

Letters

A Structure-Guided Approach to an Orthogonal Estrogen-Receptor-Based Gene Switch Activated by Ligands Suitable for *in Vivo* Studies

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Abstract: A strategy to obtain a fully orthogonal estrogen-receptor-based gene switch responsive to molecules with acceptable pharmacological properties is presented. From a series of tetrahydrofluorenones active on the wild-type estrogen receptor (ER) an inactive analogue is chosen as a new lead compound. Coevolution of receptor mutants and ligands leads to an ER-based gene switch suitable for studies in animal models.

In the field of gene therapy an important goal is to have tight temporal and dose control over the expression of an introduced gene (transgene). This can be achieved by controlling the expression of the transgene with a small molecule (inducer) with well-behaved pharmacokinetic and pharmacodynamic properties. In the presence of the inducer the ligand-dependent transcription system should be activated in a dose-dependent manner, whereas in its absence a low basal expression of the transgene should occur.^{1,2} We recently published the development of an estrogen receptor (ER) based gene switch fulfilling all the desired criteria for such a system, namely, the complete orthogonality of the artificial receptor and the exogenous ligand to the endogenous ligand/receptor pair and a low immunogenicity.³ In this paper we describe the medicinal chemistry effort of this work culminating in the identification of highly selective ligands, active on an estradiol-unresponsive ER mutant and suitable for studies *in vivo*.

The strategy applied in order to obtain an orthogonal gene switch consisted of six steps: (1) identification of inactive compounds within an active series, focusing on compounds where the activity was abolished because of minor structural changes; the compounds should also have favorable pharmacological properties; (2) determination of the residues of the protein responsible for the loss of activity of the chosen compound through molecular modeling; (3) generation of a mutant library of the ER ligand binding domain (LBD), containing mutations at the previously identified amino acid positions and selection of mutants with increased affinity for the low-affinity ligand;³ (4) optimization of the chosen lead compound in terms of selectivity profile; (5) introduction of an additional mutation to abolish the remaining affinity for

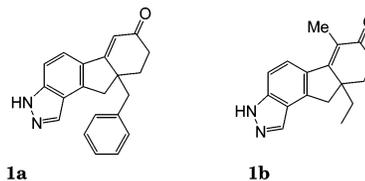


Figure 1. Inactive chosen lead compound **1a** and active analogue **1b**.

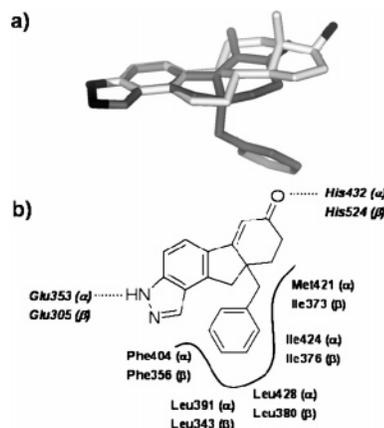


Figure 2. (a) Superposition of estradiol and **1a** and (b) residues most likely to interfere with binding of **1a** (bold). Conserved polar interactions used to define the position of the tetrahydrofluorenone are also shown (italic).

estradiol; (6) second round optimization of the ligands in terms of selectivity for the final mutant.

Compound **1a** (Figure 1) belongs to a series of tetrahydrofluorenones, which were prepared as part of our β -selective ER modulator (SERM) project.^{4,5} **1a** shows weak binding affinity for hER α and hER β (IC₅₀ > 10 μ M for both subtypes). Members of this series of benzopyrazole analogues had demonstrated good oral bioavailability and an acceptable clearance in several species (data not shown). In contrast, **1b** (Figure 1) is an active tetrahydrofluorenone with high affinity for hER β . A comparison of both compounds shows that the replacement of the ethyl group in position 9a of **1b** by the benzyl group is responsible for the loss of affinity for **1a**. This replacement increases the volume of the ligand by 54 \AA^3 .

