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# Introduction of an Electron Withdrawing Group on the Hydroxyphenylnaphthol Scaffold Improves the Potency of $17\beta$ -Hydroxysteroid Dehydrogenase Type 2 ( $17\beta$ -HSD2) Inhibitors

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# **Supporting Information**

**ABSTRACT:** Estrogen deficiency in postmenopausal women or elderly men is often associated with the skeletal disease osteoporosis. The supplementation of estradiol (E2) in osteoporotic patients is known to prevent bone fracture but cannot be administered because of adverse effect. As  $17\beta$ -hydroxysteroid dehydrogenase type 2 ( $17\beta$ -HSD2) oxidizes E2 to its inactive form estrone (E1) and has been found in osteoblastic cells, it is an attractive target for the treatment of osteoporosis. Twenty-one novel, naphthalene-derived compounds have been synthesized and evaluated for their  $17\beta$ -HSD2 inhibition and their selectivity toward  $17\beta$ -HSD1 and the estrogen receptors (ERs)  $\alpha$  and  $\beta$ . Compound **19** turned out to be the most potent and selective inhibitor of  $17\beta$ -HSD2 in cell-free assays and had a very good cellular activity in MDA-MB-231 cells, expressing naturally  $17\beta$ -HSD2. It also showed marked inhibition of the E1formation by the rat and mouse orthologous enzymes and strong inhibition of monkey  $17\beta$ -HSD2. It is thus an appropriate candidate to be further evaluated in a disease-oriented model.



# INTRODUCTION

 $17\beta$ -Hydroxysteroid dehydrogenase type 2 ( $17\beta$ -HSD2) catalyzes the conversion of the biologically active sex steroids estradiol (E2) and testosterone (T) into their inactive forms using NAD<sup>+</sup> as cofactor (Chart 1). It is a transmembrane

Chart 1. Interconversion of Estradiol (E2) to Estrone (E1) by  $17\beta$ -HSD2,  $17\beta$ -HSD4, and  $17\beta$ -HSD1 and of Testosterone (T) to Androstenedione (A-dione) by  $17\beta$ -HSD2,  $17\beta$ -HSD4,  $17\beta$ -HSD3, and  $17\beta$ -HSD5



protein,<sup>1</sup> and no crystal structure is available yet. This enzyme is abundantly expressed in placenta, liver, small intestine, and

endometrium and to a lesser extend also in kidney, pancreas, colon, uterus, breast, and prostate. It was also found in osteoblastic cells. $^{1-4}$ 

In bone physiology, osteoblastic cells<sup>5</sup> are responsible for bone formation and mineralization and osteoclastic cells (OCs) for bone resorption. The subtle remodeling balance maintained between the OB and OC activities is crucial for bone stability.

Osteoporosis<sup>6</sup> is a systemic skeletal disease characterized by low bone mass and deterioration of bone tissue. Symptoms of this disease include bone fragility<sup>6</sup> and increase in fractures,<sup>6</sup> often at the hips, spine, and wrist. Osteoporosis occurs especially in postmenopausal women and elderly men, when the levels in active sex steroids (E2 and T)<sup>7</sup> drop.

Only few drugs are efficient for the treatment of osteoporosis. Among them, the antiresorptive agent bisphosphonate alendronate<sup>8</sup> is the most potent drug but the risk of fractures in postmenopausal women<sup>8–10</sup> and elderly men<sup>11</sup> is only reduced by 50%. Estrogens stop bone loss in osteoporotic patients but are no longer used because of several nonskeletal effects such as an increased risk of deep vein thrombosis, pulmonary embolism, and breast cancer.<sup>12</sup> Furthermore, low dose of synthetic estrogens, like ethinylestradiol, has no practical

Received: June 28, 2011 Published: October 6, 2011 application. Raloxifene,<sup>13</sup> a selective estrogen receptor modulator (SERM) is frequently used to treat osteoporosis. Drawbacks of this therapy are the increased risks of thromboembolism,<sup>14</sup> hot flushes,<sup>15</sup> and leg cramps.<sup>15</sup> Therefore, there is a need to develop novel drugs that are more efficient and selective for osteoporosis.

Estrogens are known to play a key role in bone physiology. They act directly on OBs, which regulate the ability of the OC precursors to differentiate to OCs (osteoclastogenesis). The mode of action of estrogens on bone mass is not yet well understood. 17 $\beta$ -HSD2, which is expressed in OBs, can be considered as a molecular switch, as it is involved in the regulation of the level of active E2 and T in the target cell. Therefore, a local intracellular enhancement of E2 and T in bones should be feasible. This strategy has already been successfully applied for several enzymes like aromatase,<sup>16–18</sup> CYP17,<sup>19–22</sup> 17 $\beta$ -HSD1,<sup>23–35</sup> and 5 $\alpha$ -reductase.<sup>36–40</sup> Among the few 17 $\beta$ -HSD2 inhibitors described in the litera-

Among the few  $17\beta$ -HSD2 inhibitors described in the literature<sup>41-50</sup> there is only one class of nonsteroidal compounds:<sup>46-48</sup> the *cis*-pyrrolidinones. By application of a nonsteroidal inhibitor from this class in a monkey osteoporosis model,  $17\beta$ -HSD2 has been validated as a target for the treatment of osteoporosis by Bagi et al.<sup>51</sup> Treatment with this compound led to a slight decrease in bone resorption and increase in bone formation resulting in maintenance of bone balance and bone strength.<sup>50</sup> As only small effects were observed, there is a need to develop new potent  $17\beta$ -HSD2 inhibitors with better in vivo efficacy. Additionally, these compounds should show a high selectivity toward  $17\beta$ -HSD1, the isoenzyme catalyzing the reverse reaction, the reduction of estrone (E1) to E2.

In a previous work,<sup>52</sup> focusing on the design of new nonsteroidal  $17\beta$ -HSD2 inhibitors, 6-(4'-hydroxyphenyl)-1-naphthol (compound A, Table 1) has been identified as a promising scaffold with an IC<sub>50</sub> of 302 nM for  $17\beta$ -HSD2 and a selectivity factor (SF) of 8 toward 17 $\beta$ -HSD1. This hit needed to be improved regarding activity and selectivity. According to our hypothesis, 52 the OH substituted benzene ring of the naphthalene moiety in A mimics the A-ring of E2 and the OH-phenyl the D-ring, leading to a close binding of the OH-phenyl to the catalytic triad and the cofactor. As the substrate of one enzyme (17 $\beta$ -HSD2 in this case) is the product of the catalytic reaction of the other enzyme ( $17\beta$ -HSD1 in this case), it can be concluded that their active sites must be very similar too. Therefore, with analyzing  $17\beta$ -HSD1 from which in contrast to  $17\beta$ -HSD2 the crystal structure is known, it becomes apparent that there is some space available around the OH-phenyl for substituents.

In this work, first the hypothesis concerning the binding mode of hydroxyphenyl naphthol derivatives to  $17\beta$ -HSD2 was validated. Afterward, the available space for the inhibitors in the active site of the enzyme was explored by synthesis of several derivatives where the size of substituents at positions 2' and 3' of the hydroxyphenyl ring was varied. With the expectation that electronic effects play an important role, electron donating (EDG) and electron withdrawing (EWG) groups were also introduced into the OH-phenyl moiety.

