Distribution and density of α - and β -adrenergic receptor binding sites in the bovine mammary gland

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SUMMARY. Radioreceptor binding studies were designed to localize and determine the number of α - and β -adrenergic receptors in the mammary gland of lactating cows. ³H-prazosin, ³H-rauwolscine and ³H-dihydroalprenolol were used for the regional characterization of α_1 -, α_2 - and β -adrenergic receptors by competitive inhibition of binding of ³H-ligands with unlabelled adrenergic agonists and antagonists. The α_1 -, α_2 - and β_2 -adrenergic receptor subtypes could thus be demonstrated in the regions of the teats, large mammary ducts and parenchyma. Tissues of the teat wall, of the large mammary ducts above the gland cistern and of the mammary parenchyma were prepared to determine the density of α_1 , α_2 - and β receptors by saturation binding assays using ³H-prazosin, ³H-rauwolscine and ³Hdihydroalprenolol respectively. Binding to high affinity sites was reversible within minutes and saturable. Equilibrium was reached within minutes. The number of α_1 and α_2 -adrenergic receptors decreased from the teat to the mammary ducts to the parenchyma. Most of the α_1 - and α_2 -adrenergic receptors were found in the teat wall, whereas in the parenchyma α -adrenergic receptors were absent or barely detectable. The density of β -adrenergic receptors was similar in the teat wall and the large mammary ducts, but much lower in the parenchyma. Thus, α_1 -, α_2 - and β -adrenergic receptors were found mainly in the milk purging system and hardly at all in mammary parenchyma. Inhibition of milk removal by α -adrenergic stimulation is possibly due to constriction of teat wall and to constriction of the mammary ducts, whereas enhanced milk flow after β -adrenergic stimulation is possibly due to relaxation not only of the teat sphincter and teat wall, but probably also of the large mammary ducts.

The sympathetic nervous system is involved in peripheral modulation of milk removal. Milk yield was decreased during adrenalin, noradrenaline or α -specific adrenergic agonist (α -Ag) administration. The inhibition of milk removal by the α -Ag phenylephrine was shown to be dose-dependent (H. M. Hammon, unpublished observations). The effects were not mediated by inhibition of endogenous oxytocin release during milking (Lefcourt & Akers, 1984; Blum *et al.* 1989). Furthermore, the inhibition of milk let-down by adrenergic drugs could be abolished by α -, but not by β -adrenergic blockade (Blum *et al.* 1989). Contraction of both teat and gland cistern was observed using ultrasonography after α -Ag administration (Bruckmaier & Blum, 1992). Inhibition of milk removal by premilking electroshocks was abolished by

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 α -adrenergic blockade (Lefcourt & Akers, 1982; Blum *et al.* 1989). These studies support the notion that catecholamines mediate their inhibitory effects on milk removal through stimulation of α -adrenergic receptors located in the bovine mammary gland (Blum *et al.* 1989; Bruckmaier *et al.* 1991), most likely on smooth muscle cells not only in teats, but also in mammary ducts (Ziegler & Mosimann, 1960). On the other hand, enhanced milk flow after administration of β -adrenergic agonists (β -Ag) was ascribed to relaxation of the teat muscles and muscles around milk ducts leading from the alveolar tissue to the gland cistern (Hamann, 1981; Bruckmaier *et al.* 1991). Binding studies using radiolabelled adrenergic compounds as ligands indicated the existence of both α_1 - and α_2 -adrenergic receptor subtypes and of the β_2 -adrenergic receptor subtype, most likely on smooth muscles in the wall of the bovine teat, thus confirming radioligand binding studies on smooth muscles of the teat (Roets *et al.* 1984; Roets & Peeters, 1985, 1986). Adrenergic receptor density in the teat muscles was related to milkability of lactating cows (Roets *et al.* 1989).

