

Bisphenol A derivatives act as novel coactivator binding inhibitors for estrogen receptor β

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Running title

Dual activity of bisphenols as ER α -agonists and ER β -antagonists

Abstract

Bisphenol A and its derivatives are recognized endocrine disruptors based on their complex effects on estrogen receptor (ER) signaling. While the effects of bisphenol derivatives on ER α have been thoroughly evaluated, how these chemicals affect ER β signaling is not well understood. Herein, we identified novel ER β ligands by screening a chemical library of bisphenol derivatives. Many of the compounds identified showed intriguing dual activities as ER α agonists and ER β antagonists. Docking simulations suggested that these compounds act as coactivator binding inhibitors (CBIs). Direct binding experiments using wild-type and mutated ER β demonstrated the presence of a second ligand interaction position at the coactivator binding site in ER β . Our study is the first to propose that bisphenol derivatives act as CBIs, presenting a critical view point for future ER signaling-based drug development.

Keywords

endocrinology, estrogen receptor, gene transcription, humoral response, inhibitor, nuclear receptor, transcriptional coactivator

Introduction

Estrogen receptors (ERs) are members of the nuclear receptor family of transcription factors that directly bind to consensus nucleotide sequences to induce gene transcription. 48 human nuclear receptors have been identified, including those for sex steroid hormones, glucocorticoids, retinoids, and vitamin D (1, 2), with many of these receptors recognized as therapeutic targets for a wide range of diseases (3). In particular, ERs are major drug targets for breast cancer (4) and menopausal disorders. Two ER isoforms exist, ER α and ER β , that have high amino acid similarity in both the DNA-binding domains (DBDs) and ligand-binding domains (LBDs) (5). Many ER α and/or ER β -associated gene disruption experiments have been reported (6). Female mice lacking ER α are infertile, while males exhibit decreased fertility (7). Disruption of ER α in female mice leads to hypoplastic uteri, and ER α -disrupted females do not respond to estradiol treatments. ER β knockout mice present with less severe phenotypes than those with ER α knockout, even though ER β -disrupted female mice are subfertile predominantly due to reduced ovarian efficiency (8). Moreover, ER α and ER β double-knockout mice show normal reproductive tract development during the prepubertal period. However, those animals present with similar features to ER α knockout mice during adulthood. Furthermore, this diagnostic phenotype indicates that ER β plays a role in oocyte progression in the postnatal ovary (9, 10). Both ER α and ER β are activated by endogenous estrogens, however, their expression patterns and actions are different (11), with each receptor assumed to have specific biological functions.

A growing body of work in laboratory animals supports bisphenol A (BPA) as an endocrine disrupting chemical (EDC) (12) that has adverse effects on not only the female reproductive system, but also on the brain and immune system (13). BPA is used extensively as a raw material for making polycarbonate plastics and epoxy resins. However, its likely adverse effects on humans, especially infants and fetuses, has recently led to BPA being phased out of polycarbonate plastic and resin production (14). Various BPA derivatives have been developed to create more firm and stable plastics and resins, and these derivatives are now preferred as raw materials (15) (Fig. 1). However, BPA analogues have already been detected in the environment (15, 16). Fluorine-containing BPA, i.e., bisphenol AF (BPAF, 2,2-Bis(4-hydroxyphenyl)hexafluoropropane, CAS No. 1478-61-1), is seen as a practical alternative to BPA, despite reported estrogenic activity in MCF-7 breast cancer cells (17). Eight BPA derivatives, including BPAF, have been detected in sediments collected from industrialized areas (18) and indoor dust (19). In addition, BPA analogs have been found in urine samples from individuals living close to a BPAF manufacturing plant (20) and a municipal solid waste incineration plant (21). Chlorine-containing BPA, i.e., bisphenol C, (BPC, also known as bisphenol C2 or bisphenol Cl2, 1,1-Dichloro-2,2-bis(4-hydroxyphenyl)ethylene, CAS No. 14868-03-2), is a beneficial substrate for polymer production due to the high thermal stability of BPC-containing polycarbonate (22, 23, 24). Notably, BPC is structurally similar to two banned pesticides dichlorodiphenyltrichloroethane (DDT, 1,1'-(2,2,2-trichloroethylidene)bis(4-chlorobenzene), CAS No. 50-29-3) and methoxychlor (1,1'-(2,2,2-trichloroethylidene)bis(4-methoxybenzene), CAS No. 72-43-5) (25, 26). Based on its high affinity for endogenous ERs in MCF-7 cells (27), BPC was considered but ultimately not included in the list of *in vitro* endocrine disruptors by the Interagency Coordinating Committee on the Validation of Alternative Methods (ICCVAM) (NIH Publication No: 03-4503) in 2003. Historically, the designation of 2,2-bis(4-hydroxy-3-methylphenyl) propane (CAS No. 79-97-0, which does not have chlorine atoms), as BPC has led to some confusion

in the literature, however chlorine-containing BPC has been detected in human breast milk (28).

ER α and/or ER β are major targets of EDCs which interfere with their estrogen-responsive signaling pathways (29). Human ER α and ER β have almost identical DBDs, differing by only two amino acids, and both receptors bind the same estrogen-response elements in transcriptional control regions. Although ER α and ER β also have similar LBDs, they have some distinctive features in terms of ligand selectivity and target gene regulation (30). Endogenous estrogen, 17- β estradiol (E2), binds to ER α slightly stronger than to ER β . Similarly, BPA binds ER α with higher affinity than ER β , although its binding abilities are much weaker than those of E2. In contrast, BPAF and BPC display higher affinity for both ER α and ER β than BPA, with a preference for ER β over ER α binding. BPAF and BPC show antagonistic activity against ER β in reporter gene assays using HeLa cells (31, 32). BPAF and BPC show much stronger antagonist activity for ER β than ER α , (32, 33). While crystal structures have provided insight into ER α activation/inactivation mediated by of BPAF and BPC binding (32, 33), the structural changes induced by the strong antagonistic activity of BPAF and BPC against ER β are not well established. Recently, we found that the bisphenol moiety is a privileged structure for ER α . Here we describe the biphasic binding of BPAF and BPC to ER β and propose a novel two-site binding model of the ER β -BPC complex, based on the crystal structure of 4-hydroxytamoxifen (4OHT) bound to ER β . This is the first study to mechanistically associate the antagonistic actions of EDCs with interactions at the coactivator-binding site, thereby providing insight into developing safer raw materials that do not exhibit endocrine-disrupting features.

Results

The bisphenol scaffold binds both ER α and ER β

We screened a library of 119 bisphenol derivatives and related compounds using a radioligand competitive binding assay with [3 H]E2 for ER β . Some of these bisphenol derivatives have been detected in human biological samples (16). The CAS registry numbers (RNs), common names, and IUPAC names are provided in Supplementary Table 1. We found 18 bisphenol derivatives with similar or stronger ER β binding compared to BPA (Table 1 and Fig. S1). BPC showed the strongest ER β (IC₅₀: 2.99 nM), and highest ER α (IC₅₀ of 2.81 nM) binding affinity of the derivatives examined. The second strongest ER β binding was seen with Compound No.2 (4,4'-(1,3-dimethylbutylidene)bisphenol; IC₅₀: 16.1 nM), although higher affinity was measured with ER α (IC₅₀: 5.75 nM). 4,4'-(1,3-Dimethylbutylidene)bisphenol 2,2-bis(*p*-hydroxyphenyl)-1,1,1-trichloroethane (HPTE) (3) and BPAF showed comparable binding ability to ER β (IC₅₀: ~18 nM). Contrary to the results for 4,4'-(1,3-dimethylbutylidene)bisphenol (2), HPTE (3) and BPAF were preferential ER β ligands, displaying three times stronger binding to ER β than ER α . Although bisphenol Z (BPZ, 5), 4,4'-(2-ethylhexylidene)bisphenol (6) and 4,4'-(2-hydroxybenzylidene)-bis(2,3,6-trimethylphenol) (7) showed similar results to BPAF, they bound more strongly to ER α . The majority of the chemicals tested elicited comparable binding to both ER α and ER β . Of the 18 derivatives with similar or stronger ER β binding compared to BPA, 14 showed slightly stronger binding abilities to ER α than ER β (Table 1). We reported that 18 bisphenol derivatives bound to ER α more strongly than did BPA (34). Bulky functional groups at their sp³-carbon connecting two phenol groups were beneficial for ER β binding, similar to the results previously observed for ER α (34). However, ER β binding abilities did not precisely correlate with those of ER α . Fluorene derivatives, 9,9-bis(4-hydroxyphenyl)fluorene (14)

and 9,9-bis(4-hydroxy-3-methylphenyl)fluorene (15), not only bound to ER α (34, 35) but also to ER β (35), with their ER β binding ability stronger than that of BPA. Bisphenol derivatives possessing halogen atoms between two phenol groups, especially chlorine-containing derivatives, showed strong ER β binding.

To gain insight into the differences observed in ER β and ER α binding, we compared the ligand binding cavities in the deposited ER β and ER α LBD crystal structures. The sizes of the canonical binding pockets were calculated for 45 ER α and 25 ER β structures in their active conformations using SiteFinder function, and the amino acid residues surrounding the bound ligands identified (Tables S2 and S3). The average ER β pocket was smaller than for ER α , (430.9 Å³ and 369.3 Å³ for ER α and ER β , respectively; Fig. 2A). The typical ligand-binding pockets of each receptor in the active conformation is illustrated (Figs. 2C and 2D). Moreover, the average size of the ligand binding pocket in 17 β -estradiol-bound ER α and ER β structures was 419.4 Å³ and 385.0 Å³, respectively, and in genistein-bound ER α and ER β structures was 475.9 Å³ and 375.8 Å³, respectively. Although these results suggested that ER α is able to accept larger ligands than ER β , the amino acid residues surrounding the ligands differ slightly. Some of the smaller ligands fit more adequately into the ER β compared to the ER α ligand binding pocket.

