrite levels⁹⁷. Endothelial cells from donors with this mutation had lower eNOS expression/activity indicating a significant role of endothelial IRS-1 in regulation of the vascular tone in humans.

To ultimately answer the question whether insulin-mediated capillary recruitment and its trans-endothelial transport is related, or functions independently, the group of Kadowaki⁹⁸ have generated

knock out mice specifically lacking endothelial IRS-2, which is the major IRS isoform in endothelial cells. In these mice insulin signaling was impaired, as demonstrated by a decreased AKT phosphorylation after 60 minutes and a lack of Ser¹¹⁷⁷ phosphorylation of eNOS. Quantitative analysis of the data indicated that about 50% of the glucose uptake by skeletal muscle was mediated by the endothelium via insulin-mediated vasodilation and its own transport, similar to values from human subjects ⁹⁸. The observed insulin-mediated vasodilation and insulin delivery in skeletal muscle was not observed in the liver of the endothelial IRS-2 knock out mouse. This difference can be explained by the difference of the capillary structure. While capillaries in skeletal muscle have occluded conjugations in between the non-fenestrated endothelial cells, the sinusoid endothelium of the liver capillaries permits essentially free access of insulin and, thus, direct and faster action on hepatocytes. Therefore, the considerations that are valid for the skeletal muscle and adipose tissue do not apply for organs/organelles in which no tight endothelium is present, e.g. in renal glomeruli with fenestrated endothelium.

Are these insulin effects on the vasculature limited to skeletal muscle? Kadowaki's group addressed this question and investigated whether the insulin effects on vascular endothelium are also present in the pancreatic islets. Using the same mice they could show that the absence of IRS-2 in endothelial cells impairs the islet blood flow, similar to the extent that was seen in skeletal muscle ⁹⁹. Since pharmacological stimulation of islet blood flow in these animals almost completely restored insulin secretion, insulin-induced and endothelium-mediated increase in blood flow may regulate insulin distribution, in concert with other classical metabolic and hormonal stimuli. Since IRS-2 is the major IRS isoform expressed in endothelial cells and hyperinsulinemia leads to the down-regulation of IRS-2 in endothelial cells, impaired insulin action may be one of the mechanisms responsible for the decrease and/or the observed delay of insulin secretion in obese and type 2 diabetic subjects.

Impact of vascular insulin-resistance on renal hemodynamics

In rats, insulin induces NO-mediated vaso-relaxation in interlobular arteries, afferent and efferent arterioles ¹⁰⁰. In healthy subjects, insulin increased renal blood flow as analyzed by clearance of para-

aminohippuric acid during a hyperinsulinemic-euglycemic clamp ¹⁰¹. This effect was abrogated by application of the NO synthase inhibitor L-N-monomethyl-L-arginine indicating insulin-stimulated NO generation in the renal vasculature ¹⁰¹. It is expected that reduced NO signaling, that is characteristic for the insulin-resistant state, is also present in the renal vasculature. Indeed, it has been shown that renal vessels from the insulin-resistant Zucker rat failed to dilate in response to insulin and acetylcholine indicating endothelial dysfunction ¹⁰². In addition, there was a blunted myogenic vasoconstriction upon pressure, pointing to alterations in VSMC of the media. Endothelial dysfunction of renal vessels would lead to increased reno-vascular resistance and ultimately reduced renal blood flow. In a study with patients ranging from healthy to those with metabolic syndrome and type 2 diabetes, renal resistive index (RI) as index of reno-vascular resistance was gradually increased ¹⁰³. In that study, plasma concentration of adiponectin, which is known for its insulin-sensitizing effects, was an independent protective predictor, showing a negative correlation with RI.

Increased reno-vascular resistance due to reduced insulin-stimulated NO production would reduce glomerular filtration rate (GFR). However, in obese patients with metabolic syndrome or overt diabetes GFR is often increased. It is likely that mediators other than insulin account for glomerular hyperfiltration, which is also found in type 1 diabetes mellitus and defines early diabetic nephropathy. Reduced tubulo-glomerular feedback and dilation of afferent arteriole, owing to the increased reabsorption of sodium along with glucose, is thought to be one of the principal mechanism for glomerular hyperfiltration ¹⁰⁴.

Role of perivascular and renal sinus fat disposition

As schematically outlined in Figure 5 arteries of different size (except the cerebral vasculature), including small arterioles, are coated by adipose tissue. For long time it was thought that arteries were embedded in this perivascular fat tissue (PVAT) to protect and physically support the blood vessels. Recent studies have identified PVAT as an endocrine compartment releasing adipokines and other factors with vasodilatory activity. These factors may directly act on the vessel wall, because particularly in smaller arteries and microvessels there is no anatomical barrier between adventitia and PVAT^{105,106}. PVAT is not uniformly distributed along a specific vascular bed and PVAT differs in function and phenotype within the regions of a given blood vessel and also between the different PVAT compartments e.g. epicardial, periaortal, peritibial and renal sinus PVAT¹⁰⁷. Furthermore, PVAT differs from other fat compartments like subcutaneous and visceral fat¹⁰⁸. PVAT is not an innocent bystander, since these fat cells express and secrete both, pro-inflammatory cytokines like IL-6, IL-8, TNF- α and monocyte chemoattractant protein-1 (MCP-1) and the anti-inflammatory and insulin-sensitizing adiponectin and the regenerative factor hepatocyte growth factor (HGF). Extensive ex vivo studies with human PVAT showed that fetuin-A, a hepatokine released by the fatty liver, triggers expression and secretion of pro-inflammatory cytokines^{105,109}. Possibly functionally very important, but not finally characterized, is the adipocyte-derived relaxing factor (ADRF) showing anti-contractile effects on the vascular wall. This factor exerts its relaxing effects on vascular smooth muscle cells also in the absence of endothelium, indicating an independent regulation of the vascular tone by PVAT. Several reports attempting to identify the chemical nature of ADRF found evidence for low molecular weight compounds like H₂S and the palmitate methyl ester as candidates¹¹⁰.

Microvascular dysfunctions have been demonstrated to occur very early, even before clinical manifestation of vascular lesions. Therefore, Yudkin *et al.* have proposed that PVAT may be the cause of microvascular dysfunctions^{111,112}. A significant association between endothelial dysfunction and insulin resistance in young healthy first degree relatives of patients with type 2 diabetes was found,

independent of the classic cardiovascular risk factors ¹¹³. The quantity of PVAT surrounding human brachial artery is inversely associated with insulin sensitivity, but is not correlated with local endothelial dysfunction, as determined by flow-mediated dilation ¹¹⁴.

Only few studies have investigated the role of perirenal sinus adipose tissue for metabolic health and specifically for renal function ¹¹⁵. In a subcohort of the Framingham Heart Study PVAT quantity around the renal sinus was associated with hypertension and negative outcome for the patients ¹¹⁶. Furthermore, the amount of fat within this compartment predicted the outcome of diabetic renal disease. An independent study found that para- and perirenal fat thickness was an independent predictor of chronic kidney disease, increased renal resistance index and hyperuricemia in patients with type 2 diabetes¹¹⁷. Renal sinus fat mass is associated with an increased albumin excretion rate during exercise in healthy individuals who were at increased risk of type 2 diabetes ¹¹⁸. In insulin resistant individuals with fatty liver, the GFR was found to correlate inversely with increased renal sinus fat ¹⁰⁵.

Cellular dysregulation of the insulin signaling cascade in the kidney

Besides its putative actions on renal vasculature, insulin can act on virtually all other renal cell types such as mesangial cells, podocytes and tubular epithelial cells. Both isoforms of the insulin receptors are widely expressed in the kidney ³³. Unique to the kidney, insulin can access these target cells both from the lumen (in the case of podocytes) and from the basolateral side (in the case of tubular cells) as shown in a study with radiolabelled insulin ¹¹⁹. Studies published during the last years have focused on insulin signaling in these particular cells and revealed a variety of effects, ranging from glucose uptake, regulation of ion transport, prevention of apoptosis etc.