Because previous *in vivo* investigations suggested α - and β -adrenergic effects not only in the teat, but also in other regions of the udder (Bruckmaier *et al.* 1991; Bruckmaier & Blum, 1992), the goal of the present investigations was the determination of α - and β -adrenergic receptors not only in the teat, but also in the region of the large mammary ducts and in the mammary parenchyma. The possible importance of the different regions of the mammary gland for inhibition of milk removal mediated by α -adrenergic receptors and enhancement of milk flow mediated by β -adrenergic receptor interaction could thus be evaluated.

MATERIALS AND METHODS

Preparation of membranes

Tissues from three regions of lactating udders were used: the muscular layer of the teat (teat region), tissue around the gland cistern including the large mammary ducts (duct region), and mammary parenchyma from the dorsal udder in which mammary ducts were not visible macroscopically (parenchyma region). The material was removed immediately after slaughter and transferred on ice. The tissues of duct and parenchyma regions were minced into small pieces with scissors and put into ice cold 50 mm-Tris-HCl buffer, pH 7.4. The Tris-HCl buffer used in our assays contained 6 mM-MgCl₂, essential for binding of the radiolabelled ligand (Moor et al. 1988), and 1 mm-EGTA to chelate metals without affecting ligand binding (Parini et al. 1987). Teats were prepared as described by Roets et al. (1984). Internal mucosa and epidermis were removed, and the sphincter area of the teat was cut off and discarded. The tissue was suspended in $\sim 10-15$ volumes of buffer. All subsequent manipulations were carried out at 4 °C. The material was homogenized four times for 20 s at low speed (8000 rev./min) with an Ultra-Turrax homogenizer (T 25, Janke & Kunkel, Staufen, Germany). The homogenate was centrifuged at 500 g for 10 min; the supernatant was passed through two layers of cheesecloth and then centrifuged at 49000 g and 4 °C for 30 min (Moor et al. 1988). The resulting supernatant was decanted completely and the remaining pellet was suspended in ice cold buffer (as above) by a motor-driven Glass-Teflon homogenizer.

The protein concentration was determined using a kit (BCA Protein Assay Reagent; Pierce, Rockford, IL, USA) and the membrane suspension was then adjusted to 200-300 μ g protein/100 μ l. Specific ³H-ligand binding was plotted against protein concentration and was shown to increase with increasing membrane protein concentration, as expected (results not shown). Protein concentration was

kept constant (200 or 300 μ g/100 μ l) throughout. While membrane suspensions used for competitive binding assays were frozen in liquid nitrogen and stored at -80 °C, saturation binding assays were carried out immediately after preparation of the membrane suspension. Freezing of membranes did not affect binding as such, but preliminary studies demonstrated some reduction in the number of binding sites (Shi *et al.* 1989).

Binding studies

 α_1 -Adrenergic receptor binding was tested by use of ³H-prazosin, a highly α_1 -selective antagonist (Shi *et al.* 1989), α_2 -adrenergic receptor binding with ³H-rauwolscine, an α_2 -selective antagonist (Shi *et al.* 1989), and β -adrenergic binding with ³H-dihydroalprenolol (³H-DHA), a non-selective β -adrenergic antagonist (Hancock *et al.* 1979). Saturation binding assays were performed to determine binding capacity. Competitive binding assays with different specific unlabelled ligands were performed to characterize the receptor subtype.

Saturation binding assays. These assays were carried out to investigate the distribution of ³H-prazosin (n = 7), ³H-rauwolscine (n = 4) and ³H-DHA (n = 7) binding sites in teat, duct and parenchyma regions, where n is the number of udders. Membrane suspensions $(100 \ \mu$ l) were incubated with increasing concentrations of the ³H-labelled ligand with or without unlabelled ligand $(50 \ \mu$ l) at 37 °C for 15 min with constant shaking to determine total and non-specific binding. All drugs were diluted in the Tris-HCl buffer described above. The binding reaction was terminated by adding 1 ml of ice cold buffer and the material was immediately passed through glass fibre filters (diam. 20 mm; Whatman GF/C, Whatman International, Maidstone ME14 2LE, UK or MN GF-3; Macherey-Nagel, Düren, Germany), using a vacuum filtration manifold (Hölzel, Dorfen, Germany). Filters were rapidly washed three times with 5 ml saline (9 g/l), dried and placed into plastic vials. Scintillation cocktail (3 ml; OptiPhase 'HiSafe' II; FSA Laboratory, Loughborough LE11, UK) was added and the bound ³H-activity was measured in a liquid scintillation counter (BETAmatic, Kontron, Switzerland).