# CHEMISTRY

Synthesis of 1-8 and 10-20 (Scheme 1) started from the commercially available 6-methoxy-1-tetralone which was triflated and aromatized in 1a in very good yield according to the method described by us.<sup>52</sup> The cross-coupling reaction<sup>53</sup> of the triflate 1a with the appropriate methoxyphenylboronic acid led to compounds 1, 3a, 4, 6, 8, 10, 12, 14, 16–18, and 20

Table 1. Inhibition of Human  $17\beta$ -HSD2 and  $17\beta$ -HSD1 by Compounds 1–21



			inhibition of 17β-HSD2 <sup>a</sup> (%)	inhibition of 17β-HSD1 <sup>b,d</sup>	
compd	R1	R2	at 1 $\mu$ M	(%) at 1 µM	$\log P^{c}$
spiro- $\delta$ -lactone			68	ni	
Α	Н	Н	74	20	3.93
В	$CH_3$	Н	20	28	
1	$CH_3$	2'-CH3	25	20	4.68
2	Н	2'-CH3	65	25	4.41
3	Н	3'-CH3	79	48	4.41
4	$CH_3$	2'-OCH3	37	15	4.06
5	Н	2'-OH	11	10	3.54
6	$CH_3$	3'-OCH <sub>3</sub>	49	45	4.06
7	Н	3'-OCH <sub>3</sub>	36	24	3.80
8	Н	3'-OH	69	31	3.54
9	Н	3'-phenyl	84	47	5.60
10	$CH_3$	2'-F	32	41	4.35
11	Н	2'-F	87	36	4.09
12	$CH_3$	3'-F	51	66	4.35
13	Н	3'-F	78	62	4.09
14	$CH_3$	2',5'-diF	26	73	4.51
15	Н	2',5'-diF	83	74	4.24
16	Н	2',6'-diF	69	73	4.24
17	Н	2'-CF3	84	52	4.85
18	$CH_3$	3'-CF <sub>3</sub>	44	69	5.11
19	Н	3'-CF <sub>3</sub>	98	60	4.85
20	Н	3'-Cl	74	43	4.49
21	Н	3'-CN	45	ni	3.96

<sup>*a*</sup>Human placenta, microsomal fraction, substrate [<sup>3</sup>H]E2 + E2 [500 nM], NAD<sup>+</sup> [1500  $\mu$ M], mean value of three determinations, relative standard deviation of <10%. <sup>*b*</sup>Human placenta, cytosolic fraction, substrate [<sup>3</sup>H]E1 + E1 [500 nM], NADH [500  $\mu$ M], mean value of three determinations, relative standard deviation of <10%. <sup>*c*</sup>Calculated log *P* data. <sup>*d*</sup>ni: no inhibition (inhibition of <10%).

(method A or B). Ether cleavage was performed using boron tribromide for compounds 2, 7, and 13 (method C); boron trifluoride dimethylsulfide complex for compounds 3, 5, and 15 (method D); and pyridinium hydrochloride for compounds 11 and 19 (method E).

Compound 9 bearing an additional phenyl ring on the hydroxyphenyl moiety was synthesized according to the route described in Scheme 2. Compound 1a was first methylated using methyl iodide,<sup>54</sup> and subsequently the triflate group was exchanged by a boronic ester group (intermediate 9c) with bis(neopentylglycolato)diboron and a crown ether under basic conditions according to the method of Xu et al.<sup>55</sup> In parallel, compound 9b was synthesized by a Suzuki coupling with 4-bromo-2-iodo-1-methoxybenzene 9e and phenylboronic acid. Subsequent Suzuki coupling (method B) between compounds 9c and 9b led to compound 9a. Demethylation using boron tribromide (method C) afforded compound 9.

# Scheme 1. Synthesis of Compounds 1-8 and $10-20^a$



<sup>a</sup>Reagents and conditions: a. boronic acid,  $Pd(PPh_3)_4$ , aq.  $Na_2CO_3$ , DME/EtOH, 90°C, 2h for **1**, **3a**, **4**, **6**, **12** and **17**, Method A; b. boronic acid,  $Pd(PPh_3)_4$ ,  $Cs_2CO_3$ , DME/water, microwave irradiation (150°C, 150 W, 25 min) for **8**, **10**, **14**, **16**, **18** and **20**, Method B; c. BBr<sub>3</sub>,  $CH_2Cl_2$ , -78°C to rt, overnight, for **2**, **7** and **13**, Method C; d. BF<sub>3</sub>.SMe<sub>2</sub>,  $CH_2Cl_2$ , rt, overnight, for **3**, **5** and **15**, Method D; e. pyridinium hydrochloride, 220°C, 3h for **11** and **19**, Method E.

"Reagents and conditions: (a) boronic acid, Pd(PPh<sub>3</sub>)<sub>4</sub>, aq Na<sub>2</sub>CO<sub>3</sub>, DME/EtOH, 90 °C, 2 h for 1, 3a, 4, 6, 12, and 17, method A; (b) boronic acid, Pd(PPh<sub>3</sub>)<sub>4</sub>, Cs<sub>2</sub>CO<sub>3</sub>, DME/water, microwave irradiation (150 °C, 150 W, 25 min) for 8, 10, 14, 16, 18, and 20, method B; (c) BBr<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, -78 °C to room temp, overnight, for 2, 7, and 13, method C; (d) BF<sub>3</sub>·SMe<sub>2</sub>, CH<sub>2</sub>Cl<sub>2</sub>, room temp, overnight, for 3, 5, and 15, method D; (e) pyridinium hydrochloride, 220 °C, 3 h for 11 and 19, method E.

Scheme 2. Synthesis of Compounds 9 and  $21^a$ 



"Reagents and conditions: (a) MeI,  $K_2CO_3$ , 18-crown-6, acetone, reflux, overnight; (b) bis(neopentylglycolato)diboron, PdCl<sub>2</sub>(dppf), dppf, KOAc, 1,4-dioxane, 85 °C, 4 h; (c) PhI(OAc)<sub>2</sub>, I<sub>2</sub>, EtOAc, 60 °C, dark conditions, 4 h; (d) boronic acid, Pd(PPh<sub>3</sub>)<sub>4</sub>, Cs<sub>2</sub>CO<sub>3</sub>, DME/water, microwave irradiation (150 °C, 150 W, 25 min), method B; (e) BBr<sub>3</sub>, CH<sub>2</sub>Cl<sub>2</sub>, -78 °C to room temp, overnight, method C; (f) boronic acid, Pd(PPh<sub>3</sub>)<sub>4</sub>, aq Na<sub>2</sub>CO<sub>3</sub>, DME/EtOH, 90 °C, 2 h, method A; (g) BF<sub>3</sub>·SMe<sub>2</sub>, CH<sub>2</sub>Cl<sub>2</sub>, room temp, overnight, method D.

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The synthesis of **21** was carried out by palladium-catalyzed Suzuki coupling reaction using the intermediate **9c** (method A) and the commercially available 5-bromo-2-hydroxybenzonitrile, followed by ether cleavage with boron trifluoride dimethylsulfide complex (method D, Scheme 2).

# BIOLOGICAL CHARACTERIZATION

**Inhibition of Human 17\beta-HSD2.** 17 $\beta$ -HSD2 inhibitory activity of the synthesized compounds was first evaluated in a cell-free assay. Human placental microsomal enzyme was used as source. The incubation was performed with tritiated E2, cofactor, and inhibitor. The separation of substrate and product was accomplished by HPLC with a subsequent quantification by peak integration.<sup>56</sup> The percent inhibition values of compounds 1–21 are shown in Table 1, and the IC<sub>50</sub> values of the most active compounds are reported in Table 2.