To identify the residues that interfere with binding of the benzyl group, we built models of **1a** and **1b** bound to the hER α and hER β LBDs using X-ray structures of both LBDs with bound agonists.^{6,7} For this purpose the tetrahydrofluorenone scaffold was superimposed with the estradiol scaffold (Figure 2a). Severe steric clashes between the benzyl group of **1a** and five hydrophobic amino acid residues in the binding pocket were identified. These amino acids were selected as sites of replacements by other hydrophobic amino acids in the generation of an ER α -LBD mutant library (together with the mutation L384M to mimic the ER β subtype) (Figure 2b). Recently a crystal structure of a similar tetrahydrofluorenone has been published that is in good agreement with the superposition shown in Figure 2.⁸

The designed LBD-mutant library was expressed in yeast, and sequence analysis of mutants which conferred sufficient

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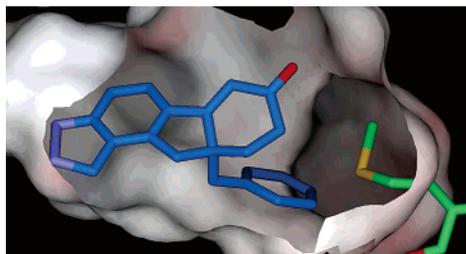
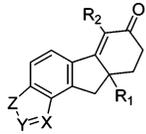


Figure 3. Model of **1a** in the estrogen pocket. The solvent accessible surface area was calculated for Gly421, showing the large pocket that becomes available to accommodate the benzyl moiety of **1a**. Met421 of the wild-type receptor is shown for comparison.

Table 1. SAR of Tetrahydrofluorenonees



No.	X	Y	Z	R ₁	R ₂	MG-ER α IC ₅₀ (nM) ^a	hER α ^b IC ₅₀ (nM)	hER β ^b IC ₅₀ (nM)
E₂ ^c	-	-	-	-	-	46 \pm 4	1.6 \pm 0.2	1.4 \pm 0.2
1a	CH	N	NH	Bn	H	3900 \pm 400	>10000	>10000
7a	CH	N	NH	Bn	Me	160 \pm 20	>10000	1200 \pm 140
7b	CH	N	NH	4-F-Bn	Me	82 \pm 58	>10000	>1000
7c	CH	N	NH	3-F-Bn	Me	440 \pm 60	>10000	1600 \pm 800
7d	CH	N	NH	2-F-Bn	Me	140 \pm 30	3400 \pm 700	560 \pm 30
7e	CH	N	NH	2-naphthyl-methyl	Me	5500 ^f	n.d. ^d	n.d.
15a	N	N	NH	Bn	Me	1200 \pm 250	>10000	5800 \pm 1400
15b	N	N	NH	4-F-Bn	Me	340 \pm 130	>10000	>10000
20a	H	-	OH	Bn	Me	14 \pm 2	1500 \pm 200	510 \pm 60
20b	H	-	OH	4-Cl-Bn	Me	22 \pm 10	6800 ^{e,f}	n.d.

^a IC₅₀ values are the mean of three or more experiments. ^b Full-length receptor. ^c Estradiol. ^d Not determined. ^e IC₅₀ on the wtER α -LBD. ^f Only one experiment.

growth in the presence of **1a** revealed the presence of one predominant mutation in the ligand binding pocket. Methionine 421 was replaced by the much smaller glycine (ER α -LBD containing mutations L384M/M421G, called MG-ER α). The IC₅₀ of **1a** on the isolated ligand binding domain MG-ER α was found to be 3.9 μ M. This increase in affinity can be rationalized by an increase of the volume of the binding pocket by 66 Å^3 , matching the 54 Å^3 increase of the volume of **1a** compared to **1b** (Figure 3). As anticipated, MG-ER α still exhibited high affinity for estradiol, although being reduced by at least 1 order of magnitude (Table 1).

Keeping in mind that MG-ER α would not be our final mutant, we started an effort to improve affinity and selectivity of lead compound **1a** for MG-ER α . Meanwhile additional mutations were evaluated for insertion into MG-ER α to abolish the remaining affinity for estradiol. We reasoned that a compound with high affinity and selectivity for MG-ER α would most likely serve as a lead for a SAR study on the final mutant.

