

# The A<sub>2</sub> Adenosine Receptor: Guanine Nucleotide Modulation of Agonist Binding Is Enhanced by Proteolysis

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Received July 26, 1990; Accepted October 30, 1990

## SUMMARY

Agonist binding to the A<sub>2</sub> adenosine receptor (A<sub>2</sub>AR) and its regulation by guanine nucleotides was studied using the newly developed radioligand <sup>125</sup>I-2-[4-(2-[2-[(4-aminophenyl)methylcarbonylamino]ethylaminocarbonyl)ethyl]phenyl]ethylamino-5'-N-ethylcarboxamidoadenosine (<sup>125</sup>I-PAPA-APEC) and its photoaffinity analog <sup>125</sup>I-azido-PAPA-APEC. A single protein of M<sub>r</sub> 45,000, displaying the appropriate A<sub>2</sub>AR pharmacology, is labeled in membranes from bovine striatum, PC12 cells, and frog erythrocytes. In DDT, MF2 cells the labeled protein has a slightly lower molecular weight. Incorporation of <sup>125</sup>I-azido-PAPA-APEC into membranes from rabbit striatum, however, reveals two specifically labeled peptides (M<sub>r</sub> ~47,000 and 38,000), both of which display A<sub>2</sub>AR pharmacology. Inhibition of protease activity leads to a decrease in the amount of the M<sub>r</sub> 38,000 protein, with only the M<sub>r</sub> 47,000 protein remaining. This suggests that the M<sub>r</sub> 38,000 peptide is a proteolytic product of the M<sub>r</sub> 47,000 A<sub>2</sub>AR protein. In membranes containing the intact undigested A<sub>2</sub>AR protein, guanine nucleotides induce a small to insignificant decrease in agonist binding, which is atypical of stimulatory G<sub>s</sub>-

coupled receptors. This minimal effect is observed in rabbit striatal membranes prepared in the presence of protease inhibitors, as well as in the other tissues studied. Binding to rabbit striatal membranes that possess the partially digested receptor protein, however, reveals a 50% reduction in maximal specific agonist binding upon addition of guanine nucleotides. Inhibition of proteolysis in rabbit striatum, on the other hand, results in a diminished ability of guanine nucleotides to regulate agonist binding. Thus, the enhanced effectiveness of guanine nucleotides in rabbit striatal membranes is associated with the generation of the M<sub>r</sub> 38,000 peptide fragment. Guanosine 5'-(β,γ-imido)triphosphate reduces photoaffinity labeling by 55% in the M<sub>r</sub> 38,000 protein, whereas the labeling is decreased by only 28% in the M<sub>r</sub> 47,000 receptor protein.

Our data, therefore, suggest that, unless proteolysis occurs, the A<sub>2</sub>AR in all tissues studied is tightly associated with the G<sub>s</sub> protein and displays minimal guanine nucleotide modulation of agonist binding, which makes the A<sub>2</sub>AR an atypical stimulatory receptor.

ARs are known to mediate a wide range of physiological effects, including vasodilatation, suppression of cardiac rate and contractility, induction of sedation, and inhibition of platelet aggregability (1). Two subtypes of ARs have been defined, based on their pharmacological profiles, and termed A<sub>1</sub>AR and A<sub>2</sub>AR (2, 3). Over the past 5 years there have been dramatic advances in our understanding of the structure, function, and regulation of the A<sub>1</sub>AR (4-6). This receptor has been (a) studied by radioligand binding, where it was found to be tightly coupled to the G<sub>i</sub> protein, both in membranes and following solubili-

zation, and to be dramatically regulated under a variety of conditions, (b) photoaffinity labeled, with its glycoprotein nature being studied, and (c) purified to homogeneity (7-12). In contrast, little is known about the A<sub>2</sub>AR. Until 1989, there had been no selective high affinity radioligands to study the A<sub>2</sub>AR. [<sup>3</sup>H]NECA had been utilized as a radioligand, but its use was associated with many artifacts, in terms of binding to both A<sub>2</sub>AR and A<sub>1</sub>AR, as well as to multiple proteins that display characteristics of neither the A<sub>1</sub>AR nor the A<sub>2</sub>AR (13-15). Recently, we (16) and Jarvis *et al.* (17) have described the synthesis of two high affinity A<sub>2</sub>-selective agonist radioligands, <sup>125</sup>I-PAPA-APEC and [<sup>3</sup>H]CGS 21680. We have used <sup>125</sup>I-PAPA-APEC and its azide derivative to define the binding subunit structure of the A<sub>2</sub>AR and its glycoprotein character-

C.N. is supported by a Postdoctoral Research Exchange Grant from the Max Kade Foundation, Inc. G.L.S. is supported in part by the National Heart, Lung and Blood Institute (Grant RO1-HL-35134) and Supplement and Grant-in-Aid (880662) from the American Heart Association and 3M Riker.

