

Pharmacological characterization of the sulphonylurea receptor in rat isolated aorta

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- 1 The binding of the sulphonylurea [3H]-glibenclamide, a blocker of adenosine 5'-triphosphate (ATP)sensitive K⁺ channels (K_{ATP} channels), was studied in endothelium-denuded rings from rat aorta.
- 2 [3 H]-glibenclamide labelled two classes of binding sites with $K_{\rm D}$ values of 20 ± 5 nM and 32 ± 1 μ M. The high affinity component, which comprised 17% of total binding at 1 nM [3H]-glibenclamide, had an estimated binding capacity of 150 fmol mg⁻¹ wet weight.
- 3 Other sulphonylureas such as glipizide and glibornuride and the sulphonylurea-related carboxylate, AZ-DF 265, inhibited high affinity [3H]-glibenclamide binding with the potencies expected from their K channel activity. At very high concentrations, AZ-DF 265 and glipizide started to interact also with the low affinity component of [3H]-glibenclamide binding.
- 4 Openers of the ATP-sensitive K+ channel belonging to different structural groups inhibited only the high affinity [3H]-glibenclamide binding; the potencies in this assay were similar to those obtained in functional (i.e. vasorelaxation) studies.
- 5 High affinity [3H]-glibenclamide binding was abolished by prolonged hypoxia combined with metabolic inhibition.
- 6 The data indicate that the high affinity component of [3H]-glibenclamide binding mediates the block of the K_{ATP} channel by the sulphonylureas in rat aorta; hence, it represents the sulphonylurea receptor in this vessel. The pharmacological properties of this binding site resemble those of the binding site for the openers of the K_{ATP} channel; present evidence suggests that these two classes of sites are negatively allosterically coupled.

Keywords: Glibenclamide binding; sulphonylurea receptor; KATP channel; KATP channel openers; [3H]-P1075; metabolic dependence of binding; rat aorta

Introduction

Adenosine 5'-triphosphate (ATP)-sensitive K⁺ channels (K_{ATP} channels) are a heterogeneous group of K+ channels that couple membrane potential to the metabolic state of the cell as given by the intracellular concentrations of ATP and adenosine 5'-diphosphate (ADP) (Ashcroft & Ashcroft, 1990; Edwards & Weston, 1993). In addition to their regulation by nucleotides, these channels are blocked by sulphonylureas like glibenclamide and tolbutamide and activated by the KATP channel openers, a structurally heterogeneous group of compounds (Ashcroft & Ashcroft, 1990; Quast, 1992; Edwards & Weston, 1993). Recent studies have shown that the K_{ATP} channels in pancreatic B-cells (Inagaki et al., 1995; Sakura et al., 1995) in heart and skeletal muscle (Inagaki et al., 1996) and the principal cells of the renal cortical collecting duct (McNicholas et al., 1996) are heteromeric proteins composed of αand β -subunits. The structure of the α -subunits is typical of inwardly rectifying K^+ channels; the β -subunits are members of the ATP-binding cassette protein superfamily and confer on the native channel the sensitivity to sulphonylureas and, perhaps with the exception of the kidney channel, also to nucleotides (Inagaki et al., 1995; 1996; Sakura et al., 1995; McNicholas et al., 1996). The molecular identity of the components making up the KATP channel in (vascular) smooth muscle is presently unknown.

One way of characterizing the components of the K_{ATP} channel is to use radiolabelled blockers, such as [3H]-glibenclamide, or openers, such as [3H]-P1075, a compound structurally related to pinacidil (Manley et al., 1993). In smooth muscle, only few such studies have been performed (Gopalakrishnan et al., 1991; Zini et al., 1991; Kovacs & Nelson, 1991; Bray & Quast, 1992; Quast et al., 1993; for re-