Table 2. IC<sub>50</sub>, Selectivity Factor, and Binding Affinities for the Estrogen Receptors  $\alpha$  and  $\beta$  for Selected Compounds

		cell-free assay			
compd	$17\beta$ -HSD2 $IC_{50}^{a}$ (nM)	$\substack{17\beta\text{-HSD1}\\\text{IC}_{50}^{b,e} \text{ (nM)}}$	selectivity factor <sup>c,e</sup>	$ER\alpha RBA^{d,e}$ (%)	$\stackrel{\text{ER}\beta \text{ RBA}^{d,e}}{(\%)}$
spiro-δ- lactone	34	nd	nd	nd	nd
Α	302	2425	8	5	5
3	275	1748	6	<0.1	0.1-1.0
9	261	919	4	<0.1	<0.1
11	132	2245	17	<0.1	<0.1
13	292	713	2	0.1-1.0	0.1-1.0
15	238	309	1	0.1-1.0	0.1-1.0
17	156	926	6	1.0-10	1.0-10
19	19	611	32	<0.1	0.1-1.0
20	307	1130	4	0.1-1.0	0.1-1.0

<sup>*a*</sup>Human placenta, microsomal fraction, substrate [<sup>3</sup>H]E2 + E2 [500 nM], cofactor NAD<sup>+</sup> [1500  $\mu$ M], mean value of three determinations, relative standard deviation of <10%. <sup>*b*</sup>Human placenta, cytosolic fraction, substrate [<sup>3</sup>H]E1 + E1 [500 nM], cofactor NADH [500  $\mu$ M], mean value of three determinations, relative standard deviation of <20%. <sup>*c*</sup>IC<sub>50</sub>(17 $\beta$ -HSD1)/IC<sub>50</sub> (17 $\beta$ -HSD2). <sup>*d*</sup>RBA: relative binding affinity. E2: 100%. <sup>*e*</sup>nd: not determined.

Compounds showing less than 10% inhibition at 1  $\mu$ M were considered to be inactive. Compound **A**, identified in a previous work,<sup>52</sup> was used as internal reference (74% 17 $\beta$ -HSD2 inhibition vs 20% 17 $\beta$ -HSD1 inhibition at 1  $\mu$ M), and a spiro- $\delta$ -lactone (compound **10** published by Poirier et al.<sup>43</sup>) was taken as external control (68% at 1  $\mu$ M in our test; 62–66% at 1  $\mu$ M in their test<sup>43</sup>).

For selected substituents (R2 = 2'-CH<sub>3</sub>, 2'-OCH<sub>3</sub>, 3'-OCH<sub>3</sub>, 2'-F, 3'-F, 2',5'-diF, 3'-CF<sub>3</sub>), both hydroxyphenyl **2**, **5**, **8**, **11**, **13**, **15**, **19** and corresponding methoxyphenyl derivatives **1**, **4**, **6**, **10**, **12**, **14**, **18** were synthesized and tested for  $17\beta$ -HSD2 inhibitory activity. As observed for the unsubstituted compound **A**, the hydroxylated molecules (**2**, **8**, **11**, **13**, **15**, and **19**) were always more potent than the respective methoxylated analogues except for the compound pair **4** (R2 = 2'-OCH<sub>3</sub>) and **5**. For compounds **3**, **9**, **16**, **17**, **20**, and **21**, the hydroxylated derivatives only were therefore tested.

Introduction of a lipophilic group like methyl (compounds 2 and 3) led to good  $17\beta$ -HSD2 inhibitory activity (65% inhibition for 2 and 79% for 3 when tested at 1  $\mu$ M), which was at the level of unsubstituted A (74% inhibition at 1  $\mu$ M). This indicates that there is space around the hydroxyphenyl moiety for introduction of a

small substituent in the 2'- or 3'-position, but the methyl is certainly not able to achieve any specific interactions, as no gain in activity is observed compared to the parent hydroxyphenylnaphthol **A**.

Exchange of the methyl group for a hydrophilic hydroxy substituent was detrimental for activity in the 2'-position (5, inactive) but tolerated in the 3'-position (8, similar activity as the unsubstituted A). It can therefore be hypothesized that there is no polar amino acid in the active site close to the 2'-position of the inhibitor. Furthermore, the 3'-OH group of compound 8 did not show specific interactions with the enzyme, as no increase in activity was observed compared to A. It is striking that an exchange of hydroxy (8) by methoxy (7) in the 3'-position decreased the inhibitory activity strongly.

A phenyl ring was also introduced into the 3'-position (9). This compound displayed a slight increase in activity (84% inhibition at 1  $\mu$ M for 9 vs 74% inhibition at 1  $\mu$ M for A), confirming our previous hypothesis that there is space available around the hydroxyphenyl core.

In order to evaluate the influence of electron withdrawing groups (EWGs) or electron donating groups (EDGs) on the  $17\beta$ -HSD2 inhibitory activity, compounds 10–21 bearing EWGs were synthesized to be compared to compounds with EDGs like methyl (3) or phenyl (9) groups. All methoxy derivatives (10, 12, 14, and 18) showed lower activity than the corresponding hydroxy compounds (11, 13, 15, and 19). Most of the hydroxy compounds exhibited identical or better potency than the parent compound A except 21. Compound 11, substituted with a fluoro group in the 2'- position, turned out to be highly active (87% inhibition at 1  $\mu$ M, IC<sub>50</sub> = 132 nM). Disubstituted fluorine compounds (2',5'-diF 15 and 2',6'-diF 16) resulted in highly active inhibitors but were slightly weaker than the monofluoro 11 (IC $_{50}$  of 238 nM for 15 and 132 nM for 11). Compounds 17 ( $R2 = 2'-CF_3$ ) and 19 ( $R2 = 3'-CF_3$ ) showed high (17) and very high (19) activity (84% and 98% inhibition at 1  $\mu$ M, respectively), resulting in IC<sub>50</sub> values of 19 nM for 19 and 156 nM for 17. However, exchange of the trifluoromethyl group for bioisosteric chloro (20) or cyano (21) groups was detrimental for the activity (74% and 45% inhibition for 20 and 21, respectively) when compared with inhibitory efficiency of 19 (98% at 1  $\mu$ M).

**Selectivity and Cellular Activity.** 17 $\beta$ -HSD1, as the main enzyme implicated in the reduction of estrone (E1) to E2 in sex steroid metabolism,<sup>57</sup> should not be affected by inhibitors of 17 $\beta$ -HSD2. Furthermore, inhibitors of 17 $\beta$ -HSD2 should have none or little affinity for the estrogen receptors  $\alpha$  and  $\beta$  (ER $\alpha$ and ER $\beta$ ), as most of the E2 effects are ER mediated. Therefore, all compounds were tested for selectivity toward 17 $\beta$ -HSD1 and a selection of the most potent inhibitors for binding affinity to ER $\alpha$  and  $\beta$ .

The assay for  $17\beta$ -HSD1 inhibition was performed in a similar way as for the  $17\beta$ -HSD2. Human placental cytosolic  $17\beta$ -HSD1 was incubated with tritiated E1, cofactor, and inhibitor. The amount of labeled E2 formed was determined after HPLC separation. For the most potent compounds, IC<sub>50</sub> values were determined and selectivity factors (SFs) calculated (SF = IC<sub>50</sub>(17\beta-HSD1)/IC<sub>50</sub>(17\beta-HSD2), Table 2). All compounds except **15** showed selectivity factor of 32, was compound **19** (R2 = 3'-CF<sub>3</sub>), which was also the most potent one exhibiting an IC<sub>50</sub> of 19 nM. The 2'-F compound **11** also showed a high SF of 17 toward  $17\beta$ -HSD1.

For the determination of ER binding affinity, a competition assay was used, with tritiated E2. Receptor bound and free E2

were separated by means of hydroxyapatite. Compounds **3**, **9**, **11**, and **19** showed a relative binding affinity (RBA) of less than 0.1% to ER $\alpha$ , compared to the affinity of E2, which is arbitrarily set to 100%. The indicated compounds are therefore classified as low affinity ligands of the ER $\alpha$ .

The log *P* values were calculated using the ChemDrawPro, version 11.0, program in order to get insight into the lipophilicity of this class of compounds and are shown in Table 1. The values for all compounds except **9** and **18** are lower than 5, fulfilling the criteria of Lipinski's rule of five. It therefore can be expected that they are able to easily permeate cell membranes.