Specific binding of the ³H-ligand was calculated. For testing α_1 - and α_2 -adrenergic receptor binding, 10 μ M phentolamine (final concentration) was used as the unlabelled ligand (or competitor), whereas for evaluation of the β -adrenergic receptor binding 1 μ M-(\pm)-propranolol (final concentration) was used as competitor. Specific binding represented 60–95% of the total binding depending on mammary tissue site and on the type of the ³H-labelled ligand.

Specific binding was expressed as fmol bound ³H-ligand per mg of membrane protein. Equilibrium binding data were plotted as a function of ³H-ligand concentration (saturation curve). Maximal binding capacities $(B_{\rm max})$ and equilibrium dissociation constants $(K_{\rm D})$ were calculated by a curve-fitting computer program (Motulsky, 1987).

Competitive binding assays. These experiments were performed by incubating membrane preparations (100 μ l) with a fixed concentration of ³H-labelled ligand (50 μ l) and a series of competitors (non-radioactive α - or β -Ag or blocking agents) at eight or ten concentrations. Final concentrations of ³H-prazosin, ³H-rauwolscine and ³H-DHA were 1, 5 and 2 nm respectively. The procedures used to determine radiolabelled binding were similar to those described for saturation binding assay but used membrane suspensions obtained from the duct region of several udders.

Binding kinetics. ³H-ligand binding kinetics were studied using a single concentration of the ³H-ligand under standard assay conditions at different

incubation times. Final concentrations of ³H-prazosin, ³H-rauwolscine and ³H-DHA were 1, 5 and 2 nm respectively. Assays with ³H-prazosin and ³H-DHA were performed at 37 °C. Assays with ³H-rauwolscine were carried out at 25 °C to maintain the reversibility of specific binding. In assays with ³H-prazosin and ³H-DHA the competitor was added after 10 min of binding reaction, in assays with ³H-rauwolscine after 6 min.

Drugs

³H-prazosin (81 Ci/mmol), ³H-rauwolscine (84 Ci/mmol) and ³H-DHA (83 Ci/mmol) were purchased from Amersham International plc (Aylesbury HP20 2TP, UK); (-)-noradrenaline-L-hydrogen tartrate, dopamine-HCl and (\pm)-isoproterenol from Fluka AG (Buchs, Switzerland); (\pm)-propranolol from Imperial Chemical Industries (Macclesfield, UK); (-)-adrenalin(+)-bitartrate, (-)-propranolol, (+)-propranolol, prazosin-HCl, (-)-alprenolol hydrate, atenolol, (-)-phenylephrine-HCl, fenoterol-HBr, yohimbine-HCl and clonidine-HCl from Sigma (St Louis, MO 63178, USA). Phentolamine mesylate and rauwolscine-HCl were purchased from Research Biochemicals Inc. (Natick, MA, USA). Dihydroergocryptine and dihydroergotamine were kindly donated by Sandoz AG (Basle) and clenbuterol by Bayer AG (Wuppertal, Germany).

Solutions were prepared freshly before each assay and substances were dissolved in Tris-HCl buffer (see above). If necessary, especially when using lipophilic agents (rauwolscine, yohimbine or prazosin), ethanol (980 ml/l) was added to promote dissolution. In this quantity (50 mM final concentration in the incubation media), ethanol does not modify binding (Roets *et al.* 1984).