Glomerular endothelial and mesangial cells

Glomerular endothelial cells express the IR and respond to insulin binding with increased NO generation in analogy the extrarenal vasculature ³³. Because of its fenestration, the glomerular endothelium is, however, freely permeable to insulin, in that it can cross the basement membrane and enter bowman's space and tubulus. Therefore, insulin-mediated endothelial NO signaling is not a limiting factor for the transport of insulin to sub-endothelial cells, such as the VSMC (Fig. 3B), suggesting that changes in insulin plasma concentrations readily affect renal mesangial cells and podocytes. Insulin-mediated NO generation could in vivo contribute to the increase in GFR after a meal ¹²⁰. NO may also be an important player of the cross-talk between the podocytes and mesangial cells, that are adjacent to each other, separated only by the glomerular basement membrane. It has been shown that in glomerular endothelial cells insulin signaling also does not stimulate glucose uptake or remodeling of the actin cytoskeleton³³.

In mesangial cells, insulin is a potent survival factor that confers protection from apoptotic stimuli via stimulation of the PI3K/Akt pathway (Fig. 1A) ¹²¹. Increased MAPK signaling by insulin has been shown to stimulate the large conductance Ca^{2+} -activated K^+ channels (BK_{Ca} channels), resulting in mesangial cell relaxation, and possibly increased proliferation ¹²². Impaired insulin signaling in mesangial cells may be associated with reduced GFR, as was shown in a cross-sectional study that included 670 individuals and investigated the effects of the Gly(972)Arg variant of the human insulin receptor substrate 1 (IRS-1) gene ¹²³. Mesangial cells transfected with this variant exhibited attenuated insulin-stimulated phosphorylation of IRS-1 and Akt. Silencing of the IR in mesangial cells resulted in formation of homodimeric IGF1-receptors and increased IGF1 signaling ¹²⁴, that may lead to cell growth and proliferation. In mesangial cells from rats with STZ-induced diabetes, insulin reversed the mitogenic action of IGF-1 by regulating STAT5/SOCS2 expression ¹²⁵. When mesangial cells were kept in insulin deficient media, Kong *et al.* noted increased IGF1 signaling, resulting in increased cell proliferation and enhanced synthesis of fibronectin and collagen IV ¹²⁶. Furthermore, mesangial cells overexpressing GLUT 1 produced fibronectin and collagen IV in the absence of insulin, indicating that elevated glucose

utilization induces mesangial matrix production ¹²⁷. Altogether, these studies suggest that reduced insulin signaling in mesangial cells could contribute to mesangial cell hypertrophy, proliferation and matrix deposition, characteristic for early diabetic nephropathy.

The essential role of insulin in both endothelial and mesangial cells *in vivo* remains to be elucidated as cell-specific knockout models targeting the IR in these cells are lacking. They have proven to be an important tool in defining the role of insulin signaling, particularly in non-classical insulin target organs.

Podocytes

Podocytes are insulin-sensitive cells and respond to insulin stimulation with an increase in PI3K and MAPK signaling, resulting in glucose uptake via GLUT4 similar to skeletal muscle cells ³⁴. Besides energy uptake, this seems to serve adaptive changes in the podocytes in response to increased insulin secretion and hyperfiltration after a meal ¹²⁸. The insulin effect is dependent on expression of the transmembrane protein nephrin, which is an essential constituent of the slit diaphragm and is involved in the regulation of the actin cytoskeleton of the podocytes ¹²⁸. Nephrin allows the insertion of GLUT1- and GLUT4-containing vesicles into the plasma membrane by interacting with the vesicular SNARE protein VAMP2 (Vesicle-associated membrane protein 2). As nephrin expression is reduced with progression of diabetic nephropathy and albuminuria, insulin signaling is expected to be reduced and, thus, to negatively affect the filtration barrier. Insulin's action on podocytes also includes up-regulation of the ion channels TRPC (transient receptor potential cation channel, subfamily C) and BK_{Ca} channels ^{129,130}. In podocytes isolated from insulin-resistant obese db/db mice insulin failed to phosphorylate PKB and this was associated with reduced viability of the cells ¹³¹. The strongest evidence for the *in vivo* relevance of insulin on podocyte function comes from mouse models with a deletion of the IR in the podocyte ¹³². Starting at 8 weeks of age, these knockout mice developed albuminuria caused by foot process loss and severe podocyte damage. After 13 weeks of age, light microscopy revealed podocyte loss, glomerulosclerosis and mesangial expansion, features characteristic of human diabetic nephropathy. As a mechanism, the authors found severe impairment of the remodeling of the actin

cytoskeleton, leading to loss of podocyte architecture and morphology. All these changes occurred in the absence of hyperglycemia, underscoring the role of normal insulin signaling and glucose uptake for podocyte health. Another mechanism could be related to dysfunction of the endoplasmic reticulum (ER) that is induced by the absence of insulin signalling¹³³. This has been found to impair PI3K-dependent translocation of transcription factor spliced X-box binding protein-1 (sXBP1) into the nucleus resulting in maladaptive ER-stress signalling.

Tubular epithelial cells

Long before the advent of the current cell biologic approach to investigate insulin signaling in the kidney, insulin effects on renal tubular function have been studied in animals since the 1970s, using microperfusion (ex vivo) or micropuncture (in vivo) methods. The results clearly showed a stimulatory effect of acutely administered insulin on tubular sodium reabsorption in the proximal tubule and Henle's loop of rabbits and dogs¹³⁴⁻¹³⁶. In addition, acutely administered i.v. insulin stimulated reabsorption of phosphate, potassium and water^{136,137}. In contrast, chronic infusion of insulin over five days in rats reduced sodium excretion only transiently on the first day, an effect that disappeared thereafter, but was associated with an increase in blood pressure¹³⁸. Extensive studies in dogs under different conditions such as insulin resistance, chronic norepinephrine infusion, reduced kidney mass, high-salt intake or chronic angiotensin II infusion could not demonstrate that chronic insulin administration causes sustained sodium retention or hypertension¹³⁹. Much of the controversy whether insulin was anti-natriuretic or not was solved when it became clear that hyperglycemia was required to induce a sodium-retaining effect of hyperinsulinemia, indicating that the euglycemic, hyperinsulinemic clamp was not the appropriate model to study the pathogenesis of hypertension in hyperglycemic diabetic patients¹³⁹. By inducing sustained hyperglycemia and hyperinsulinemia at the same time, formerly resistant dogs developed sodium retention during chronic insulin administration that was sufficient to antagonize the natriuresis caused by hyperglycemia¹⁴⁰.

Distal tubule

In cells expressing the epithelial sodium channel (ENaC), insulin was found to stimulate Na⁺ transport via up-regulation of ENaC membrane abundance, which was dependent on PI3K activity¹⁴¹. Increased PI3K signaling by insulin leads to activation of mTORC2 and the serum- and glucocorticoid kinase 1 (SGK1), which are essential serine-threonine kinases involved in the regulation of ENaC in the distal tubule (Fig. 6). SGK1 is heavily up-regulated by aldosterone on a transcriptional and activity level via K-Ras2a-activation of PI3K and PDK1. SGK1 increases ENaC membrane expression by inhibiting the ubiquitin ligase Nedd4-2 (neuronal precursor cells expressed developmentally down-regulated) mediated internalization and degradation¹⁴². Elevated glucose concentration also increases SGK1 gene expression and this may be one of the critical factors underlying the stimulation of sodium retention in type 2 diabetes¹⁴³. Increased SGK1 expression by hyperglycemia would also explain the development of sodium retention upon chronic insulin infusion in the above mentioned study in dogs¹⁴⁰.

