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K_{ATP} Channels in Vascular Smooth Muscle: Structure, Regulation and Functional Roles

N. B. Standen

K_{ATP} channels link membrane K⁺ permeability to metabolism. In vascular smooth muscle, the molecular composition of the channel isoforms expressed has not been determined definitively, though there is good evidence that a channel comprising the pore-forming subunit Kir6.1 and the sulphonylurea receptor SUR2B plays important roles. It is also likely that there is molecular diversity among K_{ATP} channels of smooth muscle, with the possible involvement of heterotetramers of pore-forming subunits. In addition to metabolic regulation, vascular K_{ATP} channels are regulated by endogenous vasoactive agents by way of cellular signalling pathways. Protein kinase A has been shown both to cause tonic channel activation, and to be important in activation by several vasodilators. These actions require PKA localization by an A-kinase anchoring protein (AKAP). Vasoconstrictors cause channel inhibition by way of protein kinase C, and vascular K_{ATP} channels have also been shown to be inhibited by intracellular calcium, an effect mediated by protein phosphatase 2B. Physiologically, K_{ATP} channels appear to be involved in the maintenance of resting blood flow in a number of vascular beds, notably the coronary circulation, as well as in vasodilation and vascular hyporeactivity of circulatory shock. Clinically, the K_{ATP} opener nicorandil is increasingly used in angina, where it may show maximum potency in ischaemic tissue, while increased understanding of these channels may offer future therapeutic opportunities. *J Clin Basic Cardiol 2003; 6: 7–14.*

Key words: K_{ATP} channel, smooth muscle, coronary blood flow, blood flow, shock, exercise, AKAP, vasodilation, vasoconstriction

t is rather over a decade since vascular ATP-sensitive potassium (KATP) channels were first identified. Since then these channels have been the subject of extensive research, and much of this work has been reviewed quite recently [1–3]. KATP channels link membrane K⁺ permeability to cellular metabolism. They are inhibited by intracellular ATP, and activated by ADP and other nucleoside diphosphates. They also form targets for therapeutic drugs that act either as channel openers or blockers. In vascular smooth muscle, opening KATP channels causes membrane hyperpolarization, decreased intracellular calcium concentration, and vasorelaxation. In addition to metabolic sensitivity, the channels are opened by vasodilators, while many vasoconstrictors close them, and such modulation represents a major component of their physiological regulation. Functional work has shown that KATP channels in arterial smooth muscle cells provide a background potassium conductance important in the regulation of membrane potential and so arterial tone and blood flow in a number of vascular beds. KATP activation can contribute both to basal blood flow, and to changes in flow in response to metabolic changes, as in exercise. Under pathological conditions, excessive KATP channel activation may play a role in the catastrophic vasodilation and vascular hyporeactivity associated with shock. This review concentrates on some recent developments in understanding the molecular structure, regulation, and some of the functional significance of vascular KATP channels.

Molecular Structure

K_{ATP} channels are heteromultimers of potassium channel subunits of the Kir6 family and sulphonylurea receptors (SUR). The pore of the channel is formed by a tetramer of Kir6 subunits, which appear to associate with SURs with 1:1 stoichiometry so that the complete channel forms as an octamer [4]. The Kir6 subunit confers sensitivity to inhibition by ATP, conductance and rectification properties. SUR modulates the sensitivity to ATP inhibition, and is also responsible for the activating effects of nucleoside diphosphates (NDPs). As their name suggests, sulphonylurea drugs inhibit KATP channels by interacting with SUR, and drugs that open KATP channels also act through this subunit [5]. Two Kir6 genes have been found: Kir6.1 and Kir6.2, while two SUR genes are also known, SUR1 and SUR2. SUR2 has two main splice variants, encoding the receptors SUR2A and SUR2B, and certain other minor variants have been identified at the RNA level [4]. Interaction of the two types of subunit is required for trafficking of correctly assembled channels to the cell membrane [6]. Consistent with this, a C-terminal truncated version of Kir6.2 that lacks the endoplasmic reticulum retention sequence can form functional channels in the absence of SUR [7]. KATP channels of different molecular composition are expressed in different tissues, and their varied properties are significant both for their physiological functions and for selective pharmacology. Correlation of the properties, regulation and pharmacology of KATP channels with their molecular structure is currently the subject of intensive research.