Mathematical and statistical evaluation

For data analysis we used the GraphPAD computer program (Motulsky, 1987). Weighted least squares curve fitting of binding values was performed to analyse saturation and competitor displacement studies. For statistical evaluation of the results, the SAS program package (SAS, 1990, release 6.04) was employed. Competitor displacement and receptor density were tested for significance of difference (P < 0.05) by means of Wilcoxon's two-sample test. All values are presented as means \pm SEM.

RESULTS

Binding kinetics

³H-prazosin, ³H-rauwolscine and ³H-DHA binding was rapid and steady state conditions were reached within minutes (Fig. 1). There was dissociation of binding within minutes after addition of competitors at 37 °C for ³H-prazosin and ³H-DHA, but only at 25 °C for ³H-rauwolscine.

Competitive binding assays

The K_i -values and Hill coefficients (h) for inhibition of specific ³H-prazosin binding by adrenergic competitors are shown in Table 1. For ³H-prazosin binding the K_i value of the unlabelled α_1 -antagonist prazosin was lowest while the α_2 -selective antagonist yohimbine and the α_2 -selective agonist clonidine had ~ 1000-fold higher K_i values than unlabelled prazosin (P < 0.05). The non-selective α -adrenergic antagonist phentolamine and the α -adrenergic agents dihydroergotamine and dihydroergocryptine were ~ 30-50 times less potent than unlabelled prazosin for inhibiting binding (P < 0.05). The K_i value of phenylephrine, an α_1 -agonist, was very high compared with the other α -adrenergic competitors. Isoproterenol, a β -

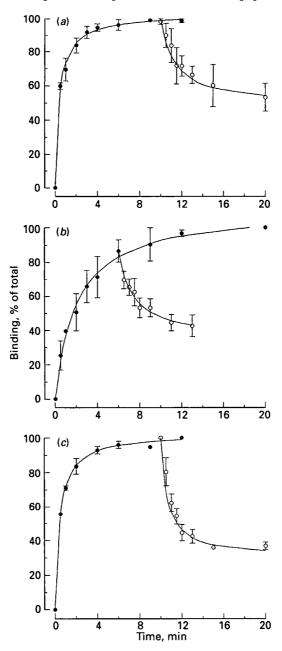


Fig. 1. Binding of (a) ³H-prazosin, (b) ³H-rauwolscine and (c) ³H-dihydroalprenolol to membranes of mammary duct tissue from the udders of three cows: \bullet , association; \bigcirc , dissociation. The competitor was (a) 10 μ M-phentolamine added at 10 min, (b) 10 μ M-phentolamine added at 6 min, (c) 1 μ M-propanolol added at 10 min; total binding of the radiolabelled ligand was taken as 100%.

adrenergic agonist, displaced ³H-prazosin binding only at very high concentrations. The order of potency for inhibition of ³H-prazosin binding was prazosin > phentolamine > dihydroergotamine > dihydroergocryptine > yohimbine > clonidine > adrenalin > noradrenaline > phenylephrine > isoproterenol > dopamine. Most of the Hill coefficients were close to one, except for phentolamine, a non-selective α -antagonist.

| | (values for \mathbf{A}_i and n are mean \pm sem for n unders) | | | |
|---------------------|---|------------------|-----------------|----------|
| | Competitor dissociation | Hill coefficient | | |
| Competitor | constant (K_i), м | Potency ratio | of plot (h) | n |
| Prazosin | $1.0 \pm 0.2 \times 10^{-9}$ n | 1.0000 | 0.90 ± 0.10 | 6 |
| Phentolamine | $2.7 \pm 0.3 \times 10^{-8}$ b | 0.0320 | 0.64 ± 0.03 | 5 |
| Dihydroergotamine | $5.0 \pm 2.0 \times 10^{-8}$ b | 0.0200 | 0.83 ± 0.07 | 6 |
| Dihydroergocryptine | $6.2 \pm 1.7 \times 10^{-8}$ b | 0.0161 | 0.87 ± 0.06 | 6 |
| Yohimbine | $1.1 \pm 0.2 \times 10^{-6}$ c | 0.0009 | 0.80 ± 0.04 | 5 |
| Clonidine | $1.3 \pm 0.1 \times 10^{-6}$ c | 0.0008 | 0.84 ± 0.11 | 5 |
| Adrenalin | $3.7 \pm 0.5 \times 10^{-6}$ d | 0.0003 | 0.80 ± 0.09 | 5 |
| Noradrenaline | 6·7±0·7×10 ^{−6} ° | 0.0001 | 0.79 ± 0.03 | 5 |
| Phenylephrine | $2.2 \pm 0.3 \times 10^{-5}$ f | 0.0000 | 0.79 ± 0.04 | 6 |
| Isoproterenol | $8.8 \pm 1.3 \times 10^{-5}$ g | 0.0000 | 0.91 ± 0.08 | 5 |
| Dopamine | $5.1 \pm 0.4 \times 10^{-4}$ h | 0.0000 | 0.87 ± 0.06 | 4 |

