# DIFFERENCES IN THE STEROID-BINDING SITE SPECIFICITIES OF RAT PROSTATE ANDROGEN RECEPTOR AND EPIDIDYMAL ANDROGEN-BINDING PROTEIN (ABP)\*

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## **SUMMARY**

The steroid specificity of androgen-binding proteins is crucial in determining their physiological functions and is important to consider for designing synthetic hormones and anti-hormones. Two proteins which bind testosterone and dihydrotestosterone with high affinity in the rat have been studied: the intracellular androgen receptor from ventral prostate and the epididymal androgen-binding protein (ABP). The binding sites were characterized in castrated animals by charcoal assay and gel electrophoresis, based on the specific binding of  $[^{3}H]$ -methyltrienolone in prostate cytosol and of  $[^{3}H]$ -dihydrotestosterone in epididymal extracts. In both tissues, kinetic and equilibrium data were compatible with a homogeneous class of sites having the expected physico-chemical properties. Competition experiments were then performed in order to assess the ability of both proteins to bind 35 steroids including androstane, pregnane and C-21 hydroxypregnane derivatives. Double-reciprocal plots suggested that all active compounds bound to the same site as the radioactive Iigand. A comparison of their apparent affinity constants indicated that the binding sites on the receptor and on ABP clearly differ in their steroid specificity. The differences involve the interaction of these sites with both extremities of the testosterone molecule since substituents either on the A-ring (e.g. 1-CH<sub>3</sub>) or on the D-ring (e.g. 17 $\alpha$ -CH<sub>3</sub>) may have opposite effects on ABP and receptor binding. Also, the presence of a pregnane side-chain is less favourable for the binding to ABP than to the androgen receptor. Thus, some compounds are very specific for the androgen receptor, while others have an affinity for ABP one order of magnitude higher than for the androgen receptor. These data may lead to a better understanding of the mechanism of the interaction of androgens and anti-androgens with their binding proteins. They also suggest that steroid analogues with selective high affinity for either the androgen receptor or ABP could be developed.

## **I. INTRODUCTION**

A variety of synthetic compounds are capable of interacting with the steroid binding site on the intracellular receptors for steroid hormones. Some compounds mimic the naturally occurring hormone and are called agonists. Others do not activate the receptor and their binding does not elicit a hormonal effect. These antagonists are therefore capable of competing with natural or synthetic agonists and of inhibiting their actions. Thus, anti-mineralocorticoids, antiestrogens and anti-androgens which act at the receptor level have become available. Development of such synthetic hormones or anti-hormones, and understanding their possible side-effects requires a knowfedge of both the steroid specificity of the receptor and the affinities of antagonists relative to the agonists fl].

Androgens control at least three distinct physiological processes perhaps via different androgen-binding proteins: development and maintenance of secondary sex organs, spermatogenesis and sperm maturation, and skeletal muscle metabolism. Apart from the sex steroid-binding proteins in plasma, two androgen-binding proteins have been characterized: the intracellular androgen receptor such as that in rat prostate and the androgen-binding protein (ABP) found in the testis and epididymis [Z, 31. ABP is produced by the Sertoli cells upon stimulation by FSH and secreted into the seminiferous tubular fluid [4]. It is transported to the epididymis by the efferent duct fluid where it seems to be degraded or reabsorbed since the level of ABP is highest in the caput epididymis and can no longer be detected beyond the cauda [5]. In mammals spermatozoa acquire fertilizing ability when passing through the epididymis. In rabbits and guinea pigs sperm maturation requires androgens acting via the epididymal tissue  $[6, 7]$ . It has been suggested that ABP because of its high androgen binding capacity (60 ng testosterone/ml tubular fluid) might serve as an androgen-concentrating factor essential for sperm maturation in the epidi-

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dymis [4]. High local concentrations of testosterone are also required for spermatogenesis in the seminiferous tubules [8]. The presence of ABP in close proximity to these androgen target cells would make testosterone available for uptake, conversion to the active metabolite Sa-dihydrotestosterone (DHT) and binding to intracellular receptors  $[4]$ .