BPC and BPAF bind but fail to activate ER β

Reporter assays using HeLa cells were performed to evaluate ER β transcriptional activity induced by BPA, BPC, BPAF, and 17 bisphenol derivatives (Fig. 2B). BPA elicited the strongest ER β agonistic activity of the derivatives, with the activity at 10 μ M comparable to that seen with the endogenous ligand E2 despite its affinity being 400 times weaker than that of E2. 4,4'-(1,3-dimethylbutylidene)bisphenol (2) and bisphenol B (8) achieved ~50% of BPA-induced transcriptional activity at the highest concentration of 10 μ M. While compound 2, found as an impurity in industrial-grade BPA, has been shown to function as an ER α agonist in yeast-two hybrid assays (36), our results reveal a high affinity for and functional activation of ER β . Compound 2 and 8 are structurally similar to BPA, possessing one methyl group on the sp³-carbon that bridges the two phenol groups, suggesting that this conformation is beneficial for ER β activation. BPC, HTPE, BPAF, BPZ, 1,1-bis(4-hydroxy-3-methylphenyl)cyclohexane (9), 9,9-bis(4-hydroxy-3-methylphenyl)fluorene (15), and 2,2-bis(4-hydroxy-3-methylphenyl)propane (18) functioned as partial agonists, inducing 20% to 30% of the E2-induced transcriptional activity. The transcriptional activity of BPC, HTPE, and BPAF was consistent with a previous report investigating ER α and ER β , in which these compounds elicited weaker activity against ER β than ER α (32, 33). Surprisingly, 4,4'-(2-ethylhexylidene)bisphenol (6), 4,4'-(2-hydroxybenzylidene)-bis(2,3,6-trimethylphenol) (7), bisphenol M (10), α , α , α' -tris(4-hydroxyphenyl)-1-ethyl-4-isopropylbenzene (12), bisphenol P (16), and α , α' -bis(4-hydroxy-3,5-dimethylphenyl)-1,4-diisopropylbenzene (20) showed no agonist activity against ER β . These findings contrast with ER α , where the majority of bisphenol derivatives with strong binding affinity also showed strong agonistic activity (34).

BPA derivatives function as ER β antagonists

The finding that many BPA derivatives with high binding affinities showed almost no agonist activity suggested that they function as ER β antagonists. To explore this possibility, the inhibitory effects of the BPA derivatives (100 pM, 1 μ M, 10 μ M) against 10 nM E2-induced ER β activation were measured (Fig. 2E). BPC showed the strongest antagonistic

activity, with additional halogen-containing bisphenols (i.e., HPTE, and BPAF) also elicited antagonistic activities, consistent with previous reports (31-33). 4,4'-(1,3-dimethylbutylidene)bisphenol (2), which had the second strongest binding ability and partial agonist activity compared to BPA, showed weak antagonist activity, contrasting with its reported ER α agonism. Bisphenol B (8) showed similar weak antagonist activity, with both Bisphenol B (8) and 4,4'-(1,3-dimethylbutylidene)bisphenol (2) inhibiting 50% of BPA-induced activation. Tricycle bisphenols (i.e., bisphenol M (10), α , α , α' -tris(4-hydroxyphenyl)-1-ethyl-4-isopropylbenzene (12), bisphenol P (16), and α , α' -bis(4-hydroxy-3,5-dimethylphenyl)-1,4-diisopropylbenzene (20)) showed antagonistic activity, presumably through the disruption of the active conformation, as reported for ER α (34). While demonstrating no agonist activity, 4,4'-(2-ethylhexylidene)bisphenol (6) and 4,4'-(2-hydroxybenzylidene)-bis(2,3,6-trimethylphenol) (7), suppressed 90% of E2-induced activation at the 10 μ M concentration. Interestingly, the fluorene derivative, 9,9-bis(4-hydroxy-3-methylphenyl)fluorene (15) functioned as a weak antagonist, demonstrating that fluorene derivatives 14 and 15 can exhibit both ER β and ER α antagonistic activity (34, 35). With the exception of the tricyclic bisphenols, these findings indicate that most bisphenol derivatives with strong ER β binding functioned as antagonists, even though they showed only agonist activities to ER α (34).

Docking analysis predicts BPC binding to the surface of ER β

To investigate the contrasting actions of BPA derivatives as ER β antagonists and ER α agonists, we performed docking simulations using the LBD of human ER β and BPC, the strongest binder among the BPA derivatives examined using a competitive binding assay with [3 H]E2. Possible ligand binding sites in 38 deposited ER β crystal structures were identified using SiteFinder, a program for binding site analysis equipped in the Molecular Operating Environment (MOE). Canonical, as well as putative binding sites were ranked according to PLB, a specific parameter in SiteFinder (37). Consistently, the top five predicted sites in each structure were the canonical ligand-binding sites. Interestingly, an actual surface 4OHT binding site close to the hydrophobic groove for the coactivator recognition surface of ER β (PDB ID: 2FSZ) was ranked 11th in the PLB order. Moreover, this location was a predicted binding site on all antagonist-bound ER β structures, based on PLB. Notably, this second site was not predicted as a binding site on over half of the agonist-bound structures (Supplementary Table S4). These predictions suggest that ER β antagonism induced by BPC and other BPA derivatives may be due to inhibition of coactivator recruitment. Next, we performed a docking simulation for ER β LBD and BPC using both its canonical and second binding sites as target rooms. BPC was able to fit and bind in both rooms, with one of its chlorine atoms interacting with the tryptophan residue (Trp335) on helix 5 via halogen interaction (Fig. 3, A and B). The obtained model structure suggested that BPC binding to the second binding site prevented recruitment of coactivators for gene transcriptions, similar to 4OHT (Fig. 3, C and D). We hypothesized that the binding affinity of BPA derivatives to this coactivator binding site would correlate with antagonistic activity. To explore this notion, docking simulations were performed for each BPA derivatives (Fig. S2) and the free energy of ligand binding evaluated using a docking simulation and the GBVI/WSA dG scoring function (larger negative scores indicate more stable ligand/receptor complexes) (38). Correlation of the GBVI/WSA dG scores with the extent of antagonism (reported as the % inhibition of 10 nM E2 induced transcriptional activity) revealed a linear relationship (correlation coefficient of - 0.83), suggesting that

inhibition of coactivator recruitment underlies the antagonism of ER β by BPA derivatives (Fig. 3E).

Biphasic 4OHT binding indicative of two ER β binding sites

To further support the presence of a second ligand binding site, competitive binding assays were performed using BPA, BPC and BPAF and tritium-labeled 4OHT ([3 H]4OHT) (Fig. 4A). Notably, a biphasic dose-response curve was observed for BPC (18.1 nM and 2281 nM IC₅₀) that was not evident in the [3 H]E2 competitive analyses. Similarly, BPAF displayed a biphasic binding curve, albeit with weaker binding at both the high- and low-affinity sites compared to BPC. Moreover, 4OHT showed a biphasic curve, consistent with the 4OHT/ER β crystal structure (PDB:ID 2FSZ). In contrast, BPA, which did not elicit antagonistic activity, showed a sigmoidal curve indicative of a single ligand binding site. Interestingly, the tri-fluorine substitution of the methyl groups in BPAF increased ER β binding ~50 fold compared to BPA. These results confirmed the presence of two distinguishable binding sites for BPC and BPAF on ER β . In contrast, the typical sigmoidal curves seen in E2 competitive binding assays using [3 H]4OHT and [3 H]E2 are indicative of single ligand binding site.

Trp335 is required for biphasic ligand binding

The docking simulations suggested that hydrophobic interactions between the BPA derivatives and the indole group of Trp335 were required for ER β binding, and identified a potential halogen interaction between the chlorine atom of BPC and the indole ring. To determine the contributions of these putative interaction to BPC binding, the corresponding tryptophan was mutated to alanine (A). Saturation binding assays revealed a typical sigmoidal dose-response curve and a K_d of 23.1 nM for E2 against ER β (W335A), indicating preservation of the canonical binding site (Fig. S3, A).

Competitive binding assays confirmed two 4OHT binding sites in ER β , with K_d values of 4.6 nM and 53.1 nM. In contrast, a single binding site was evident in ER β (W335A) (K_d 34.2 nM) (Fig. S3, B). Similarly, the biphasic binding of BPC and BPAF were lost in the ER β (W335A) mutant (Fig. 4, A and B). The IC₅₀ values of 4OHT, BPC, and BPAF were 106 \pm 51 nM, 691 \pm 29 nM, and 1249 \pm 579 nM, respectively. BPA illustrated a typical sigmoidal competitive dose-response curve against ER β (W335A), similar to the result against ER β . These results indicated that replacing Trp for Ala compromises the second 4OHT and BPA derivatives binding site on the surface of ER β LBD.

W335A reduces ER β transcription activity

Reporter assays revealed that E2-induced transcriptional activation was markedly reduced by the tryptophan to alanine substitution in ER β (Fig. 4, C and D). Given that E2 binding ability was retained, this is consistent with reduced coactivator binding. Indeed, in the active conformation, Trp335 interacts with Leu491, Met494, and Leu495 on H12 (Fig. 4E). These results indicated that Trp335 on the ER β coactivator-binding site plays an important role, not only in interacting with bisphenol derivatives, but also in recruiting coactivators on the surface of ER β by stabilizing H12 in its active conformation.