Surprisingly, mice with deletion of the IR in the distal tubule, driven by the Ksp (Kidney specific) cadherin promoter, displayed hypertension and impaired natriuresis, following a sodium load¹⁴⁴. As an explanation, the authors suggested that reduced renal NO signaling, by absent insulin signaling, could account for this finding. In a subsequent study, the authors confirmed that reduced insulin signaling in those mice conferred salt sensitivity of blood pressure by reducing renal NOS expression¹⁴⁵. In another mouse model with a deletion of the IR in the collecting duct using an aquaporin-2 promoter, the same group found that ENaC activity was reduced and associated with reduced blood pressure¹⁴⁶. The stimulating effect of insulin on ENaC currents was confirmed in isolated split-open tubules, an effect that was blunted in mice with IR deletion and abrogated by inhibition of PI3K and mTOR¹⁴⁷. Altogether, the previous and also current data confirm that insulin is per se an anti-natriuretic hormone in the distal tubule.

Proximal tubule

Insulin signaling in the proximal tubule might affect renal gluconeogenesis that accounts for up to 25% of the glucose released into the circulation ¹⁴⁸. Reduced insulin signaling would lead to disinhibition of gluconeogenesis that would in turn contribute to hyperglycemia. In mice with a deletion of the insulin receptor in the proximal tubule, driven by the glutamyltransferase promoter, fasting glucose was indeed elevated and the activity of glucose-6-phosphatase, the enzyme catalyzing the final step of gluconeogenesis and responsible for release of glucose into the circulation, was increased in renal cortex homogenates ¹⁴⁹. These data confirmed the findings of an earlier study in insulin-resistant diabetic Zucker rats, which had disinhibited renal gluconeogenesis due to increased mRNA expression and activity of rate-limiting gluconeogenic enzymes ¹⁵⁰. Another key function of the proximal tubule is the complete reabsorption of the filtered glucose load (160-180 g per day) that is accomplished by two sodium-coupled glucose transporters (SGLT1 and 2) with SGLT2 contributing to 97% of the renal glucose reabsorption ¹⁰⁴. SGLT2-mediated transport is up-regulated by insulin and involves phosphorylation of SGLT2 via IR signaling ¹⁵¹. Up-regulation of SGLT2 activity by insulin may serve to minimize glucose losses after a meal. The renal expression of SGLT2 in animal models of diabetes does not follow a clear pattern and is dependent on the model ¹⁵². However, it has been established for long time that glucose reabsorption is increased in patients with type 2 diabetes ¹⁵³, arguing against an insulin-resistance of SGLT2 up-regulation in type 2 diabetes.

Disturbances of insulin-regulated renal functions in humans

Relevance of IR deletion for understanding of human insulin resistance

Mouse models involving the deletion of IR are thought to represent the most extreme form of insulin resistance. However, the validity of the results with regards to understanding insulin resistance in humans is not always clear. First, even in conditions of severe insulin resistance in humans a complete disruption of insulin receptor signaling does not occur. Secondly, the models are often equivocal and do not fully agree with the complex situation in humans; some findings are even unexpected and

conflicting. Mice with a constitutive deletion of the IR are massively hyperglycemic and die of diabetic ketoacidosis within few days after birth, despite developing hyperinsulinemia ^{154,155}. In contrast, humans with genetic deficiency of the IR exhibit fasting hypoglycemia, retarded growth and develop ketoacidosis only after a meal ¹⁵⁶. Mice with specific deletion of the IR in the skeletal muscle, a classical insulin target tissue that is thought to be insulin resistant from the early stages of insulin resistance in humans, were normoglycemic and not hyperinsulinemic ¹⁵⁷. However, they had increased fat cell mass and obesity, suggesting redirection of energy and storage of fat similar to the human situation. Deletion of the hepatic IR resulted in severe fasting hyperglycemia, disinhibition of gluconeogenesis and glucose intolerance ¹⁵⁸. These changes resemble the human situation as the fatty liver is of central importance to insulin resistance ⁶².

Accordingly, data from mice with deletion of the IR in various renal cell types have to be interpreted with great caution. Contrary to the expectation that reduced insulin signaling may cause natriuresis, the first published mouse model using an IR deletion under the Ksp promoter showed reduced sodium excretion ¹⁴⁴. Subsequent studies with a more defined model involving the deletion of ENaC in the distal tubule fitted better with the role of insulin in promoting sodium retention ^{146,147}. Unlike in humans with diabetic nephropathy, the deleterious effect of IR deletion in podocytes has been shown to occur under normal glucose and insulin concentrations in otherwise healthy and insulin-sensitive mice. In addition, these mice did not have other typical features of diabetic nephropathy, such as enlarged kidneys, mesangial hypercellularity or Kimmelstiel–Wilson lesions, despite severe podocytopathy ¹³². In conclusion, genetic IR knock out models only highlight single distinct aspects of human insulin resistance, either within an organ, or in the whole organism, and represent only a small piece of the complex mosaic underlying systemic insulin resistance.

Insulin resistance of the kidney

Despite the compelling evidence that the kidney is an insulin responsive organ with insulin regulating different functions, it is not clear whether the kidney in general or in parts is affected by insulin,

resistance similar to the classical target organs such as liver, skeletal muscle or white adipose tissue. The human kidney abundantly expresses the IR isoform B which is the isoform found in the classically insulin-responsive organs¹⁰ and glucose uptake is insulin-dependent in some (such as podocytes), but not all cells (tubulus, mesangial cells). Studying rats with insulinopenic and insulin-resistant diabetes identified that insulin signaling, as represented by Akt phosphorylation, was reduced only in isolated glomeruli, while tubuli were not affected¹⁵⁹. In glomeruli, the phosphorylation of other downstream effectors of insulin, such as eNOS and GSK 3 α , were similarly reduced. However, there was no change of MAPK phosphorylation, indicating resistance only of the IRS-1/PI-3K driven arm of insulin signaling (Fig. 1A). The authors found reduced protein expression of IRS-1 that could be restored by inhibition of PKC β . Altogether, this important study has shown that within the kidney glomeruli, but not tubuli, can develop insulin resistance. However, the study could not elaborate whether this is confined to glomerular endothelial cells only, or also involves podocytes and mesangial cells. The study also did not address if insulin sensitivity was retained equally in the whole tubulus or only in parts.

Tubular sodium transport

If the tubulus epithelium is not affected by insulin resistance, then hyperinsulinemia will lead to a stimulation of renal sodium transport and promote sodium retention and salt-sensitive hypertension, features that are frequently encountered in obese patients with the metabolic syndrome. Indeed, insulin infusion during a euglycemic, hyperinsulinemic clamp induced a similar reduction of sodium excretion in obese adolescents¹⁶⁰ or in patients with type 2 diabetes, compared to control subjects, while the effect on peripheral glucose uptake was reduced in insulin-resistant obese and diabetic patients only¹⁶¹. The site of salt retention during hyperinsulinemia is the distal tubule and involves stimulation of ENaC by PI3K/SGK1 (Fig. 6). However, it can also involve the proximal tubule, where the stimulatory effect of insulin on sodium transport has been shown to be preserved in type 2 diabetic rat models and in patients^{162,163}. This was associated with preserved expression of IRS-2 and Akt phosphorylation, whereas expression of IRS-1 was reduced in the renal cortex. Transcription factors

such as forkhead box class O1 (FoxO1) and sterol regulatory element-binding protein 1 (SREBP 1), that are involved in the down-regulation of IRS-2 in the fatty liver, were not changed, indicating subtle differences in altered insulin signaling during insulin-resistance between liver and the kidney.