Thus, it is important to determine the steroid-binding specificities of the androgen-receptor and of ABP. Differences between the two might allow one to use steroid analogues as specific probes for further studying the physiology of ABP. Should the formation of an ABP-testosterone complex prove to be an absolute requirement for sperm maturation, preventing androgens from binding to ABP might provide a means of reversibly inducing infertility in the male. This could be achieved by competitive inhibition provided the competing drug does not bind to intracellular androgen receptors which would lead to unacceptable side effects.

fn the present report we have approached these problems by comparing the affinities of 86 steroids for rat prostate androgen receptor and epididymal ABP. We show that, although both proteins bind natural androgens with similar affinities, they differ clearly in their binding properties. We also discuss how the specificity for either protein critically depends on some substituents of the steroid molecule.

#### II. **MATERtALS AND METHODS**

## (a) Chemicals

TEG-buffer contained 2OmM Tris, I.5 mM EDTA and  $10\%$  (v/v) glycerol, pH 7.4 with or without 0.25 mM dithiothreitol.  $[^3H]$ -DHT  $[1,2,3H]$ -17 $\beta$ -hy $drows-5\alpha$ -androstan-3-one), S.A. 56 or 60 Ci/mmol, was purchased from the Radiochemical Centre, Amersham, U.K.  $[^3H]$ -Methyltrienolone (R 1881)  $[6,7 \frac{3H}{4}]$ - $17\beta$ -hydroxy-17 $\alpha$ -methyl-estra-4,9,11-triene-3-one), S.A. 55.5 Ci/mmol was generously supplied by Dr. J. P. Raynaud, Roussel-UCLAF, Romainville, France. Unlabelled steroids were either purchased from Steraloids, Pawling, N.V., U.S.A. and Sigma Co. St Louis, Missouri, U.S.A., or were gifts from Roussel-UCLAF, Romainviiie, France, Shering A.G., Berlin, Germany, Glaxo Research Ltd., Greenford, Middlesex, U.K., Merck, Sharp and Dohme, Rahway, N.J., U.S.A. and Ayerst Lab. McKenna and Harrison Ltd. Montreal, Canada.

Steroids were dissolved in absolute ethanol to a concentration (2-4mM) which was verified by spectrophotometry on the basis of molar extinetion coefficients, when available. Final dilutions were made in TEG-buffer. Al1 other chemicals were of analytic grade.

#### (b) Animals and tissues

from Wistar rats (8-14 weeks) 20-24 h after bilateral or  $[^3H]$ -MT respectively. Kapp is the apparent orchidectomy via the scrotal route under nembutal affinity constant of  $[^{3}H]$ -DHT or  $[^{3}H]$ -MT in the

anaesthesia. Epididymal tissue was minced with scissors and homogenized in 3 vol. (w/v) of TEG-buffer without dithiothreitol using a motor-driven Teflonglass homogenizer. Likewise prostate homogenates were prepared in TEG buffer containing dithiothreitol. All preparations procedures were carried out at 0°C. Homogenates were centrifuged at 200,000  $g$  for I h at  $0^{\circ}$ C to obtain the cytosol. Rat prostate cytosol was used immediately, whereas epididymis cytosol was usually stored at  $-20^{\circ}$  and incubated before use for 30 min at 50 $^{\circ}$ C [9] (in later experiments 30 $^{\circ}$ C, see below) to inactivate androgen receptors.

## (c) Binding studies bp chareoaf *assq*

*1. ABP.* Aliquots of cytosol (0.1 ml) were incubated in a total volume of  $0.4$ ml with  $[^3H]$ -DHT (0.5-5 nM) in TEG buffer *at 0°C.* Nonspecific binding was assessed in parallel incubations containing, in addition,  $10^{-5}$  M unlabelled DHT. After incubation (3 h for equilibrium studies) hormone was removed by adding 0.2 ml of activated charcoal suspension (Norit A, 50mg/ml, dextran 5 mg/ml). Tubes were briefly vortexed (3 sec) and the charcoal sedimented immediately by centrifugation  $(5 \text{ min}, 2000 \text{ g})$ . Bound radioactivity was counted in 0.3 ml of the resulting supernatant.