Discussion

Here we report the ER β transcriptional activities of BPA derivatives including BPC and BPAF using a combination of receptor binding and reporter assays. Unexpectedly, our results clearly showed that many BPA derivatives function as ER β antagonists, contrasting with their previously reported ER α agonism. Docking simulations indicated that BPA derivatives bind to a second site located near the coactivator binding site on the surface of ER β -LBD that requires interactions with Trp335. Mutation of tryptophan to alanine led to the loss of this low affinity binding site in ER β . These results indicated that some BPA derivatives act as antagonists, although most of endocrine-disrupting chemicals, including BPA, are assumed ER agonists. We previously reported that most of the BPA derivatives examined in this study act as weak agonists for ER α . The results obtained in this study demonstrate the importance of screening for both agonist and antagonist activity, especially against ER β .

We previously reported that tricyclic bisphenols, i.e., Bisphenol M, α , α , α' -tris(4-hydroxyphenyl)-1-ethyl-4-isopropylbenzene, bisphenol P, and α , α' -Bis(4-hydroxy-3,5-dimethylphenyl)-1,4-diisopropylbenzene, act as antagonists against ER α because of the steric hindrance caused by the third aromatic ring structure (34). This study showed that this feature is also valid for ER β ; tricyclic bisphenols act as antagonists not only for ER α but also ER β . In addition to tricyclic bisphenols, many BPA derivatives, including BPAF and BPC, elicit antagonist activity. Our finding for BPAF and BPC are consistent with reports that both chemicals showed partial agonism for ER α and antagonism for ER β (31, 32, 39, 40).

Several ER α - or ER β -specific agonists have been reported, including propylpyrazole triol (PPT) that selectively binds to and transcriptionally activates ER α (41). The first chemical shown to function as an ER α agonist and ER β antagonist is 2,2-bis(*p*-hydroxyphenyl)-1,1,1-trichloroethane (HPTE), a metabolite of the banned pesticide, methoxychlor [1,1,1-trichloro-2,2-bis(4-methoxyphenyl)ethane] (42, 43). Accumulated knowledge gained from protein crystal structures emphasize the importance of halogens in receptor-ligand interactions (44, 45). We found that in addition to the halogen containing BPAF and BPC, many BPA derivatives display ER α agonist activities similar to HPTE. These results indicate the complexity of establishing the mechanisms of action of environmental chemicals that activate or suppress the physiological functions of one or more nuclear receptors. In particular, antagonist activities might be overlooked if both binding affinity and transcriptional activity are not determined, as environmental chemicals are typically categorized based on the ability to active ERs.

Recent studies have indicated the value of small molecules that bind to coactivator protein-binding sites on nuclear receptors (46). Coactivator-binding inhibitors (CBIs) have been developed for ERs, an androgen receptor, a progesterone receptor, a vitamin D receptor, a thyroid hormone receptor, a pregnane X receptor, a retinoid X receptor, and peroxisome proliferator-activated receptors (47-50). This study is the first to conclude that endocrine-disrupting chemicals can function as CBIs for ER β , indicating the importance of assessing both agonist and antagonist activities of these chemicals.

In summary, we showed that tricyclic bisphenols elicit antagonistic activity against both ER α and ER β . Our results also indicate that many next-generation bisphenols are agonists and antagonists of ER α and ER β . Mutagenesis of an ER β surface amino acid indicated that these next-generation bisphenols act as CBIs. While *in silico* docking analyses support this mechanism of action, future crystallographic studies will be required to provide

more direct information on CBIs. This study highlights the mechanistic complexity of the next-generation of bisphenols acting as endocrine-disrupting chemicals.

Materials and Methods

Chemicals

17 β -estradiol (E2, CAS RN 50-28-2, >98.9%) was obtained from Research Biochemicals International (Natick, MA, USA). 4-hydroxytamoxifen (4OHT, CAS RN 68047-06-3, >98%) and 2,2-bis(*p*-hydroxyphenyl)-1,1,1-trichloroethane (CAS RN 2971-36-0, >98.9%) were obtained from Sigma-Aldrich Inc. (St. Louis, MO, USA). 4,4'-dihydroxydiphenylmethane (bisphenol F or BPF, CAS RN 620-92-8, >99.0%) and hexestrol (CAS RN 84-16-2, >99.0%) were obtained from FUJIFILM Wako Pure Chemical Corporation (Osaka, Japan); the remaining 117 chemicals were purchased from Tokyo Chemical Industry Co., Ltd. (Tokyo, Japan). Dimethyl sulfoxide (DMSO), used to dissolve each compound in a 10 mM stock solution, was obtained from Sigma-Aldrich. Tritium-labeled 17 β -estradiol ([³H]E2, 4458 GBq/mmol) and 4-hydroxytamoxifen ([³H]4OHT, 2960 GBq/mmol) were purchased from PerkinElmer (Waltham, MA, USA).

ER β expression and purification

The LBD of ER β (amino acids 263-530) was expressed as a glutathione *S*-transferase (GST)-fused protein for receptor binding assays. Human ER β cDNA was obtained from OriGene Technologies (Rockville, MD, USA). The cDNA of ER β -LBD was amplified using PCR, and subcloned into an pGEX-6p-1 expression vector (Cytiva, Marlborough, MA, USA). The expression of GST-fused ER β -LBD was induced by 1 mM isopropyl- β -D-1-thiogalactopyranoside (IPTG) in *Escherichia coli* BL21 α at 16°C for overnight. The resulting crude protein was affinity-purified using Glutathione-Sepharose 4B (Cytiva), followed by gel filtration in a Sephadex G-10 column (Cytiva).

Radioligand binding assay

Radioligand binding assays for ER β and ER β (W335A) were performed mainly according to a previously reported method (31, 34). Saturation binding assays were conducted with [³H]E2 or [³H]4OHT using GST-ER β -LBD or GST-ER β (W335A)-LBD to evaluate the binding ability of radio-labeled compounds. The reaction mixtures of each LBD (20 ng) and a series of concentrations of [³H]E2 (0.01–10 nM) or [³H]4OHT (0.1–30 nM) were incubated in a total volume of 100 μ L of binding buffer (10 mM Tris-buffered saline (pH 7.4), 1 mM ethylene glycol-bis (2-aminoethylether)-*N*, *N*, *N'*, *N'*-tetraacetic acid (EGTA), 1 mM ethylenediaminetetraacetic acid (EDTA), 10% glycerol, 0.5 mM phenylmethylsulfonyl fluoride, 0.2 mM leupeptin, and 1 mM sodium vanadate (V)) at 20°C for 2 h, to analyze total binding. Corresponding reaction mixtures, containing 10 μ M non-labeled E2 or 4OHT, were incubated to detect each non-specific binding. [³H]E2 or [³H]4OHT-specific binding was evaluated by subtracting the obtained radioactivity values of total binding from the those of non-specific binding. Following successive incubation with 100 μ L of 0.4% dextran-coated charcoal (DCC) (Sigma-Aldrich) in phosphate-buffered saline (pH 7.4) on ice for 10 min, free radioligands bound to DCC were removed using a vacuum filtration system with a 96-well filtration plate (MultiScreenHTS HV, 0.45-mm pore size, Merck KGaA, Darmstadt, Germany) for the bound/free (B/F) separation. The radioactivity of each eluent was measured using a liquid scintillation counter (LS6500; Beckman Coulter, Fullerton, CA, USA) and Clear-sol I (Nacalai Tesque Inc., Kyoto, Japan).

Calculated specific binding of [³H]E2 was assessed using Scatchard plot analysis (51). Competitive binding assays were performed to evaluate the binding ability of each test compounds using [³H]E2, for a library screening or detailed BPA binding assay. Each compound was dissolved in DMSO to prepare a 1.0 mM stock solution, and further diluted to prepare a serial dilutions (10⁻¹²M to 10⁻⁵ M) in binding buffer. To assess their binding abilities, each compound was incubated with GST-ERβ-LBD or GST-ERβ(W335A)-LBD (20 ng) and radio-labeled ligand (5 nM of [³H]E2 or 5 nM of [³H]4OHT, final concentration) for 2 h at 20°C. B/F separation was performed as described above, and the radioactivity was determined using a MicroBeta microplate counter (PerkinElmer Inc.). The IC₅₀ value of each test compounds was calculated from the dose-response curves generated via nonlinear regression analysis using Prism software (GraphPad Software Inc., La Jolla, CA, USA).

Luciferase reporter gene assay

Transcriptional activities of ERβ and ERβ(W335A) were measured as previously reported previously (31, 34). HeLa cells were maintained in Eagle's minimum essential medium (EMEM) (Nissui Pharmaceutical Co. Ltd, Tokyo, Japan) supplemented with dextran-coated charcoal treated fetal bovine serum (DCC-FBS, 10%, v/v) with at 37°C under 5% CO₂. To evaluate agonistic activity, HeLa cells were seeded at a density of 5 × 10⁵ cells per 60-mm dish and cultured for 24 h, followed by transfection of the reporter plasmid (3 μg, pGL4.23/3×ERE) and each expression plasmid (1 μg, pcDNA3.1/ERβ or pcDNA3.1/ERβ(W335A)) using Lipofectamine LTX with Plus Reagent (Thermo Fisher Scientific, Inc.), according to the manufacturer's instructions. After incubation for 24 h, cells were harvested and seeded onto 96-well plates at 5 × 10⁴ cells/well, and then treated with a series of the test compounds (10⁻¹² M to 10⁻⁵ M, final concentration) diluted with 1% bovine serum albumin/PBS (v/v). After a 24-h incubation, luciferase activity was measured using the ONE-Glo™ Luciferase Assay System (Promega Co., Madison, WI, USA) on an EnSpire multimode plate reader (Perkin Elmer, Inc.). To analyze antagonistic activity, serial concentrations of test compounds (10⁻¹² M to 10⁻⁵ M) were treated in the presence of 10 nM E2, which normally induces full transcriptional activity levels in transiently expressed ERβ.

