Identification of Endothelin Receptor Subtypes in Rat Kidney Cortex using Subtype-Selective Ligands

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SUMMARY

¹²⁵l-Endothelin (ET)-1 and ¹²⁵l-ET-3 displayed specific, saturable, and high affinity binding to membranes prepared from rat kidney cortex. Saturation binding experiments using ¹²⁵l-ET-1 and ¹²⁵l-ET-3 revealed that ¹²⁵l-ET-3 binding sites were 40–50% less abundant than ¹²⁵l-ET-1 binding sites. The dissociation constants (K_d) and maximum binding (B_{max}) for ¹²⁵l-ET-1 and ¹²⁵l-ET-3 with these membranes were 218 ± 23 pm and 275 ± 20 fmol/mg of protein and 207 ± 19 pm and 113 ± 17 fmol/mg of protein, respectively. In the presence of 10 nm sarafotoxin 6c, a selective agonist for ET_B receptors, ¹²⁵l-ET-1 binding was decreased by 45–50% and ¹²⁵l-ET-3 binding was totally abolished, suggesting

that approximately 40–50% of kidney cortex ET receptors are of the ET_B subtype and that ¹²⁵I-ET-1 binds to both ET_A and ET_B receptors with the same high affinity, whereas ¹²⁵I-ET-3 binds to only ET_B receptors with high affinity. In addition, in the presence of BQ123 [cyclo(p-Trp,p-Asp,L-Pro,p-Val,L-Leu)], a selective antagonist for ET_A receptors, ¹²⁵I-ET-1 binding was decreased by 50%, whereas ¹²⁵I-ET-3 binding was unaffected. Our results strongly suggest that rat kidney cortex contains ET_A and ET_B receptors in a 50:50 ratio and that sarafotoxin 6c and BQ123 are valuable tools in identifying the subtypes of ET receptors in various tissues.

ET-1, a 21-amino acid peptide isolated from the culture medium of endothelial cells, has been shown to have potent vasoactive properties (1). Following its discovery, two related peptides, ET-2 and ET-3, and a family of 21-amino acid peptide toxins (sarafotoxins) that show high degrees of homology to ET-1 were identified (2). The physiological actions of the ETs have been shown to be mediated by the interaction of ET with specific cell surface receptors, and ET receptors have been identified in a variety of tissues and cell lines (for review, see Refs. 3 and 4). ET-1 acts as a potent vasoconstrictor, especially in kidney, when administered in vivo. It has been shown to decrease glomerular filtration rate, renal plasma flow, and ultrafiltration coefficient (5-7). It also appears to play an important role in renal failure and cyclosporine nephrotoxicity (8-10). Based on radioligand binding studies, the existence of subtypes of ET receptors has been proposed (for review, see Ref. 11). Recently, ET receptors have been cloned from bovine (12) and rat lung (13) and designated ETA and ETB, respectively, based upon the relative binding affinity of ET-1 and ET-3. ET_A receptors are selective for ET-1, with ET-3 showing a 100-fold diminished binding affinity. ET_B receptors, termed "nonselective," display similar binding properties for ET-1 and ET-3. Williams et al. (14) have demonstrated that S6c, isolated from snake venom, binds selectively to ET_B receptors with high affinity and stimulates phosphatidylinositol turnover only in tissue slices possessing ET_B receptors (15). Recently, Ihara et

al. (16) have described a cyclic pentapeptide (BQ123) that showed >1000-fold selectivity for ET_A receptors. We have used S6c and BQ123 as tools to characterize the subtypes of ET receptors present in rat kidney cortex.

Materials and Methods

¹²⁵I-ET-1 (specific activity, 2200 Ci/mmol) and ¹²⁵I-ET-3 (specific activity, 2200 Ci/mmol) were obtained from New England Nuclear (Boston, MA). Unlabeled ET-1 and ET-3 were from American Peptides (Santa Clara, CA). S6c was from Bachem Biosciences (Philadelphia, PA) and American Peptides. BQ123 was synthesized in the Department of Peptidomimetic Research, SmithKline Beecham Pharmaceuticals (King of Prussia, PA). All other chemicals were of the highest grade available.

