Aldosterone, Dexamethasone and Triamcinolone Activate African Lungfish Mineralocorticoid

Receptor: Increased Activation After Removal of the Amino-Terminal Domain

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ABSTRACT. A distinct mineralocorticoid receptor (MR) ortholog first appears in cartilaginous fishes, such as sharks, skates, rays and chimaeras. Although aldosterone, the main physiological mineralocorticoid in humans and other terrestrial vertebrates, is a transcriptional activator of skate MR and elephant shark MR, aldosterone is not synthesized by cartilaginous fishes. Aldosterone, first appears in lungfish, which are lobe-finned fish that are forerunners of terrestrial vertebrates. Aldosterone activation of the MR regulates internal homeostasis of water, sodium and potassium, which was critical in the conquest of land by vertebrates. We studied transcriptional activation of the slender African lungfish (*Protopterus dolloi*) MR by aldosterone, other corticosteroids and progesterone and find that aldosterone, 11-deoxycorticosterone, 11-deoxycortisol and progesterone have half-maximal responses (EC50s) below 1 nM and are potential physiological mineralocorticoids. In contrast, EC50s for corticosterone and cortisol were 23 nM and 66 nM, respectively. Unexpectedly, truncated lungfish MR, consisting of the DNA-binding domain, hinge domain and steroid-binding domain, had a stronger response to aldosterone, other corticosteroids and progesterone than did full-length lungfish MR, indicating that an allosteric action of the N-terminal domain represses steroid activation of lungfish MR. This contrasts to human MR in which the N-terminal domain contains an activation function. BLAST searches of GenBank did not retrieve a GR ortholog, leading us to test dexamethasone and triamcinolone for activation of lungfish MR. At 10 nM, both synthetic glucocorticoids are about 4-fold stronger than 10 nM aldosterone in activating full-length lungfish MR, leading us to propose that lungfish MR also functions as a GR.

Keywords: Lungfish, Lobe-finned Fish, Terrestrial Vertebrates, Aldosterone evolution; mineralocorticoid receptor evolution; evolution

INTRODUCTION

The mineralocorticoid receptor (MR) and glucocorticoid receptor (GR) belong to the nuclear receptor family, a diverse group of transcription factors that arose in multicellular animals [1–3]. The MR and GR have key roles in the physiology of humans and other terrestrial vertebrates and fish [4–11]. The MR and GR evolved from an ancestral corticoid receptor (CR) in a jawless fish (cyclostome), which has descendants in modern lampreys and hagfish [12–14].

A distinct MR and GR first appear in cartilaginous fishes (Chondrichthyes) [1,13,15–17], which

diverged from bony vertebrates about 450 million years ago [18,19].

Aldosterone is the main physiological mineralocorticoid in humans and other terrestrial

vertebrates [5,6,9,20–23]. Aldosterone activation of the MR in the kidney regulates salt and

water homeostasis by promoting sodium and water reabsorption and potassium secretion, a

mechanism that conserves salt and water. Thus, it is puzzling that aldosterone is a potent

transcriptional activator of lamprey and hagfish CRs [15], skate MR [16] and elephant shark MR

[13,24] because aldosterone is not synthesized by lampreys [15], cartilaginous fishes or ray

finned fishes [25]. Aldosterone first appears in lungfish [26–28], which are lobe-finned fish that

are forerunners of terrestrial vertebrates [29–31]. The key phylogenetic position of lungfish in

the transition of vertebrates from water to land [27,29,30,32] and the role of the MR in

maintaining internal electrolyte homeostasis [5,8,33,34] motivated us to investigate the response

of the slender African lungfish MR to aldosterone, cortisol and other corticosteroids (Figure 1),

as well as activation by progestins, which also activate elephant shark MR [24], ray-finned fish

MR [35–39] and chicken MR [24,40].

Figure 1. Structures of Corticosteroids, Dexamethasone, Triamcinolone, Progesterone and Spironolactone. Aldosterone and 11-deoxycorticosterone are mineralocorticoids [41]. 11-deoxycortisol is a mineralocorticoid in lamprey [42,43]. Cortisol and corticosterone are glucocorticoids in terrestrial vertebrates and ray-finned fish [41,44]. Dexamethasone and triamcinolone are synthetic glucocorticoids. Progesterone is female reproductive steroid that also is important in male physiology [45,46]. Spironolactone is a mineralocorticoid antagonist in humans [47,48].

Our investigation also uncovered an unexpected role of the N-terminal domain (NTD) of lungfish MR in inhibiting transcriptional activation by steroids. Like other steroid receptors, lungfish MR is a multi-domain protein, consisting of an NTD (domains A and B), a central DNA-binding domain (DBD) (domain C), a hinge domain (D) and a C-terminal ligand-binding domain (LBD) (domain E) [49–51] (Figure 2). The NTD in the human MR contains an activation function domain (AF1), which is split into two segments [49–51]. (Figure 2). As described below, we find that in contrast to human MR [24,49,51,52], the NTD in full-length lungfish MR reduces steroid-mediated activation of lungfish MR, compared to truncated lungfish MR-CDE in cells transfected with a 3X-Tyrosine Amino Transferase (TAT3) promoter [53].

We also find that lungfish MR is activated by dexamethasone. At 10 nM, dexamethasone activates full-length lungfish MR and truncated lungfish MR with a signal that is 4-fold and 6-fold stronger, respectively, than that of 10 nM aldosterone. This strong response to dexamethasone and the absence of a lungfish GR sequence after a BLAST search of GenBank leads us to propose that lungfish MR also functions as a GR.

Figure 2. Comparison of the functional domains of lungfish MR to corresponding domains in selected vertebrate MRs (human, coelacanth, elephant shark, *Xenopus*, zebrafish) and GRs (human, coelacanth, elephant shark). Lungfish MR and human MR have 97% and 74% identity in DBD and LBD, respectively. Lungfish MR and elephant shark MR have 92% and 68% identity in DBD and LBD, respectively. This strong conservation of the DBD and LBD contrasts with the low sequence identity of 44% and 47% between their NTDs. There are similar % identities between corresponding domains in lungfish MR and other MRs.

RESULTS

Transcriptional activation of full-length and truncated lungfish MR by corticosteroids, progestins and dexamethasone.