Docking simulation of each antagonist onto the ERβ LBD

Three-dimensional (3D) coordinates of the compounds were obtained from the Cambridge Structural Database (CSD-Core, The Cambridge Crystallographic Data Centre, Cambridge, UK). Ligand IDs of compounds utilized for docking simulations are summarized in Supplementary Table S5. For the compounds with no corresponding entry in the CSD-System, 3D coordinates were constructed *in silico* using Gaussian 16 (Gaussian, Inc., Wallingford CT, USA), with the basis set of 6–31G. Docking simulations for the ligand/ERβ complex were performed using a Dock functions in the MOE package (Chemical Computing Group, Montreal, QC, Canada); the free energy of each complex was evaluated according to its GBVI/WSA dG score (38). Ligand-binding cavity volumes of the deposited crystal structures were analyzed and calculated using the SiteFinder function in MOE.

Statistical analysis

Significance of the data between experimental groups was determined using unpaired *t*-tests. Data are presented as the mean ± standard deviation (SD), and P values are presented in each figure legend.

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Supporting information

This article contains supporting information (Figs. S1 to S3 and Tables S1 to S5).

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Author contributions

A.M. conceived and designed the experimental approaches. A.M., M.I., and T.M. performed most of the experiments. M.H. contributed to the docking simulation analysis. K.T. and T.I. performed the experiments. A.M. wrote the manuscript. E.Y., M.D., and R.M.E. provided critical comments and contributed to the editing of the manuscript.

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Conflict of interests

Authors declare no competing interests.

Data and materials availability

All data needed to evaluate the conclusions in the paper are present in the paper and/or the Supplementary Materials.

Figures and Tables

figure 1.

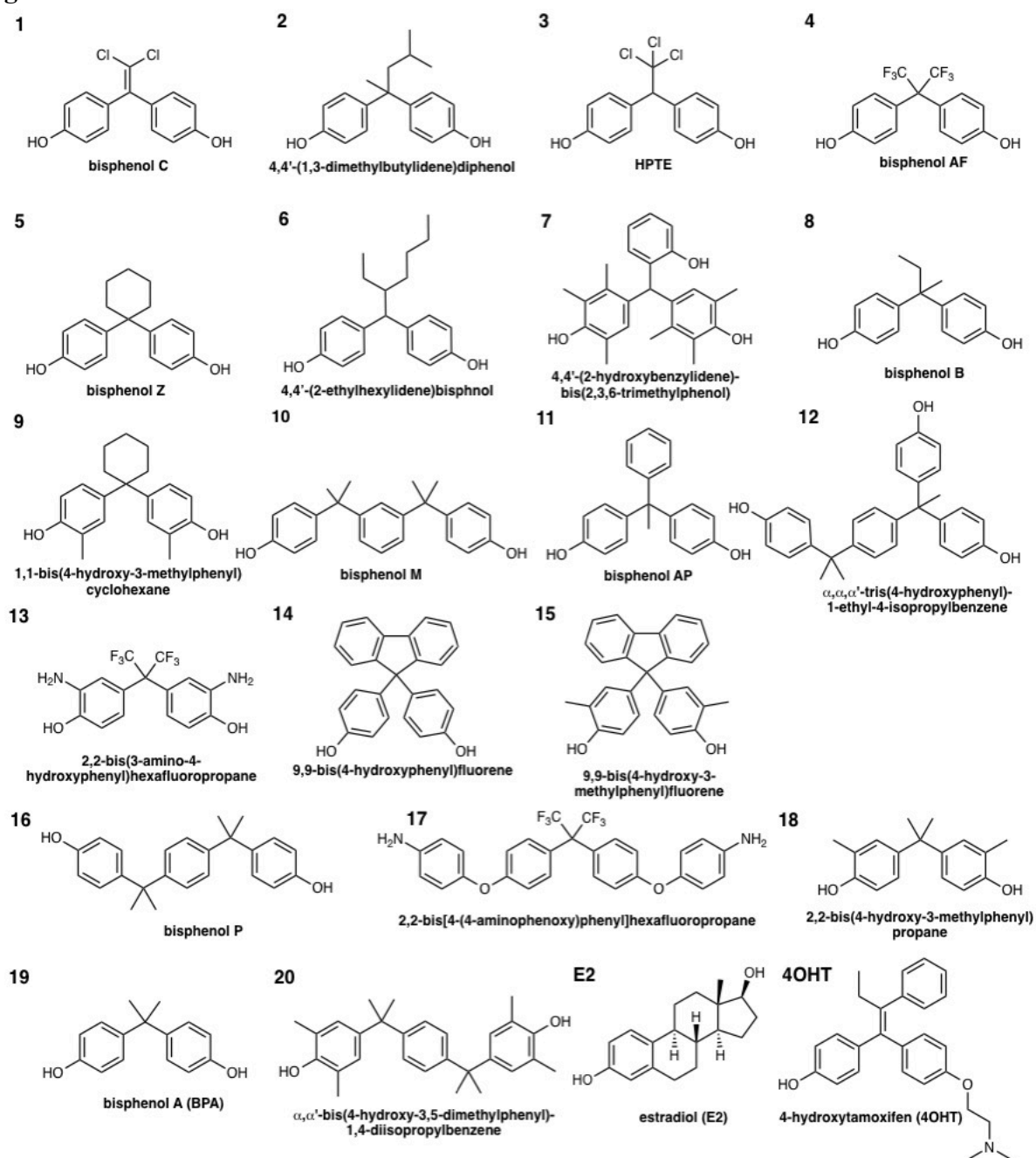


Figure 1

Fig. 1. Structures of BPA derivatives selected via screening using an ERβ binding assay. Chemical structures of E2, 4OHT, and 20 BPA-related compounds exhibited stronger binding abilities than, or comparable to, BPA; BPC had the highest binding ability to ERβ. Fluorine-containing BPA derivatives, i.e., 9,9-Bis(4-hydroxyphenyl)fluorene and 9,9-bis(4-hydroxy-3-methylphenyl)fluorene, exerted stronger binding abilities than did BPA.

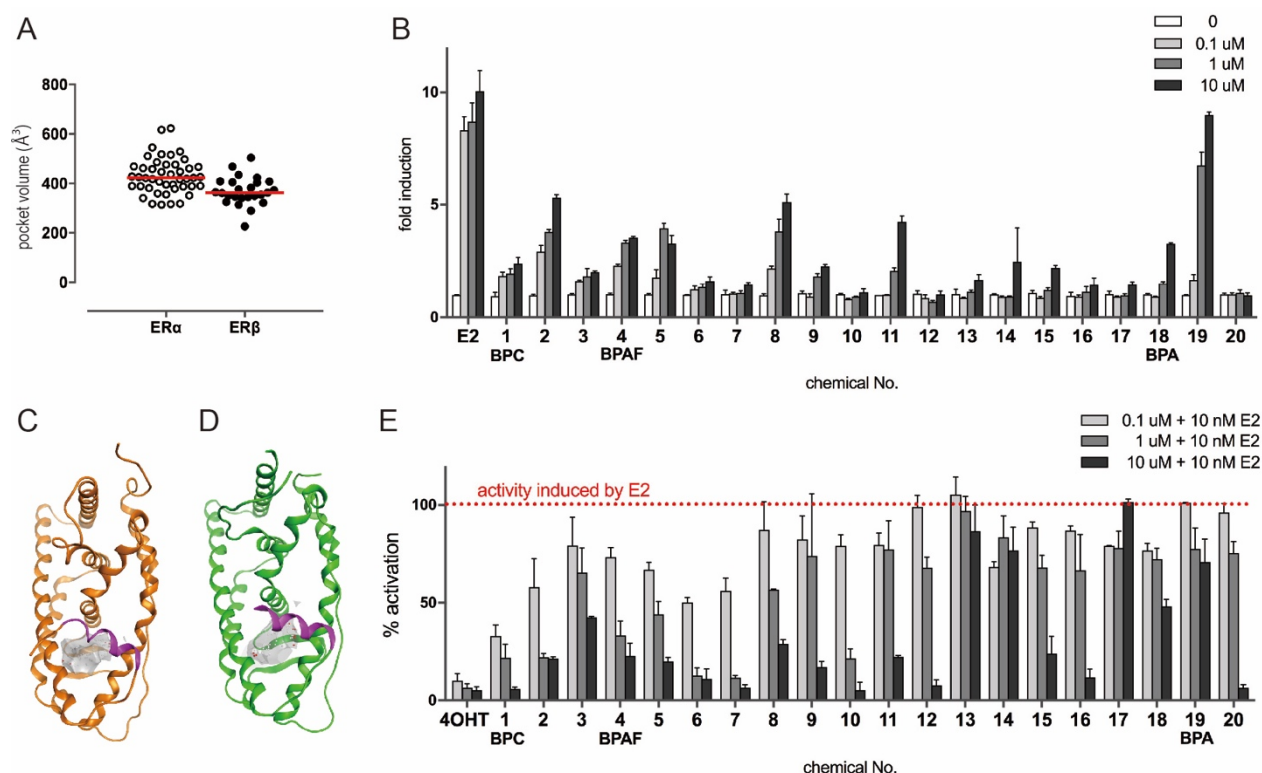


Fig. 2. Differential activities of **BPA derivatives** on **ERα** and **ERβ**. (A) Ligand-binding pocket volumes from ERα (open circles) and ERβ (filled circles) calculated from crystal structures in the presence of activating ligands; average volumes indicated by red lines. (B) Top 10 BPA derivatives binding to ERβ induced partial agonistic activity against ERβ. (C) The ligand-binding pockets of ERα (PDB ID 1QKU) and (D) ERβ (3OLL) are illustrated in gray; estradiol is bound as the ligand. (E) Sixteen chemicals, including tricyclic bisphenols, inhibited more than half of the 10 nM E2-induced transcriptional activity.