view see Ashcroft & Ashcroft, 1992). In membrane fragments prepared from guinea-pig ileum (Gopalakrishnan et al., 1991; Zini et al., 1991) including the myenteric plexus (Zini et al., 1991), [³H]-glibenclamide labelled two classes of binding sites with K_D values of 0.06 nm (Gopalakrishnan et al., 1991) and 0.4 nm (Zini et al., 1991) for the high affinity sites, and 2 nm (Gopalakrishnan et al., 1991) and 83 nm (Zini et al., 1991) for the low affinity sites, respectively. The binding capacities of the two components ranged from 20 to 3000 fmol mg⁻¹ protein (Gopalakrishnan et al., 1991; Zini et al., 1991). The potencies of different sulphonylureas in inhibiting the low affinity component of [3H]-glibenclamide binding showed some correlation with the potencies of the sulphonylureas in inhibiting openerinduced relaxation of guinea-pig small intestine; however, the openers did not inhibit [³H]-glibenclamide binding (Zini et al., 1991). In membranes from canine aorta, a single class of [³H]glibenclamide binding sites was found with $K_D = 10 \text{ nM}$. However, the pharmacological properties of this binding site were not characterized further (Kovacs & Nelson, 1991).

Binding studies with the opener [3H]-P1075 in rings of rat aorta have shown the existence of a high affinity site (Bray & Quast, 1992; Quast et al., 1993) which was abolished by metabolic inhibition of the preparation (Quast et al., 1993). [3H]-P1075 binding was inhibited by other openers with the affinity expected from functional (vasorelaxation) assays. Sulphonylureas also inhibited [3H]-P1075 binding with the correct rank order of potencies. However, the affinities determined in the binding assay were 10-20 times lower than those calculated from the inhibition of K_{ATP} channel opening (Quast et al., 1993); in these experiments, opener-induced $^{86}Rb^+$ efflux was used as a qualitative measure of channel opening (reviews Edwards & Weston, 1994; Quast, 1996). Kinetic experiments suggested a negative allosteric coupling between the sites for openers and sulphonylureas (Bray & Quast, 1992).

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In view of this coupling between the sites for openers and sulphonylureas found in the [³H]-P1075 assay (Bray & Quast, 1992; Quast *et al.*, 1993) we have now investigated the binding of [³H]-glibenclamide in metabolically intact rings of rat aorta. It was hoped that under these conditions a functionally relevant binding site for the sulphonylureas in vascular smooth muscle could be detected.

Methods

[3H]-glibenclamide binding assay in rat aortic strips

The [³H]-glibenclamide binding experiments were performed essentially according to the protocol developed for [³H]-P1075 binding (Bray & Quast, 1992; Quast *et al.*, 1993). Briefly, the thoracic aorta was removed from male Sprague-Dawley rats (350–500 g) and immersed in warm HEPES-buffered PSS containing (in mM): NaCl 139, KCl 5, CaCl₂ 2.5, MgCl₂ 1.2, glucose 11 and HEPES 5; the buffer was gassed with 95% O₂/5% CO₂ and titrated to pH 7.4 with NaOH at 37°C. Adherent fat and the adventitia were removed, the aorta was cut open, the endothelium removed by a moistened cotton bud and the organ was cut into strips of 1–2 mg wet weight.

The strips were incubated at 37°C for 90 min under light gassing with 95% $O_2/5\%$ O_2 in PSS containing [${}^3\text{H}$]-glibenclamide (1 nM) and the unlabelled test substance. They were then washed in ice-cold quench solution (154 mM NaCl, 50 mM Tris, pH 7.4) for 1 min, blotted, weighed and dissolved over 2 h in 0.3 ml of Soluene-350 at 37°C . The samples were supplemented with 50 μ l 1 M HCl and 3 ml Ultima Gold and counted for ${}^3\text{H}$. Under these conditions, B_{tot} was 42 ± 2 fmol mg $^{-1}$ wet weight and nonspecific binding, determined in the presence of unlabelled glibenclamide (300 μ M), was $14\pm3\%$ of B_{tot} .