We screened a panel of steroids (Figure 1) at 10 nM for transcriptional activation of full-length and truncated lungfish MR containing the CDE domains (MR-CDE) using two promoters: 2X-Mouse Mammary Tumor Virus (MMTV) [54,55] and TAT3 [53], which along with plasmids for both lungfish MRs were transfected into HEK293 cells.

As shown in Figure 3A, there was about 2 to 3-fold activation by 10 nM aldosterone,

other corticosteroids or progesterone of full-length lungfish MR using the MMTV-luc reporter

and less steroid activation of lungfish MR-CDE (Figure 3B).

Interestingly, compared to activation of full-length lungfish MR with the MMTV

promoter (Figure 3A), transcriptional activation of full-length lungfish MR with a TAT3

promoter and 10 nM aldosterone, other physiological corticosteroids or dexamethasone increased

by about 1.5 to 2-fold (Figure 3C). Unexpectedly, lungfish MR-CDE with the TAT3 promoter

had an additional 2-fold increase in activation by all corticosteroids (Figure 3D). Progesterone

activated lungfish MR in accord with the prediction of Fuller et al. [37,39,56]. Together, these

experiments show that removal of the NTD increases corticosteroid and progesterone activation

of lungfish MR in the presence of the TAT3 promoter.

Our results with dexamethasone, which activates human MR [52,57–59], were

unexpected. To our surprise, compared to aldosterone, dexamethasone was about 3-fold and 6-

fold more active, respectively, in activating full-length lungfish MR (Figure3C) and truncated

lungfish MR (Figure 3D) with the TAT3 promoter. Moreover, both cortisol and corticosterone

have stronger fold-activation than does aldosterone of lungfish MR using the TAT3 promoter.

Under these conditions, lungfish MR appears to have a GR-like response to steroids.

[10 nM Steroid]

Figure 3. Ligand specificity of full-length and truncated lungfish MR.
Plasmids for full-length lungfish MR or truncated lungfish MR (MR-CDE) were expressed in
HEK293 cells with an MMTV-luciferase reporter or a TAT3-luciferase reporter. Transfected
cells were treated with either 10 nM aldosterone, cortisol, 11-deoxycortisol, corticosterone, 11-
deoxycorticosterone, progesterone, dexamethasone or vehicle alone (DMSO). Results are
expressed as means ± SEM, n=3. Y-axis indicates fold-activation compared to the activity of
control vector with vehicle alone as 1. A. Full-length lungfish MR with MMTV-luciferase. B.
Truncated lungfish MR (MR-CDE) with MMTV-luciferase. C. Full-length lungfish MR with
TAT3-luciferase. D. Truncated lungfish MR (MR-CDE) with TAT3-luciferase.

 Spironolactone and eplerenone are transcriptional activators of lungfish MR.

Because spironolactone, an antagonist of human MR, activates elephant shark MR [24],

zebrafish MR [37,40,60] and trout MR [38], we investigated spironolactone for activation of full-

length lungfish MR and truncated lungfish MR-CDE. We also studied activation by eplerenone,

another antagonist of human MR [61]. As shown in Figure 4, both spironolactone and

eplerenone activated lungfish MR with a TAT3 promoter, and there was a further increase in

fold-activation by both steroids of lungfish MR-CDE.

Figure 4. Spironolactone and eplerenone activation of full-length and truncated lungfish
MR. Plasmids for full-length lungfish MR or truncated lungfish MR (MR-CDE) were expressed
in HEK293 cells with a TAT3-luciferase reporter. Transfected cells were treated with either 10
nM or 100 nM aldosterone, spironolactone or eplerenone or vehicle alone (DMSO). Results are
expressed as means ± SEM, n=3. Y-axis indicates fold-activation compared to the activity of
control vector with vehicle alone as 1. A. Full-length lungfish MR with TAT3. B. Truncated
lungfish MR (MR-CDE) with TAT3.
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Concentration-dependent activation by corticosteroids and progestins of full-length and
truncated lungfish MR.

To gain a quantitative measure of corticosteroid and progestin activation of full-length

and truncated lungfish MR, we determined the concentration dependence of transcriptional

activation by corticosteroids and progestins of full-length lungfish MR and lungfish MR-CDE

using TAT3 (Figure 5). This data was used to calculate a half maximal response (EC50) for

steroid activation of lungfish MR with a TAT3 promoter (Table 1). For full-length lungfish MR,

the four lowest EC50s were for aldosterone (0.04nM), 11-deoxycorticosterone (0.04 nM), 11-

deoxycortisol (0.17nM) and progesterone (0.03nM). These low EC50s are consistent with a

physiological role for one or more of these steroids as ligand for lungfish MR. In contrast,

corticosterone and cortisol, two physiological corticosteroids in terrestrial vertebrates, had EC50s

of 23.1nM and 66.1nM, respectively. Two synthetic glucocorticoids, dexamethasone and

triamcinolone, had EC50s of 4.7nM and 1.3nM, respectively.

For truncated lungfish MR, there were similar low EC50s for aldosterone (0.24nM), 11-

deoxycorticosterone (0.013nM), 11-deoxycortisol (0.27nM) and progesterone (0.04nM). EC50s

for corticosterone and cortisol were 85.5nM and 86.7nM, respectively. EC50s for

dexamethasone and triamcinolone were 7.7nM and 2.4nM, respectively.

Overall, these results reveal that the EC50s of aldosterone, 11-deoxycorticosterone, 11-

deoxycortisol and progesterone for full-length lungfish MR and lungfish MR-CDE are similar

and that one or more of these steroids could be a physiological mineralocorticoid in lungfish.

Although EC50s for full-length lungfish MR of triamcinolone and dexamethasone were at least

10-fold higher than that of aldosterone, deoxycorticosterone, 11-deoxycortisol and progesterone,

compared to these steroids, dexamethasone and triamcinolone have a several fold higher

activation of full-length and truncated lungfish MR (Figure 5). Consistent with data in Figure 3,

deletion of the NTD to form truncated lungfish MR-CDE increased fold-activation by

aldosterone, the other corticosteroids, progesterone, dexamethasone and triamcinolone.

However, deletion of the NTD did not have a large effect on their EC50s.