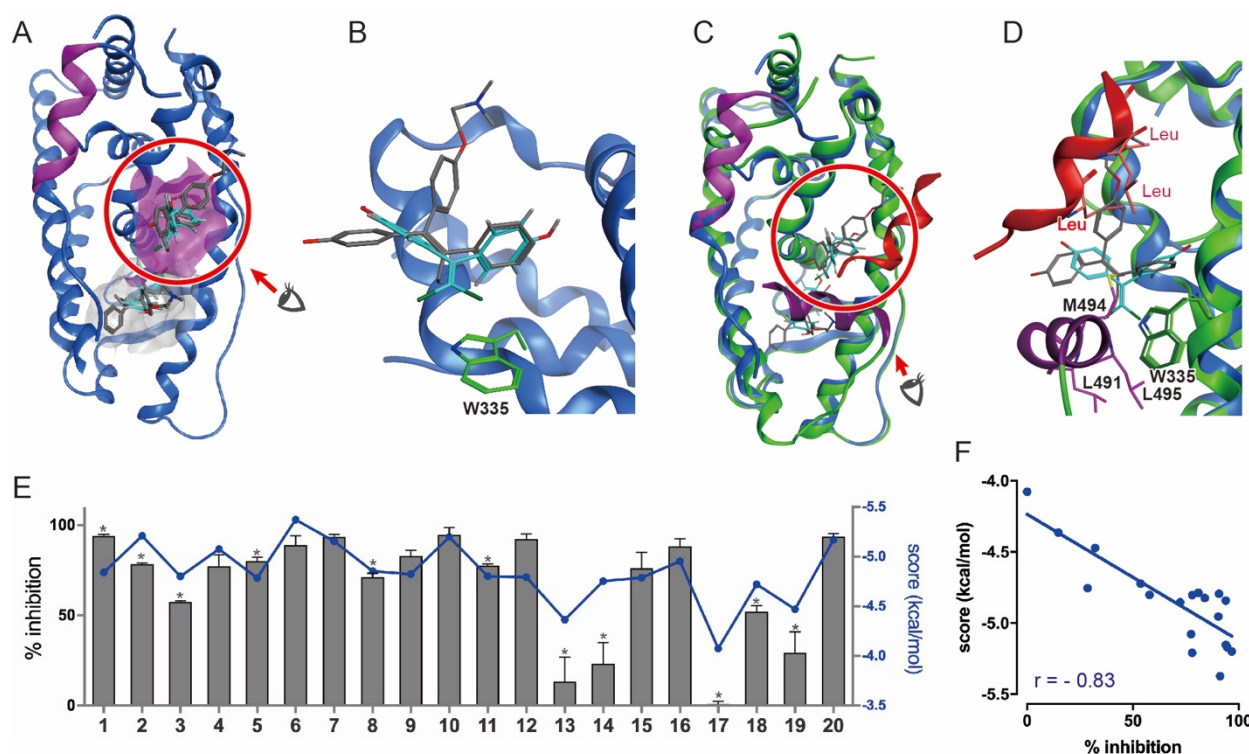


Fig. 3. ERβ harbors two ligands in its LBD. (A) Two BPC bound to ERβ during the docking simulation. The canonical binding site is indicated in gray; the second binding site, located on the surface of the receptor, is shown in magenta. The activation helix, H12, is indicated in magenta. (B) Chlorine, a halogen atom of BPC, interacted with the Trp335 side chain via halogen interaction in the second binding site. BPC and 4OHT are illustrated in blue and gray, respectively, in the stick model. (C) Superimposition of the calculated BPC-bound ERβ structure (blue) and its agonist form with the nuclear receptor coactivator 1, SRC1. (green, PDB ID, 3OLL). SRC1 is indicated as a red α-helix, H12 of its agonist form is indicated in purple, BPC is illustrated in blue, and 4OHT is shown in gray. BPC clashed with the amino acid residues on H12 in the ERβ agonist form; therefore, BPC prevented the ERβ activation. BPC and 4OHT disrupted the SRC1 binding due to steric hindrance of the amino acid residues shown in the red stick models. (D) In ERβ-agonist form, amino acid residues surrounding Trp335 within 4.5 Å on H12 are shown in the purple stick model, while leucine residues on the SRC1 LXXLL motif are indicated via the red stick model. (E, F) Correlation of the calculated binding scores and inhibitory activity for ERβ. Inhibitory activity is defined as the ratio of chemicals inhibiting transcriptional activity induced by 10 nM E2. * $p < 0.05$.

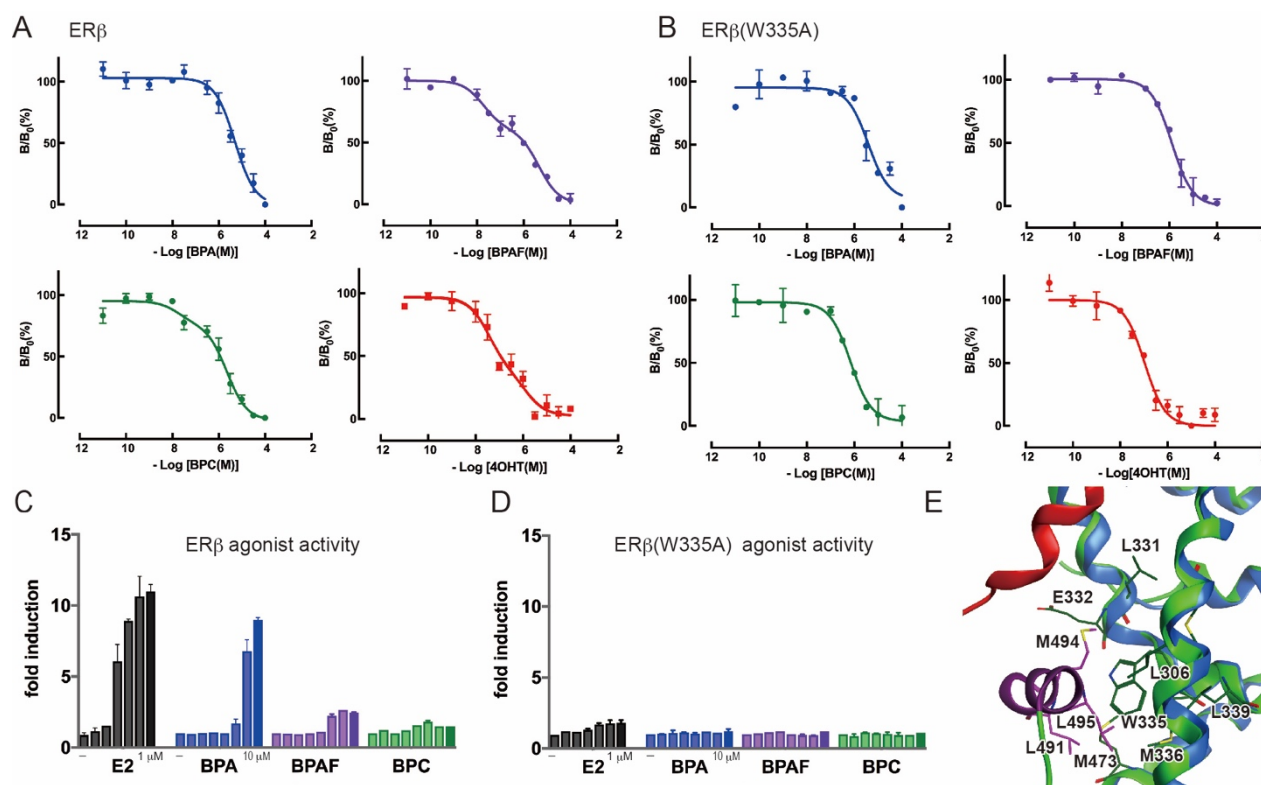


Fig. 4. Binding properties and transcriptional features of BPAF and BPC showed the importance of ERβ W335 for their receptor binding and activation. (A) Detailed competitive binding curves of BPA, BPAF, BPC, and 4OHT using [³H]4OHT illustrated a biphasic binding curve, in which chemicals compete with [³H]4OHT in two binding sites on wild type ERβ. (B) ERβ(W335A) competitive binding assays showed typical sigmoidal binding curves. (C) The reporter gene assay indicated that BPAF and BPC induced weak transcriptional activity in wild type ERβ, while E2 and BPA showed strong transcriptional activity. (D) ERβ(W335A) lost E2 or BPA-induced transcriptional activity, indicating that Trp335 substitution disrupted active conformation. (E) In ERβ agonist form, amino acid residues surrounding Trp335 within 4.5 Å are represented as green and purple stick models. (PDB ID, 3OLL).

Table 1. Receptor binding affinity (mean \pm SD) of BPA derivatives for ER β . Receptor binding affinity were evaluated by competitive binding assay using [3 H] 17 β -estradiol as a radioligand.