It must be emphasized that in addition to hyperinsulinemia, hyperglycemia stimulates sodium transport in the proximal and distal nephron by transcriptional up-regulation of SGK1 which is then activated by insulin- and PI3K-dependent phosphorylations at consecutively two sites ^{142,143,164}. Altogether, both hyperinsulinemia and hyperglycemia converge to enhance sodium transport in the distal tubule and promote net sodium retention that is able to override osmosis-driven diuresis by hyperglycemia in the proximal tubule ¹⁴⁰.

Renal gluconeogenesis

Besides periportal hepatocytes, only proximal tubular cells are capable of gluconeogenesis, which accounts for up to 25% of the glucose released into the circulation, and is suppressed by insulin ¹⁶⁵. In patients with type 2 diabetes both, renal and hepatic gluconeogenesis, was found to be increased, thus, contributing to hyperglycemia during the fasting state (IFG) ¹⁶⁶. This finding implies that the proximal tubule is resistant to the suppressive action of insulin on gluconeogenesis, as it is the case in the liver. This is in agreement with animal models of insulin-resistant rats and mice with specific deletion of the IR in the proximal tubule, showing disinhibition of gluconeogenesis ^{149,150}. In contrast, tubular sodium transport seems to be spared from insulin-resistance that can only explained by differences in intracellular signaling such as alterations of IRS-1 and IRS-2 expression ¹⁶³ and/or involvement of SGK1, regulating a variety of sodium channels. Whereas reduced IRS-1 expression may be responsible for disinhibition of gluconeogenesis ¹⁶² (and also reduced NO production ¹⁵⁹), preserved IRS-2 expression (independent of reduced IRS-1) seems to maintain stimulation of sodium transport in the proximal tubule. This evidence is supported by a study in IRS-1 and IRS-2 knockout mice showing attenuated stimulation of sodium transport exclusively in mice with deletion of IRS-2 ¹⁶⁷.

Possible therapeutic approaches to overcome renal and vascular insulin resistance

Effects of weight loss

The strong association of obesity and insulin resistance suggests that lifestyle changes, which result in weight loss and increased muscle mass, restore whole-body insulin sensitivity. Indeed, lifestyle intervention in individuals with both, impaired and also normal glucose tolerance, improved insulin sensitivity, as estimated from the relation of glucose and insulin plasma concentration, and was associated with a reduction in total-, visceral- and ectopic fat mass¹⁶⁸. MR studies showed that visceral adipose tissue and hepatic lipids can be significantly reduced during lifestyle intervention with their baseline values being predictive factors for an improvement of insulin sensitivity^{169,170}. Similar to the improvement of whole body insulin sensitivity, renal and vascular insulin resistance should similarly improve with lifestyle intervention. However, studies focusing on the insulin sensitivity of renal and vascular function are rare and can barely discriminate between systemic and organ-specific effects. The available lifestyle intervention studies with renal endpoints showed reduction of glomerular hyperfiltration and albuminuria, suggesting normalization of afferent vasodilatation, filtration pressure and possibly podocyte dysfunction¹⁷¹. They are also thought to improve salt sensitivity and to reduce blood pressure¹⁷², suggesting altered tubular sodium handling. With regard to vascular insulin resistance, lifestyle intervention improved macro- and microcirculation, as indicated by increased flow-mediated vasodilation in the brachial artery and insulin-induced vasodilation in cutaneous vessels^{173,174}. However, weight loss after lifestyle intervention is not uniformly associated with improved insulin-resistance and there is a substantially high proportion of patients not responding to lifestyle intervention that might be related to persistence of visceral obesity and, more importantly, liver steatosis^{170,175}. Compared to lifestyle intervention, bariatric surgery mostly results in a large loss of body weight, and hence, more effectively improves insulin sensitivity, but this treatment can only be considered in patients with morbid obesity¹⁷⁶.

Pharmacological treatment

Metformin and PPAR γ agonists. Pharmacologically, insulin-resistance can be improved with metformin, and with thiazolidindiones (TZD) the latter of which activate the peroxisome proliferator activated receptor gamma (PPAR γ). Whereas metformin is increasingly being recommended in the very early stages of insulin resistance and pre-diabetes, treatment with thiazolidindiones, such as pioglitazone, is indicated in patients with overt diabetes ¹⁷⁷. They lower blood glucose concentration predominantly by stimulation of lipogenesis in the white adipose, while reducing hyperinsulinemia. This has brought up their attribute to be insulin-sensitizers. Whereas metformin has predominantly effects on the liver, pioglitazone acts on both, the vasculature and the kidney, by a direct effect on various cell types such as endothelium, VSMC, podocytes or tubular cells. Pioglitazone treatment improves endothelial function and lowers blood pressure, most probably by a direct vasodilating effect ¹⁷⁸. In the kidney, several studies found a significant reduction of albuminuria, which according to a metaanalysis, amounted to 25 to 65%, depending on spot urine samples or collected urine samples ¹⁷⁹. In the randomized controlled DREAM trial involving 5269 patients with prediabetes (IFG and IGT), rosiglitazone treatment reduced renal outcomes, predominantly by reducing albuminuria (12.3 vs. 15.0 %, HR 0.82 [0.69–0.98]) ¹⁸⁰. Reduction of progression of renal function, as defined by GFR loss ≥ 30 %, was reduced with borderline significance (3.1 vs 4.0%, HR 0.77 [0.58–1.04]). However, rosiglitazone increased risk for developing heart failure (0.53 vs. 0.08%; HR 7.04 [1.60 –31.0]), which is the most severe adverse effect of TZD treatment, that has limited its wide-spread use. This complication is caused by stimulation of sodium retention by the kidney, involving activation of ENaC and up-regulation of SGK1 ¹⁸¹.

A clinical important effect of TZDs is their ability to improve liver steatosis and dysregulated hepatokine production, not only via actions on adipose tissue, but also via direct effects on the liver. In this respect pioglitazone was found to suppress mRNA and protein expression of fetuin-A in Fao hepatoma cells. Interestingly, rosiglitazone, but not metformin, also inhibited fetuin-A expression. In addition, GW

9662, an inhibitor of peroxisome proliferator-activated receptor (PPAR) gamma, reversed pioglitazone-induced suppression of fetuin-A ¹⁸². These data suggest that thiazolidinedione derivatives may have common characteristics with regard to fetuin-A suppression, possibly through PPAR gamma activation. In agreement with these studies pioglitazone, but not metformin or exercise, decreased fetuin-A levels during a 6 month intervention, although there were similar effects in the improvement of insulin sensitivity between the three groups ¹⁸³.

Insulin. One important pathophysiological factor which causes insulin resistance is the elevated plasma glucose itself. The association of increased blood glucose levels with insulin resistance is called glucotoxicity, or more accurately glucolipotoxicity ^{184,185}. As a result of chronic hyperglycemia, insulin signaling is inhibited, which can be seen also in type 1 diabetes ¹⁸⁶. Glucolipotoxicity is not only associated with insulin resistance but also with beta cell dysfunction. Therefore, one therapeutic concept of treating hyperglycemia in type 2 diabetes is to reverse glucotoxicity by insulin treatment, resulting in recovery of residual β -cell function and improvement of insulin resistance. The ORIGIN trial has examined early insulin treatment with a long-acting insulin in patients with dysglycemia including pre-diabetes and type 2 diabetes in a randomised controlled intervention study ¹⁸⁷. It has been shown that diabetes can be prevented with early insulin treatment ¹⁸⁸. In addition, early insulin therapy significantly reduces the incidence of eye and kidney disease in patients with a baseline HbA1c >6.4% ¹⁸⁹.