**ABBREVIATIONS:** AR, adenosine receptor; A<sub>1</sub>AR, A<sub>1</sub> adenosine receptor; A<sub>2</sub>AR, A<sub>2</sub> adenosine receptor; Gpp(NH)p, guanosine 5'-(β,γ-imido)triphosphate; NECA, 5'-N-ethylcarboxamidoadenosine; PAPA-APEC, 2-[4-(2-[2-[(4-aminophenyl)methylcarbonylamino]ethylaminocarbonyl)ethyl]phenyl]ethylamino-5'-N-ethylcarboxamidoadenosine; (R)-PIA, (-)-N<sup>6</sup>-(R)-1-methyl-2-phenylethyladenosine; (S)-PIA, (+)-N<sup>6</sup>-(S)-1-methyl-2-phenylethyladenosine; SANPAH, N-succinimidyl 6-(4'-azido-2'-nitrophenylamino)hexanoate; azido-PAPA-APEC, 2-[4-[2-[2-[(4-azido-phenyl)methylcarbonylamino]ethylaminocarbonyl]ethyl]phenyl]ethylamino-5'-N-ethylcarboxamido adenosine; SDS, sodium dodecyl sulfate; PAGE, polyacrylamide gel electrophoresis; CGS 21680, 2-[4-(2-carboxyethyl)phenylethylamino]-5'-N-ethylcarboxamidoadenosine; PMSF, phenylmethylsulfonyl fluoride; HEPES, N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid; G protein, guanine nucleotide-binding protein.

istics in bovine striatal membranes, as well as to demonstrate that the binding subunit has the pharmacological properties expected of the A<sub>2</sub>AR (18, 19).

Early kinetic studies from Levitzki and co-workers (20, 21) and studies on the constituents of the AR signaling pathway in striatal membranes (22) suggested that the coupling of A<sub>2</sub>AR to its effector system, adenylate cyclase, was unusual, compared with that of other receptors that act through G<sub>s</sub> to activate cyclase (20–23). These studies provided evidence that the A<sub>2</sub>AR was permanently coupled to its effector system and did not undergo an association/dissociation reaction as did the  $\beta$ -adrenergic receptor. We recently have found that guanine nucleotides have only a minimal effect on agonist binding in bovine striatal membranes (18). This is also atypical, compared with what has been described for other stimulatory receptors such as the glucagon and the D<sub>1</sub> dopamine receptors (24, 25). Therefore, we undertook the present study to answer several specific questions. First, what is the A<sub>2</sub>AR subunit structure in different tissues and species? Second, is the A<sub>2</sub>AR sensitive to endogenous proteases and what functional consequences does this proteolysis produce? Third, is the bovine brain A<sub>2</sub>AR atypical, as it relates to the inability of guanine nucleotides to substantially decrease agonist binding?

## Experimental Procedures

### Materials

(R)-PIA, (S)-PIA, adenosine deaminase, and Gpp(NH)p were obtained from Boehringer-Mannheim. Soybean trypsin inhibitor, pepstatin A, leupeptin, PMSF, chloramine T, and HEPES-Na were from Sigma. NECA was generously donated by Dr. Ray Olsson (University of South Florida). SANPAH was purchased from Pierce. Na<sup>125</sup>I (carrier-free; 100 mCi/ml) was from Amersham Corp. Electrophoresis reagents were obtained from Bio-Rad Laboratories. All other chemicals were of the highest purity grade available.

### Methods

**Preparation of membranes from rabbit striatum.** Rabbit brains were obtained from Pel-Freez or the Duke University Vivarium and were kept at  $-70^{\circ}$ . After thawing, the corpus striatum was dissected and minced in ice-cold buffer (50 mM HEPES, 10 mM MgCl<sub>2</sub>, pH 7.4). The tissue was gently homogenized using a Teflon pestle, and membranes were centrifuged at  $43,000 \times g$  for 10 min. The pellet was washed once and finally resuspended to give a concentration of 400 mg of wet weight/ml. Aliquots were stored at  $-70^{\circ}$  for use within 2 weeks. Rabbit striatal membranes prepared following the above protocol were designated as controls.

In order to limit endogenous proteolytic activity, the following buffer was used for preparation of rabbit striatal membranes: (in mM) HEPES, 50; EDTA, 5; PMSF, 0.1; (in  $\mu$ g/ml) soybean trypsin inhibitor, 100; leupeptin, 5; pepstatin A, 1; pH 7.9. Before storage, MgCl<sub>2</sub> was added at a concentration of 6 mM, which was increased to 10 mM for the binding assay. The preparation of membranes from bovine striatum as well as DDT<sub>1</sub> MF2 cells has been described and was carried out accordingly (18, 26). Membranes from frog erythrocytes were a generous gift from Dr. Marc G. Caron (Duke University Medical Center), and PC12 cell membranes were kindly provided by Dr. John W. Daly (National Institutes of Health). None of these membrane batches included protease inhibitors, except for the designated preparation from rabbit striatum.

