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Biphenyl C-Cyclopropylalkylamides: New Scaffolds for Targeting Estrogen Receptor β

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Abstract

The C-cyclopropylalkylamide scaffold was previously identified as a new structural framework for antiestrogens. A second generation library provided three compounds that bind estrogen receptor (ER) α . Further screening of this library identified ER β hits and inspired another round of SAR. A new focused library was tested for binding to the ERs, and for effects on the growth of breast cancer cell lines and protein levels of common cell cycle regulators.

The estrogen receptors (ERs) are ligand activated transcription factors. ER α and ER β demonstrate ~56% sequence homology in their ligand binding domains, but their ligand-binding pockets differ by only two amino acids: ER α Leu³⁸⁴ versus ER β Met³³⁶, and ER α Met⁴²¹ versus ER β Ile³⁷³.¹ ER β 's pocket is also about 100 Å³ smaller than that of ER α .² Additionally, the tissue distributions of these receptors are distinct.³ There is much data demonstrating the benefits of selectively targeting the ER β . ER β expression inhibits estradiol (E2)-induced proliferation in T47D and MCF-7 breast cancer cells and leads to a reduction in xenograft volume.^{4,5,6} ER β has a growth inhibitory mechanism in medullary thyroid carcinoma, prostate cancer, and ovarian cancer.^{7,8,9,10} Also, ER β -selective agonists demonstrate potential as anti-inflammatory agents.¹¹ Several ER β -selective agents have been identified (Figure 1). The plant-derived ligand genistein (GEN (**1**)) has also been shown to induce DNA strand breaks and cause chromosomal aberrations.^{3,12} The Katzenellenbogen group developed a diarylpropionitrile (DPN, **2**) that is a 100-fold selective ER β agonist.¹³ Recently, biphenyl derivatives have been investigated for ER β -selectivity. Simple 4-OH-biphenyls demonstrated 20-70 fold selectivity for ER β over ER α .¹⁴ In order to attempt to mimic the C-ring of GEN, an oxime moiety was incorporated, and altering the substitutions on the biphenyl core provided various levels of selectivity: oximes **3** and **4** were 11-fold and 43-fold selective, respectively (Figure 1).¹⁵

Previously, through screening of a library of homoallylic amides, allylic amides and C-cyclopropylalkylamides, we identified a new structural scaffold for antiestrogens.¹⁶ The synthesis of a second generation library provided three additional agents that were found to bind tightly to ER α .¹⁷ In the present analysis, we screened this second generation library with a fluorescent E2 derivative for competitive binding to ER β in a fluorescence polarization (FP)

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assay.¹⁸ The percent displacement of the E2 derivative by the test compound was calculated from the difference in measured FP values upon incubation with and without the test sample. The efficiency of displacement demonstrated for ER β by the initial screening library was significantly lower than that for ER α . In fact, only a single compound (**5**; Figure 2) demonstrated a concentration-dependent displacement over the concentration range tested and close to 50% displacement at the highest concentration. The displacement values were: 0.2 μ M, 26%; 1 μ M, 31%; and 5 μ M, 46%. Compound **5** was chosen as the lead structure for efforts to improve ER β -selectivity. Because **5** did not quite reach 50% displacement on ER β , the selection criteria were modified to include structural features previously known to promote ER β affinity.¹⁵ Screening results on ER α were used as coselection criteria, and **5** was not a hit on ER α .¹⁷