[³H]-glibenclamide bound in the presence of I (unlabelled inhibitor) was fitted to a Hill equation taking into account two binding components:

$$\begin{split} B = B_{tot} - A_1 * I^{n_{H1}} / (IC_{50,1}^{n_{H1}} + I^{n_{H1}}) - \\ A_2 * I^{n_H} / (IC_{50,2}^{n_{H2}} + I^{n_{H2}}) \end{split} \tag{1}$$

Here B and B_{tot} denote the binding of [3 H]-glibenclamide in the presence and absence of I; A₁ and A₂ are the extents (amplitudes) of the two components, IC_{50.1} and IC_{50.2} the concentrations of I producing half maximum inhibition and n_{H1} and n_{H2} the Hill coefficients of the respective components. In these experiments, the concentration of [3 H]-glibenclamide was much smaller than $K_{D,1}$ and $K_{D,2}$ of glibenclamide binding; hence, the IC₅₀ values can be equated with the respective inhibition constants, K_{i} (Tallarida, 1995). The K_{ATP} channel openers interacted only with the first component and A₂ was set to 0.

In the case of homologous displacement and a Hill coefficient of 1, the binding capacity, $B_{\rm max}$ of a component may be estimated from the amplitude, A, by taking into account the degree of saturation, y:

$$B_{\text{max}} = A/y, \text{ with } y = L/(L + K_D)$$
 (2)

where L denotes the label concentration (1 nM) and K_D the equilibrium dissociation constant (approximated by the K_i value of the component).

Drugs and solutions

[3 H]-glibenclamide (specific activities 84 and 47 Ci mmol $^{-1}$) was purchased from Amersham Buchler (Braunschweig, Germany) and NEN (Bad Homburg, Germany), respectively; the unlabelled compound was from Sigma (Deisenhofen, Germany). The following drugs were kind gifts of the pharmaceutical companies given in parentheses: AZ-DF 265 (4-([N-(α -phenyl-2-piperidino-bentyl)carbamoyl]methyl)benzoic acid,

Thomae, Biberach, Germany); glibornuride (Hoffmann-La Roche, Basel, Switzerland); glipizide (N-cyano-N'-(1,1-dimethylpropyl)-N"-3-pyridylguanidine, Pfizer, Sandwich, U.K.); aprikalim (Rhône-Poulenc Rorer, Paris, France); diazoxide (Essex Pharma, München, Germany); levcromakalim (SmithKline-Beecham, Harlow, U.K.); P1075 (Leo Pharmaceuticals, Ballerup, Denmark); pinacidil and minoxidil sulphate were synthesized by Dr W.P. Manley (Sandoz, Basel, Switzerland). Oligomycin and 2-deoxyglucose were from Fluka (Neu-Ulm, Germany). Drugs were dissolved in ethanol and dimethyl sulphoxide (1:1) and further diluted into the same solvent or into PSS; the final solvent concentration in the assays was always below 0.3%.

Calculations and statistics

Results are expressed as mean \pm s.e.mean. Individual inhibition curves were fitted to equation (1) according to the method of least squares by use of the Figure P programme (Biosoft, Cambridge, U.K.). Errors in the fitting parameters were estimated by assuming that the amplitudes were normally distributed and IC₅₀ values log-normally distributed. In calculations involving two mean values with standard errors, propagation in errors was taken into account according to Bevington (Bevington, 1969). Statistical significance of differences was assessed by ANOVA followed by the Dunnet's or the Newman-Keuls test as *post hoc* tests by use of the Instat programme (Graphpad Software, San Diego, U.S.A.).

Results

Effect of glibenclamide on [3H]-glibenclamide binding

Figure 1 illustrates the inhibition of [³H]-glibenclamide binding (1 nM) in rat aortic rings by unlabelled glibenclamide and other sulphonylureas. The inhibition curve obtained with glibenclamide was biphasic with a high affinity component comprising $17\pm1\%$ of $B_{\rm tot}$, a $K_{\rm i,1}$ value of 20 ± 5 nM and a Hill coefficient of 1.0 ± 0.2 ; the low affinity component accounted for $69\pm2\%$ of $B_{\rm tot}$ with $K_{\rm i,2}=32\pm1~\mu{\rm M}$ and $n_{\rm H,2}=2.1\pm0.2$. Homologous displacement experiments were also performed at label concentrations varying from 0.3 nM to 5 nM (not shown). $B_{\rm tot}$ increased linearly with the [³H]-glibenclamide concentration but the relative amplitudes and $K_{\rm i}$ values of the two components were not changed significantly. This would be expected if the binding parameters derived from the data in