Compound No.	Chemicals	Binding affinity (IC ₅₀ , nM)			
		ER β		ER α ^[34]	
E2	estradiol	2.17	\pm 0.6	0.88	\pm 0.13
1	bisphenol C	2.99	\pm 1.0	2.81	\pm 0.61
4OHT	4-hydroxytamoxifen	4.66	\pm 1.5	2.85	\pm 0.20
2	4,4'-(1,3-dimethylbutylidene)bisphenol	16.1	\pm 6.1	5.75	\pm 1.92
3	2,2-bis(<i>p</i> -hydroxyphenyl)-1,1,1-trichloroethane (HPTE)	18.1	\pm 4.9	59.1	\pm 1.5
4	bisphenol AF	18.9	\pm 8.4	53.4	\pm 7.3
5	bisphenol Z	21.6	\pm 1.9	56.9	\pm 0.6
6	4,4'-(2-ethylhexylidene)bisphenol	25.9	\pm 8.5	18.5	\pm 6.7
7	4,4'-(2-hydroxybenzylidene)-bis(2,3,6-trimethylphenol)	41.5	\pm 2.0	12.3	\pm 7.3
8	bisphenol B	79.8	\pm 12.6	195	\pm 44
9	1,1-bis(4-hydroxy-3-methylphenyl)cyclohexane	132	\pm 6.5	38.6	\pm 7.2
10	bisphenol M	148	\pm 80	56.8	\pm 11.7
11	bisphenol AP	158	\pm 33	259	\pm 41
12	α , α , α '-tris(4-hydroxyphenyl)-1-ethyl-4-isopropylbenzene	212	\pm 36	61.7	\pm 10.4
13	2,2-bis(3-amino-4-hydroxyphenyl)hexafluoropropane	224	\pm 113	334	\pm 112
14	9,9-Bis(4-hydroxyphenyl)fluorene	325	\pm 60	2230	\pm 202
15	9,9-bis(4-hydroxy-3-methylphenyl)fluorene	405	\pm 108	321	\pm 103
16	bisphenol P	607	\pm 28	176	\pm 35
17	2,2-bis[4-(4-aminophenoxy)phenyl]hexafluoropropane	609	\pm 81	1030	\pm 375
18	2,2-bis(4-hydroxy-3-methylphenyl)propane	744	\pm 429	368	\pm 22
19	bisphenol A	900	\pm 70	1780	\pm 764
20	α , α '-bis(4-hydroxy-3,5-dimethylphenyl)-1,4-diisopropylbenzene	10000	>	733	\pm 628

Supplementary Materials for

Bisphenol A derivatives act as novel coactivator binding inhibitors for estrogen receptor β

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This PDF file includes:

Figs. S1 to S3
Tables S1 to S5

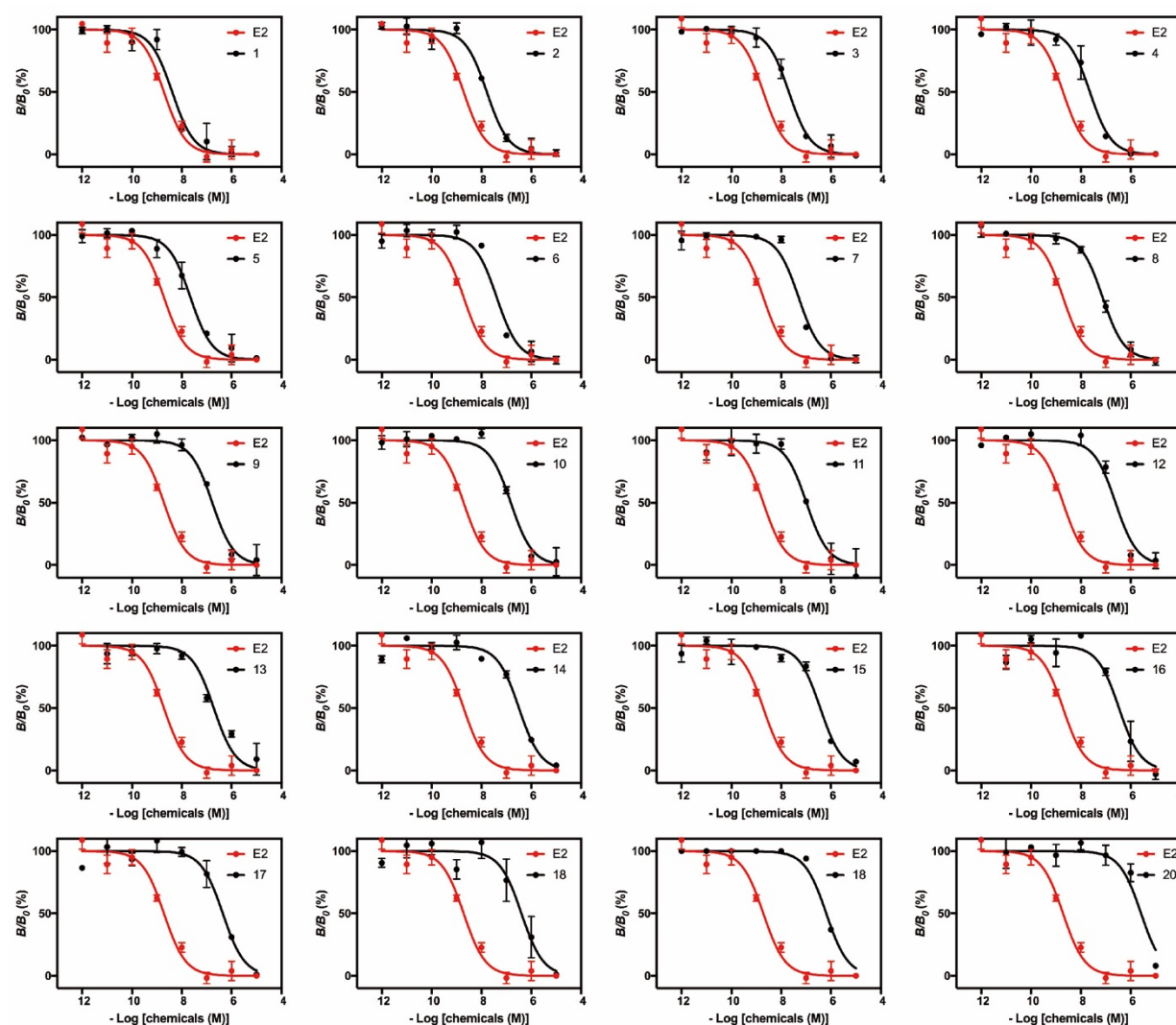


Fig. S1.

Binding curves indicated the binding ability of each bisphenol derivatives using competitive binding assays with $[^3\text{H}]\text{E2}$. B/B_0 is the ratio of displacement by the chemical tested (B) against the maximum specific binding ($B_0 = 100\%$) of $[^3\text{H}]\text{E2}$.

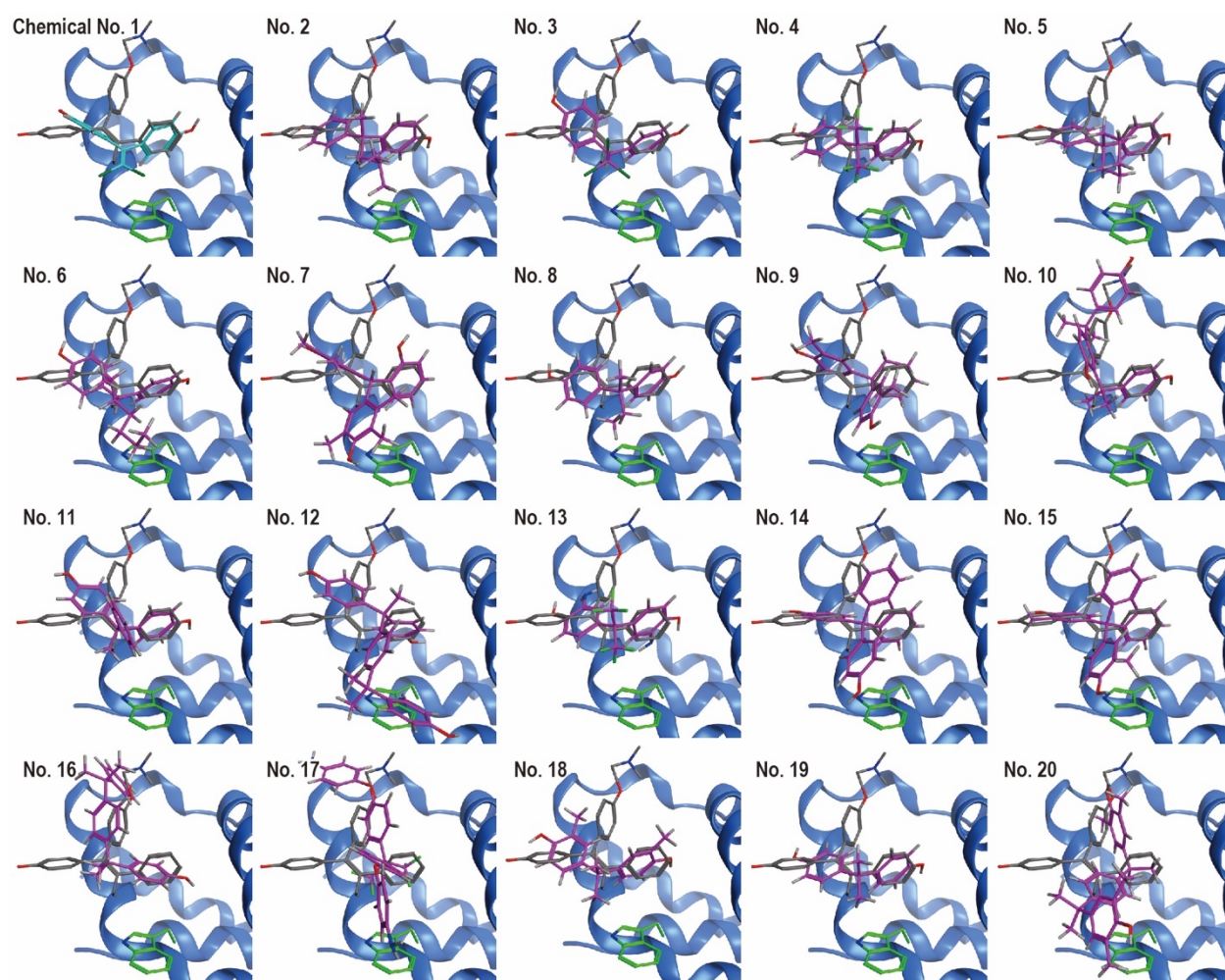


Fig. S2.