2. *Androgen receptor.* The accurate assessment of [<sup>3</sup>H]-DHT binding to prostate androgen receptors was complicated by high levels of nonspecific binding. We therefore employed the synthetic androgen methyltrienolone  $(MT)$  which has a higher affinity for the androgen receptor than the natural hormones testosterone and DHT and also exhibits lower nonspecific binding [IO]. Incubations for receptor binding of  $[^{3}H]$ -MT(0.5-5 nM) were prepared at  $0^{\circ}C$  as described for ABP. After 18-20 h (in equilibrium studies) 0.15 ml of charcoaf suspension was added and the tubes vortexed briefly. After 10 min 0.05 ml of ethanol was added [11]. The tubes were vortexed immediately and incubated for a further 20min. The charcoal was then sedimented and aliquots of the supernatant (0.3 ml) taken for counting radioactivity. Nonspecific binding of  $\lceil$ <sup>3</sup>H]-MT was determined in parallel incubations containing, in addition,  $10^{-5}$  M unlabelled MT

3. Affinity of competitors. This was determined at equilibrium using a constant concentration of unlabelled competitor against four different concentrations of radioactive ligand. Binding data were plotted according to Lineweaver-Burk and the apparent equilibrium dissociation constants of  $[^3H]$ -DHT or  $[3H]$ -MT in the absence and presence of test compounds were determined. The equilibrium dissociation constant  $(K_i)$  of competitors was calculated using the formula

$$
K_i = \frac{K_D \times F_i}{\text{Kapp} - K_D} \times \frac{K_D}{K_D}
$$

Epididymis and ventral prostate were obtained where  $K_p$  is the dissociation constant of  $[^3H]$ -DHT

presence of unlabelled competitor,  $F_i$  is the concentration of free competitor and  $\overline{K}_p$  was the mean value of the dissociation constant of the respective radioactive ligands from all our experiments.  $F_i$  was taken as the total concentration of inhibitor added as the concentration of bound inhibitor is negligible as compared to the former. Scatchard plots of specific binding data of  $[^3H]$ -DHT and  $[^3H]$ -MT were routinely done and they also demonstrated the presence of only one class of high affinity binding sites.

## (d) *Polyucrylamide gel electrophoresis (PAGE)*

Five percent Polyacrylamide gels  $(6 \times 50 \text{ mm})$  containing  $10\%$  glycerol were prepared according to Tindall *et a/.[9].* Where indicated hormones were added in glycerol to the gel solution prior to polymerisation to give a uniform concentration of hormone in the gel.

Aliquots of cytosol (0.1 ml) were layered directly over the gel and electrophoresis was run at  $1-2$  mA per tube with tubes immersed in the lower buffer cooled to  $2^{\circ}$ C. Upper and lower buffers were  $0.6\%$ Tris,  $2.9\%$  glycine in water (pH 8.6) and bromophenol blue was added to the upper buffer as an electrophoretic marker. Electrophoresis was stopped when the bromophenol blue-band had reached the bottom of the tubes and gels were sliced into 1 mm transverse sections. The sections were placed in counting vials, 10 ml of scintillation fluid added and left overnight at room temperature before counting. Relative mobilities  $(R_x)$  of proteins were expressed as the ratio of distances covered by the protein and the bromophenol blue band, respectively. Gels containing marker proteins were fixed and stained in a solution of  $45\%$  methanol and  $9.2\%$  acetic acid containing 0.25% Coomassie blue.

#### (e) *Other assays*

Protein was determined by the method of Lowry *et a/.[121* using bovine serum albumin as a standard. Radioactivity was counted by liquid scintillation spectrometry in 10 or 5 ml of scintillation fluid (Aqualuma, Lumac Systems A.G., Basel, Switzerland). Counts were corrected for quenching and systematically converted to mol of steroid using an external standard system of calibration.

### III. RESULTS AND DISCUSSION

#### (a) *Characterization of the binding proteins*

Screening a large number of compounds for androgen receptor and ABP binding called for a rapid and reproducible assay. Also, we wanted to determine whether these compounds bound to the same binding site as the natural steroid and, if so, what was their affinity constant.