The most interesting compounds **11**, **17**, and **19** with IC<sub>50</sub> values below 200 nM were further investigated in a cellular assay, using the human breast cancer cell line MDA-MB-231,<sup>58</sup> which endogenously expresses  $17\beta$ -HSD2. Cells were incubated with tritiated E2 and inhibitor for 3.5 h. Separation was performed by HPLC similar as in the cell-free inhibition assays. Compounds **17** and **19** showed very similar inhibitory activity in MDA-MB-231 cells as in the cell-free assay, i.e., IC<sub>50</sub> of 171 nM for **17** and 31 nM for **19** in cellular assay vs 156 and 19 nM, respectively, in the cell-free assay. Interestingly, compound **11** showed an IC<sub>50</sub> of 1227 nM in the cellular assay compared to 132 nM in the cell-free assay, indicating that this compound is less appropriate to permeate the cell membrane or is metabolically unstable.

Further in Vitro Assays. In order to refine the profile of the most potent compound 19, it was further tested for selectivity on other enzymes, namely,  $17\beta$ -HSD types 4 and 5.  $17\beta$ -HSD4 catalyzes the same reaction as  $17\beta$ -HSD2 (Chart 1),<sup>57</sup> and  $17\beta$ -HSD5 (beside  $17\beta$ -HSD3) is responsible for the reduction of the androgen A-dione to T.57 Both enzymes should not be inhibited by  $17\beta$ -HSD2 inhibitors. To investigate this important issue, experiments were performed with E. coli bacteria expressing human  $17\beta$ -HSD4 and  $17\beta$ -HSD5.<sup>56,59</sup> These cells were incubated with cofactor, inhibitor, and tritiated E2 in the case of  $17\beta$ -HSD4 and with A-dione in the case of 17 $\beta$ -HSD5. After RP-HPLC separation of substrate and product, the amount of labeled E1 or T formed was quantified. Compound 19 was found to be selective toward  $17\beta$ -HSD4 and  $17\beta$ -HSD5 (40%  $17\beta$ -HSD4 inhibition and 21%  $17\beta$ -HSD5 inhibition at 1  $\mu$ M).

According to the potency of compound **19** in cell-free and cellular assays and its selectivity toward  $17\beta$ -HSD1,  $17\beta$ -HSD4,  $17\beta$ -HSD5, and ERs, it can be considered as potential candidate for further development. Before evaluation in an animal model of the efficacy of this new  $17\beta$ -HSD2 inhibitor **19**, it is reasonable to determine its profile on orthologous enzymes from different species. Compound **19** was therefore tested for inhibition of  $17\beta$ -HSD2/E1-formation (activity) and  $17\beta$ -HSD1/E2-formation (selectivity) using rat, mouse, and monkey enzymes.

In rat and mouse assays, liver enzymes were used, while in monkey assay, placental enzymes were employed. Compound **19** showed good inhibition of the rat and mouse  $17\beta$ -HSD2 enzymes and very good inhibition on the monkey enzyme. In this case, the efficacy was in the same range as for the human enzyme (Table 3). In addition, selectivity toward  $17\beta$ -HSD1/E2-formation in monkey seems to be similar to that observed in human.

# DISCUSSION AND CONCLUSION

Compound **A**, identified by us in a previous study<sup>52</sup> represents a promising scaffold as  $17\beta$ -HSD2 inhibitor: substituted by two hydroxy groups (polar), the naphthalene and phenyl cores (hydrophobic) mimick the substrate E2. We have previously shown<sup>52</sup> that an exchange of the hydroxy group on the phenyl

Table 3. Inhibition of Rat, Mouse, and Monkey E1 and E2-Formation by Compound 19

	inhibiti formati	on of E1- on <sup><math>a,k</math></sup> (%)	inhibiti formatio	inhibition of E2-formation $^{b,k}$ (%)	
compd 19	$1 \ \mu M$	100 nM	$1 \ \mu M$	100 nM	
human	98 <sup>c</sup>	87 <sup>c</sup>	60 <sup>d</sup>	27 <sup>d</sup>	
rat	$77^e$	nd	52 <sup>f</sup>	nd	
mouse	72 <sup>g</sup>	nd	$26^{h}$	nd	
monkey	99 <sup>i</sup>	75 <sup>i</sup>	75 <sup>j</sup>	55 <sup>i</sup>	

<sup>a</sup>Substrate [<sup>3</sup>H]E2 + E2 [500 nM], NAD<sup>+</sup> [1500  $\mu$ M], mean value of three determinations, relative standard deviation of <10%. <sup>b</sup>Substrate [<sup>3</sup>H]E1 + E1 [500 nM], NADH [500  $\mu$ M], mean value of three determinations, relative standard deviation of <10%. <sup>c</sup>Human placenta, microsomal fraction. <sup>d</sup>Human placenta, cytosolic fraction. <sup>e</sup>Rat liver, microsomal fraction. <sup>h</sup>Mouse liver, cytosolic fraction. <sup>i</sup>Monkey placenta, microsomal fraction. <sup>j</sup>Monkey placenta, cytosolic fraction. <sup>k</sup>nd: not determined.

moiety by a sulfonamide did not affect the activity toward  $17\beta$ -HSD2 (74% vs 75% inhibition at 1  $\mu$ M, respectively), indicating that some space is available around the phenyl core. We hypothesized therefore that the phenyl ring mimics the D-ring of E2 and thus that introduction of additional substituents on this phenyl moiety would be tolerated and would allow an increase of the  $17\beta$ -HSD2 inhibitory activity. In this paper, we aimed at verifying whether this hypothesis is correct and whether introduction of various substituents on the phenyl ring could improve this activity.

The biological activity of the phenyl-substituted compound 9 confirms that the compound can indeed bind in the active site according to the previous hypothesis (the hydroxynaphthalene mimics the A-ring and the hydroxyphenyl the D-ring). The additional phenyl might be accommodated between cofactor and steroid where space is available.

The electronic effects of the substituents on the  $17\beta$ -HSD2 inhibitory activity were further elucidated. No matter where they were introduced, EDGs (Me, OH, Ph) in either the 2'- or 3'-position did not lead to significant differences in activity except for the hydroxy group where position 3' was better (compounds 5 and 8). None of the synthesized compounds (2, 3, 8, and 9) reached a better activity compared to reference A, indicating that these substituents are tolerated (there is space available) but not able to establish specific interactions with amino acid residues. The position of the substituent on the phenyl ring will induce either a pronounced rotation of the phenyl toward the naphthalene (position 2', ortho effect) or a weak rotation (position 3'). In both cases, only the 4'-hydroxy substituent stays at the same position and anchors the molecule into its binding site. Our finding that there is no difference in activity between 2'- and 3'-substituted compounds indicates that the phenyl ring is not able to establish  $\pi - \pi$  stacking interactions with amino acid side chains of the protein.

When the substituent was an EWG (F, CF<sub>3</sub>, Cl, CN), the highest potency was observed for compound 19 (R2 = 3'-CF<sub>3</sub>) in which the additional group was located at the 3'-position. Exchange of the trifluoromethyl group by the bioisosteric groups like chloro (20) and cyano (21) decreased inhibition. In the case of the CN group, the different activities might be due to the fact that CN is hydrophilic whereas CF<sub>3</sub> is lipophilic. The linear shape of CN giving steric hindrance might also explain the moderate activity of 21.

As shown in many other cases<sup>5</sup> and also observed for compound 11, the substitution with F results in an increase in inhibitory activity. However, the introduction of a second fluorine as in 15 or 16 does not further enhance activity, indicating that the effects are not additive.