Fig. 5. Concentration-dependent transcriptional activation by corticosteroids, progesterone, dexamethasone and triamcinolone of full length and truncated lungfish MR. Plasmids for full-length lungfish MR or truncated lungfish MR, were expressed in HEK293 cells with a TAT3-luciferase promoter. Cells were treated with increasing concentrations of either aldosterone, cortisol, corticosterone, 11-deoxycorticosterone, 11-deoxycortisol, progesterone, dexamethasone and triamcinolone or vehicle alone (DMSO). Results are expressed as means ± SEM, n=3. Y-axis indicates fold-activation compared to the activity of control vector with vehicle (DMSO) alone as 1. A. Aldosterone, 11-deoxycorticosterone, 11-deoxycortisol and progesterone with full-length lungfish MR with TAT3-luc. B. Aldosterone, 11-deoxycorticosterone, 11-deoxycortisol and progesterone with truncated lungfish MR (Domains CDE) with TAT3-luc. C. Cortisol, corticosterone, dexamethasone and triamcinolone with full-length lungfish MR with TAT3-luc. D. Cortisol, corticosterone, dexamethasone and triamcinolone with truncated lungfish MR (Domains CDE) with TAT3-luc.

Table 1. EC50 values for steroid activation of full-length and truncated lungfish MR with the TAT3 promoter.

	Aldosterone	11-deoxycorticosterone	11-deoxycortisol	Progesterone
	EC50	EC50	EC50	EC50
MR-full length	0.04 nM	0.04 nM	0.17 nM	0.03 nM
95% confidence interval	0.02-0.07 nM	0.02-0.09 nM	0.1-0.3 nM	0.02-0.06 nM
MR-CDE	0.24 nM	0.13 nM	0.27 nM	0.044 nM
95% confidence interval	0.17-0.35 nM	0.08-0.2 nM	0.14-0.53 nM	0.026-0.076 nM

	Corticosterone	Cortisol	Triamcinolone	Dexamethasone
	EC50	EC50	EC50	EC50
MR-full length	23.1 nM	66.1 nM	1.3 nM	4.7 nM
95% confidence interval	11.3-47.1 nM	44.5-98.4 nM	0.9-1.9 nM	3.3-6.9 nM
MR-CDE	85.5 nM	86.7 nM	2.4 nM	7.7 nM
95% confidence interval	60.0-121.8 nM	74.3-101.2 nM	2.1-2.8 nM	6.6-9.1 nM

Transcriptional activation of full-length and truncated human MR and full-length and truncated elephant shark MR by corticosteroids and progestins.

To gain an evolutionary perspective on activation of lungfish MR by steroids, we screened a panel of steroids, at 10 nM, for transcriptional activation of full-length human and elephant shark MRs and truncated human and elephant shark MR-CDEs using two reporters: MMTV-luc and TAT3-luc.

Comparison of human MR and lungfish MR.

Overall, compared to lungfish MR, fold activation of human MR was significantly higher for aldosterone and other corticosteroids. For example, compared to 2-fold activation by aldosterone of full-length lungfish MR with the MMTV promoter (Figure 3A), activation of full-length human MR by aldosterone was about 70-fold with the MMTV promoter (Figure 6A). Although fold-activation by steroids for truncated human MR (Figure 6B) decreased compared to full-length human MR (Figure 6A), activation by aldosterone and other corticosteroids of truncated human MR with the MMTV promoter (Figure 6B) was about 7-fold higher than for truncated lungfish MR (Figure 3B).

Unlike for lungfish MR, deletion of the NTD in human MR resulted in a loss of

activation by aldosterone and other corticosteroids for human MR-CDE with both promoters

(Figure 6A-D), consistent with the presence of two activation function domains in the NTD

(Figure 2) [49–52]. The relative loss of activation of human MR was greater with the MMTV

promoter than with the TAT3 promoter. For example, at 10 nM aldosterone, activation of full-

length human MR with the MMTV reporter was 70-fold (Figure 6A), which decreased to 14-fold

for human MR-CDE (Figure 6B). In contrast, at 10 nM aldosterone, fold-activation of human

MR-CDE with the TAT3 promoter was about 75% of activity for full-length human MR (Figure

6C, D). However, 11-deoxycorticosterone and 11-deoxycortisol lost substantial activity for

human MR-CDE with the MMTV and TAT3 promoters (Figure 6D).

There also was higher fold-activation by aldosterone of full-length and truncated human

MR with the TAT3 promoter (Figure 6C, D) compared to full-length and truncated lungfish MR

(Figure 3C, D). Aldosterone activation of full-length human MR with the TAT3 promoter

(Figure 6C) was about 45-fold higher than that for full-length lungfish MR with the TAT3

promoter (Figure 3C). Aldosterone activation of human MR-CDE with the TAT3 promoter

(Figure 6D) was about 15-fold higher than that for lungfish MR-CDE (Figure 3D).

The relative activation by aldosterone and dexamethasone of human MR and lungfish

MR was reversed. Aldosterone was more active than dexamethasone in stimulating transcription

by full-length human MR and human MR-CDE with the TAT3 promoter (Figure 6C, D). In

contrast, for lungfish MR dexamethasone was more active than aldosterone for full-length

lungfish MR and lungfish MR-CDE with the TAT3 promoter (Figure 3C, D).

Comparison of elephant shark MR and lungfish MR.

Activation by corticosteroids and progesterone of elephant shark MR with the MMTV

promoter has some similarities with their activation of lungfish MR. Like lungfish MR,

corticosteroids have a similar activation of about 10-fold for full-length and truncated elephant

shark MR, with little difference in potency among the corticosteroids. However, unlike lungfish

MR, aldosterone is stronger than dexamethasone in activating full-length and truncated elephant

shark MR with the MMTV promoter (Figure 6E, F).

At a 10 nM steroid concentration, aldosterone and other corticosteroids activated full-

length elephant shark MR with the TAT3 promoter by 9 to 12-fold (Figure 6G), which was

similar to activation with the MMTV promoter (Figure 6E). Activation of full-length elephant

shark MR by progesterone was about 5-fold with the TAT3 and MMTV promoters (Figure 6E,

G). Aldosterone was about 2-fold more active than dexamethasone.

However, deletion of the NTD from elephant shark MR resulted in a significant increase

is activation by steroids in the presence of the TAT3 promoter (Figure 6H). Thus, truncated

elephant shark MR with the TAT3 promoter was activated from 300 to 350-fold by aldosterone

and other corticosteroids and about 200-fold by progesterone and dexamethasone (Figure 6H),

indicating that like lungfish MR, the NTD in elephant shark inhibits activation by corticosteroids.