Firstly, the kinetics of the binding of  $[^3H]$ -DHT to ABP and of  $[^3H]$ -MT to the androgen receptor were studied to establish optimal assay conditions. Rates of association were determined by incubating

for various time-periods prostate or epididymis cytosol containing a known concentration of androgen receptor or ABP with a single concentration of their respective radioactive ligand. Dissociation rates were studied by determining the concentration of bound radioactive hormone at various times after the addition of a large excess of unlabelled hormone. Confirming earlier work [9] it was found that the androgen receptor and ABP differ greatly in their dissociation rate constant, with  $[^3H]$ -DHT-ABP complexes having a half-time for dissociation of about 4 min as compared to 23 h for  $[^3H]$ -MT-androgen receptor complex. The charcoal adsorption of free hormone in assays for ABP binding is therefore a critical step and the time of incubations with charcoal has to be the shortest to allow for efficient removal of free steroid without substantial dissociation of [<sup>3</sup>H]-DHT-ABP complexes. Consistent with the second-order association rates and the first-order dissociation rates, Scatchard plots of the equilibrium binding of  $\lceil$ <sup>3</sup>H]-MT in prostate cytosol and C3H]-DHT in epididymis cytosol were compatible with the presence of only one species of non-interacting binding sites showing high affinity and low capacity (Fig. I). Mean values for the equilibrium dissociation constants were  $0.78 \text{ nM } \pm 0.07 \text{ S.E.M.}$  and 2.32 nM  $\pm$  0.24 S.E.M. for the MT-androgen receptor complex and the DHT-ABP complex, respectively. The mean concentrations of binding sites were 42  $\pm$  4.0 S.E.M. and 850  $\pm$  170 S.E.M. fmol/mg protein, respectively.

Secondly, we verified that binding to either the androgen receptor or ABP was destroyed after preincubation of prostate or epididymis cytosol with trypsin (50  $\mu$ g/ml at 4°C for 5 h). Blocking of thiol groups by preincubation with p-chloromercuribenzoic acid (PCMB 0.5 mM) prevented the binding of  $\lceil$ <sup>3</sup>H]-MT to androgen receptors whereas binding of  $[^3H]$ -DHT to ABP in epididymis cytosol was only slightly reduced. At concentrations over 0.1 mM, dithiotreitol inhibited ABP binding, a decrease of  $70\%$  being observed at IO mM. Binding activity of prostate cytosol was completely abolished after preincubation at  $30^{\circ}$ C, whereas only a slight reduction of the binding activity in epididymis cytosol was observed at this temperature. However, binding of  $[^3H]$ -DHT in epididymis cytosol was reduced at temperatures over  $30^{\circ}$ C, being only  $6\%$  of control levels after preincubation at 50°C.

Thirdly, by PAGE, the androgen-binding component from epididymis cytosol had a relative mobility of  $R_x = 0.49{\text -}0.53$  ( $n = 6$ ). In the presence of an excess of unlabelled DHT in the gel this peak was reduced to baseline levels of radioactivity (Fig. 2). An excess of unlabelled cyproterone acetate  $(1 \mu M)$  or preincubation with PCMB had no effect. Specifically bound radioactivity in 0.1 ml epididymis cytosol was 15,700d.p.m. determined by charcoal assay as compared to 16,260d.p.m. after subtraction of baseline radioactivity under the peak with  $R_x = 0.53$  (slice



Fig. 1. Equilibrium binding of  $[^{3}H]$ -DHT in epididymis cytosol and of  $[^{3}H]$ -methyltrienolone in rat prostate cytosol. Nonspecific binding (A) was subtracted from total binding to yield specific binding (0) to ABP and androgen receptor, respectively. Insets show Scatchard plots of specific binding data. Fo, F and B refer to total free, and receptor-bound steroid concentrations.

Nos 22-28, Fig. 2). Only under steady-state PAGE conditions a second peak of radioactivity  $(R_x = 0.71)$ , undetected by charcoal assay, was observed which comigrated with albumin marker proteins and was unaffected by an excess of unlabelled DHT in the gel. A peak was occasionally observed at  $R_x = 0.3$ which may correspond to the androgen receptor described by others in rat epididymis **cytosol** [9]. When prostate cytosol prelabelled with  $[^3H]$ -MT was analysed by PAGE only one peak of radioactivity with a relative mobility of  $R_x = 0.28$  was resolved. Binding was completely abolished in the presence of a 200-fold excess of unlabelled cyproterone acetate or by preincubation with PCMB.