Docking simulation indicated the binding possibilities of BPA derivatives on the second binding site located on the coactivator-binding surface of ERβ. Calculated coordinates of BPC (blue stick model, No. 1) and the other BPA derivatives (magenta stick model, No. 2 to 20) were located close to W335. 4OHT in the crystal structure used for the docking simulation is indicated via a gray stick model.

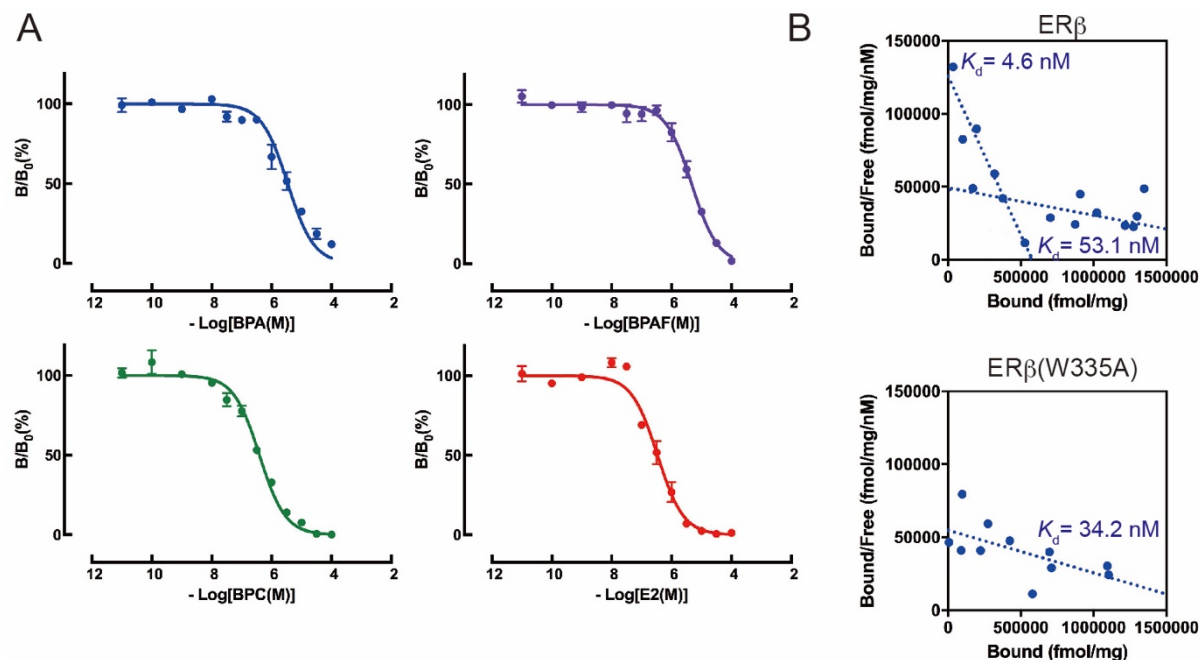


Fig. S3. Binding experiments showed that ERβ has two 4OHT binding sites, and only a single binding site for E2. (A) Competitive binding assay of ERβ(W335A) using [³H]E2 showed that ERβ(W335A) retained its binding activity for E2 and other BPA derivatives. (B) Saturation binding assays using [³H]4OHT estimated that ERβ has both a high and low binding sites for 4OHT, while ERβ(W335A) has only one binding site.

Table S1.

CAS RNs, common names, and IUPAC names of all the chemicals whose binding ability to ERβ was analyzed using competitive binding assays in this study.

CAS RN®	common name	IUPAC name
603-44-1	4,4',4"-trihydroxytriphenylmethane	4,4',4"-methanetriyltriphenol
79-95-8	tetrachloro bisphenol A	4,4'-(propane-2,2-diyl)bis(2,6-dichlorophenol)
79-94-7	tetrabromo bisphenol A	4,4'-(propane-2,2-diyl)bis(2,6-dibromophenol)
77-40-7	bisphenol B	4,4'-(butane-2,2-diyl)diphenol
1571-75-1	bisphenol AP	4,4'-(1-phenylethane-1,1-diyl)diphenol
5613-46-7	tetramethyl bisphenol A	4,4'-(propane-2,2-diyl)bis(2,6-dimethylphenol)
79-97-0	2,2-bis(4-hydroxy-3-methylphenyl)propane	4,4'-(propane-2,2-diyl)bis(2-methylphenol)
27955-94-8	1,1',1"-tris(4-hydroxyphenyl)ethane	4,4',4"-(ethane-1,1,1-triyl)triphenol
599-64-4	4-α-cumyl phenol	4-(2-phenylpropan-2-yl)phenol
2167-51-3	bisphenol P	4,4'-(1,4-phenylenebis(propane-2,2-diyl))diphenol
14868-03-2	bisphenol C	4,4'-(2,2-dichloroethene-1,1-diyl)diphenol
80-05-7	bisphenol A	4,4'-(propane-2,2-diyl)diphenol
70-30-4	hexachlorophene	6,6'-methylenebis(2,4,5-trichlorophenol)
110726-28-8	α,α,α'-tris(4-hydroxyphenyl)-1-ethyl-4-isopropylbenzene	4,4'-(1-(4-(2-(4-hydroxyphenyl)propan-2-yl)phenyl)ethane-1,1-diyl)diphenol
2716-10-1	α,α'-bis(4-aminophenyl)-1,4-diisopropylbenzene	4,4'-(1,4-phenylenebis(propane-2,2-diyl))dianiline
57100-74-0	2,2-bis(3-cyclohexyl-4-hydroxyphenyl)propane	4,4'-(propane-2,2-diyl)bis(2-cyclohexylphenol)
24038-68-4	2,2-bis(2-hydroxy-5-biphenyl)propane	5,5''-(propane-2,2-diyl)bis([1,1'-biphenyl]-2-ol))
1675-54-3	2,2-bis(4-glycidyloxyphenyl)propane	2,2'-(((propane-2,2-diylbis(4,1-phenylene))bis(oxy))bis(methylene))bis(oxirane)
10192-62-8	bisphenol A diacetate	propane-2,2-diylbis(4,1-phenylene) diacetate
4162-45-2	tetrabromobisphenol A bis(2-hydroxyethyl)ether	2,2'-((propane-2,2-diylbis(2,6-dibromo-4,1-phenylene))bis(oxy))diethanol
127-54-8	2,2-bis(4-hydroxy-3-isopropylphenyl)propane	4,4'-(propane-2,2-diyl)bis(2-isopropylphenol)
3539-42-2	4,4'-isopropylidenediphenoxyacetic acid	2,2'-((propane-2,2-diylbis(4,1-phenylene))bis(oxy))diacetic acid
13080-86-9	2,2-bis[4-(4-aminophenoxy)-phenyl]propane	4,4'-((propane-2,2-diylbis(4,1-phenylene))bis(oxy))dianiline
36395-57-0	α,α'-bis(4-hydroxy-3,5-dimethylphenyl)-1,4-diisopropylbenzene	4,4'-(1,4-phenylenebis(propane-2,2-diyl))bis(2,6-dimethylphenol)
2024-88-6	2,2-bis(4-chloroformyloxyphenyl)propane	propane-2,2-diylbis(4,1-phenylene) dicarbonochloridate
32113-46-5	2,2-bis(3-sec-butyl-4-hydroxyphenyl)propane	2-butan-2-yl-4-[2-(3-butan-2-yl-4-hydroxyphenyl)propan-2-yl]phenol
620-92-8	bisphenol F	4,4'-methylenediphenol
84-16-2	hexestrol	4,4'-(hexane-3,4-diyl)diphenol
1156-51-0	2,2-bis(4-cyanatophenyl)propane	4,4'-(propane-2,2-diyl)bis(cyanatobenzene)
479-13-0	coumestrol	3,9-dihydroxy-6H-benzofuro[3,2-c]chromen-6-one
1415-73-2	barbaloin	(10S)-1,8-dihydroxy-3-(hydroxymethyl)-10-[(2S,3R,4R,5S,6R)-3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]-10H-anthracen-9-one
961-29-5	isoliquiritigenin	(E)-1-(2,4-dihydroxyphenyl)-3-(4-hydroxyphenyl)prop-2-en-1-one
2467-25-6	4,4'-methylenebis(2-methylphenol)	4,4'-methylenebis(2-methylphenol)
17345-66-3	2,3,4-trihydroxydiphenylmethane	4-benzylbenzene-1,2,3-triol
75804-28-3	2,3-dimethyl-2,3-butanediamine	2,3-dimethylbutane-2,3-diamine
2081-08-5	bisphenol E	4,4'-(ethane-1,1-diyl)diphenol