 Table 1. Inhibition of specific ³H-prazosin binding by adrenergic agonists and antagonists (competitors) on membranes of the duct region of the udder

(Values for K_i and h are mean \pm SEM for n udders)

a, b, c, d, e, f, g, h Means without common superscript letters were significantly different (P < 0.05).

Table 2. Inhibition of specific ³H-rauwolscine binding by adrenergic agonists and antagonists (competitors) on membranes of the duct region of the udder

(Values for K_t and h are mean \pm SEM for n udders)

| | Competitor dissociation | | Hill coefficient | |
|---------------|---------------------------------|---------------|------------------|----------|
| Competitor | constant (K_i), M | Potency ratio | of plot (h) | n |
| Rauwolscine | $3.1 \pm 0.4 \times 10^{-8}$ a | 1.0000 | 1.10 ± 0.19 | 5 |
| Clonidine | $5.2 \pm 0.9 \times 10^{-8}$ ab | 0.5960 | 0.68 ± 0.06 | 5 |
| Phentolamine | $5.4 \pm 3.0 \times 10^{-8}$ ab | 0.5740 | 0.73 ± 0.08 | 5 |
| Yohimbine | $8.4 \pm 1.6 \times 10^{-8}$ b | 0.3690 | 0.70 ± 0.07 | 5 |
| Prazosin | $3.3 \pm 0.6 \times 10^{-6}$ c | 0.0090 | 1.00 ± 0.24 | 4 |
| Adrenalin | $3.5 \pm 0.7 \times 10^{-6}$ c | 0.0089 | 0.70 ± 0.05 | 5 |
| Noradrenaline | $1.0 \pm 0.2 \times 10^{-5} d$ | 0.0030 | 0.85 ± 0.05 | 5 |
| Phenylephrine | $4.1 \pm 1.6 \times 10^{-5}$ d | 0.0008 | 0.98 ± 0.11 | 5 |
| Dopamine | $1.3 \pm 0.2 \times 10^{-4}$ e | 0.0002 | 0.87 ± 0.03 | 5 |
| Isoproterenol | $2.5 \pm 0.2 \times 10^{-4}$ f | 0.0001 | 0.99 ± 0.15 | 5 |

a.b. c. d. e. f Means without common superscript letters were significantly different (P < 0.05).

Table 3. Inhibition of specific ³H-dihydroalprenolol binding by adrenergic agonists and antagonists (competitors) on membranes of the duct region of the udder

(Values for K_i and h are mean \pm SEM for n udders)