A summary of the SAR on MG-ER α with **1a** as the lead compound is given in Table 1. The introduction of a methyl group at the 6-position resulted in **7a** with a 17-fold increase in affinity for MG-ER α , being inactive on hER α and displaying an 8-fold selectivity over hER β . The exploration of the SAR around the benzyl substituent resulted in the identification of **7b** with an IC₅₀ of 82 nM on MG-ER α and a slightly improved selectivity over hER β (>12-fold). More bulky substituents such

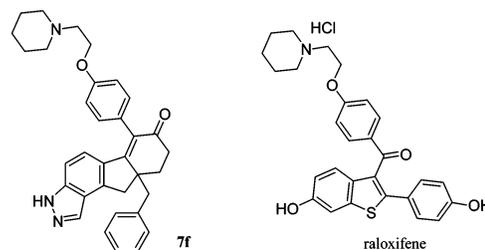


Figure 4. Compound **7f** and raloxifene.

as 2-naphthylmethyl were not tolerated (**7e**). Replacements of the pyrazole ring were investigated as well. Compound **15a**, the triazole analogue of **7a**, showed an 8-fold lower affinity, but the introduction of a 4-fluoro substituent on the benzyl ring largely restored affinity and resulted in **15b** with an IC₅₀ of 300 nM on MG-ER α and no activity on hER α and hER β up to 10 μ M. Finally, replacement of the pyrazole by a 7-hydroxy group resulted in compounds with the highest affinity and selectivity. Introduction of a chloro substituent in the 4-position on the benzyl ring yielded **20b** as the most potent and selective compound (>300-fold selective for MG-ER α over wtER α -LBD).

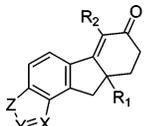
To abolish the remaining affinity of MG-ER α for estradiol, the additional mutation G521R was introduced, leading to a triple-mutated ER α -LBD (called MGR-ER α). It is known in the literature that estrogen receptors that contain the mutation G521R in the LBD exhibit a strongly reduced affinity for estradiol but remain responsive to certain ER antagonists such as 4-OH-tamoxifen and raloxifene.^{9,10}

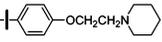
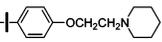
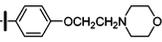
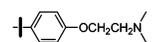
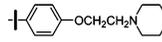
To test our compounds on MGR-ER α , a functional assay in mammalian cells was set up. (Affinity determination of our ligands on MGR-ER α by a competitive radiometric binding assay was hampered by the loss of affinity for radiolabeled estradiol.) For this purpose MGR-ER α was grafted into our previously described gene switch HEA-1¹¹ (replacement of the HEA-1-LBD with MGR-ER α), which was transfected in HeLa cells.³ In the presence of the ligands the expression of a reporter protein (SEAP, human-secreted alkaline phosphatase) was activated, protein concentrations were measured at different ligand concentrations, and the EC₅₀ values were determined. The MGR α -containing gene switch was unresponsive to estradiol up to 10 μ M, but to our disappointment, when **7a** was tested in this assay, the gene switch also remained unresponsive up to 2 μ M, suggesting a substantial loss of affinity of **7a** for MGR-ER α .

Undeterred by this result, we synthesized a hybrid molecule of **7a** and raloxifene, yielding **7f** (Figure 4).

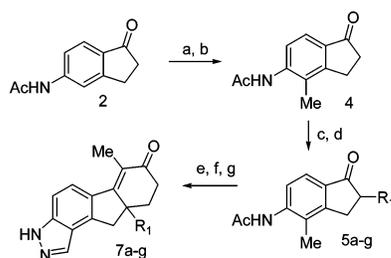
When **7f** was tested on MGR-ER α in the cell-based assay, an EC₅₀ of 300 nM was obtained. But at the same time **7f** exhibited an increased affinity for hER α and hER β compared to **7a** (Table 2). Encouraged by this result, we concentrated on compounds with bulky R₂ side chains, and a SAR study was performed to restore selectivity versus hER α /hER β .

For practical reasons we continued to measure affinity for MG-ER α , while interesting compounds were further tested in the cell-based assay on MGR-ER α . A summary of the results is given in Table 2. The hydroxy compound **20c** showed a much improved selectivity (defined as IC₅₀ ratios of MG-ER α /hER α or MG-ER α /hER β) compared to **7f** (17-fold/8-fold against 6-fold/1.2-fold), and a small SAR study around the side chain R₂ in the hydroxy series identified **20g** with still further improved selectivity (30-fold/14-fold). As expected, installation of a chloro substituent at the 4-position of the benzyl ring (**20b**) gave the most selective compound (38-fold/76-fold), confirming