Figure 2 demonstrates the three regions in **5** that were further evaluated in the next round of our SAR. The biphenyl core A was shown to be beneficial for selective targeting of ER β , in analogy to Yang *et al.*¹⁵ who used GEN as a lead structure to search for new ER β -selective ligands. The biphenyl moiety was synthetically readily accessible through diaryl Suzuki cross-couplings, and a large number of substituted phenyl precursors were commercially available. Biphenyl carbaldehyde oxime derivatives were synthesized with various substituents. Upon placement of a fluorine and/or chlorine substituent on the biphenyl core while maintaining the phenol group, agents selective for ER β were obtained that demonstrated greater than 10-fold (**3**) and up to 40-fold (**4**) selectivity.¹⁵ Among the second generation library compounds, only one compound containing the biphenyl system satisfied the criteria for hit selection in the screen on ER α ; however, this compound failed to inhibit estradiol-stimulated growth of MCF-7 ER positive breast cancer cells.¹⁷ Replacement of the biphenyl group with a smaller arene, i.e., *p*-chlorobenzene, preserved the displacement activity on ER α and recovered the cell-based activity. This substituent effect indicates that the sterically more demanding biphenyl core is less tolerated by ER α if combined with larger substituents on the nitrogen atom in the benzylic position. When the biphenyl group in **5** was replaced with a phenyl ring while keeping the remaining structure constant, the activity at ER β dropped to ~10% and activity at ER α was retained. Alterations to region B in **5** also led to detrimental consequences in the ability of the compound to displace ES2 from ER β . Compound **5** demonstrated that the best-tolerated function in sector B was a phenylamide group, while other rings were better tolerated for ER α . When the phenylamide was replaced with a biphenylamide, activity at ER β was abolished. Therefore, we concluded that only one biphenyl core was necessary and that increasing the size of the ligand would be detrimental to activity on ER β . As far as the alkyl chain in region C was concerned, an increase in its size to a cyclohexyl group while keeping the remainder of the molecule constant abolished activity at both receptors. Therefore, region C was not altered in subsequent synthetic efforts. Previous results demonstrated that the cyclopropane ring was important for cellular activity.¹⁶ When compounds containing the biphenyl core yet lacking the cyclopropane ring were tested in the FP assay, activity at both receptors was lost. Regions A and B were chosen for further elaboration and the structures were assembled with the biphenyl core, the cyclopropane ring and the amide linkage preserved. The biphenyl group was substituted with a key 4'-OH group expected to promote hydrogen bonding with ER β Glu³⁰⁵ and ER β Arg³⁴⁶. A fluorine substituent was chosen based upon published SAR for biphenyl-containing agents.¹⁵ The phenylamide was changed into a smaller acetylamide, or completely removed. Six new compounds, analogues of **5**, were synthesized, **10a**, **11b-d**, **12**, and **13** (Scheme 1). The palladium catalyst *trans*-(Cy₂NH)₂Pd(OAc)₂ (DAPCy) was used to perform Suzuki couplings in dioxane because it could be used at room temperature, thereby suggesting functional group tolerance, and its past use in similar couplings.^{19,20} This catalyst was easily prepared in yellow crystalline form from palladium (II)acetate and dicyclohexylamine.¹⁹ The cross-coupling of commercially available benzyl- or methyl-*O*-protected 4-hydroxybenzeneboronic acid to 4-bromobenzaldehyde using the DAPCy catalyst provided biphenyl aldehydes **6a-c** in good to excellent yields (65-88%). Imine

formation using titanium(IV) chloride as the Lewis acid initially gave phosphinoylimine **8a** in poor to moderate yields (30-47%).¹⁷ Changing the workup procedure to a fast filtration over a pad of silica gel and washing with ethyl acetate significantly improved the yields (80%) of the phosphinoylimines **8b-c**. One-pot hydrozirconation/transmetallation-aldimine addition in dichloromethane using a microwave (100 °C, 300 W, 10 min) immediately followed by cyclopropanation in the presence of excess diiodomethane at room temperature (16 – 20 h) gave the phosphinoyl-protected biphenyl *C*-cyclopropylalkylamides **9a-c** in moderate overall yields (43-66%). Acidic deprotection followed by acylation gave the final product **10a** (61%) or benzyl-protected compounds **10b-d** (22-75%). Benzyl ethers were easily removed by hydrogenolysis at 1 atm in the presence of palladium hydroxide on carbon to give the final products **11b-d** and **12**. Deprotections and acylations were performed in parallel using a Bradley Greenhouse reactor, and final compounds were purified individually by column chromatography. The deacetylated primary amine **13** was purified as its HCl salt.

This synthetic scheme allowed for diversification in two ways. Substituents on the biphenyl core were varied by changing the substitution on the arylboronic acid and aryl bromide coupling components. The amide linkage also allowed for diversification by altering the acylating agents, analogously to the synthesis of the second generation library. After the acidic deprotection step, the amine salt could be coupled without further purification with diverse acyl chlorides, carbamoyl chlorides, and sulfonyl chlorides in the presence of DMAP.¹⁷

3-Fluoro-4'-hydroxy-1,1'-biphenyl-4-carbaldehyde oxime **3** served as a positive control. The past synthesis of **3** was modified as shown in Scheme 2.¹⁵ Suzuki cross-coupling using DAPCy gave the desired biphenyl aldehyde **6a** in 70% yield. The aryl methyl ether was then removed with BBr₃ at low temperature, and, after simple basic aqueous work up, the resulting deprotected aldehyde was used without further purification. Biphenyl carbaldehyde oxime **3** was formed in two steps by treating the aldehyde **7** with hydroxyloxime hydrochloride in methanol at room temperature.

The six new compounds **10a**, **11b-d**, **12** and **13** and the control compound, oxime **3**, were first evaluated for their ability to compete with ES2 for binding to both ERs using the FP assay as described (Figure 3).¹⁸

Only two compounds, **11b** and **13** (Figure 4), significantly competed with ES2 as shown in Figure 3. The curves were constructed from one-site competition best-fit curves of polarization (mP) versus concentration using GraphPad Prism 4 software. The IC₅₀ values determined from this fit are listed in Table 1. Biphenyl carbaldehyde oxime **3** and E2 were used as positive controls, the former giving an IC₅₀ comparable to the previously determined value in a radioligand binding assay.¹⁵

The compounds were next evaluated for their ability to inhibit E2-induced proliferation of MCF-7 human breast cancer cells and demonstrate their potential as antiestrogens. This was done using two variants of the MCF-7 cell line. The later passage ATCC variant which expresses only ER α , and an earlier passage which expresses both ER α and ER β . The western blot demonstrating this expression is in the supplementary information. Figure 5 shows the growth inhibition curves generated for compounds determined to be selective for ER β using the MTS assay.²¹ Raloxifene (RAL), an antiestrogen known to inhibit the proliferation of MCF-7 cells, was included as a control.²² GraphPad Prism 4 was used to estimate GI₅₀ values using a non-linear best curve fit. In the ER α and ER β expressing MCF-7 cells, compounds **11b** and **13** gave GI₅₀ values of 4.1 and 1.2 μ M, respectively.