SGLT2 inhibitors. SGLT2 inhibitors might be a promising new pharmacological approach to overcome insulin-resistance. These new drugs inhibit proximal tubular glucose reabsorption in an insulin-independent manner and induce glucose loss that amounts to 50 – 150 g ¹⁹⁰ in diabetic patients, depending on the level of hyperglycemia. Although SGLT2 inhibitors lack a systemic effect, recent studies in humans and animals suggest that this new class of antidiabetic drugs improves insulin sensitivity. In a small study with 12 patients with type 2 diabetes, treatment with the SGLT2 inhibitor dapagliflozin improved peripheral glucose uptake, as analyzed with a euglycemic, hyperinsulinemic

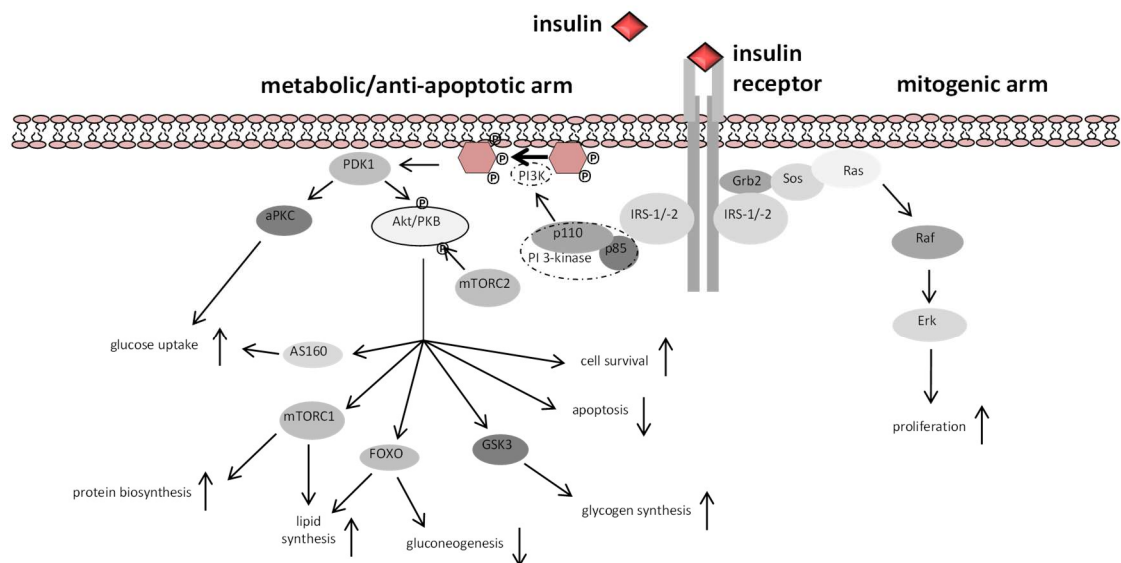
clamp, while it also reduced plasma insulin concentrations ¹⁹¹. The effects of dapagliflozin occurred rapidly, within two weeks of treatment, well before induction of significant weight loss occurred. The authors of that study explain these findings with the immediate correction of hyperglycemia by SGLT2 inhibition, leading to regression of glucotoxicity. In a larger study including 66 patients with type 2 diabetes the effects of SGLT2 inhibition on insulin secretion and peripheral glucose uptake were detectable even after a single dose of the SGLT2 inhibitor empagliflozin, which is best explained by beneficial effects of immediately lowering plasma glucose concentration and, thereby, reducing glucotoxicity ¹⁹⁰. However, the contribution of the insulin sensitizing effects to the overall beneficial effect of SGLT2 inhibitors needs to be examined and confirmed in large outcome trials, taking into account measures of insulin sensitivity such as the HOMA (homeostasis model assessment of insulin resistance), or others. In addition to possibly improving insulin sensitivity, SGLT2 inhibitors have shown beneficial effects on cardiovascular outcomes ¹⁹² and, most recently, on renal end points, by hitherto poorly-defined mechanisms. Altogether, SGLT2 inhibitors emerge as a new therapeutic approach to treat patients with insulin resistance and manifest type 2 diabetes.

Conclusion

Renal and vascular insulin resistance develops as part of the complex mosaic of systemic insulin resistance and affects organ-specific functions and cross-talk, alike. Dysregulation of insulin-regulated pathways in the kidney and vasculature culminate and sustain pathophysiological alterations found in metabolic syndrome, such reduced endothelial function, increased sodium retention and renal gluconeogenesis. These changes occur early in the course of the development of type 2 diabetes and significantly determine subsequent renal and vascular damage. To prevent micro- and macro-angiopathic diseases, intervention at the insulin resistant state is required to halt progression, or even reverse pathophysiological alterations. Identification of molecular mechanisms promoting renal and vascular insulin resistance is warranted, to better understand the organ cross-talk involving classical and non-classical insulin-responsive organs.