There is good evidence that the K_{ATP} channel of the pancreatic β -cell comprises Kir6.2/SUR1, while that of the sarcolemma of cardiac muscle is Kir6.2/SUR2A [4, 5]. Thus coexpression of the cloned subunits in *Xenopus* oocytes or mammalian cell lines reproduces the major properties of the channel in native tissue, while knockout studies have also shown the predicted abolition of channel activity [8, 9]. However, the molecular nature of the K_{ATP} channel, or channels, of vascular smooth muscle has not yet been definitively determined, and is complicated by diversity in vascular K_{ATP} channels [1, 2].

 K_{ATP} channels of vascular smooth muscle show a different pharmacology with respect to K_{ATP} opening drugs from that of pancreatic or cardiac channels, being activated by pinacidil, levcromakalim, and diazoxide [3]. This pattern of sensitivity to KCOs is conferred on recombinant channels by the

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SUR2B subunit, suggesting that this subunit might form a common component of KATP channels in VSM. In non-vascular smooth muscle cells of colon, bladder, and urethra it has been suggested on the basis of reverse transcription polymerase chain reaction (RTPCR) and mRNA measurements that KATP channels comprise Kir6.2/SUR2B [10-12]. Currently, the best evidence for the molecular composition of a particular vascular smooth muscle KATP channel is that for the nucleoside diphosphate (NDP)-activated channel, sometimes called K_{NDP} [13, 14]. K_{NDP} is activated by ATP at low concentrations, but inhibited above about 1 mmol/l, is sensitive to activation by NDPs in the presence of the K_{ATP} channel opener pinacidil, and has a relatively small unitary conductance of 30–35 pS in symmetrical high [K⁺] [13–15]. Both Kir6.1 and SUR2B show widespread expression [16, 17]. When co-expressed in heterologous systems they form a channel with ATP and NDP sensitivity and single channel conductance comparable to that of native K_{NDP}. Thus it seems likely that K_{NDP} is formed by the combination of Kir6.1 and SUR2B [18]. Recent evidence suggests that this subunit combination may also underlie KATP channels of pulmonary arterial smooth muscle. Cultured human pulmonary arterial smooth muscle cells expressed a channel with a conductance in the appropriate range (28-29 pS), and RTPCR revealed transcripts for Kir6.1 and SUR2B, but not Kir6.2, SUR2A or SUR1 [19].

There is considerable evidence that KATP channels of vascular smooth muscle can differ in terms of their single channel conductance and sensitivity to regulation by ATP and other nucleotides (reviewed in [1, 2]). Recent work on cloned KATP channels has raised the possibility that some of this diversity might arise from the co-assembly of the poreforming subunits Kir6.1 and Kir6.2 to form heterotetramers [20, 21]. Channels formed by cloned Kir6.1 or Kir6.2 alone show conductances of 30-35 pS and 70-80 pS respectively, while Kir6.1-Kir6.2 dimers form channels with an intermediate conductance [16, 22, 23]. Co-expression of Kir6.1 and Kir6.2 together with SUR2B in HEK293 cells leads to channels with a range of conductances consistent with the formation of tetramers with 1, 2, 3, or 4 of either channel subunit [20]. Further evidence for Kir6.1-Kir6.2 co-assembly in heterologous expression systems has been provided by the use of dominant negative subunits in which the conserved GFG sequence is mutated so that the subunit cannot form a functional pore; such subunits can prevent pore formation by wild-type subunits in a tetramer. In HEK293 cells, dominant negative Kir6.1 subunits can suppress currents through channels formed by wild-type Kir6.2 (or Kir6.1) subunits expressed with either SUR2B or SUR2A, and dominant negative Kir6.2 has similar effects [20, 21]. In addition, coimmunoprecipitation studies have provided evidence that Kir6.1 and Kir6.2 interact at a biochemical level in HEK293 cells [20, 21]. Co-assembly of Kir6.1 and Kir6.2 may not be possible in all systems, since Seharaseyon et al. found no evidence for coassembly in either A549 cells or cardiac ventricular myocytes [24]. With respect to vascular smooth muscle, however, it is of interest that dominant negative Kir6.2 suppressed KATP currents in the aortic-derived cell line A10 [25], where the reported properties of the KATP channel expressed are suggestive of Kir6.1 [26], consistent with co-assembly being possible in these cells. Finally, studies in Kir6.2 knockout mice show that both pinacidil-induced KATP currents of aortic smooth muscle cells and pinacidil relaxations of aortic rings were unaffected in the absence of Kir6.2, suggesting that Kir6.1 alone forms the KATP channel pore in this tissue [9]. Further investigations of the molecular basis for K_{ATP} channel diversity in smooth muscle will form an important