In the ER α expressing ATCC MCF7 cells, compound **11b** did not demonstrate inhibition, while compound **13** gave a GI₅₀ value of 51.7 μ M. In these cells, all 3 compounds tested were toxic at a concentration of 25 μ M. Additionally, the growth inhibition of ER-negative MDA-MB231

cells was also assayed.^{23,24} **11b**, **13**, and RAL did not demonstrate significant growth inhibitory activity against this cell line but the compounds again demonstrated toxicity at 25 μ M. This can also be seen in the supplementary information. Therefore, this provides support that the compounds are exerting their action through ER β , because in breast cancer cells expressing both ER β and ER α a significantly more potent anti-proliferative effect was demonstrated compared to breast cancer cells expressing only ER α or no ERs.

c-Myc is known to play a major role in cell proliferation and malignant transformation in breast cancer.²⁵ E2 has been shown to cause a rapid increase in the level of c-Myc expressed by human breast cancer cells.²⁶ Induction of the expression of ER β inhibits c-Myc at both the mRNA and protein level, while increasing the levels of its regulators, the cyclin-dependent kinase inhibitors, p21 (Cip1) and p27(Kip1).⁵ Because of these known links to breast cancer and E2, we chose to investigate how our compounds affected the protein levels of c-Myc, p21, and p27. Western blot analyses (Figure 5) from lysates of MCF-7 cells treated with the indicated compounds and densitometric analyses (Table 2) show compounds **11b** and **13** hold promise because of their ability to increase the levels of cyclin-dependent kinase inhibitors and decrease the levels of c-myc, likely inhibiting cell cycle progression and proliferation. PPT, a 410-fold ER α -selective agonist, and DPN, a 70-fold ER β -selective agonist, were included as controls.^{27,13}

In conclusion, compounds **11b** and **13** demonstrate selectivity for ER β as well as promising antiproliferative effects in breast cancer cells. The compounds are likely acting as agonists for ER β based on what has been shown about ER β 's activity in breast cancer,^{4,5,6} and this encourages future investigation. Compared to the lead compounds **3** and **5**, **11b** and **13** show no affinity for ER α and moderate affinity for ER β . This initial investigation into the biphenyl C-cyclopropylalkylamide pharmacophore demonstrates the potential of this scaffold to deliver ER-selective antiestrogens. Furthermore, the preliminary SAR provides clues for improving selectivity for ER β . A subtype-selective agent will help to understand the complex biology of the ERs and their interplay. The arrival of new ER ligands with differential subtype selectivity ratios enables the tailoring of antiestrogenic or estrogenic therapy according to the condition and level of receptor isoforms present in patients.