**Radioiodination of PAPA-APEC and synthesis of <sup>125</sup>I-azido-PAPA-APEC.** The parent compound PAPA-APEC, an A<sub>2</sub>AR-selective agonist, was synthesized as described (16). Iodination with Na<sup>125</sup>I and synthesis of the azide derivative of PAPA-APEC have also been

outlined in detail (18, 19). In brief, PAPA-APEC was iodinated by the chloramine T method and isolated by high pressure liquid chromatography, using a C18  $\mu$ Bondapak column. For isocratic elution, the mobile phase was composed of 60% methanol/40% ammonium formate (20 mM, pH 7.85). <sup>125</sup>I-PAPA-APEC was either used as a radioligand or further processed to give the azide derivative. Briefly, the substance was dried down and redissolved in acetic acid (10  $\mu$ l, 6 N). After reaction with sodium nitrite (20  $\mu$ l, 20 mg/ml), sodium azide was added (10  $\mu$ l, 5 mg/ml). In dimmed light, the incubation was run for 10 min and terminated by alkalization (8  $\mu$ l of ammonium hydroxide). Separation of <sup>125</sup>I-azido-PAPA-APEC was performed by high pressure liquid chromatography, using an isocratic protocol (75% methanol/25% 20 mM ammonium formate, pH 7.85). The radioligand was assumed to have a specific activity of 2200 Ci/mmol on the day of radioiodination.

**Photoaffinity labeling of A<sub>2</sub>AR.** Photoaffinity labeling with the agonist photoaffinity/cross-linking probe <sup>125</sup>I-azido-PAPA-APEC was performed as previously described (19). Frozen membranes were thawed, washed, and suspended in 50 mM HEPES, 10 mM MgCl<sub>2</sub> (pH 6.8), in either the absence or the presence of protease inhibitors. After pretreatment with adenosine deaminase (1 unit/mg of protein/2 ml), incubations were run for 60 min at  $37^{\circ}$ , in a total volume of 1 ml consisting of <sup>125</sup>I-azido-PAPA-APEC at  $\sim 1$  nM, competitor or Gpp(NH)p at the indicated concentrations, and a total of 0.5–1 mg of membrane protein. The binding reaction was terminated by dilution of the mixture with ice-cold incubation buffer and centrifugation at  $43,000 \times g$ . The pellet was washed once, taken up in 1 ml, and exposed to UV light for 4 min at a distance of 1 cm from the light source (UVCG-25 mineral light). The entire procedure was performed in the dark. Upon photoincorporation, membranes were washed again, solubilized in 10% SDS-glycerol buffer, and subjected to electrophoresis on a 10% acrylamide gel and autoradiography.

Alternatively, photoaffinity labeling was performed with the radioligand <sup>125</sup>I-PAPA-APEC, using SANPAH as the cross-linking agent (18).

**Radioligand binding assay.** <sup>125</sup>I-PAPA-APEC binding experiments were conducted as previously described (18). Membranes were prepared as for photoaffinity labeling assays. In saturation and competition binding experiments, incubations were for 60 min in a volume of 250  $\mu$ l, containing 25–50  $\mu$ g of membrane protein and radioligand, competitor, and/or Gpp(NH)p at the indicated concentrations. When the azide derivative was used as a radioligand, binding assays were carried out in foil-wrapped tubes. Separation of free and bound ligand was achieved by vacuum filtration over polyethylenimine (0.3%)-treated glass fiber filters and rinsing with  $3 \times 4$  ml of ice-cold buffer. The radioactivity was quantitated in a Packard Multi-Prias  $\gamma$ -counter at an efficiency of 75%.

Nonspecific binding assayed in the presence of 5 mM theophylline amounted to 50–60% of total binding in the  $K_D$  concentration range. This binding component was not altered by the inclusion of guanine nucleotides. The variability of the data from triplicate determinations was less than 5% of the mean value.

Saturation and competition curves were analyzed using a nonlinear, least-squares, curve-fitting procedure, including a weighting routine and statistical analysis, as previously published.

In dissociation experiments, equilibrium binding was attained by preincubation of rabbit striatum membranes (1 mg of protein) with <sup>125</sup>I-PAPA-APEC (1–2 nM) for 45 min at  $37^{\circ}$ . Dissociation of radioligand binding was initiated by 50-fold dilution of the preincubation volume (1 ml), using buffer at  $37^{\circ}$  with or without 0.1 mM Gpp(NH)p. At the indicated time points, 3-ml aliquots were withdrawn, poured onto membrane filters, and rinsed with ice-cold incubation buffer. Evaluation of the experimental data and estimation of a  $k_{off}$  value were achieved through fitting of the data to a monoexponential equation, based on nonlinear least-squares regression.

Protein determination was carried out according to the method of Bradford (27), with bovine serum albumin as a standard.