Thus, in epididymal extracts and prostate cytosol, the binding sites detected by our charcoal assay procedure exhibited the characteristics assigned respect-

ively to ABP and to the androgen receptor by other investigators [5, 9, 13, 14].

# (b) *Steroid-binding specificity of androgen receptor and ABP*

Unlabelled compounds were first tested at a single concentration (10  $\mu$ M) for their ability to inhibit, under equilibrium conditions, the binding of  $[^3H]$ -DHT or  $[^3H]$ -MT to ABP or androgen receptor, respectively. The affinities  $(K_i)$  or active inhibitors for the two binding proteins were then determined at an appropriate concentration by competitive equilibrium binding assay. All Lineweaver-Burk plots were compatible with competitive inhibition of binding to the same binding site. Representative experiments are shown in Figure 3. Androstane, pregnane and C-21 hydroxypregnane derivatives were tested. Their affini-





Fig. 2. Steady-state polyacrylamide gel electrophoretic profile of [<sup>3</sup>H]-DHT binding in epididymis cytosol. 0.1 ml aliquots of cytosol (400  $\mu$ g protein) were layered over 5% polyacrylamide gels containing 0.96 nM [<sup>3</sup>H]-DHT without ( $\bullet$ —— $\bullet$ ) or with ( $\circ$ —— $\circ$ ) 10  $\mu$ M unlabelled DHT. Electrophoresis was  $(-)$  or with  $(O \longrightarrow O)$  10  $\mu$ M unlabelled DHT. Electrophoresis was performed as described in Methods.  $R_x$  = mobility of bound radioactivity relative to bromophenol blue (B.B.). The electrophoretic mobility of albumin (alb.) and ovalbumin (Ov.) run on parallel gels is indicated in the shaded area.

ties for ABP, in order of decreasing importance, and for the androgen receptor are listed in Tables 1-3.

Consistent with other work [2,3, 15) we find that ABP, which reportedly is a testosterone-binding protein *in* oiuo, has a higher affinity for DHT than for testosterone (compound 1.1. vs 1.6 in Table 1). In all cases of  $5\alpha$ -reduction (1.6 vs 1.1, 1.7 vs 1.3, 1.8 vs 1.2, 1.20 vs 1.19) the affinity for ABP was improved. The only androst-4-ene derivative which bound better than testosterone was 9-dehydrotestosterone (1.5). It is interesting that 9-dehydroandrostenedione (1.13) also binds better to ABP than its reduced analogue (1.20). The 12 compounds with the highest affinity for ABP are all  $17\beta$ -OH androstane derivatives (1.1-1.12). Then comes progesterone (2.1), showing that high affinity binding to ABP is compatible with the addition of a pregnane side-chain on the testosterone molecule. However, all the 21 pregnane derivatives tested (2.2-2.22) have a lower affinity than progesterone. Thus,  $\Delta_1$  (2.1 vs 2.3),  $6\alpha$ -CH<sub>3</sub> (2.1 vs 2.10, 2.5 vs 2.13, 2.11 vs 2.16),  $11\alpha$ -OH (2.1 vs 2.17),  $11\beta$ -OH (2.1 vs 2.12),  $16\alpha$ -CH<sub>3</sub> (2.1 vs 2.11, 2.5 vs 2.21, 2.10 vs 2.16),  $16\beta$ -CH<sub>3</sub> (2.1 vs 2.15) and  $17\alpha$ -OH (2.1 vs 2.5, 2.10 vs 2.13, 2.11 vs 2.21) substitutions all decrease the affinity. Yet, some additional substituents improve somewhat the binding of pregnane derivatives. It is the case for th  $9\alpha$ -fluoro (2.14 vs 2.6) and the  $16\alpha$ -OH (2.2 vs 2.5) groups. Hydroxylation of progesterone at C-21 (deoxycorticosterone, 3.11) provokes a 12-fold decrease in the binding of progesterone, unless additional substituents are also present. Thus, a 5 $\beta$ -configuration (3.1) or a 17 (or 16)  $\alpha$ -OH substitution (3.2 and 3.3) increase the affinity of deoxycorticosterone 7-fold and 3-fold respectively. In the androstane series, we find that a  $17\alpha$ -CH<sub>3</sub> substitution has little effect on the binding of testosterone (1.7 vs 1.6, 1.9 vs 1.8) or of DHT (1.3 vs 1.1). A Cl-2 unsaturation decreases the affinity of testosterone derivatives (1.7 vs 1.8, 1.6 vs 1.9) while it increases the affinity of DHT and its analogues (1.14 vs 1.10, 1.3 vs 1.2). Addition of a lla-OH (1.6 vs 1.16) or of a  $11\beta$ -OH group (1.6 vs 1.12, 1.20 vs 1.25) decreases the affinity 5-20-fold. A 11 $\beta$ -OH group also inhibits binding in the pregnane series (2.1 vs 2.12) while it stimulates it in the C-21 hydroxypregnane series (3.11 vs 3.5). ABP binding of androstane derivatives is reduced by the presence of a  $1$ -CH<sub>3</sub> group whether in quasi-equatorial (1.8 vs 1.15, 1.4 vs 1.22) or in  $\beta$ -configuration (1.1 vs 1.14). Finally, oxidation of the  $17\beta$ -OH group into a ketone (1.1 vs 1.19, 1.5 vs 1.13, 1.6 vs 1.20, 1.12 vs 1.25) or loss of either the 3-ketone  $(1.17 \text{ vs } 1.1)$  or the 17 $\beta$ -OH  $(1.6 \text{ vs } 1.18)$  lead to a 5@-2OO-fold decrease in affinity for ABP.