References and notes

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18. The ER competitor assays were performed according to the manufacturer's (Invitrogen, Carlsbad, CA) recommendations with some modifications. Human recombinant ER was used at the recommended concentration of 15 nM and the Fluormone™ ES2 was used at a concentration of 1 nM. Aliquots (20 µL) of the mixture of ER and Fluormone™ ES2 were distributed in 384-well, black flat-bottom plates and serial dilutions of test compounds in DMSO were added. Each compound was tested in duplicate, and 1 µM E2 was used as a positive control. The DMSO concentration was kept at 1% (v/v) throughout the experiment. After 2 h, the fluorescence polarization (FP) was measured using an Analyst™ AD & HT Assay Detection Systems reader (Molecular Devices, Sunnyvale, CA) equipped with 485 nm excitation and 530 nm emission filters with the appropriate FL505 dichroic mirror. Data was then analyzed by linear regression using GraphPad Prism's one-site competition method. The fit was constrained by the high polarization control, which was the ER/ES2 complex with no competition, and the low polarization control, which was the ER/ES2 complex with 1 µM E2 for 100% competition.
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21. MCF-7 breast cancer cells were seeded in 96-well plates at a density of 4×10^3 cells per well in phenol redfree RPMI-1640 medium containing 10% charcoal-dextran stripped FBS and allowed to adhere overnight. Test compounds were then added (concentrations ranging from 25 pM to 50 µM) along with 1 nM E2, and cells were incubated for 6 days. Effect of the test compounds on growth was determined using the CellTiter Aqueous One assay system (Promega, Madison, WI), which utilizes the 3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium (MTS) dye reduction assay with phenazine methosulfate as the electron acceptor. Absorbance was measured at 490 nm using an M5 plate reader (Perkin Elmer, Boston, MA) and the background absorbance at 630 nm was subtracted after 2 h incubation with the reagents. A plate of control cells (16 wells) was measured at day 0 to establish 0% growth. E2-stimulated growth at day 6 was set to 100% growth. GraphPad Prism 4 (GraphPad Software Inc, La Jolla, CA) software was used for constructing dose-response curves and calculating GI₅₀ values.
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24. MDA-MB231 cells were plated in 96-well plates at 1×10^3 cells/well and allowed to adhere for 72 h in phenol redfree RPMI-1640 containing 10% FBS. Test agents were added (concentrations ranging from 3.2 nM to 50 µM) and cells were incubated for 72 h. Cell density was determined using the MTS assay as described above.
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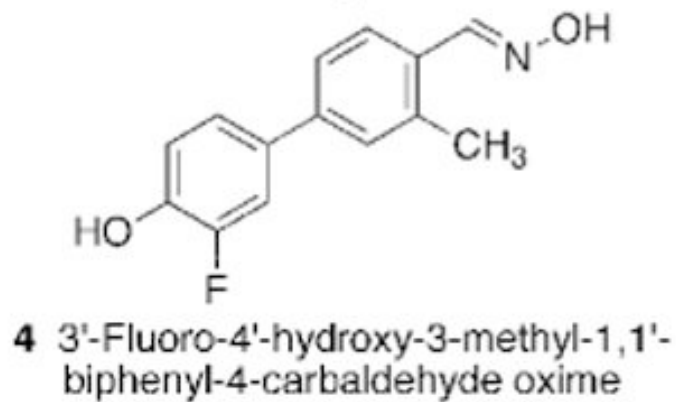
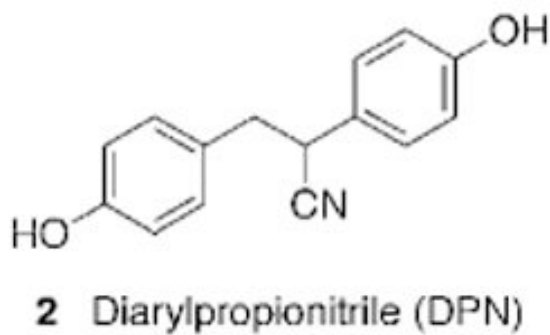
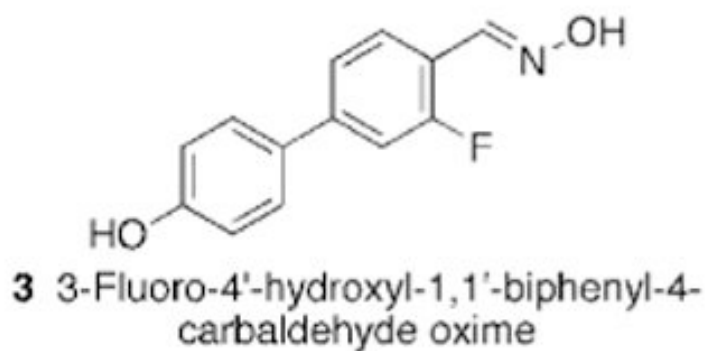
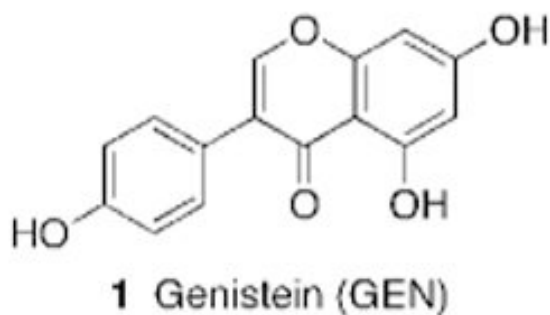


Figure 1. Structures of several known ER β -selective ligands: GEN (**1**), DPN (**2**), and biphenyl carbaldehyde oximes **3** and **4**.

