



COMMENTARY

Intracellular Diadenosine Polyphosphates

A NOVEL FAMILY OF INHIBITORY LIGANDS OF THE ATP-SENSITIVE K^+ CHANNEL

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ABSTRACT. Intracellular diadenosine polyphosphates (Ap_nA) are signal molecules that alert the cell under stress conditions. Herein, we review evidence that has recently identified a novel target for Ap_nA , namely the ATP-sensitive K^+ (K_{ATP}) channel. These channels are abundant in pancreatic β -cells and cardiac myocytes where they are essential in coupling the cellular metabolic state with membrane excitability. The potency and efficacy of Ap_nA to inhibit K_{ATP} channel activity were first described in cardiac K_{ATP} channels, and appear similar to those of intracellular ATP, the primary ligand of K_{ATP} channels. Also, the inhibitory ligand action of Ap_nA is dependent upon the operative condition of K_{ATP} channels and the presence of nucleotide diphosphates. In addition to a direct antagonism of channel opening, an indirect effect of Ap_nA on K_{ATP} channel activity has been associated with inhibition of adenylate kinase, a catalytic system believed essential for the regulation of channel opening. At present, the precise role for Ap_nA -induced K_{ATP} channel inhibition remains to be established. Yet, it is known that, under glucose challenge of pancreatic β -cells, intracellular concentrations of Ap_nA do increase to micromolar levels necessary to block K_{ATP} channels, leading to insulin secretion. Thus, the Ap_nA -mediated K_{ATP} channel gating represents a previously unrecognized pathway of channel regulation. *BIOCHEM PHARMACOL* 54;2:219–225, 1997. © 1997 Elsevier Science Inc.

KEYWORDS. adenylate kinase; alarmone; dinucleotide polyphosphates; glucose; K_{ATP} channels; stress

Intracellular Ap_nA^\dagger (where $n = 2-6$; Fig. 1) have emerged as important intracellular signal molecules in cells under stress [1, 2]. As Ap_nA are synthesized during metabolic challenge and can act homeostatically under stress conditions, these molecules, particularly Ap_4A , have been termed putative “alarmones” [3]. Specifically, during oxidative and heat shock stress, cytosolic Ap_4A rapidly rises in concentration [4, 5] and binds to several heat shock and oxidative stress proteins to inhibit their activities [6]. In several cell types, a direct intracellular effect of Ap_nA has been demonstrated on enzymes with nucleotide-binding domains, which are associated with cellular metabolism [1, 2].

Recently, a novel target for Ap_nA has been uncovered—the K_{ATP} channel [7]. Thus far, an inhibitory effect of Ap_nA has been determined on cardiac [8, 9] and pancreatic K_{ATP} channels [10], where these signaling molecules may serve a novel role in the regulation of K_{ATP} channel behavior.

It is well established that K_{ATP} channels are present at high density in cardiac myocytes and pancreatic β -cells where they link membrane excitability with the cellular bioenergetic state [11–15]. A defining attribute of K_{ATP}

channels is their regulation by intracellular mononucleotide polyphosphates, primarily ATP [13, 16, 17]. In pancreatic β -cells, K_{ATP} channels mediate glucose-induced insulin secretion. When the glucose concentration in serum is high, intracellular ATP inhibits the opening of K_{ATP} channels, which in turn depolarizes the cellular membrane, and promotes Ca^{2+} influx, triggering insulin release [18, 19]. In the heart, K_{ATP} channels have been implicated in the shortening of the action potential duration and in cellular loss of K^+ that occurs during metabolic stress and hypoxia [20, 21]. During early ischemia, opening of K_{ATP} channels may promote cardiac arrhythmias [22, 23], although opening of K_{ATP} channels has also been implicated in the cardioprotective mechanism of ischemic preconditioning [24, 25].

This synopsis summarizes the actions of Ap_nA on K_{ATP} channels, as these intracellular molecules now emerge as a novel regulator family of channel gating.

AP_nA : DIRECT INHIBITORS OF K_{ATP} CHANNEL OPENING

Ligand Effect of Ap_nA on K_{ATP} Channels in Excised Membrane Patches

That Ap_nA act as inhibitory ligands of K_{ATP} channels was first described for Ap_6A in membrane patches excised from cardiac cells [7] (Fig. 2). Further investigation, using the

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† Abbreviations: Ap_nA , diadenosine polyphosphates; Ap_4A , diadenosine tetraphosphate; Ap_6A , diadenosine hexaphosphate; Ap_3A , diadenosine triphosphate; Ap_5A , diadenosine pentaphosphate; and K_{ATP} , ATP-sensitive K^+ .