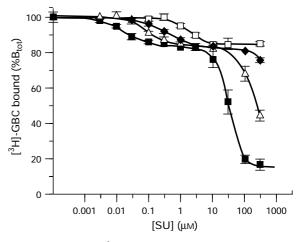


Figure 1 Inhibition of [3 H]-glibenclamide (GBC) binding (1 nM) in rat aortic rings by the sulphonylureas (SU) glibenclamide (\blacksquare), glipizide (\spadesuit), glibornuride (\square) and the sulphonylurea-related carboxylate, AZ-DF 265 (\triangle). Pooled data from 4−8 experiments for each compound are shown; individual inhibition curves were fitted to equation (1) yielding the fitting parameters given in the text and in Table 1. B_{tot} was $42\pm2\,\mathrm{fmol\,mg}^{-1}$ wet weight.

Figure 1 were correct since even at 5 nm [³H]-glibenclamide, the high affinity component would be saturated only to 20%.

Effect of other sulphonylureas on $[^3H]$ -glibenclamide binding

Other sulphonylureas such as glipizide (long chain sulphonylurea) and glibornuride (short sulphonylurea) as well as the sulphonylurea-related carboxylate AZ-DF 265 (Garrino & Henquin, 1988), inhibited high affinity [3H]-glibenclamide binding; at very high concentrations, AZ-DF 265 and glipizide also started to interact with the low affinity component (Figure 1). Analysis of the inhibition curves shows that the amplitudes of the high affinity component for these compounds are not different from that obtained with glibenclamide (Table 1); the corresponding pK_i (= $-\log K_i$) values and the Hill coefficients are listed in Table 1.

Openers interact with high affinity [3H]-glibenclamide binding

Representatives of the major structural families of the openers (Edwards & Weston, 1994) like P1075 and pinacidil (cyanoguanidines), levcromakalim (a benzopyran), aprikalim (a thioformamide), minoxidil sulphate (a pyrimidine) and diazoxide (a benzothiadiazine, chemically related to the thiazide diuretics) inhibited the high affinity component of glibenclamide binding completely but did not interfere with the low affinity component even at the highest concentrations used (Figure 2). Analysis of the curves yielded the pK_i values compiled in Table 1; in general, Hill coefficients were not different from 1 with the exception of aprikalim and diazoxide.

Modulation of [3H]-glibenclamide binding by metabolic inhibition

We have shown previously that under hypoxia and metabolic poisoning, binding of the opener [3H]-P1075 in rat aortic strips disappears concomitantly with the reduction in intracellular ATP levels (Quast et al., 1993). Severe metabolic blockade was produced by inhibition of glycolysis (substitution of glucose by deoxyglucose) combined with hypoxia (gassing with 95% N₂/ 5% CO₂) or with the inhibition of oxidative phosphorylation by oligomycin for 90 min. Figure 3 shows that under these conditions, total binding of [3H]-glibenclamide was reduced by 15-16%, i.e. by exactly the amount of high affinity binding.

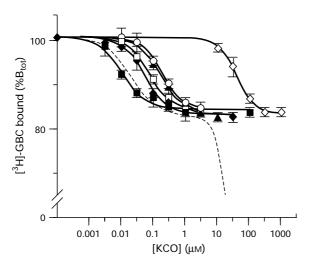


Figure 2 Inhibition of [³H]-glibenclamide (GBC) binding (1 nm) by K_{ATP} channel openers: (\blacksquare) P1075, (\spadesuit) minoxidil sulphate, (\square) pinacidil, (\triangle) levcromakalim, (\bigcirc) aprikalim and (\diamondsuit) diazoxide. Data are means and vertical lines s.e.mean from 4-6 experiments; individual data were fitted to equation (1) with $A_2 = 0$; for fitting parameters see Table 1. The dashed curve represents glibenclamide.