In our previous study,<sup>52</sup> it was observed that the hydroxy group bound to the phenyl moiety was important for activity (74% at 1  $\mu$ M for A with OH vs 20% inhibition for B with OMe). This is confirmed in this study, as all methoxy compounds displayed lower 17 $\beta$ -HSD2 inhibitory activity than their corresponding hydroxy analogues. Contrary to our compounds, the other described nonsteroidal 17 $\beta$ -HSD2 inhibitors,<sup>46–48</sup> the *cis*-pyrrolidinones, do not need any hydrophilic group to be active, allowing us to presume that they might have a different binding mode compared to ours.

As other  $17\beta$ -HSD enzymes of the short-chain dehydrogenase/reductase (SDR) and the aldo-keto reductases (AKR) family participate in steroid interconversions,<sup>60</sup> the development of enzyme-specific inhibitors is a challenge. We therefore tested the compounds on other  $17\beta$ -HSDs, like  $17\beta$ -HSD types 1, 2, 4, and 5. Selectivity toward  $17\beta$ -HSD1 was low with nearly all of the tested hydroxyphenylnaphthol derivatives; however, the most potent  $17\beta$ -HSD2 inhibitors 11 and 19 showed the highest selectivity toward  $17\beta$ -HSD1, exhibiting promising selectivity factors (SFs) of 17 and 32, respectively. Although some inhibition of  $17\beta$ -HSD4 and -5 was observed with the best inhibitor compound 19, the substance can be assumed to be selective toward both enzymes. We were surprised by the fact that compound 19 showed some minor inhibition of  $17\beta$ -HSD4. In our previous projects we have not seen any effect on this particular enzyme with steroidal or nonsteroidal inhib-itors.<sup>56,61</sup> Nevertheless, the observed inhibition of compound 19 is fairly low and the impact on the ubiquitously expressed 17 $\beta$ -HSD4 contributing to the peroxisomal  $\beta$ -oxidation of fatty acids<sup>62</sup> is most probably negligible.

Compounds 11 and 19 also displayed very little binding affinity to ER $\alpha$ , which is responsible for cell proliferation. Only compound 19 showed a very good cellular activity (IC<sub>50</sub> = 31 nM in MDA-MB-231 cells). Thus, it is able to permeate the cell membrane and is not quickly metabolized.

An osteoblastic cell-line expressing  $17\beta$ -HSD2 and the ERs like SV-HFO should be promising to prove our concept that  $17\beta$ -HSD2 inhibition is able to induce bone formation and inhibit osteoclastogenesis. For further in vivo experiments, species differences have to be taken into consideration,<sup>63,64</sup> and compound **19** should inhibit the enzyme of the chosen species. In this pursuit, **19** was tested on rat, mouse, and monkey enzymes and revealed a high (72–77% inhibition at 1  $\mu$ M in mouse and rat enzymes) to very high (99% inhibition at 1  $\mu$ M in monkey enzyme) inhibitory activity for  $17\beta$ -HSD2/E1-formation. Showing a better activity and selectivity in marmoset (*Callithrix jacchus*) enzymes, this species should be appropriate for proving our concept.

In the present study, we describe the synthesis of substituted 6-phenyl-1-naphthols as inhibitors of  $17\beta$ -HSD2 and the evaluation of their biological activities. A new highly potent inhibitor of  $17\beta$ -HSD2 has been discovered (compound **19**) with very good cellular activity and good selectivity toward  $17\beta$ -HSD1,  $17\beta$ -HSD4,  $17\beta$ -HSD5, and ERs. Compound **19** is also able to inhibit E1-formation in rat, mouse, and monkey. Further biological analyses still need to be investigated to get information on the efficacy of the compound to demonstrate

that  $17\beta$ -HSD2 inhibitors are able to inhibit osteoclastogenesis and induce bone formation.

# EXPERIMENTAL SECTION

**Chemical Methods.** <sup>1</sup>H NMR and <sup>13</sup>C NMR spectra were recorded on a Bruker DRX-500 instrument in CDCl<sub>3</sub> and acetone- $d_6$ . Chemicals shifts are reported in  $\delta$  values (ppm). The hydrogenated residues of deuteriated solvent were used as internal standard (CDCl<sub>3</sub>,  $\delta$  = 7.26 ppm in <sup>1</sup>H NMR and  $\delta$  = 77.0 ppm in <sup>13</sup>C NMR; acetone- $d_6$ ,  $\delta$  = 2.05 ppm in <sup>1</sup>H NMR and  $\delta$  = 30.8 ppm and 206.3 ppm in <sup>13</sup>C NMR). Signals are described as s, br, d, t, dd, dt, qt, and m for singlet, broad, doublet, triplet, doublet of doublets, doublet of triplets, quintuplet, and multiplet, respectively. All coupling constants (*J*) are given in Hertz. IR spectra were measured neat on a Bruker Vector 33FT infrared spectrometer. Melting points (mp) were determined in open capillaries on a Stuart Scientific SMP3 apparatus and are uncorrected.

Mass spectra (GC/MS) were measured on a GCD series G1800A (Hewlett-Packard) instrument with an Optima-5-MS (0.25  $\mu$ M, 30 m) column (Macherey Nagel). All microwave irradiation experiments were carried out in a CEM-Discover monomode microwave apparatus.

Flash chromatography was performed on silica gel 40 (35/40 to 63/70  $\mu$ M) with hexane/ethyl acetate mixtures as eluents and the reaction progress was determined by thin-layer chromatography analyses on Alugram SIL G/UV254 (Macherey-Nagel). Visualization was accomplished with UV light.

Tested compounds are  $\geq$ 95% chemical purity as measured by LC/ MS. Data for all tested compounds are provided in the Supporting Information. The Surveyor LC system consisted of a pump, an autosampler, and a PDA detector. Mass spectrometry was performed on a MSQ electrospray mass spectrometer (ThermoFisher, Dreieich, Germany). The system was operated by the standard software Xcalibur. A RP-C18 NUCLEODUR 100-5 (125 mm × 3 mm) column (Macherey-Nagel GmbH, Dühren, Germany) was used as stationary phase. All solvents were HPLC grade. In a gradient run the percentage of acetonitrile (containing 0.1% trifluoroacetic acid) in 0.1% trifluoroacetic acid was increased from an initial concentration of 0% at 0 min to 100% at 15 min and kept at 100% for 5 min. The injection volume was 15  $\mu$ L, and flow rate was set to 800  $\mu$ L/min. MS analysis was carried out at a spray voltage of 3800 V and a capillary temperature of 350 °C and a source CID of 10 V. Spectra were acquired in positive mode from 100 to 1000 m/z at 254 nm for the UV trace.

Starting materials (different boronic acids and compounds 1e, 9f, and 21b) were purchased from Aldrich, Alfa Aeser, Acros, and Combi-Blocks and were used without further purification. No attempts were made to optimize yields.

The following compounds were prepared according to previously described procedures: 5-hydroxy-2-naphthyl trifluoromethanesulfonate **1a**, <sup>52</sup> 4-bromo-2-iodo-1-methoxybenzene **9e**.<sup>27</sup>

General Procedures for Suzuki Coupling. Method A. To a mixture of aryl bromide (1 equiv) and tetrakis(triphenylphosphine)palladium(0) (0.02 equiv) in DME was added a 2 N aqueous solution of sodium carbonate (2 equiv), and the mixture (oxygen free) was purged with N<sub>2</sub>. The resultant solution was stirred at room temperature for 5 min, and a solution of boronic acid (1.3 equiv) in EtOH was added. The mixture was heated to 90 °C and stirred for 2 h. The reaction mixture was cooled to room temperature, quenched by water, and extracted with ethyl acetate. The organic layer was washed with brine, dried over sodium sulfate, filtered, and concentrated to dryness. The product was purified by column chromatography.