area of future research, and it may be that such diversity may also offer potential for future selective pharmacology.

Regulation by Cellular Signalling Pathways

Arterial KATP channels are regulated by multiple mechanisms. In general, regulation is consistent with their metabolic sensitivity and a role in adjusting blood vessel diameter and so blood flow to tissue metabolic demand. Like KATP channels of other tissues, they can be inhibited by intracellular ATP, though their reported sensitivity varies in different smooth muscle preparations. The reported effects of ATP may be complicated by the role of MgATP in maintaining channel function, and by the effects of ATP levels on tonic channel activation by protein kinase A, discussed below. Since intracellular ATP levels usually change little except in relatively severe metabolic compromise, it is more likely that inhibition by ATP provides a background level of activity against which other regulatory mechanisms operate. Among these, activation by nucleoside diphosphates seems important for many smooth muscle KATP channels [14, 15], and may account for some of the metabolic sensitivity of such channels. The role of nucleotides in KATP channel regulation has been reviewed recently [1]. KATP channels of vascular smooth muscle are also regulated by a wide range of vasodilators and vasoconstrictors. It is likely that such regulation forms an important part of their control under physiological conditions, and such regulation is discussed below, with an emphasis on experiments where currents through KATP channels have been measured directly from vascular smooth muscle cells using the patch clamp technique.

Activation by Vasodilators

Several vasodilators have been shown to activate vascular KATP channels, including calcitonin gene-related peptide, adenosine, and b-adrenoceptor agonists [27-30]. There is good evidence that activation by these vasodilators can occur through the classical adenylyl cyclase-protein kinase A pathway [30-32] (see [1] for review). In addition, intracellular application of either cyclic AMP or the catalytic subunit of protein kinase A (PKA) has been shown to activate KATP channels [30-32]. Recent work, discussed below, suggests that in addition to activation in response to binding of vasodilators to their receptors, PKA exerts a tonic drive on KATP channel activity in the absence of receptor activation. The molecular mechanism by which PKA activates vascular KATP channels remains to be determined. The simplest hypothesis is that the channel is phosphorylated directly, and both Kir and SUR subunits have consensus sites for phosphorylation by PKA [4]. In Kir6.2, PKA phosphorylates T224 to cause channel activation [33]. Little is known of Kir6.1 or SUR2B in this respect, though the consensus PKA site is conserved in Kir6.1 at T234.

Tonic Channel Activation by PKA

Most experimental work using patch clamp to study vascular K_{ATP} channels has used a low intracellular ATP concentration (usually 0.1 mmol/l) to reduce channel inhibition by ATP and so increase K_{ATP} current. It has been shown recently, however, that low intracellular ATP obscures a tonic activation of the channels by PKA, which is revealed by increasing ATP towards physiological levels. Thus K_{ATP} current is reduced by block of PKA when intracellular ATP is 1 mmol/l, but not when it is 0.1 mmol/l [34]. Such tonic activation by PKA does not require the presence of receptor agonists. Further, block of phosphatases with calyculin A leads to small but significant increases in K_{ATP} current. Such tonic channel acti-

vation by PKA may be important for the tonic contribution of the channels to resting blood flow discussed below, and inhibition of tonic PKA activation may also provide a new pathway for vasoconstrictor inhibition of the channels.