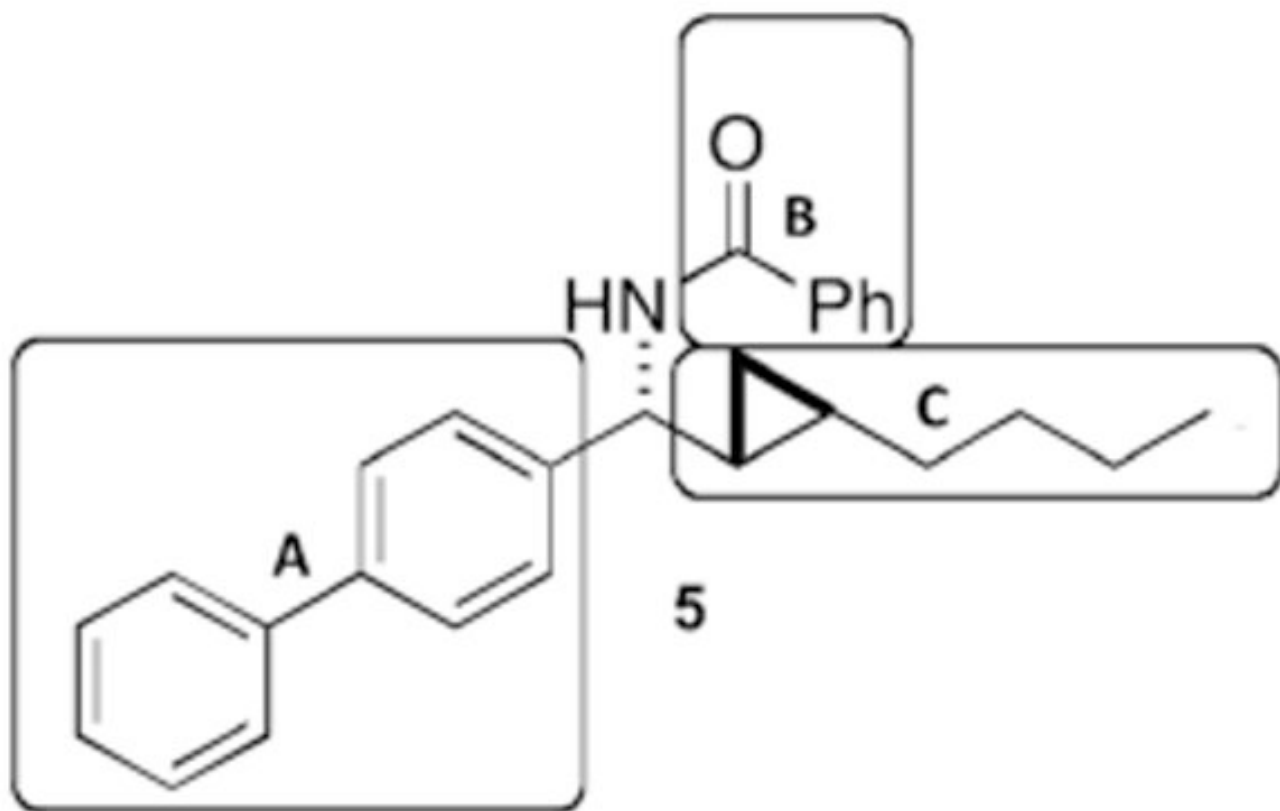


Figure 2.
Lead compound **5**, and subunits **A**, **B** and **C** used in the SAR.