Addition of glibenclamide (1 µM) or P1075 (100 µM), concentrations which saturate the high affinity component, did not decrease B_{tot} further; in addition, non-specific binding was not affected. This indicated that severe metabolic inhibition abolished the high affinity component of [3H]-glibenclamide binding but had no effect on low affinity binding.

Discussion

Low affinity glibenclamide binding

To our knowledge, this study presents the first investigation of [³H]-glibenclamide binding in intact vascular smooth muscle. In homologous competition experiments two binding components were identified with K_i values of 20 nm and 32 μ m. The low affinity component was detectable only at glibenclamide concentrations $\geq 10 \, \mu \text{M}$; in this concentration range glibenclamide is known to bind to several proteins, be they membraneous or cytosolic (Panten et al., 1989; Ashcroft & Ashcroft, 1992; Sheppard & Welsh, 1992; Rabe et al., 1995). This heterogeneity of binding sites in the low affinity component is in obvious contradiction to the observed Hill coefficient

Table 1 Inhibition of high affinity [3H]-glibenclamide binding by K_{ATP} channel modulators

Substance	pK_i $(-\log M)$	$\begin{array}{c} A_I^2 \\ (\% \ \mathbf{B}_{tot}) \end{array}$	$n_{H,I}$
Glibenclamide	7.69 ± 0.10	17 ± 1	1.0 ± 0.2
AZ-DF 265	7.00 ± 0.10	17 ± 2	1.2 ± 0.3
Glipizide	6.44 ± 0.08	18 ± 1	0.9 ± 0.1
Glibornuride	5.75 ± 0.05	15 ± 1	1.6 ± 0.2
P1075	7.95 ± 0.05	16 ± 1	1.1 ± 0.1
Minoxidil sulphate	7.27 ± 0.05	17 ± 1	1.2 ± 0.1
Pinacidil	7.09 ± 0.02	16 ± 1	1.1 ± 0.1
Levcromakalim	6.74 ± 0.05	17 ± 1	1.2 ± 0.2
Aprikalim	6.74 ± 0.02	15 ± 1	1.4 ± 0.1
Diazoxide	4.38 ± 0.03	16 ± 1	1.5 ± 0.1

¹Individual inhibition curves for the K_{ATP} channel modulators were fitted to equation (1) yielding the pK_i $(=-\log K_i)$ values, amplitudes (A_1) , and Hill coefficients (n_{H,1}) listed below. Pooled data are shown in Figures 1 and 2 (n=4-8).²A₁ values not different from that for glibenclamide (ANOVA followed by Dunnet's test).

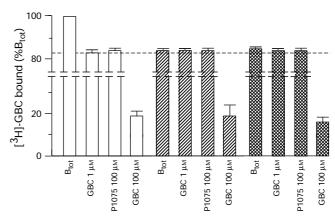


Figure 3 Effects of metabolic inhibition on [3H]-glibenclamide binding. Binding is given as % Btot under control conditions. Open columns: control; hatched columns: hypoxia combined with inhibition of glycolysis (i.e. bubbling with $95\%~N_2/5\%~CO_2$ in the presence of 2-deoxyglucose (1 mm) and absence of glucose); cross-hatched columns: inhibition of oxidative phosphorylation (oligomycin (1 μ M) in the presence of 2-deoxyglucose (1 mm)). The dashed line represents the contribution of the high affinity component to Btot under control conditions.

Table 2 Correlation of the potencies of K_{ATP} channel modulators in the [³H]-glibenclamide binding assay with those obtained in functional and [3H]-P1075 assays1

	x vs y	Correlation coefficient	Slope	d^2
Blockers:	pA_2 (⁸⁶ Rb ⁺ efflux) ³ vs pK_i ([³ H]-GBC binding)	0.992	1.1 ± 0.1	-0.14
	pK_i ([³ H]-P1075 binding) ⁴ vs pK_i ([³ H]-GBC binding)	0.989	1.0 ± 0.1	-1.22
Openers:	pD_2 (vasorelaxation) ⁵ vs pK_i ([³ H]-GBC binding)	0.979	0.9 ± 0.1	0.43
pk	pK_i ([³ H]-P1075 binding) ⁴ vs pK_i ([³ H]-GBC binding)	0.993	1.0 ± 0.1	0.32