| Competitor | Competitor dissociation constant (K_i) , M | Hill coefficient Potency ratio of plot (h) | | |
|-------------------|--|---|-----------------|----------|
| (—)-Propranolol | $3.5 \pm 0.5 \times 10^{-9}$ a | 1.0000 | 0.72 ± 0.08 | 5 |
| Alprenolol | $3.7 \pm 0.7 \times 10^{-9}$ a | 1.0000 | 0.91 ± 0.13 | 5 |
| (±)-Propranolol | $8.6 \pm 0.8 \times 10^{-9}$ b | 0.4444 | 0.98 ± 0.08 | 5 |
| Clenbuterol | $4.9 \pm 0.7 \times 10^{-8}$ c | 0.0816 | 0.99 ± 0.06 | 5 |
| (+)-Propranolol | $3.2 \pm 0.5 \times 10^{-7}$ d | 0.0124 | 1.22 ± 0.13 | 5 |
| Fenoterol | $4.8 \pm 1.1 \times 10^{-7}$ d | 0.0084 | 0.94 ± 0.05 | 5 |
| Isoproterenol | $9.1 \pm 2.6 \times 10^{-7}$ d | 0.0044 | 0.88 ± 0.03 | 5 |
| Adrenalin | $2.2 \pm 0.3 \times 10^{-6}$ e | 0.0018 | 0.86 ± 0.14 | 5 |
| Noradrenaline | $1.7 \pm 0.3 \times 10^{-5}$ f | 0.0002 | 1.24 ± 0.28 | 5 |
| Atenolol | $3.9 \pm 1.3 \times 10^{-5}$ f | 0.0001 | 0.73 ± 0.08 | 4 |
| Dopamine | $3.8 \pm 1.9 \times 10^{-4}$ g | 0.0000 | 0.95 ± 0.10 | 4 |

a, b, c, d, e, f, g Means without common superscript letters were significantly different (P < 0.05).

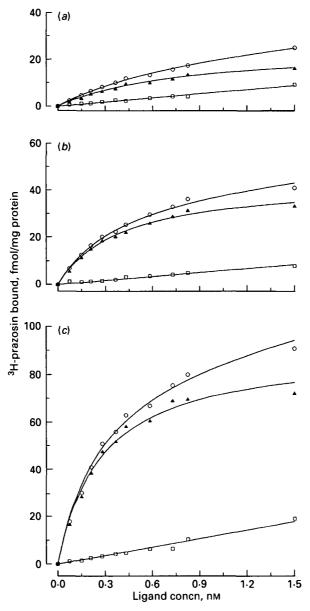


Fig. 2. Representative saturation plot for binding of the α -antagonist ³H-prazosin to suspensions of membrane preparations of (a) parenchyma, (b) mammary duct and (c) teat regions of one bovine udder: \bigcirc , total binding; \square , non-specific binding; \blacktriangle , specific binding. Each point is the mean of triplicate determinations.

The α_2 -adrenergic competitors rauwolscine, clonidine, and yohimbine had low, and similar, K_i values for displacement of specific ³H-rauwolscine binding (Table 2). Inhibition potency of the non-selective α -adrenergic antagonist phentolamine was of the same order of magnitude, whereas the α_1 -antagonist prazosin was about 100 times less potent than unlabelled rauwolscine for inhibition of ³H-rauwolscine binding (P < 0.05). Displacement by isoproterenol of ³H-rauwolscine binding was in agreement with displacement of ³H-prazosine from its binding sites. The order of potency for inhibition of ³H-rauwolscine binding was rauwolscine > clonidine >

| Ligand | Region | Maximal binding capacity (B _{max}), fmol/mg protein | Dissociation constant $(K_{\rm D})$, nm | Hill coefficient (h) |
|----------------------------|--------------|---|--|-------------------------|
| ³ H-prazosin | Teat | $126.4 \pm 25.4^{ m a}$ | 0.38 ± 0.03 | 0.92 ± 0.06 |
| (7 udders) | Mammary duct | 38.8 ± 6.3^{b} | 0.34 ± 0.02 | 0.97 ± 0.03 |
| | Parenchyma | 7·8±3·4° | 0.58 ± 0.13 | 1·01 ± 0·01 |
| ³ H-rauwolscine | Teat | 130.1 ± 21.6^{a} | 4.42 ± 0.68 | 1.05 ± 0.08 |
| (4 udders) | Mammary duct | $23 \cdot 4 \pm 9 \cdot 4^{\mathrm{b}}$ | $4\cdot38\pm1\cdot37$ | 1.09 ± 0.04 |
| | Parenchyma | < 3.0 | | |
| ³ H-DHA | Teat | 123.3 ± 21.2^{a} | 1.13 ± 0.23 | 0.96 ± 0.001 |
| (7 udders) | Mammary duct | 119.1 ± 21.8^{a} | 1.41 ± 0.12 | 0.97 ± 0.01 |
| | Parenchyma | 23.7 ± 5.3^{b} | 1.78 ± 0.43 | 1·07 ± 0·03 |