**Method B.** A mixture of aryl bromide (1 equiv), boronic acid (1 equiv), cesium carbonate (3 equiv), and tetrakis(triphenylphosphine)palladium(0) (0.02 equiv) was suspended in an oxygen-free DME/ water (2:1) solution. The reaction mixture was exposed to microwave irradiation (25 min, 150 W, 150 °C, 15 bar). After the mixture reached room temperature, water was added and the aqueous layer was extracted with ethyl acetate. The combined organic layers were washed with brine, dried over sodium sulfate, filtered, and concentrated to dryness. The product was purified by column chromatography. General Procedures for Ether Cleavage. Method C. To a solution of methoxy derivative (1 equiv) in dry dichloromethane cooled at -78 °C under nitrogen was slowly added boron tribromide (1 M solution in dichloromethane, 5–10 equiv per methoxy function). The reaction mixture was stirred at -78 °C for 1 h and then allowed to warm to room temperature overnight. The reaction was quenched by water, and the aqueous layer was extracted with ethyl acetate. The combined organic layers were washed with brine and dried over sodium sulfate, evaporated to dryness under reduced pressure, and purified by column chromatography.

**Method D.** To a solution of methoxy derivative (1 equiv) in dry dichloromethane, boron trifluoride-dimethylsulfide complex (75 equiv per methoxy function) was added dropwise at room temperature. The reaction mixture was stirred at room temperature overnight. Water was added to quench the reaction, and the aqueous layer was extracted with ethyl acetate. The combined organic layers were washed with brine, dried over sodium sulfate, evaporated to dryness under reduced pressure, and purified by column chromatography.

**Method E.** To pyridinium hydrochloride (100 equiv) at 190  $^{\circ}$ C was added methoxy derivative (1 equiv), and the solution was stirred for 3 h. The reaction mixture was cooled to room temperature and was stirred with 1 N HCl. The mixture was dissolved in ethyl acetate, and the combined organic layers were washed with water, dried over sodium sulfate, filtered, evaporated to dryness under reduced pressure, and purified by column chromatography.

**6-(4-Hydroxy-3-methylphenyl)-1-naphthol (3).** The title compound was prepared by reaction of 6-(4-methoxy-3-methylphenyl)-1-naphthol **3a** (20 mg, 0.08 mmol, 1 equiv) with boron trifluoride–dimethylsulfide complex (6.00 mmol, 75 equiv) according to method D. The product was purified by column chromatography (hexane/ethyl acetate, 1:1) to give 12 mg (60%) of the analytically pure compound as a brown solid. C<sub>17</sub>H<sub>14</sub>O<sub>2</sub>; MW 250; mp 164–166 °C; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>) δ 8.90 (s, br, 1H), 8.26 (s, br, 1H), 8.23 (d, *J* ≈ 8.5 Hz, 1H), 7.96 (d, *J* ≈ 1.9 Hz, 1H), 7.69 (dd, *J* ≈ 2.0 Hz, *J* ≈ 8.5 Hz, 1H), 7.53 (d, *J* ≈ 1.9 Hz, 1H), 7.43 (dd, *J* ≈ 2.0 Hz, *J* ≈ 8.5 Hz, 1H), 7.40–7.37 (m, 1H), 7.28–7.24 (m, 1H), 6.91 (d, *J* ≈ 8.5 Hz, 1H), 6.84 (dd, *J* ≈ 0.8 Hz, *J* ≈ 7.4 Hz, 1H), 2.27 (s, 3H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>) δ 156.1, 154.0, 139.6, 136.4, 133.0, 130.4, 127.4, 126.4, 125.6, 124.9, 124.7, 123.5, 120.2, 118.2, 116.0, 108.7, 16.3; IR 3337, 2952, 1506, 1274 cm<sup>-1</sup>; GC/MS *m*/*z* 250 (M)<sup>+</sup>.

**5-(5-Hydroxy-2-naphthyl)biphenyl-2-ol (9).** The title compound was prepared by reaction of 2-methoxy-5-(5-methoxy-2-naphthyl)biphenyl **9a** (75 mg, 0.22 mmol, 1 equiv) with boron tribromide (2.2 mmol, 10 equiv) according to method C. The product was purified by column chromatography (hexane/ethyl acetate, 8:2), then by preparative HPLC (gradient acetonitrile/water from 20:80 to 100:0) to give 10 mg (15%) of the analytically pure compound as an orange oil. C<sub>22</sub>H<sub>16</sub>O<sub>2</sub>; MW 312; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>) δ 8.16 (d, *J* ≈ 8.5 Hz, 1H), 7.96 (d, *J* ≈ 2.0 Hz, 1H), 7.67 (dd, *J* ≈ 2.0 Hz, *J* ≈ 8.5 Hz, 1H), 7.61 (d, *J* ≈ 2.0 Hz, 1H), 7.59–7.55 (m, 2H), 7.51 (dd, *J* ≈ 2.0 Hz, *J* ≈ 8.5 Hz, 1H), 6.76 (d, *J* ≈ 7.6 Hz, 1H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>) δ 154.8, 153.9, 139.7, 139.2, 136.4, 133.7, 130.3, 128.9, 128.1, 127.7, 127.5, 125.2, 124.7, 124.6, 123.6, 120.2, 117.5, 108.7; GC/MS *m*/z 312 (M)<sup>+</sup>.

**6-(2-Fluoro-4-hydroxyphenyl)-1-naphthol (11).** The title compound was prepared by reaction of 6-(2-fluoro-4-methoxyphenyl)-1-naphthol **10** (100 mg, 0.09 mmol, 1 equiv) with pyridinium hydrochloride (9 mmol, 100 equiv) according to method E. The product was purified by column chromatography (hexane/ethyl acetate, 7:3) to afford 48 mg (54%) of the analytically pure product as a beige solid. C<sub>16</sub>H<sub>11</sub>FO<sub>2</sub>; MW 254; mp 145–146 °C; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>)  $\delta$  9.15 (s, br, 1H), 9.07 (s, br, 1H), 8.28 (d, *J*  $\approx$  8.5 Hz, 1H), 7.83 (d, *J*  $\approx$  8.5 Hz, 2H), 7.62 (dt, *J*  $\approx$  1.8 Hz, *J*  $\approx$  8.5 Hz, 1H), 7.50–7.46 (m, 1H), 7.42 (d, *J*  $\approx$  8.5 Hz, 1H), 7.32–7.31 (m, 1H), 6.92 (dd, *J*  $\approx$  1.0 Hz, *J*  $\approx$  7.3 Hz, 1H), 6.84–6.81 (m, 1H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>)  $\delta$  162.7, 154.0, 148.1, 136.0, 132.4, 130.1, 127.8, 127.5, 126.5, 123.1, 120.2, 115.1, 115.0, 112.9, 109.1, 104.2; IR 3225, 1623, 1214 cm<sup>-1</sup>; GC/MS *m*/z 254 (M)<sup>+</sup>.

**6-(3-Fluoro-4-hydroxyphenyl)-1-naphthol (13).** The title compound was prepared by reaction of 6-(3-fluoro-4-methoxyphenyl)-1-naphthol **12** (20 mg, 0.08 mmol, 1 equiv) with boron tribromide (0.80 mmol, 10 equiv) according to method C. The product was purified by column chromatography (hexane/ethyl acetate, 1:1) to give 8 mg (40%) of the analytically pure compound as a yellow oil. C<sub>16</sub>H<sub>11</sub>O<sub>2</sub>F; MW 254; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>) δ 8.99 (s, br, 1H), 8.74 (s, br, 1H), 8.29 (d, *J* ≈ 8.8 Hz, 1H), 8.05 (d, *J* ≈ 2.0 Hz, 1H), 7.74 (dd, *J* ≈ 2.1 Hz, *J* ≈ 8.7 Hz, 1H), 7.57 (dd, *J* ≈ 2.1 Hz, *J* ≈ 8.7 Hz, 1H), 7.57 (dd, *J* ≈ 0.9 Hz, *J* ≈ 7.6 Hz, 1H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>) δ 154.2, 136.3, 127.6, 125.4, 124.4, 124.0, 123.7, 120.3, 119.1, 115.5, 109.0; IR 3247, 1692, 1524, 1277 cm<sup>-1</sup>; GC/MS *m*/z 254 (M)<sup>+</sup>.