¹K_{ATP} channel blockers were glibenclamide (GBC), AZ-DF 265, glipizide and glibornuride; openers were P1075, aprikalim, diazoxide, leveromakalim, minoxidil sulphate and pinacidil. ²d denotes the mean logarithmic distance of the correlation line from the line of identity in the region of interest; d>0: shift upwards. ³pA₂ values were obtained from inhibition of P1075 (60 nM)-induced ⁸⁶Rb⁺ efflux from rat aorta (Quast et al., 1993); this efflux is taken as a qualitative measure of KATP channel activation (Edwards & Weston, 1994; Quast, 1996). pKi values were calculated from the IC₅₀ values in (Quast et al., 1993) after correction for the presence of the agonist, i.e. by dividing the IC₅₀ values by a factor of 4 (Quast, 1996); in view of the allosteric coupling between openers and sulphonylureas (Bray & Quast, 1992), the values calculated in this manner should be regarded as pseudo-pA₂ ('pA₂')-values. ⁴Data from Bray & Quast (1992). ⁵pD₂ values for opener-induced vasorelaxation were determined in endothelium-denuded rat aortic rings precontracted by noradrenaline (0.1 µM); data from Quast et al. (1993).

of 2.1. However, glibenclamide is known to partition into the membrane (Zünkler et al., 1989) and, at the high concentrations needed to saturate the low affinity component, nonspecific effects of glibenclamide are likely to occur. At concentrations $\geq 100 \ \mu M$, glipizide and AZ-DF 265 also started to interact with these sites whereas the K_{ATP} channel openers did not. The low affinity component was not investigated further.

High affinity glibenclamide binding: pharmacological characterization

Although the high affinity component of glibenclamide binding was only 17% of total binding it was possible to characterize this component in detail. The Hill coefficient was close to unity $(n_{H,1} = \bar{1}.0 \pm 0.2)$ and the $K_{i,1}$ value of glibenclamide was 20 nm. This K_i value is in excellent agreement with the potency of glibenclamide for block of the vascular KATP channel opened by guanosine 5'-diphosphate (GDP) (Beech et al., 1993) or by low concentrations of KATP channel openers (Xu & Lee, 1994; Quast, 1996). It is also close to the inhibition constant of glibenclamide against the vasorelaxant effect of the openers calculated from the observed rightward shifts $(K_i = 10 - 30 \text{ nM}; \text{ Bray & Quast, 1991}; \text{ Quast } et \ al., 1993). Si$ milarly, other sulphonylureas and the carboxylate derivative, AZ-DF 265, inhibited high affinity glibenclamide binding completely. The $K_{i,1}$ values determined from these curves agree within a factor of 2 with the respective inhibition constants determined previously against P1075-induced 86Rb+ efflux (Table 2). These data strongly suggest that the high affinity component of [3H]-glibenclamide binding mediates the block of the K_{ATP} channel by the sulphonylureas.

Previous binding studies with [3H]-glibenclamide in smooth muscle membranes have failed to show a meaningful inhibition by K_{ATP} channel openers (Zini et al., 1991). However, in the intact aortic rings representatives of the major structural classes of K_{ATP} channel openers inhibited high affinity glibenclamide binding completely. Table 2 shows that the midpoints of these inhibition curves agree within a factor of 3-4 with those of the vasorelaxation-concentration curves determined previously in rat aorta (Quast et al., 1993). Similarly, the potencies of the openers in the binding assay correlate very well with the potencies to induce 86Rb+ efflux from rat aorta (data from Bray & Quast, 1992; Quast et al., 1993; correlation not shown). Hence, the high affinity sulphonylurea binding site described here also recognizes the K_{ATP} channel openers in a pharmacologically relevant way.