However, unlike lungfish MR, compared to aldosterone, dexamethasone was less active for full-

length and truncated elephant shark MR with the MMTV promoter and truncated elephant shark

MR with the TAT3 promoter.

A. Full-length human MR with MMTV

B. Truncated human MR with MMTV

C. Full-length human MR with TAT3

D. Truncated human MR with TAT3

E: Full-length elephant shark MR with MMTV

F. Truncated elephant shark MR with MMTV

G: Full-length elephant shark MR with TAT3

H. Truncated elephant shark MR with TAT3

Figure 6. Ligand specificity of full-length and truncated human MR and elephant shark

MR. Plasmids for full-length human and elephant shark MR or truncated human and elephant

shark MR (MR-CDE) were expressed in HEK293 cells with an MMTV-luciferase reporter or a

TAT3-luciferase reporter. Transfected cells were treated with either 10 nM aldosterone, cortisol,

11-deoxycortisol, corticosterone, 11-deoxycorticosterone, progesterone, dexamethasone or

vehicle alone (DMSO). Results are expressed as means ± SEM, n=3. Y-axis indicates fold-

activation compared to the activity of control vector with vehicle alone as 1. A. Full-length

human MR with MMTV-luciferase. B. Truncated human MR (MR-CDE) with MMTV-

luciferase. C. Full-length elephant shark MR with TAT3-luciferase. D. Truncated elephant

shark MR (MR-CDE) with TAT3-luciferase.

Does lungfish contain a separate GR gene?

We used sequences of human GR, coelacanth GR and elephant shark GR as probes in a

BLAST search of GenBank and did not retrieve a lungfish GR sequence. The absence of a

lungfish GR ortholog coupled with the strong response of lungfish MR to dexamethasone and

triamcinolone (Figure 5) leads us to propose that lungfish MR also functions as a GR.

Discussion

Dobzhansky's aphorism "Nothing in Biology Makes Sense Except in the Light of

Evolution" [62] explains the importance of the evolution of aldosterone in lungfish because

aldosterone activation of the kidney MR in terrestrial vertebrates regulates sodium, potassium

and water transport, which is critical in maintaining internal electrolyte homeostasis in terrestrial

vertebrates [30,34,63–65] an activity that was important in the transition from water to land.

Here we report that aldosterone, 11-deoxycorticosterone, and progesterone have EC50s below 1

nM for lungfish MR (Table 1), which makes these steroids potential physiological ligands for

lungfish MR. Another potential physiological steroid is 11-deoxycortisol, which is a steroid for

the CR in Atlantic sea lamprey [42,43]. 11-deoxycortisol has EC50 of 0.17 nM for full-length

lungfish MR (Table 1).

A functional advantage of 11-deoxycorticosterone, 11-deoxycortisol and progesterone as

ligands for the MR is that they lack an 11β-hydroxyl group, and thus, like aldosterone, they are

inert to 11β-hydroxysteroid dehydrogenase-type 2, unlike cortisol and corticosterone [66–69].

Indeed, this inertness to 11β-hydroxysteroid dehydrogenase-type 2 and the low EC50s of these

steroids for lungfish MR suggests that more than one corticosteroid and progesterone [39] may

be physiological mineralocorticoids.

Like ray-finned fish MRs [24,37,38,60] and elephant shark MR [24,39], lungfish MR is

activated by spironolactone (Figure 4), and, as reported here, by eplerenone [47,48,61].

We also find important differences between the response of lungfish MR and human MR

to aldosterone, 11-deoxycorticosterone, 11-deoxycortisol and progesterone, indicating that

further selectivity for aldosterone in human MR occurred during the evolution of terrestrial

vertebrates [13,15,37,40,52,69,70].

An unexpected difference between lungfish MR and human MR is the substantial

increase in fold-activation by steroids of lungfish MR after deletion of the NTD, in contrast to

human MR in which the NTD contains an activation function domain (Figure 2) [49–52].

Deletion of the NTD in elephant shark MR also resulted in a substantial increase in fold-

activation by corticosteroids and progesterone using the TAT3 promoter (Figure 6), but not for

the MMTV promoter [17]. These data with lungfish MR and elephant shark MR suggest that

early in the evolution of the MR there was an allosteric interaction between the LBD and NTD

[71,72] that repressed steroid activation of the MR, and that the activation function in the NTD

as found in human MR [49–52] evolved later in terrestrial vertebrates, along with changes in

steroid specificity, such loss of MR activation by progesterone [37,39,70]. The different

responses of full-length and truncated lungfish MR, human MR and elephant shark MR with the

MMTV and TAT3 promoters indicate that the NTD and the promoter are important regulators of

steroid activation of these MRs. Corticosteroid activation of these MRs in the presence of other

promoters merits investigation.

The stronger response of lungfish MR to dexamethasone compared to aldosterone and the

absence a lungfish GR ortholog sequence are puzzling. At a 10 nM concentration, fold-

activation by dexamethasone and triamcinolone is substantially higher than that of cortisol,

corticosterone, as well as aldosterone, 11-deoxycorticosterone, 11-deoxycortisol and

progesterone for lungfish MR (Figure 5). One explanation is that lungfish MR also has a GR

function.

Materials and Methods

Chemical reagents

Aldosterone, cortisol, corticosterone, 11-deoxycorticosterone, 11-deoxycortisol and

progesterone, spironolactone and eplerenone were purchased from Sigma-Aldrich. For reporter

gene assays, all hormones were dissolved in dimethyl-sulfoxide (DMSO); the final DMSO

concentration in the culture medium did not exceed 0.1%.

Animal

A slender spotted African lungfish, *Protopterus dolloi,* was purchased from a local

commercial supplier. Lungfish were anesthetized in freshwater containing 0.02% ethyl 3-

aminobenzoate methane-sulfonate from Sigma-Aldrich, and tissue samples were quickly

dissected and frozen in liquid nitrogen. Animal handling procedures conformed to the guidelines

set forth by the Institutional Animal Care and Use Committee at the University of Tokyo.