Membrane preparation. Whole kidneys of male Sprague-Dawley rats (175–200 g) were removed, and the cortex was dissected out. This cortical tissue was homogenized (1 g/10 ml) in 20 mM Tris·HCl, pH 7.4, 5 mM EDTA, 0.25 M sucrose, 100 μ g/ml phenylmethylsulfonyl fluoride, 10 μ g/ml aprotinin, 10 μ g/ml leupeptin (buffer A), using a tissue homogenizer (15 strokes). The homogenate was centrifuged for 15 min at 4° at 1000 \times g. The supernatant was decanted, filtered through cheesecloth, and centrifuged for 30 min at 4° at 40,000 \times g. The resulting pellets were resuspended in buffer B (50 mM Tris·HCl, pH 7.5, 20 mM MgCl₂).

¹²⁶I-ET-1 binding. ¹²⁵I-ET-1 binding to membranes from rat kidney cortex was performed in buffer containing 50 mm Tris·HCl, pH 7.5, 10 mm MgCl₂, and 0.05% bovine serum albumin, as described (10). Membrane protein (2-6 µg/tube) was added to tubes containing either

buffer (total binding) or 100 nm unlabeled ET-1 (nonspecific binding). The reaction was started by the addition of 0.3 nm 125I-ET-1 and continued for 60 min at 30°. After the incubation, the reactions were stopped with 3 ml of cold buffer containing 50 mm Tris. HCl, pH 7.5, and 10 mm MgCl₂. Membrane-bound radioactivity was separated from free ligand by filtering through Whatman GF/C filter paper that had been presoaked in 0.1% bovine serum albumin. The filters were washed five times with 3 ml of buffer, using a Brandel cell harvester, and the filter papers were counted in a γ counter with an efficiency of 75%. Saturation binding experiments were performed using increasing concentrations of ¹²⁵I-ET-1 (30-500 pM), in the absence and presence of 100 nm unlabeled ET-1. Competition binding experiments were performed using 0.3 nm 125I-ET-1, in the absence and presence of increasing concentrations of the indicated compounds. 125I-ET-3 binding was performed in a fashion similar to that described above. Each experiment was performed three or four times; the data shown are from one experiment and are representative of those obtained in similar experi-

Data analysis. Saturation binding experiments were analyzed by nonlinear regression analysis of untransformed data, using the Lundon 1 program (Lundon Software Inc., Cleveland, OH). Competition curves were analyzed using the Lundon 2 program. Single- and multiple-site models were statistically compared to determine the best fit, and differences among models were tested by comparing the residual variance, using an F test and a significance level of p < 0.05.

Results

 125 I-ET-1 and 125 I-ET-3 bound to rat kidney cortex membranes in a specific and saturable manner, as shown in Fig. 1, A and B. Scatchard transformation of the specific binding from the saturation binding experiments revealed linear Scatchard plots, suggesting a single class of high affinity binding sites, with a dissociation constant (K_d) of 218 ± 23 pM and maximum binding $(B_{\rm max})$ of 275 ± 20 fmol/mg of protein for 125 I-ET-1 and a K_d of 207 ± 19 pM and a $B_{\rm max}$ of 113 ± 17 fmol/mg of protein for 125 I-ET-3 (Fig. 1C). The membrane protein concentrations chosen were in the linear range of the protein concentration curve. Time course experiments suggested that both 125 I-ET-1 and 126 I-ET-3 binding reached steady state by 60 min at 30°, at the ligand and protein concentrations used (data not shown).