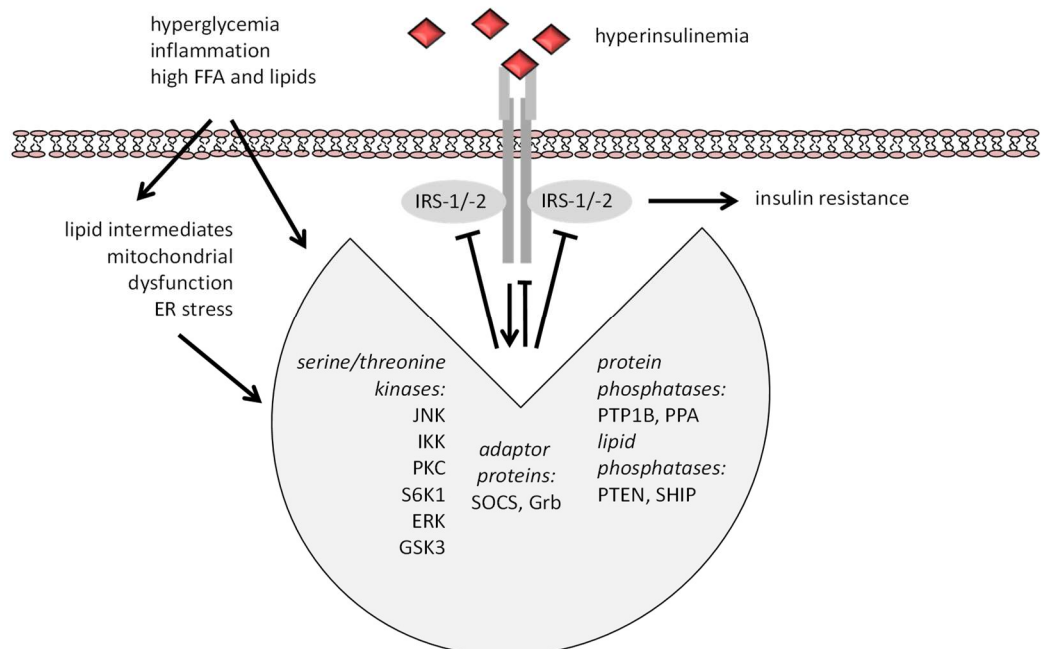
Figures with legends

Figure 1A



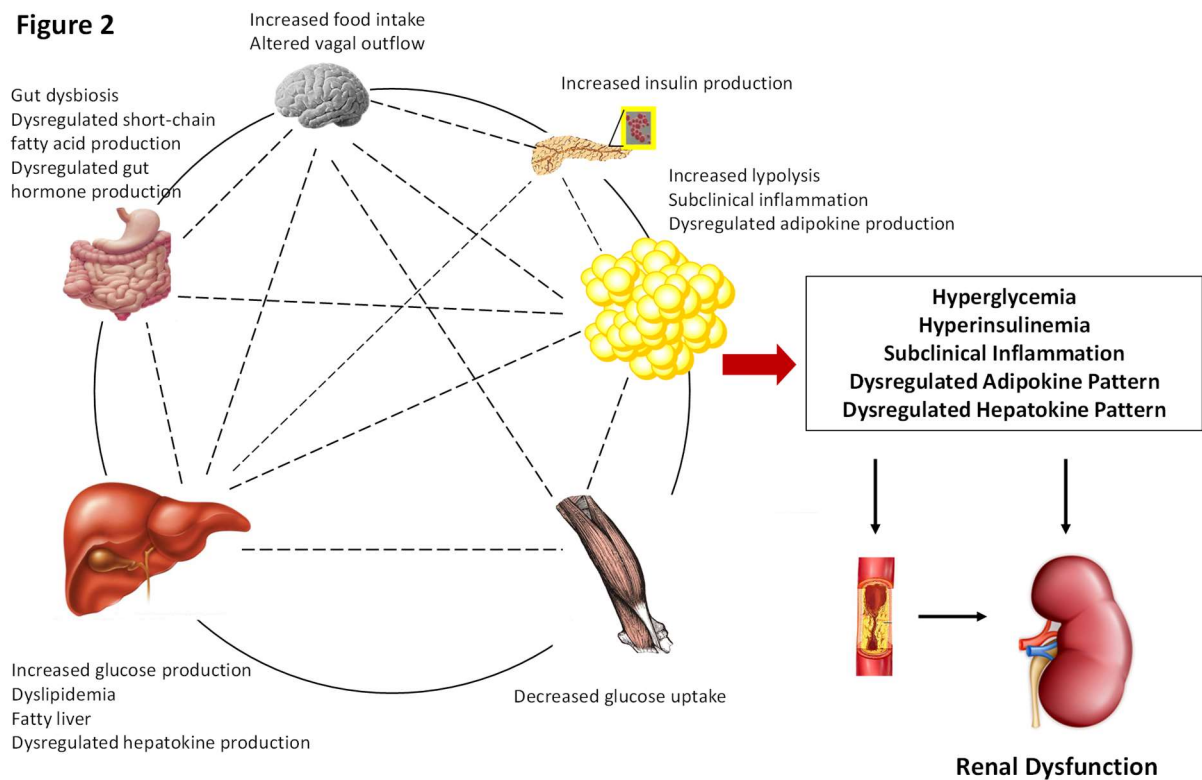
Scheme of the insulin signaling pathway. Binding and activation of the insulin receptor leads to recruitment of IRS isoforms and subsequently to activation of the PI3-kinase-Akt/PKB pathway (left arm) and regulates glucose and lipid metabolism, protein biosynthesis, cell survival and apoptosis. Activation of the Ras/Erk pathway increases proliferation (right arm).

Figure 1B



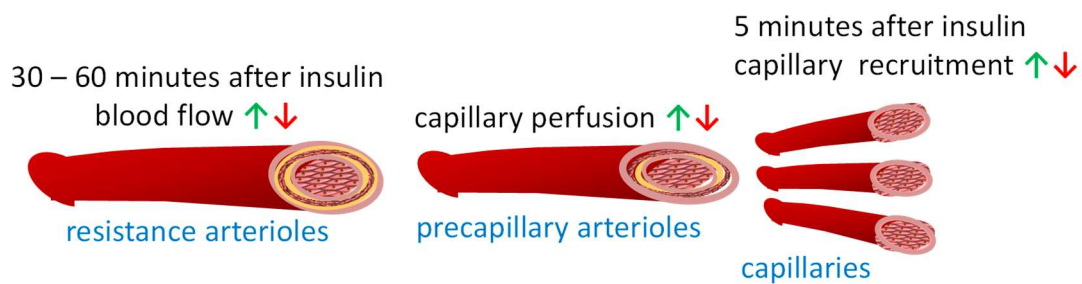
Negative regulation of the insulin signaling pathway. Hyperglycemia, hyperinsulinemia, high plasma free fatty acids (FFA) and inflammation activate via several mechanisms serine/threonine kinases, adaptor proteins and phosphatases resulting in a chronically reduced cellular response to insulin

Figure 2



Organ cross-talk in the insulin-resistant state. Insulin exerts its metabolic actions classically on liver, fat and skeletal muscle. Non-classical insulin actions include the brain, gut, pancreas, vasculature and the kidney. The physiological organ cross-talk is impaired in the insulin resistant state