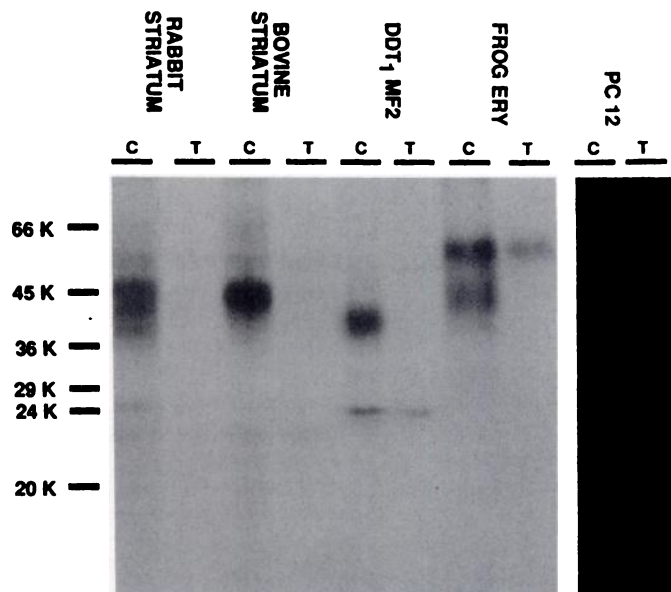
## Results

Photoaffinity labeling of the A<sub>2</sub>AR was performed in membranes from rabbit and bovine striatum, rat PC12 and DDT<sub>1</sub> MF-2 cells, and frog erythrocytes. The photoaffinity radioligand <sup>125</sup>I-azido-PAPA-APEC labels distinct proteins, as shown by autoradiography after separation on SDS-PAGE (Fig. 1). Specific incorporation into the A<sub>2</sub>AR is completely suppressed by addition of 5 mM theophylline. As estimated from their migration profile, most of the A<sub>2</sub>AR species examined migrate with apparent *M<sub>r</sub>* 44,000–47,000, whereas in DDT<sub>1</sub> MF2 cells the A<sub>2</sub>AR has *M<sub>r</sub>* 40,000. Specific labeling is confined to a single protein, except in the case of rabbit striatum membranes, wherein a doublet is observed at *M<sub>r</sub>* 47,000 and 38,000 (Figs. 1 and 2).

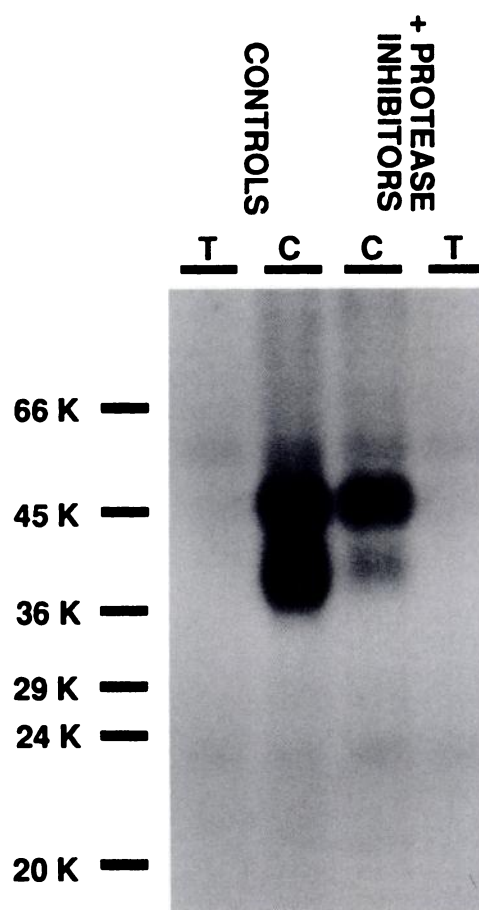
This particular feature of the A<sub>2</sub>AR from rabbit striatum likely arises from partial enzymatic digestion of the receptor protein during cell disruption. Modification of the membrane preparation procedure to inhibit protease activity markedly reduces the amount of the *M<sub>r</sub>* 38,000 protein, whereas labeling of the *M<sub>r</sub>* 47,000 protein is similar or enhanced, compared with control (Fig. 2). The relative amount of radioactivity incorporated into each protein was quantified and gave identical results regardless of whether the azide derivative was used for direct photoaffinity labeling or, alternatively, <sup>125</sup>I-PAPA-APEC was cross-linked into the receptor proteins using SANPAH as a cross-linking agent (data not shown).

Thus, inclusion of protease inhibitors and omission of Mg<sup>2+</sup> induces a considerable shift in the ratio of upper to lower bands, from 1.6 (0.7–3.6, 95% confidence intervals) in control to 5.3 (2.9–9.4) under conditions in which proteolysis is inhibited.

Further evidence that the <sup>125</sup>I-azido-PAPA-APEC-labeled *M<sub>r</sub>* 38,000 protein originates from enzymatic cleavage of the *M<sub>r</sub>* 47,000 A<sub>2</sub>AR comes from displacement experiments with AR



**Fig. 1.** Photoaffinity labeling of A<sub>2</sub>ARs in membranes from rabbit and bovine striatum, DDT<sub>1</sub> MF2 and PC12 cells, and frog erythrocytes (ERY). Membranes were incubated with the agonist photoaffinity probe <sup>125</sup>I-azido-PAPA-APEC (1 nM) in the absence (C) or presence of 5 mM theophylline (T), as described in Experimental Procedures. After photoaffinity labeling, samples were solubilized and aliquots were subjected to SDS-PAGE and autoradiography. This autoradiograph is representative of two separate experiments.