Table 4. Values from saturation assays of ³H-prazosin, ³H-rauwolscine and ³Hdihydroalprenolol binding on membranes of different regions of the udder

(Values are means \pm sem for no. of udders shown)

^{a, b, c} Means without common superscript letters were significantly different between regions within ligand (P < 0.05).

phentolamine > yohimbine > prazosin > adrenalin > noradrenaline > phenylephrine > dopamine > isoproterenol. Hill coefficients were in the same range as in 3 H-prazosin binding studies.

The unlabelled non-selective β -antagonists alprenolol and propranolol had the lowest K_i values for specific ³H-DHA binding sites, followed by the β_2 -adrenergic agonist clenbuterol (Table 3). The β_2 -agonist fenoterol and the non-selective β -agonist isoproterenol were about 100-fold more potent than the β_1 -antagonist atenolol for inhibition of ³H-DHA binding (P < 0.05). The order of potency for inhibition of ³H-DHA binding was alprenolol > propranolol > clenbuterol > fenoterol > isoproterenol > adrenalin > noradrenaline > atenolol > dopamine. Inhibition of ³H-DHA binding was investigated with different isomers of propranolol. The order of potency was dependent on stereospecificity: (-)-propranolol > (\pm)-propranolol (P < 0.05) > (+)-propranolol (P < 0.05; Table 3). Most of the substances competed for ³H-DHA binding with Hill coefficients close to one, except for (-)-propranolol and atenolol.

Saturation binding assays

The specific binding of ³H-prazosin to membrane suspensions was a saturable process (Fig. 2). There was a decrease in the density of ³H-prazosin binding sites from the teat region to the duct region to the parenchyma region (P < 0.05; Table 4). K_D values of the three udder regions were in the same range. Hill coefficients were all close to one.

The specific binding of ³H-rauwolscine to membrane suspensions was a saturable process. There was a decrease in the number of ³H-rauwolscine binding sites from the teat to the duct region (P < 0.05; Table 4). In the parenchyma region no specific ³H-rauwolscine binding could be measured. $K_{\rm D}$ values of the teat and duct region were in the same range. Hill coefficients were all close to one.

The specific binding of ³H-DHA was saturable for each udder region. Numbers of binding sites for ³H-DHA were similar in the teat and duct regions, but markedly lower in the parenchyma region (P < 0.05; Table 4). K_D values of the three udder regions were of the same order. However, K_D values for ³H-prazosin, ³H-rauwolscine and ³H-DHA binding were significantly different (P < 0.05). All Hill coefficients were close to one.

DISCUSSION

Kinetic experiments demonstrated the reversibility of ³H-prazosin, ³H-rauwolscine and ³H-DHA binding. Association of ³H-rauwolscine had to be studied at 25 °C and an excess of unlabelled phentolamine had to be added after only 6 min because ³H-rauwolscine binding was no longer reversible with incubation for longer times or at 37 °C. This was surprising, because competitive binding assays with simultaneous addition of rauwolscine and phentolamine at 37 °C caused inhibition of ³H-rauwolscine binding.

Three different adrenergic receptor subtypes $(\alpha_1, \alpha_2, \beta_2)$ could be identified by inhibition of specific ³H-prazosin, ³H-rauwolscine and ³H-DHA binding by different subtype-specific drugs.