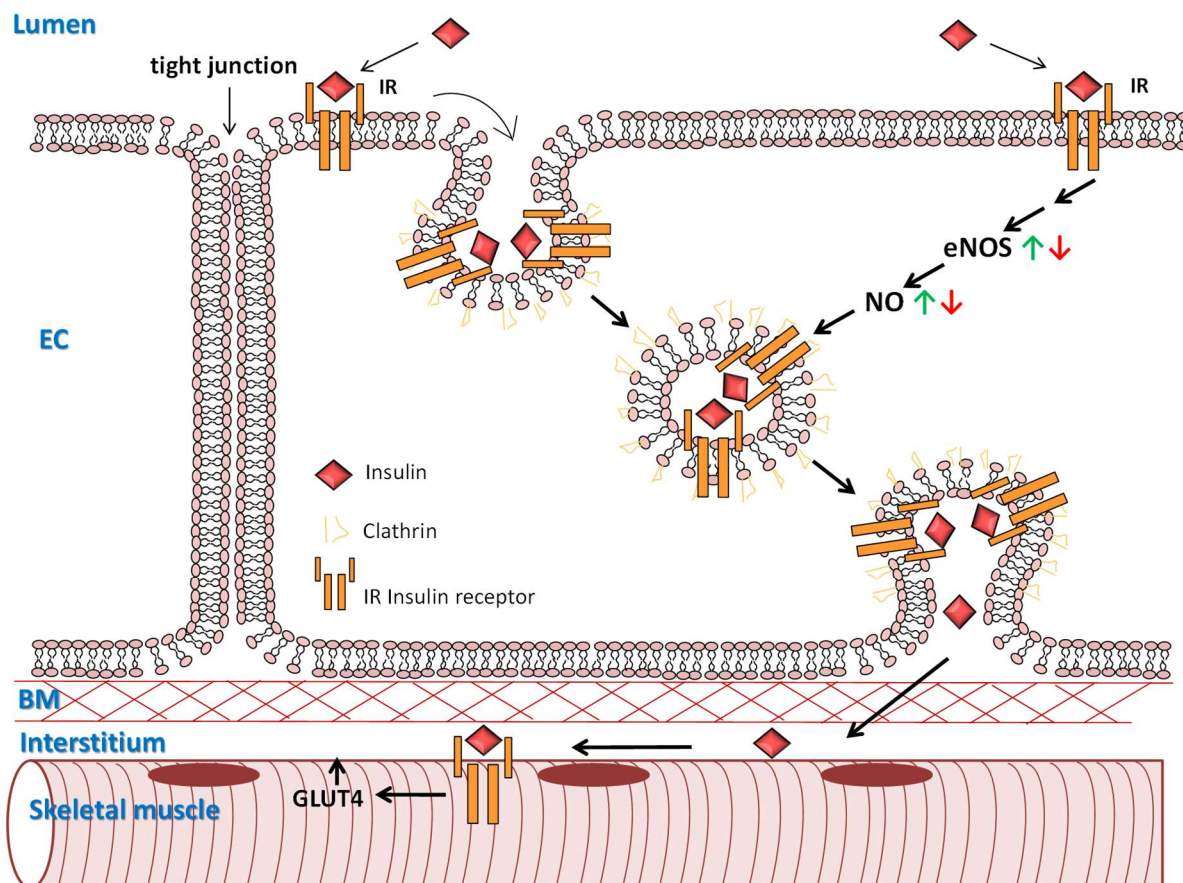
Figure 3A



Physiologic and impaired insulin effects on the arterial vasculature

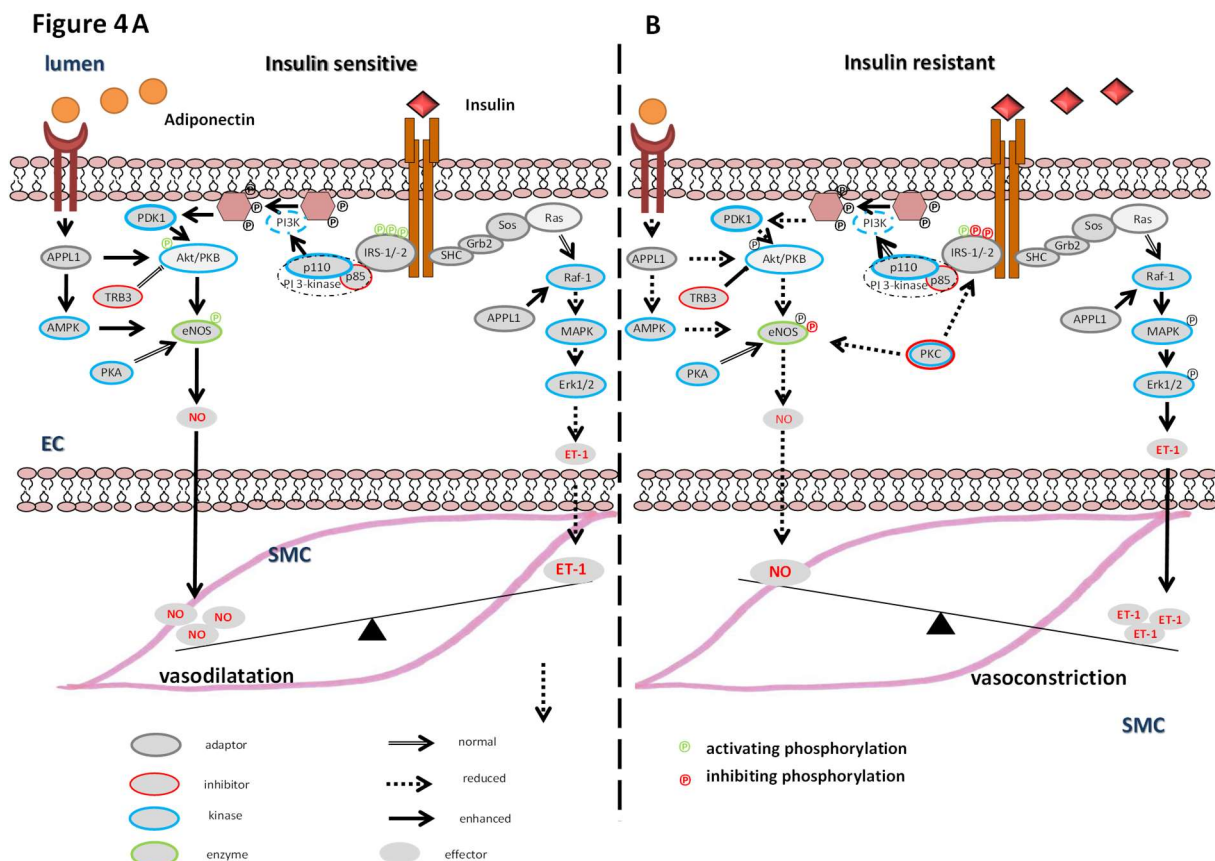
Insulin acts at all three levels of the vascular tree. In the skeletal muscle, after a few minutes, insulin induces capillary recruitment by vasodilation of the precapillary arterioles, followed by relaxation of the larger resistance arterioles, leading to an enhanced blood flow (green arrows). Accordingly, insulin improves endothelial function in conduit arteries. These actions of insulin are reduced in obese subjects and patients with type 2 diabetes (red arrows).

Figure 3B



Insulin stimulates its own trans-endothelial transport by clathrin-dependent NO-mediated endo- and transcytosis. In skeletal muscle and in adipose tissue the non-fenestrated microvascular endothelial monolayer forms a tight barrier, restricting free access of plasma constituents to the subendothelial interstitium and, if not activated, preventing the interaction of blood cells with the endothelium. For delivery to subendothelial target cells insulin binds to its receptor (IR) on the endothelial plasma

membrane and the resulting complex is clathrin-dependently engulfed, transported through the endothelial cell and released by vesicle-mediated exocytosis. After passage through the basement membrane (BM) insulin stimulates glucose-uptake at the target cells. Transcytosis of insulin is stimulated by NO which in turn is generated by insulin-stimulated activation of eNOS (green arrows). All these insulin actions are impaired in insulin resistant subjects (red arrows).



Physiologic and impaired insulin signal transduction in endothelial cells

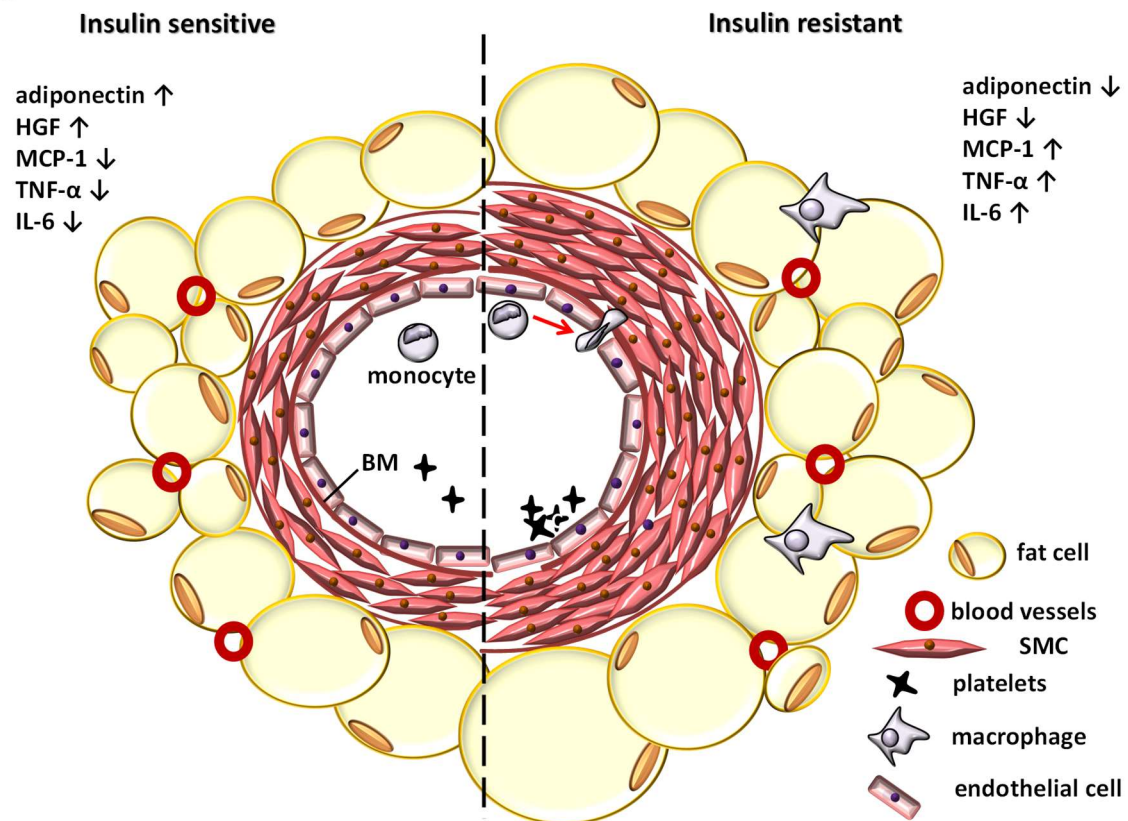
Insulin plays an essential role in maintaining vascular function by regulating both endothelial NO (left arm) and endothelin-1 (ET-1) production (right arm) acting on vascular smooth muscle cells (VSMC) causing vasodilation and vasoconstriction, respectively. Insulin resistance induces endothelial dysfunction