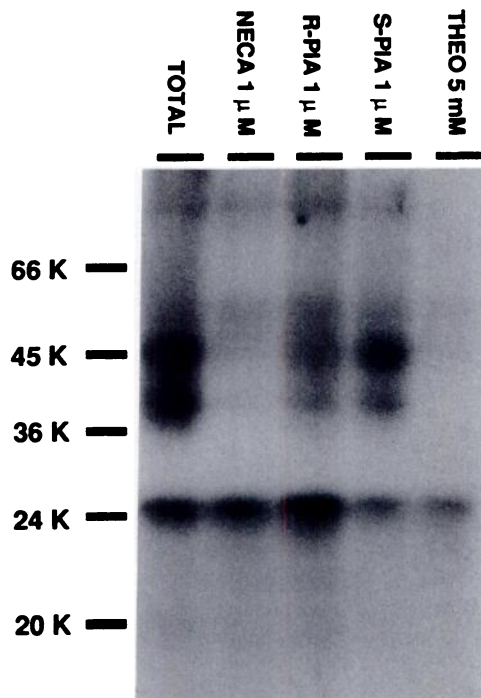
**Fig. 2.** <sup>125</sup>I-Azido-PAPA-APEC labeling of A<sub>2</sub>ARs in control membranes from rabbit striatum and in membranes prepared with protease inhibitors (see Experimental Procedures). The inner lanes (C) show the result of photoaffinity labeling in the absence and the outer lanes (T) in the presence of 5 mM theophylline. Equal amounts of protein from each sample (200 µg) were loaded onto the gel. This experiment was repeated several times.

agonists. The incorporation of the photoaffinity probe into both proteins is antagonized to a similar extent by NECA, (R)-PIA, and (S)-PIA, revealing a rank order of potency that is characteristic of the A<sub>2</sub>AR (Fig. 3). Competition binding curves with <sup>125</sup>I-PAPA-APEC accordingly reflect the A<sub>2</sub>AR pharmacology. *K<sub>i</sub>* values (nM) derived from experiments in control membranes are NECA, 18.0 ± 1.2; (R)-PIA, 343 ± 104; (S)-PIA, 7170 ± 2600; and theophylline, 7623 ± 5228 (means ± standard errors). Thus, the receptor fragment retains the binding affinity for agonists as well as antagonists, suggesting that the binding subunit remains functionally intact during enzymatic alteration of the receptor protein.

A differentiating characteristic, however, is observed for the effect of guanine nucleotides on the incorporation of <sup>125</sup>I-azido-PAPA-APEC into the doublet of A<sub>2</sub>AR proteins. Fig. 4 demonstrates the effect of the nonhydrolyzable guanine nucleotide Gpp(NH)p on photoaffinity labeling in membranes prepared in either the presence or the absence of protease inhibitors. In the smaller molecular weight peptide, labeling is clearly reduced by Gpp(NH)p (to 45 ± 12% of specific incorporation), whereas the effect appears small to insignificant in the larger protein (to 72 ± 10% of specific incorporation).

It has been well established that guanine nucleotides reduce

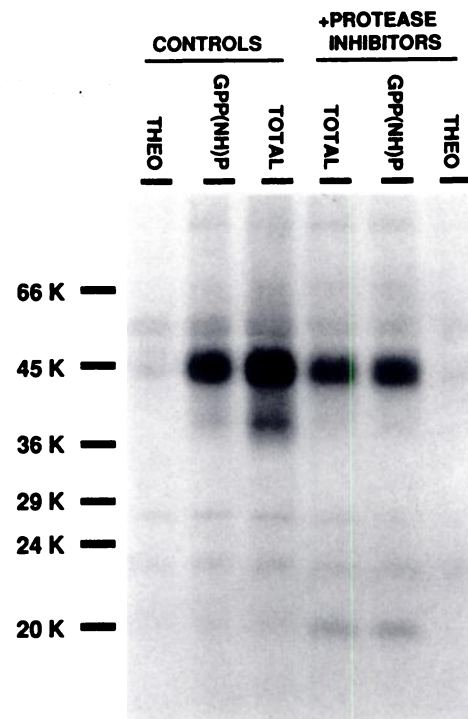




**Fig. 3.** Inhibition of  $^{125}\text{I}$ -azido-PAPA-APEC photoaffinity labeling in control rabbit striatum membranes by AR agonists. Membranes were photoaffinity labeled with  $^{125}\text{I}$ -azido-PAPA-APEC alone (TOTAL) or in the presence of the indicated concentrations of the competing ligands. Autoradiography of the samples after separation on SDS-PAGE is shown, with the positions of molecular weight markers to the left. The results are representative of three similar experiments. THEO, theophylline.