From the characteristics of Scatchard and Hill plots, ³H-prazosin, ³H-rauwolscine and ³H-DHA saturation binding indicated interaction with only a single binding site, identified as α_1 -, α_2 - and β_2 -adrenergic receptor subtypes respectively. K_D values calculated for ³H-prazosin, ³H-rauwolscine and ³H-DHA binding were significantly different from each other. On the other hand, K_D values of the same ³H-ligand for teat, duct and parenchyma region were of the same order. Numbers of α_1 -, α_2 - and β_2 -adrenergic receptors of membranes obtained from the teat region agreed well with the results of Roets *et al.* (1984) and Roets & Peeters (1985, 1986). Our studies demonstrate that there are additional regions in the bovine mammary gland containing α_1 -, α_2 - and β_2 -adrenergic receptors, localized in the area around the gland cistern including the large mammary ducts. However, it is important to note that the parenchyma region contained only very small numbers of adrenergic receptors. Thus, adrenergic receptors were mainly present in the milk purging system of the udder.

That α -adrenergic receptors were specifically localized on smooth muscle cells of mammary ducts cannot be stated with absolute certainty. If we assume their presence, several *in vivo* studies on inhibition of milk removal after α -Ag administration and in stress situations may be explained. The connection between mammary parenchyma and gland cistern may be interrupted completely or in part by α -adrenergic receptor-stimulated smooth muscle contractions of the mammary ducts. Although oxytocin is released (Blum *et al.* 1989) and may cause myoepithelial contraction, no milk or only small amounts are transported to the gland cistern and only milk already available in the teat and gland cistern can be removed during milking, a view supported by studies on effects of the α -Ag phenylephrine on milk removal and intramammary pressure before and after teat stimulation (Bruckmaier *et al.* 1991). Because the number of α -adrenergic receptors in particular was very low in mammary parenchyma, the influence of the adrenergic system on this region of the mammary gland is probably relatively small.

It has been proposed that under stress conditions or after α -Ag administration oxytocin may not reach the receptors of myoepithelial cells because of decreasing blood flow to the udder due to vascular contraction after catecholamine administration through interaction with α -adrenergic receptors in the vascular bed (Dhondt *et al.* 1976; Lefcourt & Akers, 1984; Gorewit & Aromando, 1985). However, only a very small increase of oxytocin concentration in blood is necessary to elicit milk ejection (Schams *et al.* 1984). Furthermore, vasoconstriction under stress conditions is hardly the main reason for peripheral inhibition of milk removal, because our study demonstrated only small numbers of α - and β -adrenergic receptors in the mammary parenchyma. From *in vivo* studies (Bruckmaier *et al.* 1991), it also appears unlikely that α -Ag have an effect on the responsiveness of myoepithelial cells to oxytocin. Because of the significantly higher density of α -adrenergic receptors in teat tissue and the region of the large mammary ducts, peripheral disturbed milk removal by α -Ag is primarily caused by constriction of the milk purging system of the udder.

It has been shown that administration of β -Ag has a favourable effect on milk flow as a consequence of teat muscle relaxation (Bernabé & Peeters, 1980; Hamann, 1981; Vandeputte-Van Messom *et al.* 1986). Through interaction with the numerous β -adrenergic receptors in the large mammary ducts, probably localized on smooth muscles, relaxation of the large mammary ducts is likely. Thus, after β -Ag administration, transport of milk from parenchyma tissue to the cistern is facilitated by enlarged mammary ducts and milk withdrawal accelerated after milk ejection (Blum *et al.* 1989; Bruckmaier *et al.* 1991). Mielke *et al.* (1991) have suggested the possibility of milk ejection after administration of β -Ag. However, lack of increased intramammary pressure by the administration of a β -Ag contradicts these findings (Bruckmaier *et al.* 1991).

We conclude that α - and β -adrenergic receptors are present not only in teat walls, but also around the gland cistern and large milk ducts. Thus, the modulation of milk ejection and milk removal by the adrenergic system occurs not only during the milk's passage through the teat cavity, but also very probably during its passage from gland parenchyma to the gland cistern.

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