As to the methyltrienolone-binding component in prostate cytosol, its properties are similar to those of the androgen receptor described by others in this tissue [10, 16]. Of all the steroids tested, only methyltrienolone had a higher affinity than DHT for the androgen receptor. Testosterone had a slightly lower affinity than DHT. This is in agreement with other work using similar techniques [17]. The 11 steroids with the highest affinities are found in the series of androstane derivatives. As a rule,  $5\alpha$ -reduction increases the affinity (1.6 vs 1.1, 1.7 vs 1.3, 1.8 vs 1.2,



Fig. 3. Competitive inhibition of  $[^{3}H]$ -DHT binding to ABP and of  $[^{3}H]$ -methyltrienolone binding to the androgen receptor by various steroids. Specific binding of the radioactive steroids was determined in absence and presence of a constant concentration of nonradioactive competitors as indicated on the figure.

1.20 vs 1.19). However, **progesterone** (2.1) and four of its analogues (2.10, 2.7, 2.8, and 2.13) also bind with high affinity. Unfavourable substitutions on progesterone are  $11\beta$ -OH (2.1 vs 2.12),  $16\alpha$ -OH (2.2 vs 2.51 17a-OH (2.1 vs 2.5, 2.10 vs 2.13, 2.11 vs 2.21),  $16\alpha$ -CH<sub>3</sub> (2.1 vs 2.11, 2.10 vs 2.16, 2.5 vs 2.21) and  $16\beta$ -CH<sub>3</sub> (2.1 vs 2.15) groups as well as a Cl-2 unsaturation (2.1 vs 2.3). Transition to the C2l-hydroxypregnane series leads to a 3-fold decrease in affinity  $(3.11 \text{ vs } 2.1)$ . Desoximethasone  $(3.10)$  has a surprisingly high affinity as compared to the negligible binding of other C-21 hydroxypregnane derivatives. Addition of a  $17\alpha$ -OH group on desoximethasone (3.32) virtually inhibits the binding. In the androstane derivatives, an interesting finding is the unfavourable effect of the  $17\alpha$ -CH<sub>3</sub> (1.6 vs 1.7, 1.9 vs 1.8, 1.1 vs 1.3). The  $C1-2$  unsaturation either decreases (1.6 vs 1.9, 1.7 vs 1.8), does not change (1.3 vs 1.2) or increases  $(1.14 \text{ vs } 1.10)$  the affinity. An equatorial-  $(1.8$ vs 1.15, 1.4 vs 1.22) or a  $\beta$ -methyl group (1.1 vs 1.14) on Cl reduces the affinity considerably. Thus. one would predict that  $\Delta 1$ -DHT (not tested) binds with





Table 2. Affinity (equilibrium dissociation constant, nM) of pregnane derivatives for the androgen-binding protein (ABP) and the androgen receptor (AR)