Molecular cloning of lungfish mineralocorticoid receptor

Two conserved amino acid regions, GCHYGV and LYFAPD of vertebrate MRs were

selected and degenerate oligonucleotides were used as primers for PCR. First-strand cDNA was

synthesized from 2 µg of total RNA isolated from the liver after amplification, and an additional

primer set (CKVFFK and LYFAPD) was used for the second PCR. The amplified DNA

fragments were subcloned with TA-cloning plasmid pGEM-T Easy vector, sequenced using a

BigDye terminator Cycle Sequencing-kit with T7 and SP6 primers, and analyzed on the 3130

Genetic Analyzer (Applied Biosystems). The 5'- and 3'-ends of the mineralocorticoid receptor

cDNAs were amplified by rapid amplification of the cDNA end (RACE) using a SMART RACE

cDNA Amplification kit. Genbank accessions for this lungfish MR are: Nucleotide ID:

LC630795 and Protein ID: BCV19931.

Construction of plasmid vectors

The full-length and truncated MRs were amplified by PCR with KOD DNA polymerase. The

PCR products were gel-purified and ligated into pcDNA3.1 vector (Invitrogen). The truncated

MR proteins were designed to possess methionine and valine residues at the N-terminus and

contain a DNA-binding domain, a hinge-region, and a ligand-binding domain. The truncated

MRs were amplified by PCR with KOD DNA polymerase by using the following primers:

lungfish MR forward primer (5'-

C<u>AAGCTT</u>ACCATGGTGTGTCTGGTGTGTGGTGACGAAG-3' containing *Hind*III site) and

lungfish MR reverse primer (5'-CCTACTTCCTGTGAAAGTACAATGAC -3' containing stop

codon), human MR forward primer (5'-

C<u>GGATCC</u>ACCATGGTGTGTTTGGTGTGTGGGGATGAG-3' containing *Bam*HI site) and

human MR reverse primer (5'-CTCACTTCCGGTGGAAGTAGAGCGGC -3' containing stop

codon). The amplified DNA fragments were subcloned with TA-cloning plasmid pGEM-T Easy

vector and sequenced, and then subcloned into pcDNA 3.1 vector by using *Hind*III-*Not*I sites for

lungfish MR truncated form or *Bam*HI-*Not*I sites for human MR truncated form. Mouse

mammary tumor virus-long terminal repeat (MMTV-LTR) was amplified from pMSG vector by

PCR, and inserted into pGL3-basic vector containing the *Photinus pyralis* lucifease gene. 3X-

Tyrosine Amino Transferase (TAT3) promoter containing reporter vector named pGL4.23-

TAT3-Luc was constructed as described previously [53]. All cloned DNA sequences were

verified by sequencing.

Transactivation assay and statistical methods

Transfection and reporter assays were carried out in HEK293 cells, as described previously

[40,73]. All experiments were performed in triplicate. The values shown are mean ± SEM from

three separate experiments, and dose-response data, which were used to calculate the half

maximal response (EC50) for each steroid, were analyzed using GraphPad Prism.

DECLARATION OF INTEREST

We have no conflict of interest.

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AUTHOR CONTRIBUTIONS

Y.K., S.O., and M.E.B. carried out the research and analyzed data. S.H. aided in the collection

of animals. X.L. constructed plasmid DNAs used in this study. Y.K. and M.E.B. conceived and

designed the experiments. Y.K. and M.E.B. wrote the paper. All authors gave final approval for

publication.

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REFERENCES

1. Baker ME, Nelson DR, Studer RA. Origin of the response to adrenal and sex steroids: Roles of promiscuity and co-evolution of enzymes and steroid receptors. J Steroid Biochem Mol Biol. 2015;151:12-24. doi:10.1016/j.jsbmb.2014.10.020.

2. Evans RM. The steroid and thyroid hormone receptor superfamily. Science. 1988;240(4854):889-895. doi:10.1126/science.3283939.

3. Bridgham JT, Eick GN, Larroux C, et al. Protein evolution by molecular tinkering: diversification of the nuclear receptor superfamily from a ligand-dependent ancestor. PLoS Biol. 2010;8(10):e1000497. Published 2010 Oct 5. doi:10.1371/journal.pbio.1000497.

4. Baker ME, Funder JW, Kattoula SR. Evolution of hormone selectivity in glucocorticoid and mineralocorticoid receptors [published correction appears in J Steroid Biochem Mol Biol. 2014 Jan;139:104]. J Steroid Biochem Mol Biol. 2013;137:57-70. doi:10.1016/j.jsbmb.2013.07.009.

5. Lifton RP, Gharavi AG, Geller DS. Molecular mechanisms of human hypertension. Cell. 2001;104(4):545-556. doi:10.1016/s0092-8674(01)00241-0.

6. Shibata S. 30 YEARS OF THE MINERALOCORTICOID RECEPTOR: Mineralocorticoid
receptor and NaCl transport mechanisms in the renal distal nephron. J Endocrinol.
2017;234(1):T35-T47. doi:10.1530/JOE-16-0669.

7. Jaisser F, Farman N. Emerging Roles of the Mineralocorticoid Receptor in Pathology:
Toward New Paradigms in Clinical Pharmacology. Pharmacol Rev. 2016;68(1):49-75.
doi:10.1124/pr.115.011106.

8. Hanukoglu I, Hanukoglu A. Epithelial sodium channel (ENaC) family: Phylogeny,
structure-function, tissue distribution, and associated inherited diseases. Gene.
2016;579(2):95-132. doi:10.1016/j.gene.2015.12.061.

9. Joëls M, de Kloet ER. 30 YEARS OF THE MINERALOCORTICOID RECEPTOR: The
brain mineralocorticoid receptor: a saga in three episodes. J Endocrinol. 2017
Jul;234(1):T49-T66. doi: 10.1530/JOE-16-0660. PMID: 28634266.

10. Gross KL, Cidlowski JA. Tissue-specific glucocorticoid action: a family affair. Trends
Endocrinol Metab. 2008;19(9):331-339. doi:10.1016/j.tem.2008.07.009.

11. Weikum ER, Knuesel MT, Ortlund EA, Yamamoto KR. Glucocorticoid receptor control of
transcription: precision and plasticity via allostery. Nat Rev Mol Cell Biol. 2017;18(3):159-
174. doi:10.1038/nrm.2016.152.

12. Thornton JW. Evolution of vertebrate steroid receptors from an ancestral estrogen receptor
by ligand exploitation and serial genome expansions. Proc Natl Acad Sci U S A. 2001 May
8;98(10):5671-6. doi: 10.1073/pnas.091553298. Epub 2001 May 1. PMID: 11331759;
PMCID: PMC33271.