Severe metabolic inhibition selectively and completely abolished the high affinity component of [3H]-glibenclamide binding. The manoeuvres used here have previously been shown to reduce ATP levels in rat aortic rings by >90% (Quast et al., 1993); this could possibly result in inadequate phosphorylation of the drug receptor and/or in the disruption of the cytoskeleton (Hall & Haworth, 1996). Phosphorylation has indeed been shown to be necessary for functional binding of [3H]-glibenclamide to membraneous or solubilized preparations from an pancreatic B-cell line (Niki & Ashcroft, 1991; Schwanstecher et al., 1992a) and from cerebral cortex (Schwanstecher et al., 1992b); an intact cytoskeleton has been shown to be necessary for proper functioning of the K_{ATP} channel in the principal cells of the rat cortical collecting duct (Wang et al., 1994). It is well possible that both inadequate phosphorylation of the drug receptor and disassembly of the actin cytoskeleton prevent meaningful binding of K_{ATP} channel modulators in membranes from smooth muscle (Gopalakrishnan et al., 1991; Zini et al., 1991; Quast et al., 1993).

Comparison of high affinity [3H]-glibenclamide binding with $[^3H]$ -P1075 binding

In their ability to recognize both sulphonylureas and openers and in their sensitivity to metabolic blockade, the high affinity binding sites for [3H]-glibenclamide strongly resemble the binding sites labelled by the opener, [3H]-P1075, in rat aorta (Bray & Quast, 1992; Quast et al., 1993). When the results of the binding assays were compared, the openers were two times more potent in the [3H]-P1075 assay whereas the sulphonylureas were 17 times more potent in the [3H]-glibenclamide binding assay (Table 2). Collectively, the data of Table 2 show that the high affinity sulphonylurea sites recognize binding of the openers with almost the correct potency whereas the openers sense binding of the sulphonylureas in a more remote fashion. These shifts may reflect a negative allosteric coupling between the binding sites for openers and sulphonylureas. A negative allosterism between the sites for openers and blockers is supported by our earlier observation that glibenclamide, at high concentrations, enhanced the dissociation of the [3H]-P1075-receptor complex (Bray & Quast, 1992).

A requirement for such an allosteric coupling is that the densities of binding sites for openers and sulphonylureas are of similar magnitude. The B_{max} value for [${}^{3}H$]-P1075 in rat aortic strips has been determined to be 21 ± 3 fmol mg⁻¹ wet weight (Bray & Quast, 1992). However, the lability of the binding, probably due to the strong dependence on cell metabolism, and more recent experiments suggest that this represents a lower limit and that the real value may be considerably higher. The B_{max} value of the [3H]-glibenclamide sites may be estimated from the amplitude of the high affinity component $(7.1 \pm 0.5 \text{ fmol mg}^{-1} \text{ wet weight)}$ by considering that at the label concentration used ([3H]-glibenclamide = 1 nm) the high affinity sites are only saturated to 5%. By use of equation 2 in the Methods section and respecting propagation of errors, a $B_{\text{max}} = 150 \pm 38 \text{ fmol mg}^{-1}$ wet weight was obtained. Considering the uncertainty in these numbers and taking into account that the stoichiometry between the two classes of sites is unknown, these estimates are compatible with the notion of an allosteric coupling between the sites for sulphonylureas and openers.

In conclusion, we have characterized a high affinity [³H]-glibenclamide binding component in rat aortic rings which recognizes both sulphonylureas and K_{ATP} channel openers with the potencies expected from functional assays; hence, these sites may represent the sulphonyl-urea receptor. The properties of this binding component, including its sensitivity to metabolic inhibition, resemble the properties of sites for the

openers, indicating a strong coupling between the two classes of sites.

Note added in proof

While this paper was in press, cloning of a smooth muscle type sulphonylurea receptor was reported [Isomoto *et al.* (1996). *J. Biol. Chem.*, **271**, 24321–24324].

We thank F. Metzger and Dr U. Ruß for helpful discussion. The study was supported by Deutsche Forschungsgemeinschaft, grants Qu 100/1-1 and Qu 100/2-1.

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