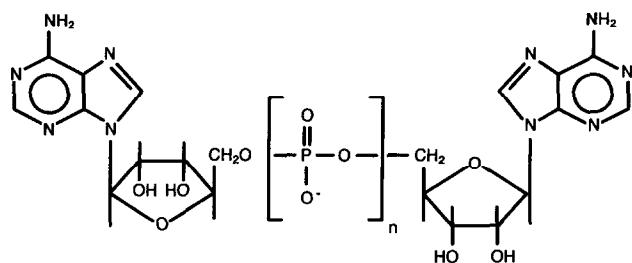


FIG. 1. General structure of Ap_nA , which are dinucleotides composed of two adenosine moieties linked through ribose 5'-carbons to the phosphate group chain. The number of phosphate groups (n) commonly varies from 2 to 6.

inside-out configuration of the patch-clamp technique, indicated that, in addition to Ap_6A , micromolar concentrations of other Ap_nA , such as Ap_4A , as well as Ap_3A and Ap_5A inhibit K_{ATP} channel opening in a reversible manner [8–10, 26] (Fig. 2 and Table 1).

The idea that led to the discovery of Ap_nA -mediated inhibition of K_{ATP} channels was based on the previous observation that these molecules inhibit the activity of certain cytosolic nucleotide-binding enzymes through interaction with the intracellular ATP and/or AMP binding sites of targeted proteins [2, 27]. As the K_{ATP} channel structure also possesses nucleotide-binding domains [28–32], it was conceivable that Ap_nA could affect the behavior of K_{ATP} channels. Indeed, application of Ap_nA to the intracellular side of myocardial and pancreatic β -cell membrane patches decreased the probability of K_{ATP} channel opening [7–10, 26].

The site of action of Ap_nA is most likely intracellular, as

TABLE 1. Summary of the IC_{50} and Hill coefficient values defining the concentration-dependent inhibition of cardiac and pancreatic K_{ATP} channels by Ap_nA *

	Cardiac		Pancreatic	
	IC_{50} (μM)	Hill coefficient	IC_{50} (μM)	Hill coefficient
Ap_3A	ND†	ND	74	ND
Ap_4A	17	1.2	17	1.2
Ap_5A	16	1.6	ND	ND
Ap_6A	14	1.1	ND	ND

*Data from Refs. 7–10.

†ND = not determined.

these molecules are poorly membrane-permeable [2]. In addition, the effect of Ap_nA occurs in the absence of GTP [7–10], eliminating the possibility of the involvement of a GTP-binding protein in the transduction of the inhibitory effect of Ap_nA on K_{ATP} channels. The concentration-dependence and saturable nature of the action of Ap_nA on K_{ATP} channel opening (Fig. 3) suggest the existence of specific binding sites for Ap_nA within K_{ATP} channel subunits or a closely associated protein.

Regardless of the length of the phosphate chain, the potency of various Ap_nA , e.g. Ap_4A , Ap_5A , and Ap_6A , to inhibit K_{ATP} channels is within a close micromolar range (Table 1). This is in contrast with a variable affinity displayed by members of the Ap_nA family in binding to other nucleotide-binding proteins, such as adenosine kinase [33], adenylate kinase [27, 34, 35], terminal deoxynucleotidyl transferase and carbamyl phosphate synthase [36]. For