13. Baker ME, Katsu Y. 30 YEARS OF THE MINERALOCORTICOID RECEPTOR:
Evolution of the mineralocorticoid receptor: sequence, structure and function. J Endocrinol.
2017;234(1):T1-T16. doi:10.1530/JOE-16-0661.

14. Osório J, Rétaux S. The lamprey in evolutionary studies. Dev Genes Evol. 2008
May;218(5):221-35. doi: 10.1007/s00427-008-0208-1. Epub 2008 Feb 15. PMID:
18274775.

15. Bridgham JT, Carroll SM, Thornton JW. Evolution of hormone-receptor complexity by
molecular exploitation. Science. 2006;312(5770):97-101. doi:10.1126/science.1123348.

16. Carroll SM, Bridgham JT, Thornton JW. Evolution of hormone signaling in elasmobranchs
by exploitation of promiscuous receptors. Mol Biol Evol. 2008;25(12):2643-2652.
doi:10.1093/molbev/msn204.

17. Katsu Y, Shariful IMD, Lin X, Takagi W, Urushitani H, Kohno S, Hyodo S, Baker ME. N-
terminal Domain Regulates Steroid Activation of Elephant Shark Glucocorticoid and
Mineralocorticoid Receptors. J Steroid Biochem Mol Biol. 2021 Feb 27:105845. doi:
10.1016/j.jsbmb.2021.105845. Epub ahead of print. PMID: 33652098.

18. Inoue JG, Miya M, Lam K, et al. Evolutionary origin and phylogeny of the modern
holocephalans (Chondrichthyes: Chimaeriformes): a mitogenomic perspective. Mol Biol
Evol. 2010;27(11):2576-2586. doi:10.1093/molbev/msq147.

19. Venkatesh B, Lee AP, Ravi V, et al. Elephant shark genome provides unique insights into
gnathostome evolution [published correction appears in Nature. 2014 Sep
25;513(7519):574]. Nature. 2014;505(7482):174-179. doi:10.1038/nature12826.

20. Funder JW. Aldosterone and Mineralocorticoid Receptors-Physiology and Pathophysiology.
Int J Mol Sci. 2017 May 11;18(5):1032. doi: 10.3390/ijms18051032. PMID: 28492512;
PMCID: PMC5454944.

21. Hawkins UA, Gomez-Sanchez EP, Gomez-Sanchez CM, Gomez-Sanchez CE. The
ubiquitous mineralocorticoid receptor: clinical implications. Curr Hypertens Rep.
2012;14(6):573-580. doi:10.1007/s11906-012-0297-0.

22. Rossier BC, Baker ME, Studer RA. Epithelial sodium transport and its control by
aldosterone: the story of our internal environment revisited. Physiol Rev. 2015;95(1):297-
340. doi:10.1152/physrev.00011.2014.

23. Arriza JL, Weinberger C, Cerelli G, et al. Cloning of human mineralocorticoid receptor
complementary DNA: structural and functional kinship with the glucocorticoid receptor.
Science. 1987;237(4812):268-275. doi:10.1126/science.3037703.

24. Katsu Y, Kohno S, Oka K, et al. Transcriptional activation of elephant shark
mineralocorticoid receptor by corticosteroids, progesterone, and spironolactone. Sci Signal.
2019;12(584):eaar2668. Published 2019 Jun 4. doi:10.1126/scisignal.aar2668, (n.d.).

25. Jiang JQ, Young G, Kobayashi T, Nagahama Y. Eel (Anguilla japonica) testis 11beta-
hydroxylase gene is expressed in interrenal tissue and its product lacks aldosterone
synthesizing activity. Mol Cell Endocrinol. 1998;146(1-2):207-211. doi:10.1016/s0303-
7207(98)00147-6.

26. Joss JM, Itahara Y, Watanabe TX, Nakajima K, Takei Y. Teleost-type angiotensin is
present in Australian lungfish, Neoceratodus forsteri. Gen Comp Endocrinol.
1999;114(2):206-212. doi:10.1006/gcen.1999.7246.

27. Joss JM. Lungfish evolution and development. Gen Comp Endocrinol. 2006 Sep
15;148(3):285-9. doi: 10.1016/j.ygcen.2005.10.010. Epub 2005 Dec 7. PMID: 16337631.

28. Uchiyama M, Maejima S, Yoshie S, Kubo Y, Konno N, Joss JM. The epithelial sodium
channel in the Australian lungfish, Neoceratodus forsteri (Osteichthyes: Dipnoi). Proc Biol
Sci. 2012;279(1748):4795-4802. doi:10.1098/rspb.2012.1945.

29. Meyer A, Schloissnig S, Franchini P, Du K, Woltering JM, Irisarri I, Wong WY,
Nowoshilow S, Kneitz S, Kawaguchi A, Fabrizius A, Xiong P, Dechaud C, Spaink HP,
Volff JN, Simakov O, Burmester T, Tanaka EM, Schartl M. Giant lungfish genome
elucidates the conquest of land by vertebrates. Nature. 2021 Feb;590(7845):284-289. doi:

10.1038/s41586-021-03198-8. Epub 2021 Jan 18. PMID: 33461212; PMCID:
PMC7875771.

30. Biscotti MA, Gerdol M, Canapa A, Forconi M, Olmo E, Pallavicini A, Barucca M, Schartl
M. The Lungfish Transcriptome: A Glimpse into Molecular Evolution Events at the
Transition from Water to Land. Sci Rep. 2016 Feb 24;6:21571. doi: 10.1038/srep21571.
PMID: 26908371; PMCID: PMC4764851.

31. Brinkmann H, Venkatesh B, Brenner S, Meyer A. Nuclear protein-coding genes support
lungfish and not the coelacanth as the closest living relatives of land vertebrates. Proc Natl
Acad Sci U S A. 2004 Apr 6;101(14):4900-5. doi: 10.1073/pnas.0400609101. Epub 2004
Mar 22. PMID: 15037746; PMCID: PMC387346.

32. Wang K, Wang J, Zhu C, Yang L, Ren Y, Ruan J, Fan G, Hu J, Xu W, Bi X, Zhu Y, Song
Y, Chen H, Ma T, Zhao R, Jiang H, Zhang B, Feng C, Yuan Y, Gan X, Li Y, Zeng H, Liu
Q, Zhang Y, Shao F, Hao S, Zhang H, Xu X, Liu X, Wang D, Zhu M, Zhang G, Zhao W,
Qiu Q, He S, Wang W. African lungfish genome sheds light on the vertebrate water-to-land
transition. Cell. 2021 Feb 4:S0092-8674(21)00090-8. doi: 10.1016/j.cell.2021.01.047. Epub
ahead of print. PMID: 33545087.