high affinity agonist binding to G protein-coupled receptors (28). An increase in the agonist dissociation rate is considered to cause a shift to the low affinity binding state. In order to characterize the effect of guanine nucleotides on binding parameters, saturation experiments were performed with  $^{125}\text{I}$ -PAPA-APEC in rabbit striatal membranes. Fig. 5 displays saturable and monophasic binding isotherms in the absence as well as the presence of 0.1 mM Gpp(NH)p. As predicted from the results of photoaffinity labeling experiments, rabbit striatal membranes containing the intact receptor reveal almost superimposable binding curves. This is also observed in membranes that are demonstrated in Fig. 1 to harbor the undigested  $\text{A}_2\text{AR}$  protein, i.e., from frog erythrocytes, DDT<sub>1</sub> MF-2 cells, and PC12 cells (data not shown). In unprotected control membranes from rabbit striatum, however,  $^{125}\text{I}$ -PAPA-APEC binding is subject to G protein modulation. Upon addition of Gpp(NH)p, no change in receptor affinity becomes evident, whereas the receptor number decreases by 50% of maximum specific binding (Table 1). Because  $^{125}\text{I}$ -PAPA-APEC is a full agonist radioligand, it is not surprising that we do not detect a low affinity binding state in saturation experiments.

This finding is further confirmed in dissociation experiments. The resulting curves are shown in Fig. 6 and demonstrate a monophasic dissociation, according to an exponential decay that transforms to a straight line on a semilogarithmic plot. This indicates dissociation from a single affinity state of the receptor, giving a uniform  $k_{\text{off}}$  of  $0.16 \pm 0.02 \text{ min}^{-1}$  (six experiments). Gpp(NH)p creates a small, statistically nonsignificant,  $k_{\text{off}}$  increase ( $0.20 \pm 0.05 \text{ min}^{-1}$ ), without altering the monophasic characteristics of the dissociation curve. In order to avoid interference from a time-dependent onset of guanine



**Fig. 4.** Effect of the guanine nucleotide analog Gpp(NH)p on the incorporation of the agonist photoaffinity probe  $^{125}\text{I}$ -azido-PAPA-APEC into the partially digested and the intact  $\text{A}_2\text{AR}$ . Rabbit striatum membranes were prepared as controls or in the presence of protease inhibitors (see Experimental Procedures). Membranes were incubated with  $^{125}\text{I}$ -azido-PAPA-APEC (1 nM) alone (TOTAL) or in the presence of Gpp(NH)p (0.1 mM) or theophylline (5 mM) (THEO). Autoradiography of the gel is shown, with the positions of molecular weight markers to the left. Three additional experiments gave similar results.

nucleotide effects, additional experiments were performed with Gpp(NH)p being present also during the preincubation period (data not shown). This modification would allow for dissociation from a supposedly low affinity state of the receptor but revealed  $k_{\text{off}}$  values identical to those estimated in control experiments, providing evidence for the inability of the agonist radioligand  $^{125}\text{I}$ -PAPA-APEC to discern a low affinity state of the  $\text{A}_2\text{AR}$ .

## Discussion

In this paper we have provided information on the structure and function of  $\text{A}_2\text{AR}$ s from different tissues and species. We have documented that (a) with a single exception the  $\text{A}_2\text{AR}$  binding subunit resides on a  $M_r \sim 45,000$  protein, (b) either the  $\text{A}_2\text{AR}$  from the rabbit striatum is very sensitive to the effects of endogenous proteases or there are high levels of proteases in this species, and (c) this endogenous proteolysis has profound effects on the ability of guanine nucleotides to regulate agonist binding.

The  $\text{A}_2\text{AR}$  of rabbit striatum is shown to be unstable in the presence of endogenous proteolytic enzymes (Fig. 2). Photoaffinity labeling with an  $\text{A}_2\text{AR}$ -selective probe demonstrates two proteins are specifically labeled, both of which display the appropriate  $\text{A}_2\text{AR}$  pharmacology. Inhibition of protease activity largely reduces the doublet to a single labeled protein of  $M_r 47,000$ . This protein, i.e., the undigested  $\text{A}_2\text{AR}$ , reveals a molecular weight and pharmacological properties similar to those of the  $\text{A}_2\text{AR}$  in bovine striatal membranes (18). Although the