KATP Channel Inhibition by Vasoconstrictors

Vasoconstrictors cause an increase in the intracellular Ca²⁺ concentration of vascular smooth muscle and so in contractile force, by increasing influx of extracellular Ca²⁺ and by releasing Ca²⁺ from intracellular stores [35–37]. Closure of K^+ channels will contribute to membrane depolarization, and therefore to increased Ca^{2+} entry through voltage-dependent Ca²⁺ channels [36]. Many studies have shown that vasoconstrictors can inhibit a variety of potassium channels [38], and a number of vasoconstrictors have been shown to inhibit KATP channels of vascular smooth muscle. These include angiotensin II, endothelin, vasopressin, serotonin, phenylephrine, neuropeptide Y and histamine [39-42], (see [1] for review). Several of these vasoconstrictors have been shown to inhibit KATP channels through protein kinase C (PKC) [41, 42], and in mesenteric artery such inhibition is entirely prevented by the peptide translocation inhibitor of PKC epsilon, suggesting that this is the isoform involved [34]. Constitutively active PKC has also been shown to inhibit nucleotide diphosphate-sensitive KATP channels when applied to inside-out membrane patches excised from rabbit or rat pulmonary vein [43]. As for PKA, the means by which PKC affects channel activity at the molecular level remains to be determined. As for PKA, one possibility is through phosphorylation of Kir6.1 itself. PKC has been shown to modulate the activity of cardiac KATP channels by phosphorylation of threonine 180 on Kir6.2 [44], and this consensus site is conserved at T190 in Kir6.1.

Initial studies that implicated PKC in the actions of vasoconstrictors showed that block of PKC abolished KATP channel inhibition by vasoconstrictors, suggesting that this was the sole pathway for their action [41, 42]. Recent work, however, has provided intriguing new possibilities for the mechanisms by which vasoconstrictors may inhibit KATP channels. The tonic KATP activation by PKA described above raises the possibility that vasoconstrictors might inhibit such activation. The earlier studies that showed entire dependence on PKC were done under conditions of low intracellular ATP, where tonic activation would have been absent, and so would not have revealed such a pathway. When intracellular ATP is higher, however, about half the KATP channel inhibition by angiotensin II can be attributed to reduction in tonic activation by PKA [34]. Whether this effect involves inhibition of adenvlyl cyclase or occurs elsewhere in the pathway is currently unknown. It will also be of great interest to determine whether inhibition of tonic PKA drive forms a widespread pathway for the action of vasoconstrictors.

Recent work in aortic smooth muscle cells has provided evidence that K_{ATP} channel activity can also be regulated by the intracellular Ca^{2+} concentration by way of the calciumsensitive protein phosphatase 2B (calcineurin) [45]. K_{ATP} currents were maximally activated when $[Ca^{2+}]_i$ was 10 nM, but were strongly inhibited at 300 nM. Inhibition of calcineurin (with cyclosporin A, FK-506, or calcineurin autoinhibitory peptide) increased K_{ATP} currents, even when Ca^{2+} was high. It is presently unclear how calcineurin exerts its action on the channel, but it is likely to regulate the degree of phosphorylation of a site that causes channel activation, either on the channel itself or on an accessory regulatory protein [45]. One possible such target is the site at which PKA activates the channel. These findings suggest another pathway that might contribute to vasoconstrictor inhibition of K_{ATP} channels: vasoconstrictor-induced increases in $[Ca^{2+}]_i$ may increase the activity of calcineurin to reduce channel activity.

Anchoring proteins and signalling complexes

The work discussed above has provided evidence that vascular KATP channels are influenced by multiple regulators, including protein kinases A and C and protein phosphatase 2B. The tonic activation of KATP channels by protein kinase A suggests an intimate spatial relationship between kinase and channel. In recent years subcellular targeting of PKA through association with A-kinase anchoring proteins, or AKAPs, has been shown to account for the specificity of PKA phosphorylation in a number of cellular pathways [46-48]. AKAPs have a conserved helical region that binds the regulatory (RII) subunits of the PKA holoenzyme, and an anchoring domain that tethers the PKA-AKAP complex close to its site of action. Membrane targeting of PKA by AKAPs has been shown to be involved in the regulation of several types of ion channel (eg [49-51]). We have recently shown that an AKAP is involved in PKA regulation of arterial KATP channels [52]. Ht31 peptide binds to the RII subunit of PKA with nanomolar affinity, thus competing for PKA with native AKAPs, and so disrupting PKA anchoring within cells [53]. Intracellular application of Ht31 peptide blocked KATP current activation either by CGRP, which activates adenylyl cyclase, or by cAMP itself [52]. In addition, it prevented tonic KATP channel activation by PKA. However, intracellularly applied PKA catalytic subunit, which should act independently of native PKA localisation, was still able to activate KATP channels. These findings provide strong evidence for a key role of an AKAP in PKA-KATP channel signalling.