33. McCormick SD, Bradshaw D. Hormonal control of salt and water balance in vertebrates.
Gen Comp Endocrinol. 2006 May 15;147(1):3-8. doi: 10.1016/j.ygcen.2005.12.009. Epub
2006 Feb 2. PMID: 16457828.

34. Rossier BC. Osmoregulation during Long-Term Fasting in Lungfish and Elephant Seal: Old
and New Lessons for the Nephrologist. Nephron. 2016;134(1):5-9. doi: 10.1159/000444307.
Epub 2016 Feb 23. PMID: 26901864.

35. Sugimoto A, Oka K, Sato R, Adachi S, Baker ME, Katsu Y. Corticosteroid and
progesterone transactivation of mineralocorticoid receptors from Amur sturgeon and
tropical gar. Biochem J. 2016;473(20):3655-3665. doi:10.1042/BCJ20160579.

36. Katsu Y, Baker ME. Progesterone activation of zebrafish mineralocorticoid receptor may
influence growth of some transplanted tumors. Proc Natl Acad Sci U S A.
2018;115(13):E2908-E2909. doi:10.1073/pnas.1802441115.

37. Fuller PJ, Yao YZ, Jin R, et al. Molecular evolution of the switch for progesterone and
spironolactone from mineralocorticoid receptor agonist to antagonist. Proc Natl Acad Sci U
S A. 2019;116(37):18578-18583. doi:10.1073/pnas.1903172116, (n.d.).

38. Sturm A, Bury N, Dengreville L, et al. 11-deoxycorticosterone is a potent agonist of the
rainbow trout (Oncorhynchus mykiss) mineralocorticoid receptor. Endocrinology.
2005;146(1):47-55. doi:10.1210/en.2004-0128.

39. Baker ME, Katsu Y. Progesterone: An enigmatic ligand for the mineralocorticoid receptor.
Biochem Pharmacol. 2020;177:113976. doi:10.1016/j.bcp.2020.113976.

40.   Katsu Y, Oka K, Baker ME. Evolution of human, chicken, alligator, frog, and zebrafish
mineralocorticoid receptors: Allosteric influence on steroid specificity. Sci Signal.
2018;11(537):eaao1520. Published 2018 Jul 3. doi:10.1126/scisignal.aao1.

41.   Baker ME, Funder JW, Kattoula SR. Evolution of hormone selectivity in glucocorticoid and
mineralocorticoid receptors [published correction appears in J Steroid Biochem Mol Biol.
2014 Jan;139:104]. J Steroid Biochem Mol Biol. 2013;137:57-70.
doi:10.1016/j.jsbmb.2013.07.009.

42.   Close DA, Yun SS, McCormick SD, Wildbill AJ, Li W. 11-deoxycortisol is a corticosteroid
hormone in the lamprey. Proc Natl Acad Sci U S A. 2010 Aug 3;107(31):13942-7. doi:
10.1073/pnas.0914026107. Epub 2010 Jul 19. PMID: 20643930; PMCID: PMC2922276.

43.   Shaughnessy CA, Barany A, McCormick SD. 11-Deoxycortisol controls hydromineral
balance in the most basal osmoregulating vertebrate, sea lamprey (Petromyzon marinus).
Sci Rep. 2020;10(1):12148. Published 2020 Jul 22. doi:10.1038/s41598-020-69061-4.

44.   Bridgham JT, Carroll SM, Thornton JW. Evolution of hormone-receptor complexity by
molecular exploitation. Science. 2006;312(5770):97-101. doi:10.1126/science.1123348.

45.   Conneely OM, Mulac-Jericevic B, DeMayo F, Lydon JP, O'Malley BW. Reproductive
functions of progesterone receptors. Recent Prog Horm Res. 2002;57:339-355.
doi:10.1210/rp.57.1.339.

46.   Publicover S, Barratt C. Reproductive biology: Progesterone's gateway into sperm
[published correction appears in Nature. 2011 Mar 31;471(7340):589]. Nature.
2011;471(7338):313-314. doi:10.1038/471313a.

47.   Epstein M, Calhoun DA. Aldosterone blockers (mineralocorticoid receptor antagonism) and
potassium-sparing diuretics. J Clin Hypertens (Greenwich). 2011 Sep;13(9):644-8. doi:
10.1111/j.1751-7176.2011.00511.x. Epub 2011 Aug 9. PMID: 21896143.

48.   Kolkhof P, Bärfacker L. 30 YEARS OF THE MINERALOCORTICOID RECEPTOR:
Mineralocorticoid receptor antagonists: 60 years of research and development. J Endocrinol.
2017 Jul;234(1):T125-T140. doi: 10.1530/JOE-16-0600. PMID: 28634268; PMCID:
PMC5488394.

49.   Fuse H, Kitagawa H, Kato S. Characterization of transactivational property and coactivator
mediation of rat mineralocorticoid receptor activation function-1 (AF-1). Mol Endocrinol.
2000 Jun;14(6):889-99. doi: 10.1210/mend.14.6.0467. PMID: 10847590.

50.   Fischer K, Kelly SM, Watt K, Price NC, McEwan IJ. Conformation of the
Mineralocorticoid Receptor N-Terminal Domain: Evidence for Induced and Stable
Structure. Mol Endocrinol. 2010 Oct;24(10):1935-48. Doi: 10.1210/Me.2010-0005. Epub
2010 Aug 4. PMID: 20685853; PMCID: PMC5417395.".

51. Pascual-Le Tallec L, Lombès M. The mineralocorticoid receptor: a journey exploring its
diversity and specificity of action. Mol Endocrinol. 2005 Sep;19(9):2211-21. doi:
10.1210/me.2005-0089. Epub 2005 Mar 31. PMID: 15802372.

52. Rupprecht R, Arriza JL, Spengler D, et al. Transactivation and synergistic properties of the
mineralocorticoid receptor: relationship to the glucocorticoid receptor. Mol Endocrinol.
1993;7(4):597-603. doi:10.1210/mend.7.4.8388999.