Number	Trivial name	Systematic name	<b>ABP</b>	<b>AR</b>
2.1	Progesterone	4-Pregnene-3,20-dione	120	90
2.2	$16\alpha$ , $17\alpha$ -Dihydroxyprogesterone	$16\alpha$ , 17-Dihydroxy-4-pregnene-3,20-dione	425	> 10,000
2.3	1-Dehydroprogesterone	1,4-Pregnadiene-3,20-dione	600	430
2.4	$20\alpha$ -Hydroxyprogesterone	20x-Hydroxy-4-pregnen-3-one	645	535
2.5	$17\alpha$ -Hydroxyprogesterone	17-Hydroxy-4-pregnene-3,20-dione	645	1590
2.6	$6\alpha$ , 9 $\alpha$ -Difluoro-11 $\beta$ -Hydroxy-proges- terone acetonide	$6\alpha$ , 9 $\alpha$ -Difluoro-11 $\beta$ -hydroxy 16 $\alpha$ -17-isopro- pylidene-dioxy-4-pregnene-3,20-dione	800	2570
2.7	Chlormadinone acetate	6-Chloro-4,6-pregnadiene-3,20-dione-17 $\alpha$ -yl acetate	990	25
2.8	Cyproterone acetate	$6$ -Chloro-1 $\alpha$ , $2\alpha$ -methylene-4, 6-pregna- diene-3,20-dione-17 $\alpha$ -yl acetate	1060	37
2.9	$6.17\alpha$ -Dimethyl-6-dehydroprogesterone (Medrogestone)	$6,17\alpha$ -Dimethyl-1,6-pregnadiene-3,20-dione	1380	330
2.10	$6\alpha$ -Methylprogesterone	$6\alpha$ -Methyl-4-pregnene-3,20-dione	1470	19
2.11	16x-Methylprogesterone	$16\alpha$ -Methyl-4-pregnene-3,20-dione	2310	600
2.12	$11\beta$ -Hydroxyprogesterone	$11\beta$ -Hydroxy-4-pregnene-3,20-dione	2340	350
2.13	$6\alpha$ -Methyl-17-hydroxyprogesterone	$6\alpha$ -Methyl-17-hydroxy-4-pregnene-3,20- dione	2340	115
2.14	$6\alpha$ -Fluoro-11 $\beta$ -hydroxyprogesterone acetonide	6α-Fluoro-11β-hydroxy-16α, 17-iso-propyl idene-dioxy-4-pregnene-3,20-dione	2680	4940
2.15	$16\beta$ -Methylprogesterone	168-Methyl-4-pregnene-3.20-dione	2720	4320
2.16	$6\alpha$ , $16\alpha$ -Dimethylprogesterone	$6\alpha$ , $16\alpha$ -Dimethyl-4-Pregnene-3, 20-dione	3920	220
2.17	$11\alpha$ -Hydroxyprogesterone	$11\alpha$ -Hydroxy-4-pregnene-3,20-dione	5590	790
2.18	6x-Methyl-16-dehydroprogesterone	6x-Methyl-4,16-pregnadiene-3,20-dione	6610	1630
2.19	Alphaxalone	$3\alpha$ -Hydroxy-5 $\alpha$ -pregnane-11,20-dione	7250	>10,000
2.20	6x-Fluoro-11x-Hydroxyprogesterone acetonide	$6\alpha$ -Fluoro-11 $\alpha$ -hydroxy-16 $\alpha$ ,17-isopropyl idene-dioxy-4-pregnene-3,20-dione	>10,000	>10,000
2.21	16x-Methyl-17-hydroxyprogesterone	$16\alpha$ -Methyl-17-hydroxy-4-pregnene-3,20- dione	>10,000	4810
2.22	6x-Progesterone Ethylcarbamate	$6\alpha$ -Ethylcarbamate-4-Pregnene-3,20-dione	>10,000	>10,000







Fig. 4. Comparison of the affinity of various steroids for the androgen receptor and for **ABP.** Numbers in the symbols refer to compound numbers in Table 1 (O), Table 2 ( $\Delta$ ) and Table 3 ( $\square$ ). Compounds which have a negligible affinity  $(K_D > 1 \mu M)$  for both binding sites are not shown.