A Binding of insulin to its cognate receptor stimulates the PI3K-Akt axis of the cascade via IRS-1/2 as shown in detail in Figure 1A. Activated Akt phosphorylates Ser¹¹⁷⁷ of eNOS for NO-dependent vasodilation. Ser¹¹⁷⁷ may also be phosphorylated by activated AMP-kinase (AMPK) or protein kinase A (PKA). The insulin sensitizing hormone adiponectin supports insulin action by activating AMPK via the adaptor protein phosphotyrosine interacting with PH domain and leucine zipper 1 (APPL1). APPL1 competes with the Akt inhibitor TRB3 both actions further stimulate eNOS activity and subsequent NO production. Since insulin-mediated stimulation of the MAPK/ERK axis is reduced by APPL1-mediated inhibition of Raf-1 ET-1 production and action is overbalanced by NO resulting in vasodilation. (enhanced actions = green arrow; reduced actions = red arrow), MAPK = mitogen-activated protein kinase; ERK = extracellular signal-regulated kinase

B In insulin resistant state post-receptor interferences reduce insulin signal transduction. Various metabolic and inflammatory factors inhibit the action of IRS1/2 through serine phosphorylation (Fig. 1B) leading to reduced activation of the PI3 kinase-Akt axis and NO production. Furthermore, since adiponectin levels are reduced in insulin resistant and obese subjects phosphorylation of APPL1 Ser⁴⁰¹

and subsequently its function is reduced further shifting the insulin signal from the PI3 kinase-Akt axis to the MAPK/ERK axis which is independent of IRS1/2. Since the inhibitory effect of APPL1 on RAF-1 is reduced the MAPK/ERK axis is even enhanced. Furthermore, activated PKC in insulin resistant individuals (see Fig. 1B) phosphorylated the inhibitory Ser⁴⁹⁵ of eNOS further reducing NO production. Together the reduced NO production and increased ET-1 causes vasoconstriction of the small artery. Furthermore since the expression of adhesion molecules is under the control of the MAPK/ERK axis overexpression of vascular cell adhesion molecule-1 (VCAM-1) and E-selectin is stimulated.

Figure 5

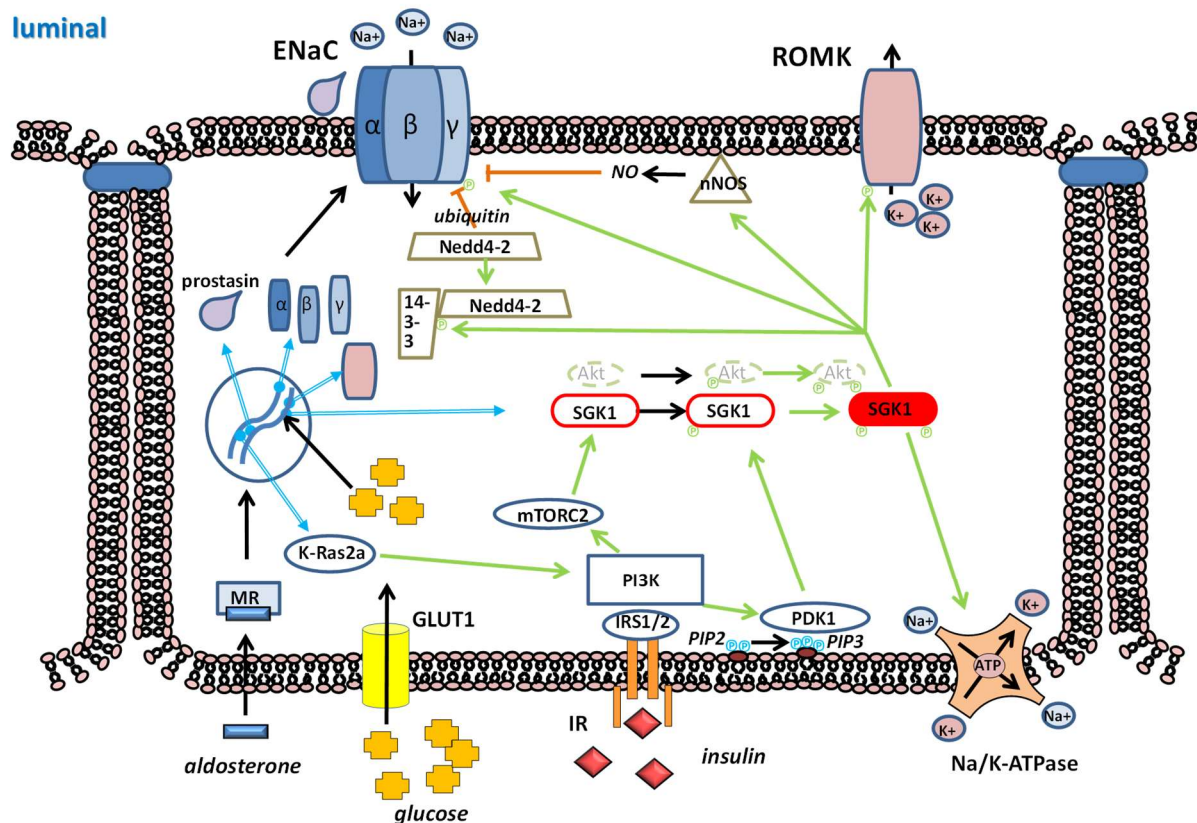


Perivascular adipose tissue influences vascular function

Large and smaller arteries are surrounded by adipose tissue. In healthy insulin sensitive individuals adiponectin levels are increased and the perivascular fat cells secrete vasodilating factors e.g. ADRF which are still ill-defined. In the insulin resistant state perivascular fat cells increase in size and in number and secrete more proinflammatory factors like IL-6 and TNF-α together with monocyte chemoattractive factors (MCP-1) and less regenerative factors like hepatocyte growth factor (HGF) particularly under the influence of fatty liver derived fetuin-A. Furthermore these proinflammatory factors increase endothelial permeability and enhance insulin resistance by inhibition of the IRS-PI3K-Akt- axis of insulin signal transduction (see Fig. 1). Factors like MCP-1 foster the attraction and infiltration of monocyte/macrophages into the vascular wall and the surrounding adipose tissue and in turn may further increase the proinflammatory state (vicious cycle). The activated endothelium expresses adhesion factors thus attracting leukocytes and activating platelets.

Figure 6

luminal



basolateral

Insulin signaling in the principal cell of the aldosterone-sensitive distal nephron.

Serum-and Glucocorticoid-Inducible Kinase 1 (SGK1) is the central effector of both insulin and aldosterone. SGK1 expression is rapidly induced on a transcriptional level by aldosterone/MR and also by hyperglycemia. It is activated after sequential phosphorylation at Ser⁴²² by mTORC2¹⁶⁴ and Thr²⁵⁶ by PDK1. Active SGK1 increases ENaC membrane abundance and currents via direct and indirect effects, among them inhibition of the ubiquitin ligase Nedd4-2. It also stimulates renal outer medullary potassium channel (ROMK) and the Na/K-ATPase. Akt1 is of minor importance as effector of insulin, although it is similarly phosphorylated and has the same downstream targets. (MR=mineralocorticoid receptor)..

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Author contributions

All authors contributed equally to researching the data for the article, discussing the article's content, writing the article and review/editing of the manuscript before submission.

Competing interests statement

No conflict to declare.