AKAPS do not only bind PKA. The prototypic AKAP, AKAP79, can bind both PKC and PP2B in addition to PKA [54]. Thus AKAPs can assemble multi-enzyme complexes that act in concert to regulate the phosphorylation state of cellular substrates. Such AKAP complexes may themselves be part of large signalling complexes that may involve both upstream activators and downstream targets [55]. Since PKA, PKC and PP2B are all involved in K_{ATP} regulation, it is an attractive idea that all of these enzymes might be localised by



Figure 1. Schematic diagram of factors that regulate vascular K_{ATP} channels. Activation pathways are shown with solid arrows and inhibitory pathways with dotted arrows. The A-kinase anchoring protein (AKAP) may be important in localising key regulatory enzymes close to the channel to form a regulatory complex. Many vasodilators act through adenylyl cyclase (AC) and PKA to cause channel activation, while PKA also provides tonic channel activation. Vasoconstrictors acting through phospholipase C (PLC) inhibit the channel through PKC and might also do so by activation of PP2B or inhibition of AC. Whether these kinases and phosphatases act directly on the channel proteins as indicated remains to be determined, as do the detailed interactions between the regulatory pathways and channel into a signalling complex.

the same AKAP, and that their functions in regulating the K_{ATP} channel might be integrated by anchoring into a signalling complex. Figure 1 shows a hypothetical scheme for such a complex. The identity of the AKAP involved remains to be determined, as do possible scaffolding links between receptors for vasoactive agonists, AKAP, and K_{ATP} channel.

Physiological and Pathophysiological Roles

Opening arterial K_{ATP} channels causes membrane hyperpolarization, decreased intracellular [Ca²⁺], and vasorelaxation, and the mechanisms involved have been reviewed previously [1, 36]. The following sections consider functional roles played by K_{ATP} channels in circulatory responses under physiological and pathophysiological conditions.

Resting Arterial Tone and Blood Flow

The initial evidence that vascular KATP channels have the potential to contribute to vasodilation came from studies that preceded the identification of the channels. A series of pharmacological agents including nicorandil, cromakalim, and pinacidil proved to be effective vasorelaxants of systemic and coronary blood vessels [56-62]. These drugs caused membrane hyperpolarization of smooth muscle, increased ⁸⁶Rb or ⁴²K efflux, and were ineffective at relaxing vessels constricted with high extracellular [K⁺], consistent with their acting by opening K⁺ channels. They are now generally named K⁺ channel openers. Following the demonstration that KATP channels occurred in smooth muscle, it became clear that these openers exert their hyperpolarizing and vasorelaxant effects by activating these KATP channels (see [1] for review). More recently, pharmacological KATP channel openers have been shown to increase blood flow or decrease vascular tone in the systemic [63, 64], coronary [63, 65, 66], and pulmonary [64, 67, 68] circulations. These studies with KATP openers demonstrate the potential of KATP channel activation to contribute to vasodilation under physiological conditions.

Over the past ten years substantial evidence has accumulated that KATP channels contribute a tonic vasodilator component in the coronary and systemic circulations, so contributing to blood flow at rest. Much of this evidence has come from studies of the effects of KATP channel inhibition with the sulphonylurea blocker glibenclamide on tone, vascular resistance or blood flow. Thus glibenclamide has been shown to increase vascular resistance in mesenteric and renal vascular beds [69-71]. KATP channels have also been reported to influence resting vascular tone in the skeletal muscle bed in some preparations [70, 72–74], but not in others [64, 75–77]. In the pulmonary circulation, evidence for functional KATP channels comes from vasodilator responses to KATP channel agonists in a number of species [64, 67, 68, 78, 79], and glibenclamide has been reported to decrease pulmonary vascular conductance in newborn piglets [68]. However, KATP channel blockade did not affect pulmonary vascular conductance in rats, cats, dogs or adult pigs, suggesting that these channels make little contribution to the regulation of basal pulmonary vascular tone [67, 76, 79].