very high affinity since its 1-CH<sub>3</sub> analogue (1.10) binds almost as well as testosterone itself. This was found to be the case by other investigators [IS]. Finally, substitution of the  $17\beta$ -OH by a ketone group (1.1 vs 1.19, 1.5 vs 1.13, 1.6 vs 1.20, 1.12 vs 1.25) decreases the affinity dramatically while loss of the  $17\beta$ -OH group (1.6 vs 1.18) virtually abolishes binding. Loss of the 3-ketone (1.1 vs 1.17) is still compatible with binding. Addition of a  $11\beta$ -OH (1.6 vs 1.12, 1.20 vs 1.25) or of a  $11\alpha$ -OH (1.6 vs 1.16) can reduce the affinity by 1O-20-fold. The binding to rat prostate androgen receptor of a few of the steroids examined here has been studied by other investigators  $[17-22]$  who measured their relative inhibitory effect on the binding of radioactive DHT or MT. Although the "percentage of inhibition" data found in the literature are not always easy to compare, except in [19], with our  $K_i$  values, they are in general agreement.

If we now compare the steroid-binding specificity of the androgen receptor and of ABP, we find that both proteins have in common a high affinity for the natural androgens DHT and testosterone. However, the mechanisms involved in the binding must somehow differ for the two proteins, since the very fast off-rate of DHT from ABP is compatible with high affinity only because it is compensated by a very high rate of association. These rapid kinetics are in keeping with the proposed androgen-carrier function of ABP. By contrast, androgens associate with and dissociate from their intracellular receptor very slowly and this accounts for the selective retention of the hormone in target cells. Thus, it is not surprising that differences in the binding site properties become also apparent when discrimination between different steroid structures is at stake. This is illustrated in Fig. 4. It can be seen that several androstane derivatives have an affinity higher for ABP than for the androgen receptor. On the other hand, some compounds bind better to the androgen receptor than to ABP. The most striking example is methyltrienolone (1.11) with a ratio of 60-fold, Others have ratios of 7-14-fold. Here, substituents unfavourable to ABP are the 1-CH<sub>3</sub> (1.10, 1.14) and  $11\alpha$ -OH (1.16). The presence of a 17-ketone (1.19) is also tolerated better by the androgen receptor than by ABP. As to the pregnane derivatives tested, seventy percent bind better to the androgen receptor than to ABP. They include the known anti-androgens chlormadinone acetate  $(2.7)$ , cyproterone acetate  $(2.8)$  and medrogestone  $(2.9)$ . The anti-androgenic activity of spironolactones has been ascribed to their binding to the androgen receptor [23]. We find that aldactone (1.23) has a low affinity for this receptor and binds better to ABP. The  $1 - CH_3 - \Delta 1$ -substitution appeared interesting because the corticoid derivative tested (3.20) bound neither to the glucocorticoid receptor [24] nor to the androgen receptor and yet it interacted somewhat with ABP. We thought that  $1 - CH_3 - \Delta 1$ -androgens might bind with high affinity to ABP while losing

**their affinity for intracellular receptors. Such compounds however, not only bind with high affinity 10 ABP; they even bind better to the androgen receptor**  (1.10). **Finally, the results obtained with compound**  3.1 suggest that a 5*ß*-configuration may selectively **favor ABP binding** 

**In conclusion. we have shown that ABP and the androgen receptor clearly differ in their steroid specificity. Among the steroids tested, 35 compounds including pregnane and C-21 hydroxypregnane derivatives could bind to the androgen receptor at concentrations achievable in** *rivo* ( $K<sub>D</sub> < 1 \mu M$ ). Our data do **not allow distinction between androgens and antiandrogens. However, they should lay the ground for a comprehensive study of the structural requirements for highly specific binding of steroid analogues to the**  androgen receptor. Some substitutions inhibit binding **to ABP. Thus, androgen agonists or antagonists could be designed which would not bind to the circulating sex hormone binding globulin (TeBG) since in species. such as man. where this protein occurs it resem**bles [25] or may be identical to ABP [26]. On the **other hand. twelve compounds among those which**  had a good affinity for ABP  $(K_D < 1 \mu M)$  did bind **better to this protein than to the androgen receptor. Derivatives of such compounds which would preferentially or exclusively bind to ABP might be helpful for further studying the role of this protein in physiology and in pathology.** 

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