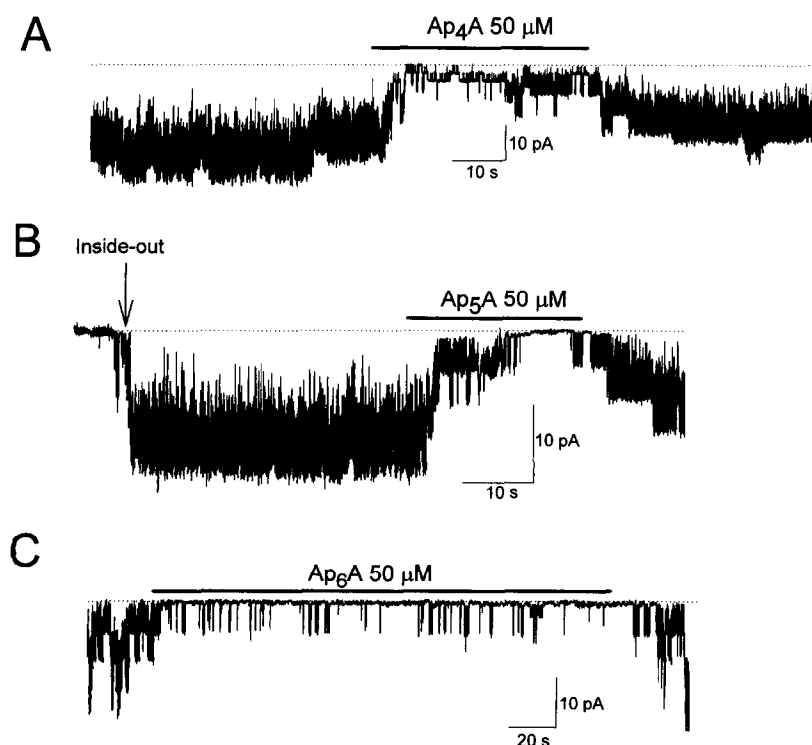


FIG. 2. Diadenosine tetra- (A), penta- (B), and hexaphosphate (C) (Ap_nA , $n = 4-6$)-induced inhibition of myocardial K_{ATP} channels. The patch-clamp technique was employed in the inside-out configuration. Holding potential: -60 mV. Modified from Refs. 7–9.

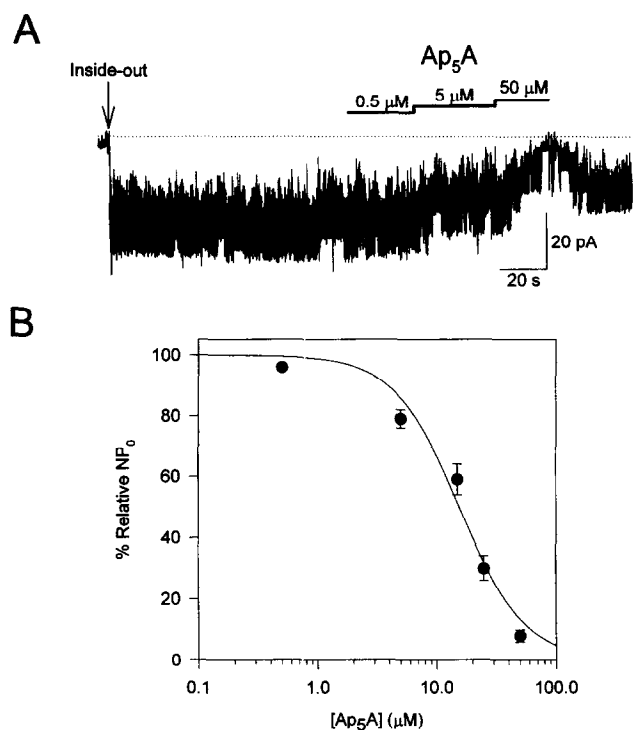


FIG. 3. Concentration-dependent effect of Ap_5A -induced inhibition of myocardial K_{ATP} channels. The patch-clamp technique was employed in the inside-out configuration. Holding potential: -60 mV. (A) Original record. (B) Concentration-response curve fitted by the Hill equation. Channel activity is expressed as NP_o , where N represents the number of channels and P_o the probability for channels to be open. Values, expressed as means \pm SEM, are from 3–6 membrane patches for each data point. Slightly modified from Ref. 9. Reprinted with permission from Naunyn-Schmiedeberg's Arch Pharmacol 353: 241–244, 1996. Copyright (1996) Springer-Verlag GmbH & Co. KG.

example, Ap_4A inhibits the activity of adenosine kinase [33], a property not shared by other members of the Ap_nA family. Despite their different selectivity toward other nucleotide-binding proteins and their different affinities for the ATP-binding site located on these proteins, it appears that the property of Ap_nA to inhibit K_{ATP} channels is common to all Ap_nA so tested.