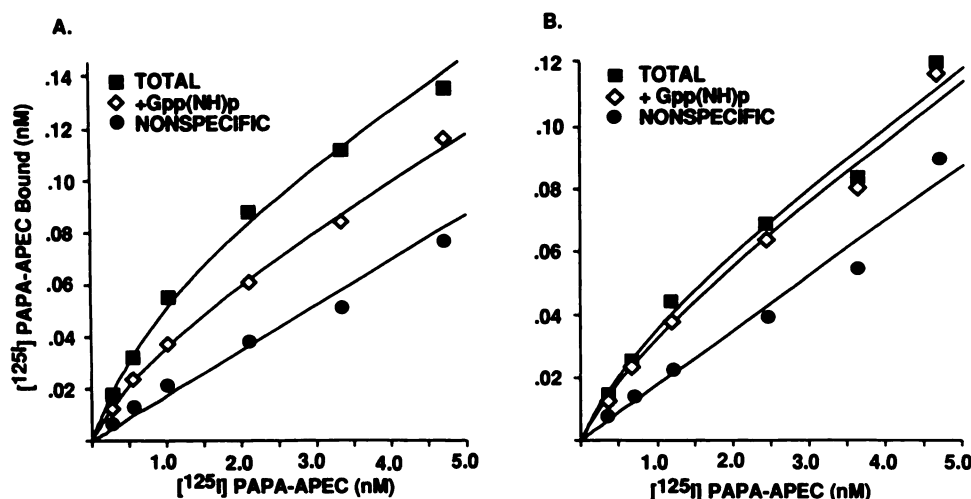


Fig. 5. Effect of the guanine nucleotide analog Gpp(NH)p on reversible binding of the agonist radioligand  $^{125}\text{I}$ -PAPA-APEC to rabbit striatum membranes harboring either the intact or the partially digested  $\text{A}_2\text{AR}$  protein. Control or protease-inhibited membrane preparations were used in the binding assay. Incubations were run in a volume of 250  $\mu\text{l}$  containing 25–50  $\mu\text{g}$  of membrane protein and  $^{125}\text{I}$ -PAPA-APEC, at the indicated concentrations, alone (TOTAL) or in the presence of Gpp(NH)p (0.1 mM) or theophylline (5 mM). Gpp(NH)p had no effect on the displacement of radioligand binding by theophylline. Experiments were performed in triplicate, and the data were fitted to an equation describing the interaction with a single binding site.

TABLE 1

Parameter estimates for  $^{125}\text{I}$  PAPA-APEC binding to rabbit striatum  
Values are mean  $\pm$  standard error.

	Controls		+Protease inhibitors	
	$K_D$	Receptor number	$K_D$	Receptor number
	nM	fmoI/mg	nM	fmoI/mg
No addition	$1.94 \pm 0.54$	$248 \pm 54$	$1.03 \pm 0.08$	$197 \pm 43$
+Gpp(NH)p (0.1 mM)	$1.43 \pm 0.35$	$125 \pm 23^*$ ( $50 \pm 11\%^b$ )	$1.57 \pm 0.27$	$189 \pm 40$ ( $96 \pm 8\%^b$ )

\* Significantly different from the value in the absence of Gpp(NH)p ( $p < 0.01$ ).

<sup>b</sup> Expressed as percentage of the total receptor number in the same experiment.

receptor protein from rabbit striatum is susceptible to endogenous proteolysis,  $\text{A}_2\text{AR}$ s from other species and tissues, i.e., those in membranes from bovine striatum, frog erythrocytes, and PC 12 and DDT<sub>1</sub> MF2 cells, have a structurally stable receptor protein. Thus, the rabbit striatum  $\text{A}_2\text{AR}$  species appears to be unique among ARs, because neither  $\text{A}_2\text{AR}$ s nor  $\text{A}_1\text{AR}$ s have been reported to be susceptible to proteolysis (1, 18, 26).

It should be remembered that many transmembrane receptors, such as the  $\alpha_1$ - and the  $\beta$ -adrenergic receptors, have been found to be labile in the presence of tissue-specific proteases, a feature that has been described in several tissues (29, 30). Similarly, their degradation is inhibited by addition of protease inhibitors, particularly upon the removal of divalent metal cations.

Furthermore, our data suggest that alteration of the  $\text{A}_2\text{AR}$  protein by proteolysis leads to a markedly enhanced modulation of agonist radioligand binding by guanine nucleotides. Treatment of membranes with protease inhibitors, on the other hand, results in an almost complete lack of effect of guanine nucleotides on agonist binding. Thus, the effectiveness of guanine nucleotides may be associated with the generation of the  $M_r$  38,000 peptide fragment, as shown in Fig. 4. Prevention of proteolysis of the  $\text{A}_2\text{AR}$  itself or of other membrane proteins appears to diminish the ability of guanine nucleotides to decrease agonist binding. This finding may be interpreted as tight coupling between receptor and G protein or, alternatively, as a uniform conformation of the receptor binding subunit regardless of G protein modulation. The latter concept could explain why the receptor does not display a low affinity state for the agonist PAPA-APEC unless the receptor structure is enzymatically altered.