Table 2. SAR of Tetrahydrofluorenones with Bulky Side-chains R₂


No.	X	Y	Z	R ₁	R ₂	MG-ER α IC ₅₀ (nM) ^a	MGR-ER α EC ₅₀ (nM)	hER α IC ₅₀ (nM) ^b	hER β IC ₅₀ (nM) ^b
7f	C	NH	Bn			110 ±10	300 ±40	620 ±90	130 ±10
20c	H	-	OH	Bn		22 ±3	14 ±2	380 ±40	180 ±20
20d	H	-	OH	Bn	n-butyl	25 ±4	n.d. ^c	260 ±40	310 ±40
20e	H	-	OH	Bn	4-OH-phenyl	29 ±4	11 ±3	220 ±20	100 ±20
20f	H	-	OH	Bn		25 ±4	n.d.	710 ±80	230 ±35
20g	H	-	OH	Bn		40 ±5	n.d.	1200 ±110	560 ±60
20h	H	-	OH	4-Cl-Bn		29 ±3	41 ±2	1100 ±90	2200 ±140

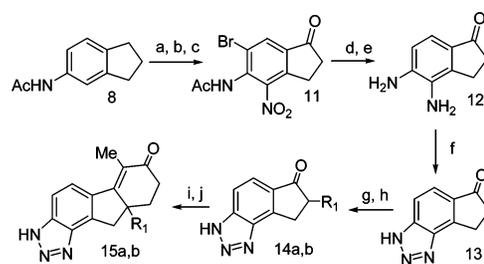
^a IC₅₀ values are the mean of three or more experiments. ^b Full-length receptor. ^c Not determined.

Scheme 1^a

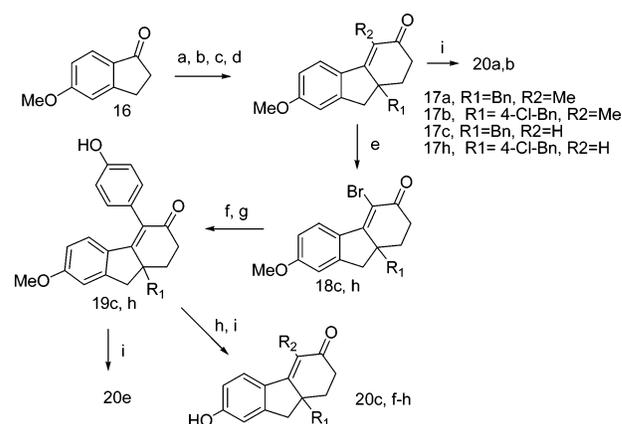
^a Reagents and conditions: (a) NBS, CH₃CN, 60 °C; (b) PdCl₂(PPh₃)₂, LiCl, SnMe₄, DMF, 100 °C; (c) ArCHO, NaOMe, MeOH; (d) Pd/C, H₂; (e) ethyl vinyl ketone, NaOMe, MeOH, 60 °C; (f) 6 N aqueous HCl/HOAc, 1:1, 80 °C; (g) (i) NOBF₄, CH₂Cl₂, -35 to 4 °C; (ii) KOAc, db-18-crown-6, CH₂Cl₂, -35 °C to room temp.

the results obtained in the series lacking the bulky side chain. **20c**, **20e**, and **20h** were tested in the cell-based assay on MGR-ER α (Table 2). Their EC₅₀ values correlated well with their IC₅₀ values measured on MG-ER α , suggesting that the additional mutation present in MGR-ER α has no detrimental effect on the affinity of ligands with bulky side chains R₂. The data obtained in mammalian cells also suggest a good cell penetration of the molecules.

Compounds **20c**, **20e**, and **20h** were evaluated for their suitability for in vivo studies: **20c** and **20e** were dosed iv in rats (dose 3 mg/kg). **20c** displayed a moderate clearance of 22 mL min⁻¹ kg⁻¹ and a terminal half-life of 2 h. Meanwhile, **20e** displayed a clearance of 39 mL min⁻¹ kg⁻¹ and a terminal half-life of 1.5 h. The plasma concentration of **20c** after 6 h was more than 10-fold its EC₅₀ in mammalian cells, and that of **20e** was more than 4-fold. Compound **20h** was dosed in male Balb-c mice (4 mg/kg, ip). The plasma concentration after 4 h was 1.5-fold its EC₅₀ in mammalian cells with a C_{max} of 0.5 μ M after 30 min. These data demonstrate that the obtained class of compounds is suitable for in vivo studies in animal models, ideally using ip dosing.