**6-(2, 5-Difluoro-4-hydroxyphenyl)-1-naphthol (15).** The title compound was prepared by reaction of 6-(2,5-difluoro-4-methoxyphenyl)-1-naphthol 14 (40 mg, 0.14 mmol, 1 equiv) with boron trifluoride–dimethylsulfide complex (10.5 mmol, 75 equiv) according to method D. The product was purified by column chromatography (hexane/ethyl acetate, 8:2) to afford 20 mg (50%) of the analytically pure product as a beige solid. C<sub>16</sub>H<sub>10</sub>F<sub>2</sub>O<sub>2</sub>; MW 272; mp 178–181 °C; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>)  $\delta$  9.22 (s, br, 1H), 9.05 (s, br, 1H), 8.29 (d, *J* ≈ 8.8 Hz, 1H), 7.44 (d, *J* ≈ 8.4 Hz, 1H), 7.42–7.39 (m, 1H), 7.36–7.32 (m, 1H), 6.96–6.90 (m, 2H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>)  $\delta$  162.4, 159.6, 154.4, 153.2, 135.2, 134.5, 126.6, 125.9, 125.1, 123.5, 122.0, 120.8, 117.9, 117.5, 108.9, 104.6; IR 3427, 1641, 1522, 1268 cm<sup>-1</sup>; GC/MS *m/z* 272 (M)<sup>+</sup>.

**6-[4-Hydroxy-2-(trifluoromethyl)phenyl]-1-naphthol (17).** The title compound was prepared by reaction of 5-hydroxy-2-naphthyl trifluoromethanesulfonate **1a** (100 mg, 0.34 mmol, 1 equiv) with 4-hydroxy-2-trifluoromethylphenylboronic acid (98 mg, 0.44 mmol, 1.3 equiv) according to method A. The product was purified by column chromatography (hexane/ethyl acetate, 8:2) to give 70 mg (68%) of the analytically pure compound as a brown solid. C<sub>17</sub>H<sub>11</sub>F<sub>3</sub>O<sub>2</sub>; MW 304; mp 140–142 °C; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>) δ 9.03 (s, br, 1H), 9.01 (s, br, 1H), 8.25 (d, *J* ≈ 8.8 Hz, 1H), 7.73 (s, 1H), 7.42–7.39 (m, 2H), 7.36–7.31 (m, 2H), 7.28 (d, *J* ≈ 2.5 Hz, 1H), 7.18 (dd, *J* ≈ 2.5 Hz, *J* ≈ 8.6 Hz, 1H), 6.96–6.93 (m, 1H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>) δ 157.9, 154.3, 138.8, 135.4, 134.9, 128.7, 127.8, 127.2, 125.0, 122.6, 120.4, 119.4, 113.7, 109.5, 60.5; IR 3361, 1691, 1322 cm<sup>-1</sup>; GC/MS *m/z* 304 (M)<sup>+</sup>.

**6-[4-Hydroxy-3-(trifluoromethyl)phenyl]-1-naphthol (19).** The title compound was prepared by reaction of 6-[4-methoxy-3-(trifluoromethyl)phenyl]-1-naphthol **18** (100 mg, 0.15 mmol, 1 equiv) with pyridinium hydrochloride (15 mmol, 100 equiv) according to method E. The product was purified by column chromatography (hexane/ethyl acetate, 7:3) to afford 40 mg (42%) of the analytically pure product as a brown solid. C<sub>17</sub>H<sub>11</sub>F<sub>3</sub>O<sub>2</sub>; MW 304; mp 158–160 °C; <sup>1</sup>H NMR (acetone-*d*<sub>6</sub>) δ 8.28 (d, *J* ≈ 8.5 Hz, 1H), 8.04 (d, *J* ≈ 2.0 Hz, 1H), 7.91 (d, *J* ≈ 2.0 Hz, 1H), 7.85 (dd, *J* ≈ 2.0 Hz, *J* ≈ 8.5 Hz, 1H), 7.29 (m, 1H), 7.22–7.19 (m, 1H), 6.89 (dd, *J* ≈ 0.9 Hz, *J* ≈ 7.3 Hz, 1H); <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>) δ 156.3, 154.1, 138.0, 136.3, 133.0, 127.7, 125.5, 124.9, 124.4, 123.9, 120.2, 118.6, 109.1; IR 3229, 3066, 2986, 1619, 1263 cm<sup>-1</sup>; GC/MS *m/z* 304 (M)<sup>+</sup>.

**6-(3-Chloro-4-hydroxyphenyl)-1-naphthol (20).** The title compound was prepared by reaction of 5-hydroxy-2-naphthyl trifluoromethanesulfonate **1a** (200 mg, 0.68 mmol, 1 equiv) and 3-chloro-4-hydroxyphenylboronic acid (153 mg, 0.89 mmol, 1.3 equiv) according to method B. The product was purified by column chromatography (hexane/ethyl acetate, 7:3) to give 117 mg (64%) of the analytically pure compound as a beige solid.  $C_{16}H_{11}ClO_2$ ; MW 270; mp 112–114 °C; <sup>1</sup>H NMR (acetone- $d_6$ )  $\delta$  9.10 (s, br, 2H), 8.31 (d,  $J \approx 8.5$  Hz, 1H), 8.07 (d,  $J \approx 2.0$  Hz, 1H), 7.80 (d,  $J \approx 2.0$  Hz, 1H), 7.75 (dd,  $J \approx 2.0$  Hz,  $J \approx 8.5$  Hz, 1H), 7.64 (dd,  $J \approx 2.0$  Hz,  $J \approx 8.5$  Hz, 1H), 7.47 (d,  $J \approx 8.4$  Hz, 1H), 7.36–7.32 (m, 1H), 7.19 (d,  $J \approx 8.4$  Hz, 1H), 6.94 (dd,  $J \approx 1.0$  Hz,  $J \approx 7.4$  Hz, 1H); <sup>13</sup>C NMR (acetone- $d_6$ )  $\delta$ 154.0, 153.5, 137.9, 136.2, 134.5, 129.2, 127.7, 127.6, 125.4, 125.3, 124.4, 123.8, 121.7, 120.2, 118.1, 109.0; IR 3330, 1599, 1283, 1187 cm<sup>-1</sup>; GC/MS m/z 270–272 (M)<sup>+</sup>.

**Biological Assays.** [2,4,6,7-<sup>3</sup>H]E2 and [2,4,6,7-<sup>3</sup>H]E1 were bought from Perkin-Elmer, Boston, MA. Quickszint Flow 302 scintillator fluid was bought from Zinsser Analytic, Frankfurt, Germany. Cytosolic (17 $\beta$ -HSD1) and microsomal (17 $\beta$ -HSD2) fractions were obtained from human and *Callithrix jacchus* placenta according to previously described procedures<sup>29,65,66</sup> and from rat and mouse liver tissues.<sup>67</sup> Fresh tissue was homogenized and centrifuged. The pellet fraction contains the microsomal 17 $\beta$ -HSD2 and was used for the determination of E1-formation, while 17 $\beta$ -HSD1 was obtained after precipitation with ammonium sulfate from the cytosolic fraction for use of testing of E2-formation.