Much experimental work has concentrated on the contribution of K_{ATP} channels to basal coronary blood flow, and here there is good evidence that a substantial component of resting flow depends on K_{ATP} channel activity. In anaesthetised dogs, glibenclamide can increase coronary resistance by up to 67 % [80–83], while equivalent responses have been reported in the coronary circulation of isolated hearts from rabbits, rats and mice [84–86]. Experiments on chronically implanted awake dogs and pigs at rest are consistent with these findings: here glibenclamide also increased coronary vascular

resistance [66, 79]. Together, this work suggests that K_{ATP} channels in the coronary vasculature are tonically active under basal conditions, and contribute a degree of vasodilation that is important for resting blood flow.

Studies on Genetically Altered Mice

Two recent studies on mice in which genes encoding KATP channel subunits were disrupted have given important insights into both the role and composition of arterial KATP channels. Disruption of the gene encoding Kir6.1 abolished pinacidil-activated currents in aortic smooth muscle cells and also abolished vasodilator responses to pinacidil, suggesting that this subunit is an essential component of arterial KATP channels [87]. Kir6.1-null mice had a high rate of sudden cardiac death associated with spontaneous elevation of the ST segment in the electrocardiogram followed by atrioventricular node block, indicative of periods of cardiac ischaemia resulting from coronary arterial vasospasm. The vasoconstrictor ergometrine induced ST elevation followed by cardiac death in Kir6.1 mice, but not in wild-type animals [87]. Thus the Kir6.1-null animals have hypercontractile coronary arteries, consistent with a critical role for the KATP channel in regulating vascular tone and in protecting against vasospasm. Very similar effects are seen in mice with disruption of the gene encoding SUR2 [88]. SUR2-null mice also lack vascular KATP channels and show ST segment elevation, coronary vasospasm and sudden cardiac death. Focal narrowing of coronary arteries was also observed in these animals. In addition, SUR2-null mice have significantly elevated resting blood pressure [88]. Together, these genetic studies confirm the results of pharmacological studies in intact animals in indicating the importance of KATP channels for regulating vascular tone and so blood pressure and blood flow.

Vasodilator Responses to Metabolic Demand and Exercise In many vascular beds, blood flow is closely correlated to metabolic demand. Thus local vasodilator signals released by metabolically active tissue lead to adjustments of blood vessel diameter to provide appropriate blood flow. These signals may include falls in oxygen tension, falls in pH and release of local regulators such as adenosine and prostacyclins. KATP channel activation appears important in hypoxic vasodilation in the coronary circulation and several other vascular beds. Recent studies of reactive hyperaemia in both the guinea-pig heart and human forearm suggest that KATP channels play an important role in post-ischaemic vasodilation [89, 90]. The role of KATP channels in the cerebral circulation has been reviewed recently by Faraci and Heistad [91]. Here, though KATP channels are clearly present and functional in many vessels, a number of studies have provided little evidence for a role of KATP channels in resting tone. However, several endogenous substances, including EDHF, CGRP, prostacyclin, opioids, and adenosine, that hyperpolarize and relax cerebral vascular smooth muscle, appear to do so via activation of KATP channels. Similarly, several studies have suggested that KATP channels may contribute to cerebral vasodilation in response to hypoxia [92-97] and KATP channels may also contribute to cerebral vascular autoregulation [98].