Operative Condition-Dependent Regulation of K_{ATP} Channels by Ap_nA

K_{ATP} channels can function under two operative conditions: operative condition 1, when K_{ATP} channel activity is sustained spontaneously and not enhanced by nucleotide diphosphates, versus operative condition 2, when after the decline of spontaneous channel activity ("run-down") nucleotide diphosphates restore channel activity [15, 17, 37]. The effect of Ap_4A and Ap_5A on K_{ATP} channels under different operative conditions was tested in membrane patches excised from ventricular cardiac cells [26]. Under operative condition 1, application of nucleotide diphosphates prevented the Ap_4A - and Ap_5A -induced inhibition of spontaneous K_{ATP} channel activity (Fig. 4). In contrast, under operative condition 2, the nucleotide diphosphate-induced K_{ATP} channel openings were inhibited by Ap_4A (Fig. 4) or Ap_5A . Thus, the outcome of the interaction between an Ap_nA and nucleotide diphosphates on K_{ATP} channel opening is not constant but changes depending on the operative condition of the channel. The mechanism responsible for the switch in the responsive behavior of K_{ATP} channels toward a ligand is unknown. Phosphorylation of the channel protein has been suggested as a possible candidate [26, 37], as it is for the mechanism underlying the switch between operative conditions 1 and 2 [15, 16]. The K_{ATP} channel protein complex possesses several nucleotide binding and phosphorylation sites [29, 30] that could be involved in regulating the interaction of Ap_nA with K_{ATP} channels.

Comparison of the Effects of ATP and Ap_nA on K_{ATP} Channels

ATP and Ap_nA inhibit the opening of K_{ATP} channels with a similar micromolar potency (Table 2). The relationship between the concentration of ATP or Ap_nA on channel activity could be fitted by a Hill equation with similar values for the apparent Hill coefficient (Table 2). However, the precise number of molecules required to close the channel remains to be determined. Neither ATP nor Ap_nA affect the amplitude of the single channel current, but they decrease the probability

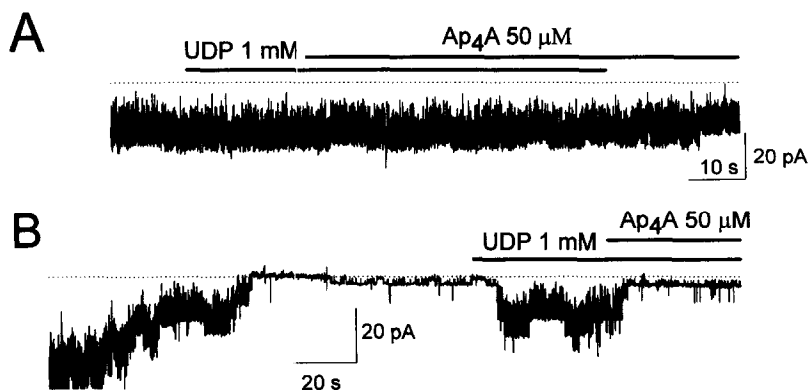


FIG. 4. Operative condition-dependent responsiveness of myocardial K_{ATP} channels to Ap_4A . The patch-clamp technique was employed in the inside-out configuration. Holding potential: -60 mV. (A) Operative condition 1 (spontaneous channel opening). (B) Operative condition 2 (nucleotide diphosphate-UDP-induced channel opening after "run-down"). Under operative condition 1, Ap_4A did not inhibit K_{ATP} channel activity in the presence of UDP. Under operative condition 2, Ap_4A did inhibit K_{ATP} channel activity in the presence of UDP. Modified from Ref. 26.

TABLE 2. Comparison of the effects of ATP and $A_{p_n}A$ ($n = 4-6$) on myocardial K_{ATP} channels*

	Channel inhibition			Channel maintenance
	IC_{50} (μM)	Hill coefficient	Operative condition-dependent	
ATP	24–30	1.7–1.8	Yes (with UDP)	Yes (with Mg^{2+})
$A_{p_n}A$	14–17	1.1–1.6	Yes (with UDP)	No

*Data from Refs. 7–9, 26, and 37–40.

of channel opening, most likely by apparently reducing the time the channel spends in a burst [8–10, 13].

As described above for $A_{p_n}A$, the outcome of interaction between ATP and K_{ATP} channels is governed by the operative condition of the channel and the presence of nucleotide diphosphates [15, 37]. Thus, the interaction of K_{ATP} channels with both $A_{p_n}A$ and ATP provides further evidence that the basal state of the effector system governs the outcome of a stimulus [37, 41, 42].

In contrast to ATP, which serves not only as a ligand to close the K_{ATP} channel but also maintains channel activity [12–16], $A_{p_n}A$ did not promote sustained channel activity. The maintenance of channel activity by ATP can be observed after removal of this nucleotide, and is the exclusive property of the Mg^{2+} bound form of ATP ($MgATP$), and is not shared by non-hydrolyzable analogs of ATP. Since both Mg^{2+} and ATP hydrolysis are necessary for kinase activity, maintenance of K_{ATP} channel activity may relate to an enzyme-dependent action of ATP, such as phosphorylation. When K_{ATP} channels are in “run-down,” treatment of membrane patches with $MgATP$ reactivates spontaneous channel activity [13, 15, 16]. In contrast, $A_{p_n}A$ do not reverse channel activity after “run-down” [8–10]. Thus, $A_{p_n}A$ possess a “pure” inhibitory property on K_{ATP} channel opening, without having the ability to maintain channel opening. In this regard, the action of $A_{p_n}A$ on K_{ATP} channels mostly resembles that observed with non-hydrolyzable analogs of ATP, such as $ATP\gamma S$.