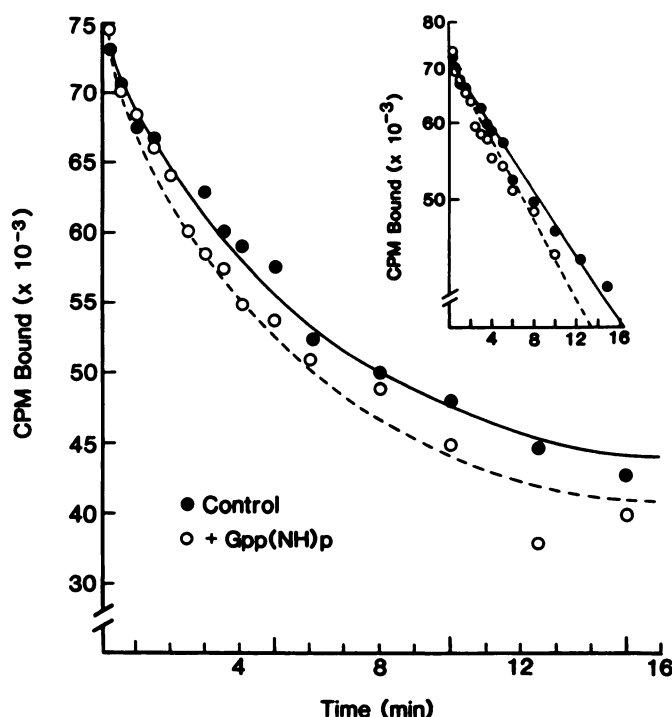


Fig. 6. Dissociation of the agonist radioligand  $^{125}\text{I}$ -PAPA-APEC from the  $\text{A}_2\text{AR}$  in control membranes from rabbit striatum. Membranes prepared in the absence of protease inhibitors were preincubated with 1 nM  $^{125}\text{I}$ -PAPA-APEC, and dissociation was initiated by 50-fold dilution, without or with the addition of 0.1 mM Gpp(NH)p. Aliquots were withdrawn and filtered at the indicated time points. After fitting of the data, the lines were drawn according to an equation for monoexponential decay. Inset, data have been linearized for illustrative purposes. The results are representative of four experiments.

Nevertheless, it might be argued that treatment of membranes with EDTA in the absence of  $\text{Mg}^{2+}$  to inactivate metal ion-dependent proteases might potentially perturb the  $\text{A}_2\text{AR}$ - $\text{G}_i$  interaction and, thus, account for the loss of sensitivity to guanine nucleotides. Experiments with different conditions for the preparation of membranes from rabbit striatum (i.e., with and without  $\text{Mg}^{2+}$ ), however, led us to conclude that proteolysis and not alterations in the ion concentrations are responsible for the varying effects of guanine nucleotides (data not shown).

Provided that proteolysis is inhibited, the minimal effect of

guanine nucleotides observed in rabbit striatal membranes is reminiscent of the results reported for <sup>125</sup>I-PAPA-APEC binding in membranes from bovine striatum, wherein no proteolysis appears to occur and agonist binding is reduced by only ~10% in the presence of guanine nucleotides. This has also been found in DDT<sub>1</sub> MF-2, PC12, and frog erythrocyte membranes and makes the A<sub>2</sub>AR distinctly unusual, compared with other receptors that activate adenylate cyclase. Although the common paradigm has been that guanine nucleotides totally ablate high affinity agonist binding (33), our data suggest that what initially appeared anomalous, i.e., a minimal effect of Gpp(NH)p on agonist binding in the bovine striatal A<sub>2</sub>AR, is actually typical for a number of A<sub>2</sub>ARs if endogenous proteolysis is inhibited.

Although our current understanding of the mode of coupling between the A<sub>2</sub>AR and the stimulatory G protein, G<sub>s</sub>, is still rudimentary, there are data to suggest that there may be a very tight coupling between A<sub>2</sub>AR and G<sub>s</sub> (20–23). For example, in turkey erythrocyte membranes it has been proposed that the A<sub>2</sub>AR is permanently coupled to the catalytic unit via the G<sub>s</sub> protein, a model that is opposite to the transient nature of the β-adrenergic receptor-G<sub>s</sub> interaction in the same membrane. Both models are, however, potentially in accord with the findings on the guanine nucleotide regulation of agonist binding. Whereas the β-adrenergic receptor-G<sub>s</sub> complex apparently dissociates (at least functionally) in the presence of a GTP analog, which is associated with a complete loss of receptors in the agonist-specific high affinity state, there is virtually no effect of guanine nucleotides on agonist binding to the A<sub>2</sub>AR, as demonstrated in these studies. Taken together, these findings imply that A<sub>2</sub>AR and G<sub>s</sub> do not functionally uncouple even in the presence of guanine nucleotides.

We and others have documented that the A<sub>1</sub>AR and G<sub>i</sub> are tightly coupled, in that guanine nucleotides do not totally eliminate agonist-specific high affinity binding in membranes but do decrease binding by 40–60% (A<sub>1</sub>ARs are also not susceptible to endogenous proteolysis) (10, 34). Thus, ARs appear to be atypical in their responses to guanine nucleotides, with the A<sub>2</sub>AR being even more resistant to their effects than is the A<sub>1</sub>AR. In addition, it is clear that proteolysis of the membranes containing A<sub>2</sub>AR promotes guanine nucleotide sensitivity and, therefore, great care must be taken to avoid proteolysis, which can induce binding artifacts.

#### Acknowledgments

We would like to thank Linda Scherich for her assistance in the preparation of this manuscript.

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