53. Iñiguez-Lluhí JA, Pearce D. A common motif within the negative regulatory regions of
multiple factors inhibits their transcriptional synergy. Mol Cell Biol. 2000
Aug;20(16):6040-50. doi: 10.1128/mcb.20.16.6040-6050.2000. PMID: 10913186; PMCID:
PMC86080.

54. Beato M, Arnemann J, Chalepakis G, Slater E, Willmann T. Gene regulation by steroid
hormones. J Steroid Biochem. 1987;27(1-3):9-14. doi: 10.1016/0022-4731(87)90288-3.
PMID: 2826895.

55. Cato AC, Skroch P, Weinmann J, Butkeraitis P, Ponta H. DNA sequences outside the
receptor-binding sites differently modulate the responsiveness of the mouse mammary
tumour virus promoter to various steroid hormones. EMBO J. 1988 May;7(5):1403-10.
PMID: 2842149; PMCID: PMC458390.

56. Fuller PJ, Yao Y, Yang J, Young MJ. Mechanisms of ligand specificity of the
mineralocorticoid receptor. J Endocrinol. 2012 Apr;213(1):15-24. doi: 10.1530/JOE-11-
0372. Epub 2011 Dec 12. PMID: 22159507.

57. Lombes M, Kenouch S, Souque A, Farman N, Rafestin-Oblin ME. The Mineralocorticoid
Receptor Discriminates Aldosterone from Glucocorticoids Independently of the 11 Beta-
Hydroxysteroid Dehydrogenase. Endocrinology. 1994;135(3):834-840.
Doi:10.1210/Endo.135.3.8070376.

58. Hellal-Levy C, Couette B, Fagart J, Souque A, Gomez-Sanchez C, Rafestin-Oblin M.
Specific hydroxylations determine selective corticosteroid recognition by human
glucocorticoid and mineralocorticoid receptors. FEBS Lett. 1999;464(1-2):9-13.
doi:10.1016/s0014-5793(99)01667-1.

59. Mani O, Nashev LG, Livelo C, Baker ME, Odermatt A. Role of Pro-637 and Gln-642 in
human glucocorticoid receptors and Ser-843 and Leu-848 in mineralocorticoid receptors in
their differential responses to cortisol and aldosterone. J Steroid Biochem Mol Biol. 2016
May;159:31-40. doi: 10.1016/j.jsbmb.2016.02.017. Epub 2016 Feb 22. PMID: 26907965.

60. Pippal JB, Cheung CM, Yao YZ, Brennan FE, Fuller PJ. Characterization of the zebrafish
(Danio rerio) mineralocorticoid receptor. Mol Cell Endocrinol. 2011 Jan 30;332(1-2):58-66.
doi: 10.1016/j.mce.2010.09.014. Epub 2010 Oct 12. PMID: 20932876.

61. Kiilerich P, Triqueneaux G, Christensen NM, et al. Interaction between the trout
mineralocorticoid and glucocorticoid receptors in vitro. J Mol Endocrinol. 2015;55(1):55-
68. doi:10.1530/JME-15-0002.

62.   Dobzhansky.T (1973) Nothing in Biology Makes Sense except in the Light of Evolution.
Am Biol Teach 35(3):125-129.

63.   Fournier D, Luft FC, Bader M, Ganten D, Andrade-Navarro MA. Emergence and evolution
of the renin-angiotensin-aldosterone system. J Mol Med (Berl). 2012 May;90(5):495-508.
doi: 10.1007/s00109-012-0894-z. Epub 2012 Apr 14. PMID: 22527880; PMCID:
PMC3354321.

64.   Vize PD, Smith HW. A Homeric view of kidney evolution: A reprint of H.W. Smith's
classic essay with a new introduction. Evolution of the kidney. 1943. Anat Rec A Discov
Mol Cell Evol Biol. 2004;277(2):344-354. doi:10.1002/ar.a.20017.

65.   Xanthakis V, Vasan RS. Aldosterone and the risk of hypertension. Curr Hypertens Rep.
2013 Apr;15(2):102-7. doi: 10.1007/s11906-013-0330-y. PMID: 23378101; PMCID:
PMC3619380.

66.   Funder JW. Apparent mineralocorticoid excess. J Steroid Biochem Mol Biol. 2017
Jan;165(Pt A):151-153. doi: 10.1016/j.jsbmb.2016.03.010. Epub 2016 Mar 5. PMID:
26956190.

67.   Frey FJ, Odermatt A, Frey BM. Glucocorticoid-mediated mineralocorticoid receptor
activation and hypertension. Curr Opin Nephrol Hypertens. 2004 Jul;13(4):451-8. doi:
10.1097/01.mnh.0000133976.32559.b0. PMID: 15199296.

68.   Chapman K, Holmes M, Seckl J. 11β-hydroxysteroid dehydrogenases: intracellular gate-
keepers of tissue glucocorticoid action. Physiol Rev. 2013;93(3):1139-1206.
doi:10.1152/physrev.00020.2012.

69.   Funder J. 30 YEARS OF THE MINERALOCORTICOID RECEPTOR: Mineralocorticoid
receptor activation and specificity-conferring mechanisms: a brief history. J Endocrinol.
2017 Jul;234(1):T17-T21. doi: 10.1530/JOE-17-0119. Epub 2017 May 22. PMID:
28533421.

70.   Fuller PJ, Yao YZ, Yang J, Young MJ. Structural determinants of activation of the
mineralocorticoid receptor: an evolutionary perspective. J Hum Hypertens. 2021
Feb;35(2):110-116. doi: 10.1038/s41371-020-0360-2. Epub 2020 May 28. PMID:
32467588.

71.   Hilser VJ, Thompson EB. Structural dynamics, intrinsic disorder, and allostery in nuclear
receptors as transcription factors. J Biol Chem. 2011;286(46):39675-39682.
doi:10.1074/jbc.R111.278929.

72.   Changeux JP, Christopoulos A. Allosteric Modulation as a Unifying Mechanism for
Receptor Function and Regulation. Cell. 2016 Aug 25;166(5):1084-1102. doi:
10.1016/j.cell.2016.08.015. PMID: 27565340.

73.   Oka K, Hoang A, Okada D, Iguchi T, Baker ME, Katsu Y. Allosteric role of the amino-
terminal A/B domain on corticosteroid transactivation of gar and human glucocorticoid

receptors. J Steroid Biochem Mol Biol. 2015;154:112-119.
doi:10.1016/j.jsbmb.2015.07.025.