Competition of ¹²⁵I-ET-1 binding by unlabeled ET-1, ET-3, S6c, and BQ123 is shown in Fig. 2A. Unlabeled ET-1, ET-3, S6c, and BQ123 displaced ¹²⁵I-ET-1 binding in a concentration-dependent manner. ET-1 gave a monophasic curve, with K_d and $B_{\rm max}$ values of 0.3 ± 0.04 nM and 260 fmol/mg of protein (represented as 100% in Table 1), respectively. ET-3 and S6c displayed shallow competition curves (Fig. 2A). ET-3 blocked 48% of ¹²⁵I-ET-1 binding sites with high affinity (0.7 \pm 0.06 nM) and 52% with low affinity (540 \pm 260 nM). Similarly, S6c blocked 34% of ¹²⁵I-ET-1 binding with high affinity (0.9 \pm 0.1 nM) and 66% with low affinity (930 \pm 240 nM) (Table 1A; Fig. 2A). In addition, BQ123, an ET_A subtype-selective antagonist, displaced only 50% of ¹²⁵I-ET-1 binding, even at concentrations as high as 1 μ M (Fig. 2A), suggesting the presence of subtypes of ET receptors.

Next, we attempted to quantitate the proportion of these subtypes by performing $^{125}\text{I-ET-1}$ competition binding experiments in the presence of the ET_B-selective ligand S6c (to block all ET_B receptors) or the ET_A-selective antagonist BQ123 (to block all ET_A receptors). Fig. 2B shows the competition curves for ET-1, ET-3, and S6c performed in the presence of 1 μM BQ123. In the presence of 1 μM BQ123, $^{125}\text{I-ET-1}$ binding was

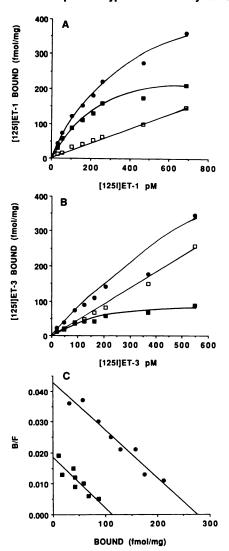


Fig. 1. A and B, Saturation binding of ¹²⁵I-ET-1 (A) and ¹²⁵I-ET-3 (B) to membranes prepared from rat kidney cortex. Assays were performed in duplicate at 30° for 60 min. Increasing concentrations of ¹²⁵I-ET-1 or ¹²⁵I-ET-3 were added to membranes in the absence (total binding) (●) or presence (nonspecific binding) (□) or 100 nm unlabeled ligand. Specific binding (■) was the difference between total and nonspecific binding bound and free ligands were separated as described in Materials and Methods. C, Scatchard transformations of the specific binding of ¹²⁵I-ET-1 (●) and ¹²⁵I-ET-3 (■) from A and B.

decreased by 48% (135 fmol/mg of protein), compared with control (260 fmol/mg of protein); the competition curves for ET-3 and S6c were monophasic, like that for ET-1, with K_d values of 0.2 ± 0.03 , 3.5 ± 0.3 , and 0.3 ± 0.04 nm, respectively (Fig. 2B; Table 2B). These data further indicated that 48% of ¹²⁵I-ET-1 binding sites were ET_A receptors. Similar competition experiments were performed in the presence of 100 nm S6c, and the data are shown in Fig. 2C and Table 1C. Total binding in the presence of S6c was decreased by 56% (115 fmol/mg of protein), compared with control. Whereas the ET-1 curve did not change, compared with control (Fig. 2A), the ET-3 curve was shifted to the right and predominantly displayed low affinity binding, with a K_d of 308 \pm 160 nM, suggesting that, when ET_B receptors were blocked by S6c, ET-3 bound to ET_A receptors with low affinity. On the other hand, the BQ123 curve was shifted to the left (Fig. 2C), compared with control (Fig. 2A), and was monophasic, with a K_d of 0.4 \pm 0.06 nm. In addition,