$A_{p_n}A$: INDIRECT REGULATORS OF K_{ATP} CHANNEL BEHAVIOR?

The possibility that certain $A_{p_n}A$ could regulate K_{ATP} channel activity through an indirect mechanism has been indicated recently in cardiomyocytes [43]. As has been already mentioned, the defining property of K_{ATP} channels is their inhibition by ATP, as is the effect of ADP to reverse the ATP-inhibited state [12, 13, 15–17]. However, the mechanism by which opening of K_{ATP} channels is governed *in situ* remains undetermined, and the conventional assumption that a fall in the cytosolic concentration of ATP determines channel opening has been contested [20]. The cytosolic concentration of ATP remains, even under metabolic stress, over 100-fold higher than the value required for K_{ATP} channel closure. Therefore, additional mecha-

nisms must allow K_{ATP} channels to open in the presence of millimolar concentrations of cytosolic ATP. Recent studies suggest that the microenvironment surrounding K_{ATP} channels may regulate the ATP-dependent gating of these channels [38, 44]. Furthermore, evidence has been presented that transition from the closed (ATP-liganded) to the open (ADP-liganded) state of K_{ATP} channels is possible through catalytic transformation of ATP into ADP within the microenvironment surrounding K_{ATP} channels [43]. Such transition could be mediated through adenylate kinase activity, known to catalyze the transformation of ATP into ADP [43, 45–47]. Thus, adenylate kinase emerges as an important determinant of the composition of adenine nucleotide species at the K_{ATP} channel site. Of importance, $A_{p_n}A$, in particular $A_{p_5}A$, are potent blockers of adenylate kinase activity [27]. Indeed, $A_{p_5}A$ inhibited the adenylate kinase-mediated transition of K_{ATP} channels from the closed to the open state [43] (Fig. 5). Such an effect of $A_{p_5}A$ was observed at concentrations that were not sufficient to directly inhibit K_{ATP} channel activity, in accord with the high affinity of adenylate kinase for $A_{p_5}A$ [27, 43]. Therefore, it is apparent that besides a direct action on K_{ATP} channels, $A_{p_5}A$ may modulate the opening behavior of these channels through an indirect mechanism, i.e. through inhibition of adenylate kinase-dependent channel gating. At present, such indirect regulation of K_{ATP} channel behavior has been reported only for $A_{p_5}A$.

ROLE FOR $A_{p_n}A$ IN REGULATING K_{ATP} CHANNEL BEHAVIOR

Under resting conditions, the intracellular concentration of $A_{p_n}A$ ($<1 \mu M$) is at least 10,000-fold lower than the concentration of ATP [1, 2]. However, it is established that after cell exposure to various oxidants or metabolic stress, cytosolic concentrations of $A_{p_n}A$ could rise to micromolar levels [1, 2]. Furthermore, as described above, $A_{p_n}A$ may act as alarmones to alert the cell to the onset of metabolic stress [3]. In pancreatic β -cells, evidence has been obtained that the $A_{p_n}A$ -mediated regulation of K_{ATP} channels may be of importance in glucose-triggered insulin secretion [10]. It was shown that a glucose challenge of a pancreatic β -cell induces an important increase of the cytosolic concentra-