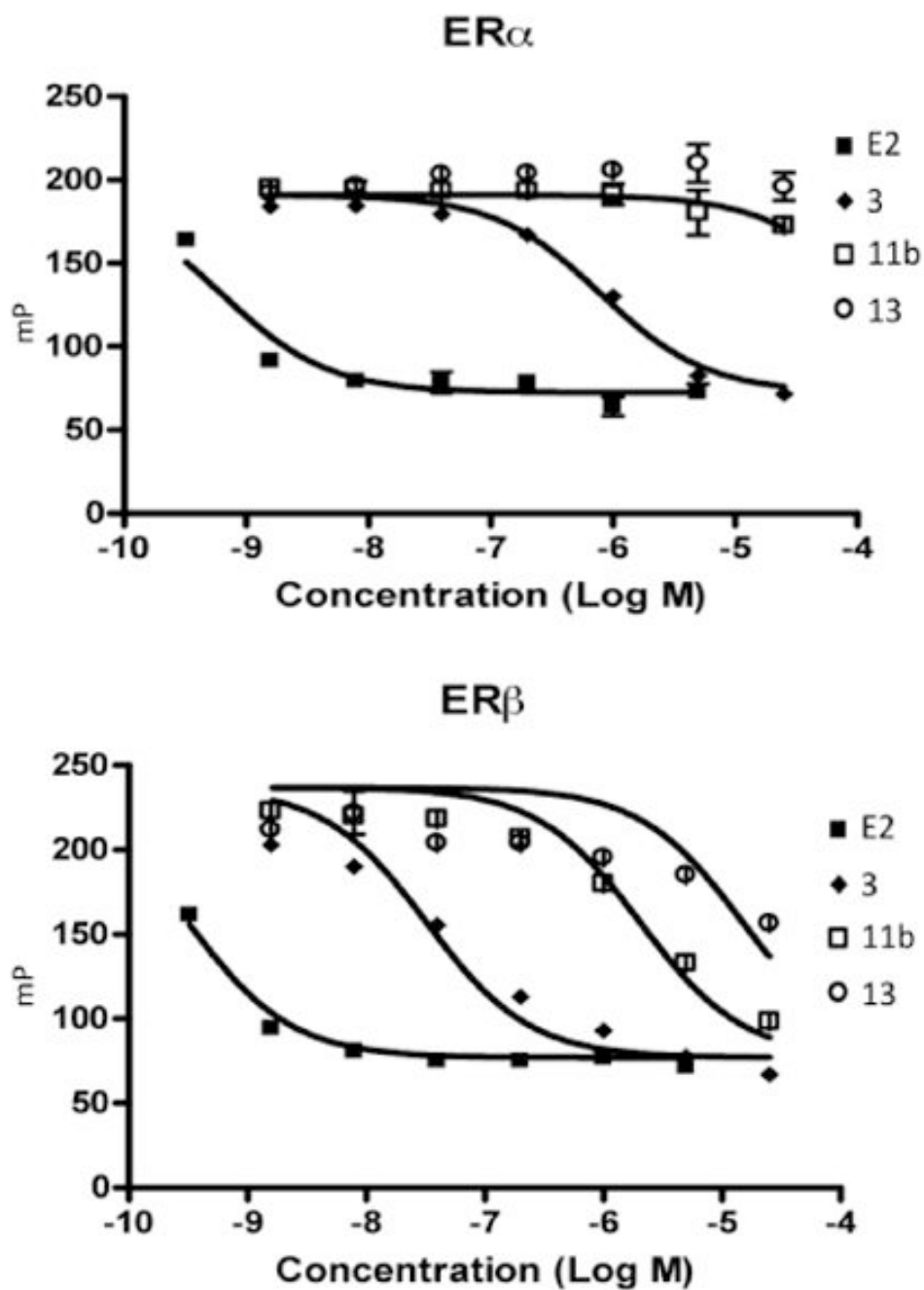


Figure 3. Competition FP assay for **11b** and **13** with E2 for binding to ER α or ER β . E2 and **3** were used as controls.

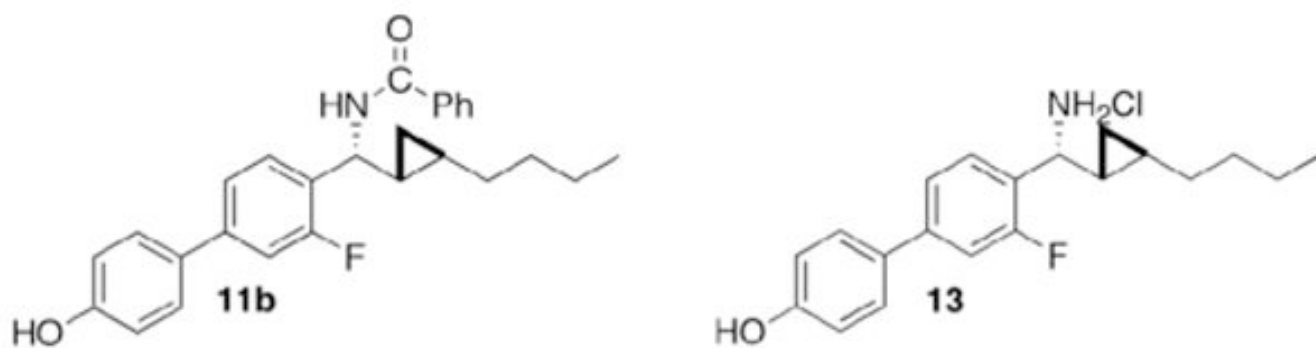


Figure 4.
Lead compounds **11b** and **13**.