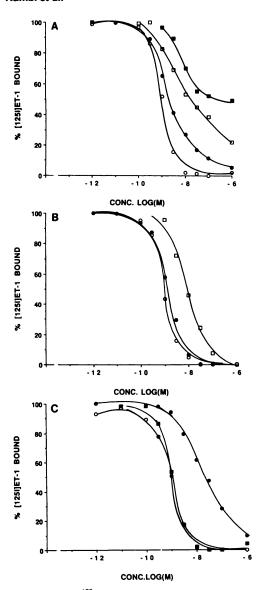


Fig. 2. A, Competition of ¹²⁵I-ET-1 binding to membranes prepared from rat kidney cortex, by unlabeled ET-1 (O), ET-3 (●), S6c (□), and BQ123 (■). Membranes were incubated in triplicate with 0.2 nm ¹²⁵I-ET-1 and increasing concentrations of unlabeled ligands, for 60 min, at 30°. Bound and free ligands were separated as described. Data obtained in the absence of competitor were normalized to 100% and those obtained in the presence of 1 μM ET-1 were 0%; data are expressed as the percentage of ¹²⁵I-ET-1 binding. B, Competition of ¹²⁵I-ET-1 binding to cortex membranes by unlabeled ET-1 (O), ET-3 (●), and S6c (□) in the presence of 1 μM BQ123. The experiment was performed as described for A. C, Competition of ¹²⁵I-ET-1 binding to cortex membranes by unlabeled ET-1 (O), ET-3 (●), and BQ123 (■) in the presence of 100 nm S6c. The experiment was performed as described for A.

BQ123 displaced >90% of ¹²⁵I-ET-1 binding (Fig. 2C), compared with control, where the maximum displacement was 50% (Fig. 2A).

The percentage distribution of ET_A and ET_B receptors in this tissue was further quantitated by the binding of $^{125}\text{I-ET-1}$ and $^{125}\text{I-ET-3}$, in the presence and absence of S6c and BQ123, and the data are presented in Fig. 3. $^{125}\text{I-ET-1}$ binding was 193 \pm 5 fmol/mg of protein (100%) and $^{125}\text{I-ET-3}$ binding was 92 \pm 17 fmol/mg of protein (48%), suggesting that 48% of $^{125}\text{I-ET-1}$ binding was to the ET_B receptor subtype. In the presence of 10 nm S6c, $^{125}\text{I-ET-1}$ binding was inhibited by 52%, to 90.0 \pm 2

TABLE 1

Percentage distribution and dissociation constants for ET-1, ET-3, S6c, and BQ123 in competing for ¹²⁵IET-1 binding to membranes from rat kidney cortex

 126 IET-1 competition binding experiments using unlabeled ET-1, ET-3, S6c, and BQ123 were performed as described in Materials and Methods and the Fig. 2 legend. The data presented are mean \pm standard error of three determinations and were analyzed using the Lundon 2 program. $R_{\rm H}$ and $R_{\rm L}$ represent high and low affinity sites for the respective ligands. $K_{\rm H}$ and $K_{\rm L}$ represent dissociation constants for high and low affinity sites for the respective ligands. For A, B, and C, 100% represents 260 \pm 15,135 \pm 10, and 115 \pm 9 fmol/mg of protein, respectively.

	R _H	KH	R _L	K,
	%	ПМ	%	пм
A. Control				
ET-1	100	0.3 ± 0.04		
ET-3	48 ± 4	0.7 ± 0.06	52 ± 5	540 ± 260
S6c	34 ± 3	0.9 ± 0.10	66 ± 4	930 ± 240
BQ123	50 ± 8	9.3 ± 2.0	50 ± 12	
B. In the presence of 1				
μ м BQ123				
EŤ-1	100	0.3 ± 0.04		
ET-3	100	0.2 ± 0.03		
S6c	100	3.5 ± 0.30		
C. In the presence of				
100 nм S6c				
ET-1	100	0.4 ± 0.04		
ET-3	10 ± 2	3.0 ± 0.70	90 ± 6	308 ± 160
BQ123	100	0.4 ± 0.06		

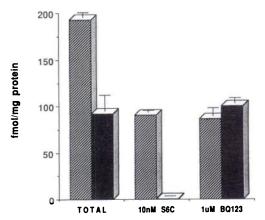


Fig. 3. 125 I-ET-1 and 125 I-ET-3 binding to rat kidney cortex membranes in the absence and presence of 10 nm S6c or 1 μ m BQ123. 125 I-ET-1 (22) or 125 I-ET-3 (125 I) (0.3 nm) was added to membranes in the absence of S6c or BQ123 (total) or the presence of 10 nm S6c or 1 μ m BQ123, and membranes were incubated for 60 min at 30°. Bound and free ligands were separated as described in Materials and Methods. Nonspecific binding was measured in the presence of 100 nm unlabeled ET-1 or ET-3 and was 10 and 20% of total binding for 125 I-ET-1 and 125 I-ET-3, respectively. The data shown are mean \pm standard error of three determinations.