The roles of K_{ATP} channels in metabolic vasodilation suggests that they might be involved in circulatory responses to the increased metabolic demands induced by exercise. There have been relatively few studies in this area, but work on exercising dogs has provided interesting results. K_{ATP} channel blockade with glibenclamide reduced resting coronary blood flow and the reactive increase in blood flow in response to ischaemia, but did not prevent exercise-induced vasodilation [66]. However, when adenosine receptors were inhibited,

blockade of KATP channels blunted exercise-induced coronary vasodilation [99], an effect that became very severe when nitric oxide (NO) synthase was inhibited as well [100]. Neither adenosine nor nitric oxide-dependent mechanisms appear obligatory for maintaining either resting coronary flow or the increase in coronary flow on exercise, since these parameters were unaffected by blockade of both pathways [100]. The findings suggest that KATP channels are important for maintaining coronary vasodilation during exercise under normal conditions, but that adenosine and NO can act to increase coronary blood flow in exercise when KATP channels are blocked [100]. Interestingly, hypertrophied hearts seem to have an increased dependence on KATP channel opening to increase coronary blood flow in exercise, since here glibenclamide alone blunted the exercise-induced increase in flow [101]. A recent study has extended these investigations to the systemic and pulmonary circulations in exercising pigs. This work showed that KATP channel activation contributed to vasodilation both at rest and during exercise in the systemic as well as coronary circulations, but was not essential for exercise-induced vasodilation. In contrast, pulmonary vascular conductance was unaffected by glibenclamide, either at rest or in exercise [79].

Shock

Circulatory shock is characterized by hypotension, low systemic resistance and inadequate tissue perfusion, and vascular hyporesponsiveness to vasoconstrictors. Activation of potassium channels in the peripheral circulation seems important in the refractory vasodilation of shock, since K⁺ channel blockers have been shown to reduce such vasodilation [2, 102]. In particular, there is considerable evidence that increased activity of K_{ATP} channels occurs in both endotoxic and haemorrhagic shock.

In 1992 Landry and Oliver provided the first evidence for a role of KATP channels in the vasodilation induced by shock, showing that glibenclamide caused vasoconstriction and restored blood pressure in dogs with endotoxic hypotension [103]. Since then studies in rat and pig models in which shock is induced with bacterial endotoxins (lipopolysaccharides) have also shown at least partial restoration of arterial pressure by glibenclamide [104-108]. In the conscious rat, the major effect of glibenclamide on haemodynamics appears to be in the hindquarters vascular bed, where vascular resistance is increased and flow reduced [107]. Glibenclamide also abolished endotoxin-induced hyporesponsiveness to phenylephrine in rats [108]. Additional evidence for involvement of KATP channels comes from experiments showing that vasorelaxations to the KATP channel openers cromakalim and pinacidil were enhanced in endotoxic shock, consistent with an increase in KATP channel activity under such conditions [108]. Aortic smooth muscle from endotoxaemic rats has recently been shown to have a hyperpolarized membrane potential compared to control animals, and this hyperpolarization was partially reversed by blockers of either KATP or large-conductance Ca²⁺-activated K⁺ channels [109].

Blockade of K_{ATP} channels has also been reported to be beneficial in haemmorhagic shock. In anaesthetized rats subjected to severe haemorrhagic shock, glibenclamide increased arterial pressure [110, 111] and also improved survival rate [110]. In similar experiments using pigs, glibenclamide caused a sustained recovery of blood pressure, improved a number of haemodynamic variables, and restored tissue perfusion and metabolism to pre-shock levels [112]. In rats, glibenclamide has also been reported to reverse shock-induced arteriolar smooth muscle hyperpolarization and to improve intracellular Ca²⁺ responses and vascular hyporesponsiveness to noradrenaline [113], but not to affect hyporesponsiveness to angiotensin II [111].