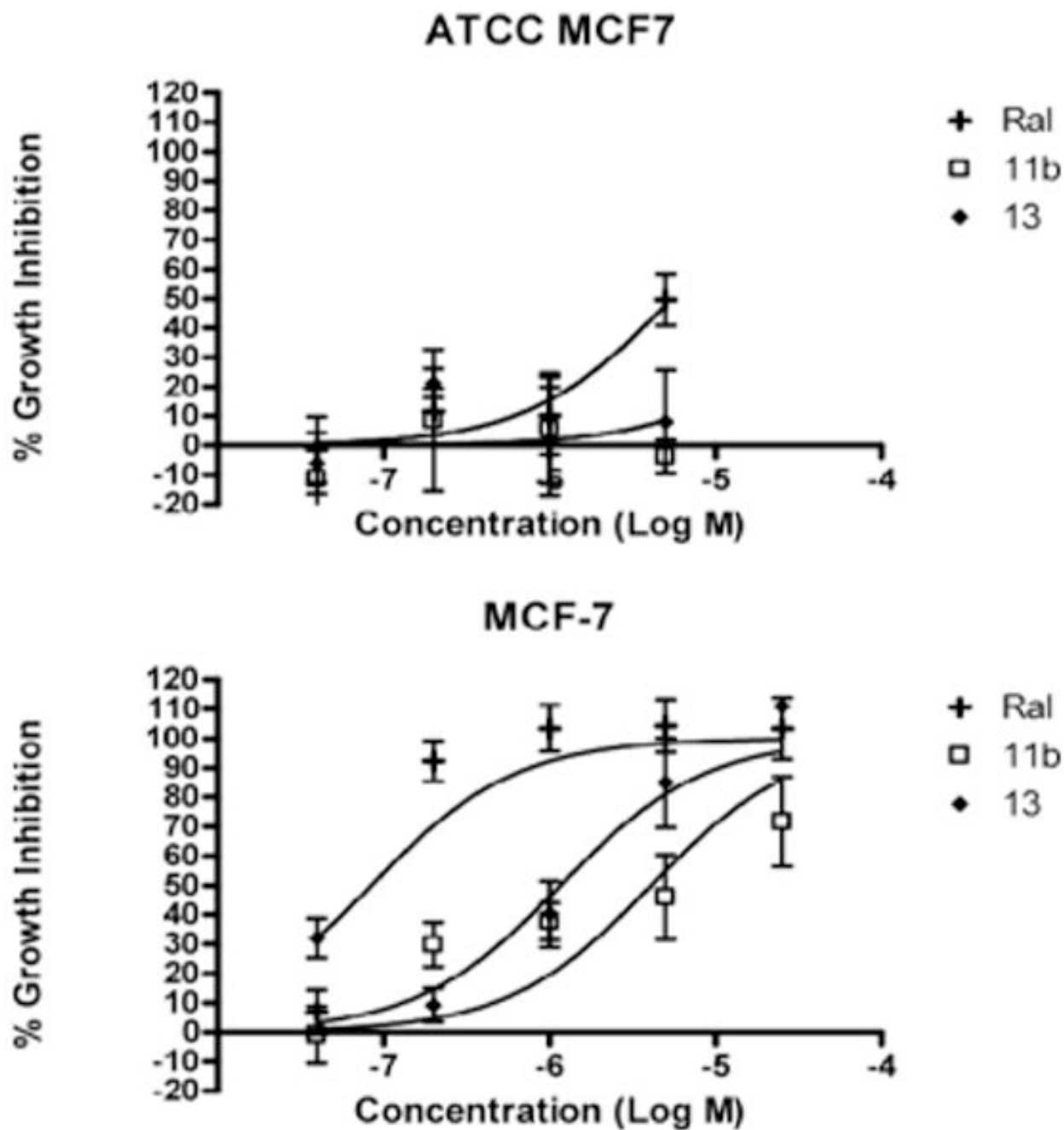


Figure 5. Inhibition of E2-stimulated growth of ATCC MCF7 (*Top*) or MCF-7 (*Bottom*) cells by biphenyl C-cyclopropylalkylamides **11b** and **13**. RAL was used as a positive control. Data represents % growth inhibition (mean \pm SD, N = 3).

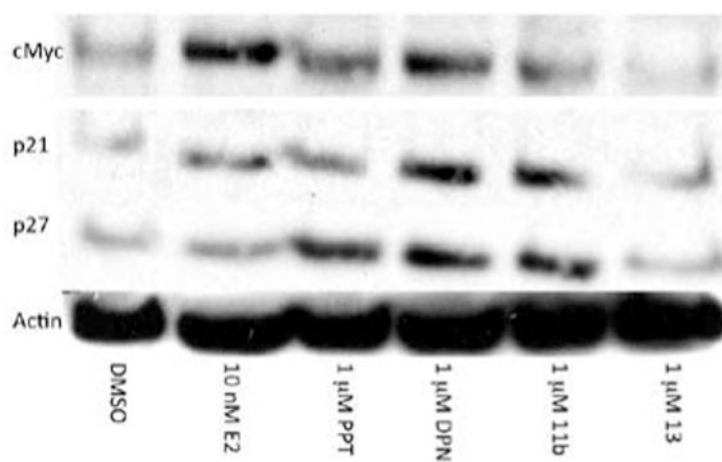
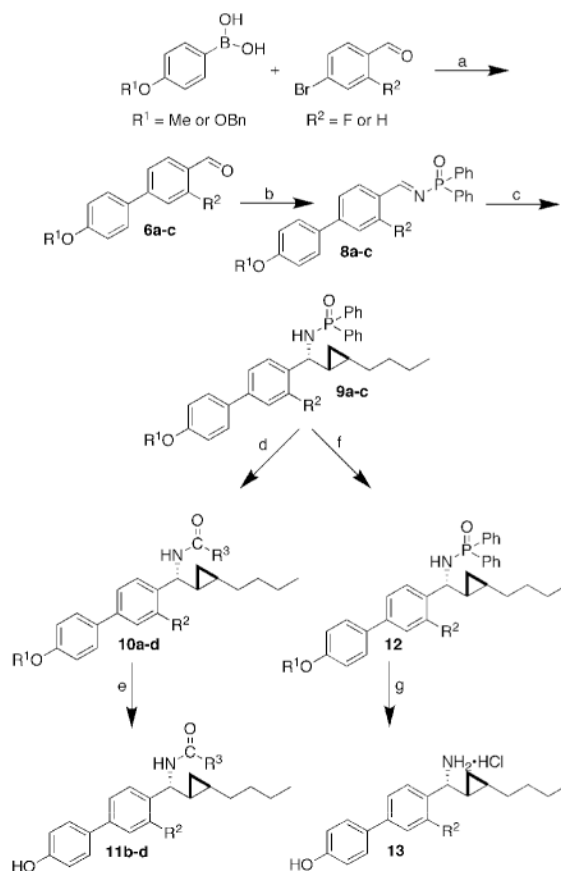
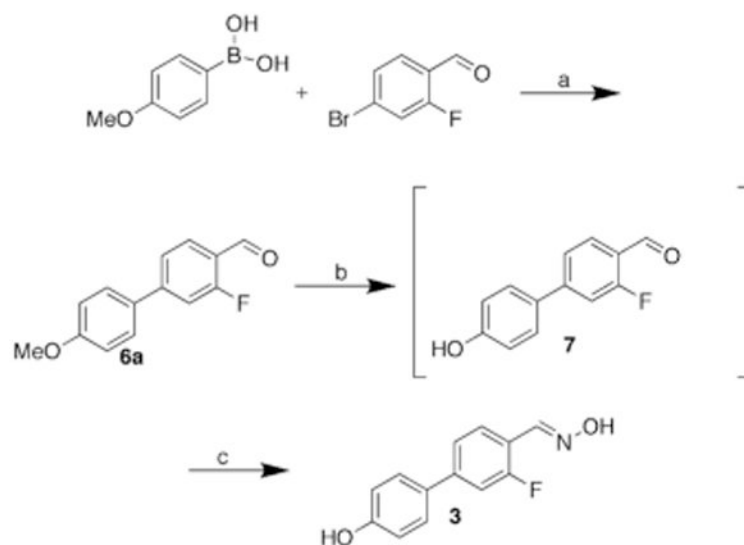