fmol/mg of protein, whereas $^{125}\text{I-ET-3}$ binding was totally abolished, further confirming that 50–55% of $^{125}\text{I-ET-1}$ binding was to ET_B receptors and that, at the concentration used (0.3 nm), $^{125}\text{I-ET-3}$ bound only to ET_B receptors (Fig. 3). Addition of 1 μ M BQ123 did not have any effect on $^{125}\text{I-ET-3}$ binding (100 \pm 5.7 fmol/mg of protein), whereas it inhibited $^{125}\text{I-ET-1}$ binding by 53% (86.7 \pm 8.3 fmol/mg of protein), further confirming that 50% of $^{125}\text{I-ET-1}$ binding was to ET_A receptors. In the presence of BQ123 and S6c, $^{125}\text{I-ET-1}$ binding was totally abolished.

Discussion

There are many reports in the literature suggesting the presence of subtypes of ET receptors in various tissues (17-

21), and the recent cloning of ETA and ETB receptors from bovine and rat lung (12, 13) confirmed these suggestions. The present study demonstrates the presence of high affinity ET_A and ET_B receptors in membranes prepared from rat kidney cortex. Whereas 126I-ET-1 bound to both ETA and ETB receptors with the same high affinity, 125I-ET-3 bound to only ETB receptors with high affinity. Analysis of 125I-ET-1 competition binding experiments with unlabeled ET-1, ET-3, S6c, and BQ123 suggested that the ET-1 competition curve was monophasic, with high affinity, in the absence or presence of subtypeselective ligands. On the other hand, the ET-3 competition curve was shallow in the absence of any subtype-selective ligands and was monophasic and shifted to the left (high affinity) in the presence of BQ123 or to the right (low affinity) in the presence of S6c. Similar results were obtained for S6c and BQ123. In the presence of BQ123, the S6c competition curve was shifted to the left and was monophasic, whereas in the absence of BQ123 the S6c competition curve was shallow and of low affinity. In the presence of S6c the BQ123 competition curve was monophasic and shifted to the left, whereas in its absence BQ123 inhibited only 50% of ¹²⁵I-ET-1 binding. even at 1 µM. Also, in the presence of S6c or BQ123 125 I-ET-1 binding was decreased by ~50%, and in the presence of both together ¹²⁵I-ET-1 binding was totally abolished, suggesting that kidney cortex displayed only ETA and ETB receptors. In addition, 125I-ET-3 binding was totally abolished by S6c and was unaffected by BQ123, further confirming that BQ123 is selective for ET_A receptors and that ¹²⁵I-ET-3 bound to only ET_B receptors with high affinity.

At present, we do not know the physiological functions of these two subtypes of receptors, although both ET_A and ET_B receptors, when transiently expressed in COS cells or Xenopus oocytes, mediated calcium release when challenged with the agonists (12, 13, 22). Whereas ET-1 binds to both ETA and ET_B receptors with the same affinity, ET-3 is more selective for ET_B receptors. The physiological function of ET-3 is not known at this time. Both ETs and sarafotoxins display prolonged vasoconstrictor actions in many vascular beds (23). In addition to the vasopressor effects, ETs produce a characteristic short-lived vasodepressor effect before the pressor effect (24). Warner et al. (25) and Douglas and Hiley (26) reported endothelium-dependent relaxation in response to ET-3, ET-2, ET-1, and S6b. The order of potency of these peptides for relaxation was different from that for contraction. ET-3 was the most potent relaxant but the least potent constrictor (26), suggesting that different receptors may be mediating these two responses. Recently, MacCumber et al. (27) and Matsumoto et al. (28) have reported the tissue distribution of various ET-related peptides in rat. Their data showed that both ET-1 and ET-3 were identified in kidney. This distribution of ET-1 and ET-3 is consistent with the ET receptor subtypes identified in this study and might suggest a role for ET-1 and ET-3 as local hormones in renal physiology.

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