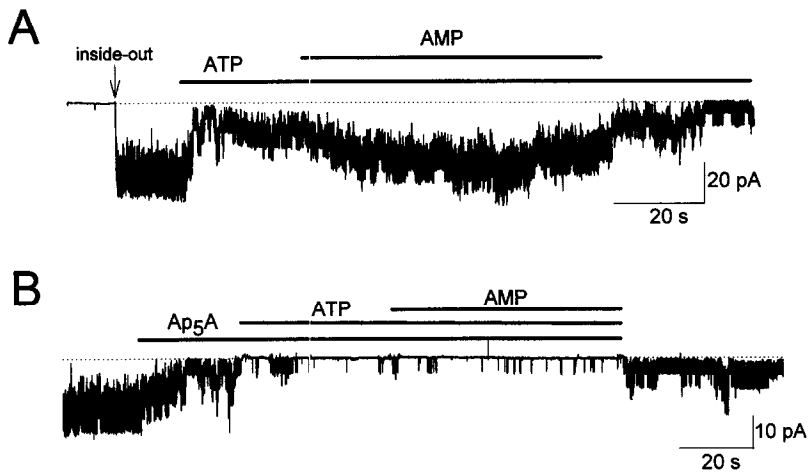


FIG. 5. Prevention by Ap_5A of the opening of myocardial K_{ATP} channels induced by substrates (AMP and ATP) of adenylate kinase. The patch-clamp technique was employed in the inside-out configuration. Holding potential: -60 mV. (A) Adenylate kinase-mediated activation of ATP-inhibited channels by AMP. (B) Prevention by Ap_5A of the opening of the K_{ATP} channel by the two substrates of adenylate kinase (ATP and AMP). Modified from Ref. 40.

tion of Ap_nA to micromolar concentrations sufficient to inhibit K_{ATP} channel opening [10]. By analogy to pancreatic β -cells, it is conceivable that under stress conditions the concentration of Ap_nA may also rise in other tissues sufficient to gate K_{ATP} channel opening. This would be of importance due to the critical cellular functions with which K_{ATP} channels have been associated, including vasodilation, skeletal muscle contraction, and neuroprotection [12–17, 48–51]. In particular, a role for Ap_nA in the regulation of K_{ATP} channels in the heart may be significant since the myocardium is under constant metabolic stress. Activation of K_{ATP} channels does appear to be protective against myocardial infarction [24, 25, 52], while it has also been reported that opening of K_{ATP} channels is associated with cellular loss of K^+ during ischemia and hypoxia with possible proarrhythmic consequences [20, 23].

An important question to be resolved is why the K_{ATP} channel system would rely on two inhibitory signaling pathways, the conventional ATP-dependent and the now described Ap_nA -dependent inhibition. It is tempting to speculate that under resting conditions in cardiac cells, for example, K_{ATP} channels would be closed by high cytosolic levels of ATP, while the concentration of Ap_nA would be too low to affect channel activity. Under metabolic stress with the decrease in the cytosolic concentration of ATP, as well as an associated adenylate kinase-mediated opening of K_{ATP} channels [43], intracellular levels of Ap_nA may rise to directly impede channel opening and/or block adenylate kinase-mediated channel opening. Thus, the role of Ap_nA may lie in counteracting the consequence of decreased intracellular ATP concentration under metabolic stress.

CONCLUDING REMARKS

Intracellular Ap_nA are signal molecules proposed to act homeostatically under stress conditions. Herein, we present evidence identifying Ap_nA as inhibitors of K_{ATP} channels, an important ion conductance that couples the metabolic state of a cell with membrane excitability.

The overall inhibitory profile of Ap_nA appears similar to that of intracellular ATP, the primary ligand of K_{ATP} channels. However, the dual ability of ATP to both block as well as maintain K_{ATP} channel activity is not shared by Ap_nA . Thus, the inhibitory activity of Ap_nA mostly resembles that reported for non-hydrolyzable analogs of ATP.

At present, the mechanism through which Ap_nA inhibit K_{ATP} channels is unknown. Two possible mechanisms for Ap_nA -induced channel inhibition have been proposed: (1) a direct ligand inhibitory effect, probably mediated through binding to a specific binding site on a K_{ATP} channel subunit or on an associated protein, and/or (2) an indirect effect mediated through inhibition of adenylate kinase-mediated channel regulation. A particular feature of the direct inhibitory action of Ap_nA is its dependence upon the operative condition of K_{ATP} channels and the presence of nucleotide diphosphates. The indirect effect through inhibition of adenylate kinase may occur at a slightly lower concentration of Ap_nA than the direct ligand inhibitory effect.

The precise role related to the Ap_nA -induced K_{ATP} channel inhibition remains to be determined. It is known, however, that the intracellular concentration of Ap_nA does increase to levels sufficient to gate K_{ATP} channels under cellular challenges. Of particular importance is the ability of glucose to induce a rise in the cytosolic levels of Ap_nA within pancreatic β -cells, which in turn may play an important role in the K_{ATP} channel-dependent regulation of insulin secretion. Other roles for the Ap_nA -dependent K_{ATP} channel gating await to be identified.

In summary, this synopsis reviews the evidence for a previously unrecognized property of Ap_nA , as well as the discovery of a novel family of endogenous ligands gating K_{ATP} channel behavior.

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