Together, these findings provide good evidence that the activity of KATP channels is increased in circulatory shock, but the mechanism by which this occurs is still unclear. There is substantial evidence that increased synthesis of NO contributes to the vascular hypotension and hyporeactivity of shock [102, 114]. One possibility is that this leads to KATP channel activation via nitric oxide (NO) and cGMP. NO has been shown to cause glibenclamide-sensitive membrane potential hyperpolarization in rat and rabbit mesenteric arteries [115, 116], and may also play a role in hypoxic dilation of pial arteries [97, 117]. NO has also been reported to activate KATP channels in cultured smooth muscle cells [118]. However, studies in freshly isolated cells from rabbit mesenteric or pig coronary arteries failed to find KATP channel activation by NO or NO donors [30, 31]. In addition, glibenclamide did not inhibit NO donor-induced vasodilation in normal animals [103, 119]. A common theme in these studies has been the use of glibenclamide to define the involvement of KATP channels. Two recent brief reports suggest that KATP channels may have an altered pharmacology in endotoxaemia. Thus the compound PNU-37883A, which blocks KATP channels by interaction with the pore-forming subunit rather than the sulphonylurea receptor, inhibited both hyporeactivity to phenylephrine and relaxations to L-arginine in lipopolysaccharidepretreated rat mesenteric artery or aortic rings under conditions where glibenclamide was ineffective [120, 121]. The suggestion that endotoxaemia can reduce KATP channel sensitivity to glibenclamide, maybe through an action of NO, indicates that the possibility of KATP channel activation by NO requires further experimental study using pore-blocking agents.

CGRP levels are also increased in shock, and can contribute to shock-induced hypotension [122–124]. Since CGRP is a potent activator of K_{ATP} channels, it may also contribute to their activation in shock. A further possibility is that tissue metabolic changes might contribute to K_{ATP} channel activation, since lactic acidosis-induced hypotension in dogs could also be reversed by glibenclamide [103]. Finally, in addition to acute activation of K_{ATP} channels by one or all of these mechanisms, it is also possible that they affect gene regulation to upregulate K_{ATP} channel expression in shock.

Clinical Pharmacology

Both KATP channel opening and blocking drugs are used clinically, though vascular KATP channels are not always their intended target. Openers include the antihypertensives diazoxide and minoxidil sulphate, and the anti-anginal drug nicorandil which is currently in increasing use in Europe. In theory, at least, KATP channel openers might be expected to have advantageous properties in the treatment of ischaemic conditions like angina. The intrinsic metabolic sensitivity of vascular KATP channels should increase channel open probability in ischaemic tissue. If the action of KATP channel openers is synergistic with such intrinsic activation, such openers should have a self-targetting property, being more effective as vasodilators in ischaemic tissue. Studies on isolated coronary arteries suggest that this is the case for nicorandil [125]. Nicorandil also combines a nitrovasodilator action with its KATP channel opening properties, and this may contribute to a combination of rapid onset of action with lack of the development of the tolerance seen with classical nitrovasodilators [126]. Further interest in the drug has come from recent evidence that it is able to confer cardioprotection similar to that induced by ischaemic preconditioning [126], an effect that appears to be mediated by cardiac rather than vascular KATP channels.

 K_{ATP} channels of pancreatic beta cells form the therapeutic target for a widely-used family of drugs, the antidiabetic sulphonylureas. While such drugs show selectivity for the beta-cell form of the channel, they can block cardiac and vascular K_{ATP} channels at higher concentrations, and there has been a long-standing controversy as to whether this may contribute to possible adverse cardiovascular effects associated with sulphonylurea therapy for type 2 diabetes [127, 128].

Conclusions and Future Perspectives

We have accumulated a large amount of information about vascular KATP channels since they were discovered. However, important questions remain to be answered. Definitive information on molecular structure and the relationship between structure and the variations in KATP channel properties in smooth muscle will be an important goal. Achieving a full understanding of KATP channel regulation by cellular signalling pathways will be a major task, given the complexity that is already becoming apparent. It is likely that such an understanding will require information about the structural as well as functional relationships between the components of the signalling complexes involved. Functional studies in intact tissues and whole animals will also continue to be important in further defining the functional roles of vascular KATP channels. These should benefit from developing understanding of the channels and their regulation at a molecular and cellular level. Recent evidence suggests that the use of pore blocking drugs as well as KATP channel inhibitors will be important to help define these functions. As information about channel regulation becomes more precise, it is very likely that transgenic animals developed in the light of such knowledge will also play an important apart in elucidating channel function. Increased understanding of the regulation and functions of these channels should also have the potential to improve clinical practice in situations where they are involved, either to positive effect as in angina, or negatively as in shock. In the longer term, it is possible that new drugs designed to target aspects of KATP channel regulation may prove to be valuable therapeutic agents.

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