Human  $17\beta$ -HSD4 and  $17\beta$ -HSD5 were cloned into the modified pGEX-2T vector.<sup>56</sup> For the multidomain enzyme  $17\beta$ -HSD4, only the steroid converting SDR domain was subcloned.<sup>56</sup>

Inhibition of  $17\beta$ -HSD2/E1-Formation. Inhibitory activities were evaluated by a well established method with minor modifications.<sup>68-70</sup> Briefly, the enzyme preparation was incubated with NAD<sup>+</sup> [1500  $\mu$ M] in the presence of potential inhibitors at 37 °C in a phosphate buffer (50 mM) supplemented with 20% of glycerol and 1 mM EDTA. Inhibitor stock solutions were prepared in DMSO. Final concentration of DMSO was adjusted to 1% in all samples. The enzymatic reaction was started by addition of a mixture of unlabeled E2 and  $[2,4,6,7^{-3}H]E2$  (final concentration of 500 nM, 0.11  $\mu$ Ci). After 20 min at 37 °C, the incubation was stopped with HgCl<sub>2</sub> and the mixture was extracted with ether. After evaporation, the steroids were dissolved in acetonitrile. E1 and E2 were separated using acetonitrile/ water (45:55) as mobile phase in a C18 RP chromatography column (Nucleodur C18 Gravity, 3 µm, Macherey-Nagel, Düren, Germany) connected to a HPLC system (Agilent 1100 series, Agilent Technologies, Waldbronn, Germany). Detection and quantification of the steroids were performed using a radioflow detector (Berthold Technologies, Bad Wildbad, Germany). The conversion rate was calculated according to following equation:

% conversion = 
$$\frac{\%(E1)}{\%(E1) + \%(E2)} \times 100$$

Each value was calculated from at least three independent experiments.

**Inhibition of 17** $\beta$ **-HSD1/E2-Formation.** The 17 $\beta$ -HSD1 inhibition assay was performed similarly to the 17 $\beta$ -HSD2 procedure. The cytosolic fraction was incubated with NADH [500  $\mu$ M], test compound, and a mixture of unlabeled E1 and [2,4,6,7-<sup>3</sup>H]E1 (final concentration of 500 nM, 0.15  $\mu$ Ci) for 10 min. Further treatment of the samples and HPLC separation were carried out as mentioned above.

Inhibition of  $17\beta$ -HSD4. Inhibitory activity was assessed as originally described<sup>71,72</sup> with minor modifications.<sup>56</sup> Briefly, bacteria containing recombinant  $17\beta$ -HSD4 were resuspended in PBS and enzymatic assay was performed at pH 7.7. The enzyme preparation was incubated with NAD<sup>+</sup> [7.5 mM]. Inhibitor (dissolved in DMSO) was added in a final concentration of 1  $\mu$ M. Final concentration of DMSO was adjusted to 1% in all samples. The enzymatic reaction was started with the addition of [6,7-<sup>3</sup>H]E2 at 21 nM. The incubation at 37 °C was stopped with 0.21 M ascorbic acid in methanol/acetic acid (99:1) after the time needed to convert approximately 30% of the substrate in a control assay without inhibitor. Steroids were extracted from the assay mixture by SPE using Strata C18-E columns (Phenomenex), eluted with methanol, and separated by RP-HPLC (column, Luna, 5  $\mu$ m C18(2), 150 mm; Phenomenex) at a flow rate of 1 mL/min acetonitrile/water (43:57). Radioactivity was detected by scintillation counting after mixing with ReadyFlowIII (Beckman). Conversion was calculated from integration of substrate and product peaks. Assay was run in triplicate of three independent experiments.

**Inhibition of 17\beta-HSD5.** The 17 $\beta$ -HSD5 inhibition assay was performed similarly to the 17 $\beta$ -HSD4 procedure. The recombinant enzyme was incubated with NADPH [6 mM], inhibitor [1  $\mu$ M], and

 $[1,2,6,7^{-3}H]$ A-dione at 21 nM. Further treatment of the samples and HPLC separation were carried out as mentioned above.<sup>56</sup>

Inhibition of 17 $\beta$ -HSD2 in a Cellular Assay. Cellular 17 $\beta$ -HSD2 activity is measured using the breast cancer cell-line MDA-MB-231<sup>58</sup> (17 $\beta$ -HSD1 activity negligible). [<sup>3</sup>H]E2 (200 nM) is taken as substrate and is incubated with the inhibitor for 3.5 h at 37 °C. After ether extraction, substrate and product are separated by HPLC and detected with a radioflow detector. Potency is evaluated as percentage of inhibition (inhibitor concentration of 1  $\mu$ M) and as IC<sub>50</sub>.

ER Affinity.<sup>29</sup> The binding affinity of selected compounds to the ER $\alpha$  and ER $\beta$  was determined according to Zimmermann et al.<sup>73</sup> using recombinant human proteins. Briefly, 0.25 pM ER $\alpha$  or ER $\beta$  was incubated with [3H]E2 (10 nM) and test compound for 1 h at room temperature. The potential inhibitors were dissolved in DMSO (5% final concentration). Evaluation of nonspecific binding was performed with diethylstilbestrol (10 µM). After incubation, ligandreceptor complexes were selectively bound to hydroxyapatite (5 g/60 mL TE buffer). The formed complex was separated, washed, and resuspended in ethanol. For radiodetection, scintillator cocktail (Quickszint 212, Zinsser Analytic, Frankfurt, Germany) was added and the samples were measured in a liquid scintillation counter (Rack Beta Primo 1209, Wallac, Turku, Finland). From these results the percentage of [<sup>3</sup>H]E2 displacement by the compounds was calculated. The plot of percent displacement versus compound concentration resulted in sigmoidal binding curves. The compound concentration to displace 50% of the receptor bound [<sup>3</sup>H]E2 was determined. Unlabeled E2 was used as a reference. For determination of the relative binding affinity (RBA) the ratio was calculated according to the following equation:

$$RBA(\%) = \frac{IC_{50}(E2)}{IC_{50}(compound)} \times 100$$

This results in an RBA of 100% for E2. After the assay was established and validated, a modification was made to increase throughput. Compounds were tested at concentrations of 1000 IC<sub>50</sub>(E2) and 10000 IC<sub>50</sub>(E2). Results were reported as RBA ranges. Compounds with less than 50% displacement of [<sup>3</sup>H]E2 at a concentration of 10000 IC<sub>50</sub> (E2) were classified with RBA < 0.01%. Compounds that displace more than 50% at 10000 IC<sub>50</sub> (E2) but less than 50% at 10000 IC<sub>50</sub>(E2) were classified with 0.01% < RBA < 0.1%.

**log** P **Determination.** The log P values were calculated from CambridgeSoft Chem & Bio Draw, version 11.0, using the ChemDrawPro, version 11.0, program.

#### ASSOCIATED CONTENT

# Supporting Information

Chemical synthesis, characterization of all compounds, and HPLC purity determination. This material is available free of charge via the Internet at http://pubs.acs.org.

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# ABBREVIATIONS USED

17β-HSD1, 17β-hydroxysteroid dehydrogenase type 1; 17β-HSD2, 17β-hydroxysteroid dehydrogenase type 2; A-dione, 4androstene-3,17-dione; AKR, aldo-keto redutase; APCI, atmospheric pressure chemical ionization; DME, dimethoxyethane; DMSO, dimethylsulfoxide; E1, estrone; E2, 17βestradiol; EDG, electron donating group; equiv, equivalent; EDTA, ethylenediaminetetraacetic acid; ER, estrogen receptor; ESI, electronspray interface; EtOH, ethanol; EWG, electron withdrawing group; HPLC, high pressure liquid chromatography; NAD(P)(H), nicotinamide adenine dinucleotide (phosphate); OB, osteoblast; OC, osteoclast; PBS, phosphate buffered saline; RBA, relative binding affinity; RP, reversed phase; SAR, structure-activity relationship; SDR, short chain dehydrogenase/reductase; SERM, selective estrogen receptor modulator; SF, selectivity factor; T, testosterone

# ADDITIONAL NOTE

For the sake of clarity, IUPAC nomenclature is not strictly followed except for the experimental part where the correct IUPAC names are given.

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