Figure 6. Western blot analysis of the levels of c-Myc, p21, p27, and actin in lysates of MCF-7 cells treated with 10 nM E2, 1 μM PPT, 1 μM **11b** and 1 μM **13** for 12 h.

**Scheme 1.**

Overview of the synthesis of six new biphenyl substituted C-cyclopropylalkylamides. Reagents and conditions: *a*, DAPCy (2 mol%), Cs₂CO₃, dioxane, r.t.; *b*, Ph₂P(O)NH₂, TiCl₄, Et₃N, r.t.; *c*, i) Cp₂Zr(H)Cl, 1-hexyne, CH₂Cl₂, ii) Me₂Zn, toluene, -78 to 0°C, iii) μW 100 °C, 300 W, 10 min, iv) CH₂I₂, r.t.; *d*, (1) HCl, MeOH, (2) R³COCl, DMAP, *i*Pr₂EtN, r.t.; *e*, H₂, Pd(OH)₂/C, THF, MeOH, r.t.; *f*, H₂, Pd(OH)₂/C, THF, MeOH, r.t.; *g*, HCl, MeOH. Percent yields: **6a**, 70; **6b**, 88; **6c**, 50; **8a**, 47; **8b**, 67; **8c**, 40; **9a**, 30; **9b**, 68; **9c**, 40; **10a**, 61; **10b**, 57; **10c**, 22; **10d**, 75; **11b**, 69; **11c**, 92; **11d**, 94; **12**, 80; **13**, 43. R¹ = Me, R² = F (**6a**, **8a**, **9a**); R¹ = Bn, R² = F (**6b**, **8b**, **9b**); R¹ = Bn, R² = H (**6c**, **8c**, **9c**); R¹ = Me, R² = F, R³ = Ph (**10a**); R¹ = H, R² = F, R³ = Ph (**10b**, **11b**); R¹ = H, R² = H, R³ = Ph (**10c**, **11c**); R¹ = H, R² = F, R³ = Me (**10d**, **11d**); R² = F (**12**, **13**).

**Scheme 2.**

Synthesis of positive control agent, 3-fluoro-4'-hydroxy-1,1'-biphenyl-4-carbaldehyde oxime **3**. Reagents and conditions: *a*, DAPCy (2 mol%), Cs_2CO_3 , dioxane, r.t, 70%; *b*, BBR_3 ; *c*, $\text{NH}_2\text{OH}\cdot\text{HCl}$, 42%.

Table 1
IC₅₀ values for competition at both ERs from the FP assay.

Compounds	ER α IC ₅₀ , nM ^a	ER β IC ₅₀ , nM ^a
E2	0.627	0.316
3	795	32.3
11b	> 100,000	2048
13	Na	15030

^aValues are means of three experiments, (na = not active).

Table 2

Relative fold changes of protein levels induced by the indicated treatment in MCF-7 cells compared to DMSO vehicle treatment.

	10 nME2	1 μ M PPT	1 μ M DPN	1 μ M 11b	1 μ M 13
cMyc ^a	3.84	2.17	2.67	0.77	0.59
p21 ^a	1.15	4.05	3.47	3.64	1.10
p27 ^a	1.76	2.47	3.32	3.90	1.51

^aImageJ was used to determine band intensity, and levels were standardized to actin.