Scheme 2^a

^a Reagents and conditions: (a) Br₂, HOAc, 10 °C; (b) CrO₃, HOAc, 15–17 °C; (c) HNO₃ (100%), -40 °C; (d) 6 N HCl, MeOH, reflux; (e) Pd/C, H₂, NaOAc, EtOAc; (f) NaNO₂, HCl, EtOH, 0–5 °C; (g) ArCHO, NaOMe, MeOH; (h) Pd/C, H₂; (i) ethyl vinyl ketone, NaOMe, MeOH, 60 °C; (j) 6 N aqueous HCl/HOAc, 1:1, 80 °C.

Scheme 3^a

^a Reagents and conditions: (a) ArCHO, NaOMe, MeOH; (b) Pd/C, H₂, EtOAc for **17a,c**; Se, NaBH₄, EtOH/THF, 0–50 °C for **17b,h**; (c) ethyl vinyl ketone, NaOMe, MeOH, 60 °C for **17a,b**; methyl vinyl ketone, DBU, THF, 55 °C for **17c,h**; (d) 6 N aqueous HCl/HOAc, 1:1, 80 °C, for **17a,b**; pyrrolidine, HOAc, THF, 55 °C, for **17c,h**; (e) Br, NaHCO₃, CCl₄, 0 °C; (f) aryl-SnBu₃, Pd(PPh₃)₄, toluene, 100 °C; (g) MeOH/2 N aqueous HCl, 5:1, 60 °C; (h) amino alcohol, DIAD, PPh₃, THF, 0–25 °C; (i) AlCl₃, 2-propanethiol, CH₂Cl₂.

The synthesis of the pyrazole-containing ligands followed the strategy developed for the synthesis of the hER- β -active compounds^{4,5} and is summarized in Scheme 1. Selective bromination of N-protected 5-amino-1-indanone (**2**) at the 4-position with NBS and subsequent replacement of the bromo substituent by a methyl group via Stille cross-coupling led to intermediate **4**. Introduction of the alkyl substituents R₁ was achieved by an aldol condensation with the corresponding arylaldehydes and a subsequent catalytic reduction of the double bond. A two-step Robinson annulation with ethyl vinyl ketone and a concomitant N-deacetylation led to intermediates **5a–g**. The final pyrazole ring formation was achieved under mild conditions by formation of the diazonium tetrafluoroborate intermediates, which were then cyclized under phase-transfer conditions.¹²

The triazole-containing ligands were synthesized as shown in Scheme 2. To obtain the key intermediate **12**, a transient protection of the 6-position of **8** with a bromo substituent was necessary because the alternative direct nitration of **8** or **2** gave mixtures of regioisomers that were difficult to separate. After formation of the triazole ring the introduction of the alkyl substituents R₁ was achieved in the same manner as in the pyrazole series by aldol condensation with the corresponding arylaldehydes and subsequent reduction of the double bond. A two-step Robinson annulation led to the final triazole com-

pounds, although in low yields because of a partial conjugate addition of ethyl vinyl ketone to the unprotected triazole ring.

The series of hydroxytetrahydrofluorenones was prepared mostly according to the synthesis of the hER- β -active compounds (Scheme 3).^{4,5}

In summary we have successfully demonstrated an approach for the construction of an orthogonal gene switch, responsive to molecules with acceptable pharmacological properties. The essence of this approach lies in the early consideration of the properties of the ligand and the subsequent construction of a suitable receptor for the chosen ligand. Similar to our work, directed evolution approaches have been successfully applied in the creation of highly orthogonal gene switches.^{13–15} In these studies though the pharmacological properties of the ligands were not taken into consideration. By contrast, our concept consists of the coevolution of the receptor and of the ligand. To our knowledge the work presented here is the first example where the ligands of a completely orthogonal estrogen-receptor-based gene switch are small molecules with modest clearance and acceptable half-life in vivo, suitable for studies in animal models.

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Supporting Information Available: Experimental procedures and characterization of intermediates and final compounds. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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