2971-36-0	HPTE	4,4'-(2,2,2-trichloroethane-1,1-diyl)diphenol
83558-87-6	2,2-bis(3-amino-4-hydroxyphenyl)hexafluoropropane	4,4'-(perfluoropropane-2,2-diyl)bis(2-aminophenol)
47250-53-3	2,2-bis(3-aminophenyl)hexafluoropropane	3,3'-(perfluoropropane-2,2-diyl)dianiline
116325-74-7	2,2-bis(3-amino-4-methylphenyl)hexafluoropropane	5,5'-(perfluoropropane-2,2-diyl)bis(2-methylaniline)
1095-78-9	2,2-bis(4-aminophenyl)hexafluoropropane	4,4'-(perfluoropropane-2,2-diyl)dianiline
69563-88-8	2,2-bis[4-(4-aminophenoxy)phenyl]hexafluoropropane	4,4'-(((perfluoropropane-2,2-diyl)bis(4,1-phenylene))bis(oxy))dianiline
1478-61-1	bisphenol AF	4,4'-(perfluoropropane-2,2-diyl)diphenol
1107-00-2	4,4'-(hexafluoroisopropylidene)diphthalic anhydride	5,5'-(perfluoropropane-2,2-diyl)bis(isobenzofuran-1,3-dione)
1171-47-7	2,2-bis(4-carboxyphenyl)hexafluoropropane	4,4'-(perfluoropropane-2,2-diyl)dibenzoic acid
10224-18-7	2,2-bis(4-isocyanatophenyl)hexafluoropropane	4,4'-(perfluoropropane-2,2-diyl)bis(isocyanatobenzene)
83558-76-3	hexafluoro-2,2-diphenylpropane	(perfluoropropane-2,2-diyl)dibenzene
4221-68-5	1,1-bis(3-cyclohexyl-4-hydroxyphenyl)cyclohexane	4,4'-(cyclohexane-1,1-diyl)bis(2-cyclohexylphenol)
15499-84-0	9,9-bis(4-aminophenyl)fluorene	4,4'-(9H-fluorene-9,9-diyl)dianiline
184355-68-8	4,4'-(2-hydroxybenzylidene)-bis(2,3,6-trimethylphenol)	4,4'-((2-hydroxyphenyl)methylene)bis(2,3,6-trimethylphenol)
6807-17-6	4,4'-(1,3-dimethylbutylidene)diphenol	4,4'-(4-methylpentane-2,2-diyl)diphenol
3236-71-3	9,9-bis(4-hydroxyphenyl)fluorene	4,4'-(9H-fluorene-9,9-diyl)diphenol
88938-12-9	9,9-bis(4-hydroxy-3-methylphenyl)fluorene	4,4'-(9H-fluorene-9,9-diyl)bis(2-methylphenol)
74462-02-5	4,4'-(2-ethylhexylidene)diphenol	4,4'-(2-ethylhexane-1,1-diyl)diphenol
117344-32-8	9,9-bis[4-(2-hydroxyethoxy)phenyl]fluorene	2,2'-(((9H-fluorene-9,9-diyl)bis(4,1-phenylene))bis(oxy))diethanol
2362-14-3	1,1-bis(4-hydroxy-3-methylphenyl)cyclohexane	4,4'-(cyclohexane-1,1-diyl)bis(2-methylphenol)
3282-99-3	1,1-bis(4-aminophenyl)cyclohexane	4,4'-(cyclohexane-1,1-diyl)dianiline
843-55-0	bisphenol Z	4,4'-(cyclohexane-1,1-diyl)diphenol
13595-25-0	1,3-bis[2-(4-hydroxyphenyl)-2-propyl]benzene	4,4'-(1,3-phenylenebis(propane-2,2-diyl))diphenol
20601-38-1	4,4'-bicyclohexanol	[1,1'-bi(cyclohexane)]-4,4'-diol
1980-4-69	2,2-bis(4-hydroxycyclohexyl)propane	4,4'-(propane-2,2-diyl)dicyclohexanol
2433-14-6	4-cyclohexylcyclohexanol	[1,1'-bi(cyclohexan)]-4-ol
119-42-6	2-cyclohexylphenol	2-cyclohexylphenol
947-42-2	diphenylsilanediol	diphenylsilanediol
20714-70-9	4-(phenylazo)phenol	(E)-4-(phenyldiazenyl)phenol
501-36-0	resveratrol	(E)-5-(4-hydroxystyryl)benzene-1,3-diol
3127-14-8	spirobiromane	4,4,4',4'-tetramethyl-2,2'-spirobi[chroman]-7,7'-diol
2246-46-0	4-(2-Thiazolylazo)resorcinol	(E)-4-(thiazol-2-yl diazenyl)benzene-1,3-diol
32737-35-2	6,6',7,7'-tetrahydroxy-4,4,4',4'-tetramethyl-2,2'-spirobichroman	4,4,4',4'-tetramethyl-2,2'-spirobi[chroman]-6,6',7,7'-tetraol
269409-97-4	4-(4,4,5,5-Tetramethyl-1,3,2-dioxaborolan-2-yl)phenol	2-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)phenol
611-99-4	4,4'-dihydroxybenzophenone	bis(4-hydroxyphenyl)methanone
90-96-0	4,4'-dimethoxybenzophenone	bis(4-methoxyphenyl)methanone
61445-50-9	2,3',4,4'-tetrahydroxybenzophenone	(2,4-dihydroxyphenyl)(3,4-dihydroxyphenyl)methanone
131-55-5	2,2',4,4'-tetrahydroxybenzophenone	bis(2,4-dihydroxyphenyl)methanone
345-92-6	4,4'-difluorobenzophenone	bis(4-fluorophenyl)methanone
131-54-4	2,2'-dihydroxy-4,4'-dimethoxybenzophenone	bis(2-hydroxy-4-methoxyphenyl)methanone
90-98-2	4,4'-dichlorobenzophenone	bis(4-chlorophenyl)methanone
2421-28-5	3,3',4,4'-benzophenonetetracarboxylic dianhydride	5,5'-carbonylbis(isobenzofuran-1,3-dione)
85-58-5	benzophenone-2,4'-dicarboxylic acid monohydrate	2-(4-carboxybenzoyl)benzoic acid

342-25-6	2,4'-difluorobenzophenone	(2-fluorophenyl)(4-fluorophenyl)methanone
611-98-3	4,4'-diaminobenzophenone	bis(4-aminophenyl)methanone
83846-85-9	4-benzoyl 4'-methyldiphenyl sulfide	phenyl(4-(p-tolylthio)phenyl)methanone
964-68-1	benzophenone-4,4'-dicarboxylic acid	4,4'-carbonyldibenzoic acid
131-53-3	2,2'-dihydroxy-4-methoxybenzophenone	(2-hydroxy-4-methoxyphenyl)(2-hydroxyphenyl)methanone
85-29-0	2,4'-dichlorobenzophenone	(2-chlorophenyl)(4-chlorophenyl)methanone
1470-79-7	2,4,4'-trihydroxybenzophenone	(2,4-dihydroxyphenyl)(4-hydroxyphenyl)methanone
835-11-0	2,2'-dihydroxybenzophenone	bis(2-hydroxyphenyl)methanone
21222-05-9	3,3'-dinitrobenzophenone	bis(3-nitrophenyl)methanone
611-79-0	3,3'-diaminobenzophenone	bis(3-aminophenyl)methanone
2958-36-3	2-amino-2',5-dichlorobenzophenone	(2-amino-5-chlorophenyl)(2-chlorophenyl)methanone
33077-87-1	2,2',4-trimethoxybenzophenone	(2,4-dimethoxyphenyl)(2-methoxyphenyl)methanone
3708-39-2	4,4'-bis(methylamino)benzophenone	bis(4-(methylamino)phenyl)methanone
119-61-9	benzophenone	benzophenone
31127-54-5	2,3,4,4'-tetrahydroxybenzophenone	(4-hydroxyphenyl)(2,3,4-trihydroxyphenyl)methanone
118-82-1	4,4'-methylenebis(2,6-di-tert-butylphenol)	4,4'-methylenebis(2,6-di-tert-butylphenol)
122-25-8	methylenedisalicylic acid	5,5'-methylenebis(2-hydroxybenzoic acid)
105391-33-1	bis(3-ethyl-5-methyl-4-maleimidophenyl)methane	1,1'-(methylenebis(2-ethyl-6-methyl-4,1-phenylene))bis(1H-pyrrole-2,5-dione)
97-23-4	2,2'-methylenebis(4-chlorophenol)	2,2'-methylenebis(4-chlorophenol)
13676-54-5	4,4'-bismaleimidodiphenylmethane	1,1'-(methylenebis(4,1-phenylene))bis(1H-pyrrole-2,5-dione)
19900-72-2	4,4'-methylenebis(2-ethyl-6-methylaniline)	4,4'-methylenebis(2-ethyl-6-methylaniline)
838-88-0	4,4'-diamino-3,3'-dimethyldiphenylmethane	4,4'-methylenebis(2-methylaniline)
101-77-9	4,4'-diaminodiphenylmethane	4,4'-methylenedianiline
101-61-1	Bis[4-dimethylamino-phenyl]methane	4,4'-methylenebis(N,N-dimethylaniline)
88-24-4	2,2'-methylenebis(6-tert-butyl-4-ethylphenol)	6,6'-methylenebis(2-(tert-butyl)-4-ethylphenol)
101-14-4	4,4'-methylenebis(2-chloroaniline)	4,4'-methylenebis(2-chloroaniline)
5384-21-4	4,4'-methylenebis(2,6-dimethylphenol)	4,4'-methylenebis(2,6-dimethylphenol)
19430-83-2	3,4'-diaminodiphenylmethane	3-(4-aminobenzyl)aniline
119-47-1	2,2'-methylenebis(6-tert-butyl-p-cresol)	6,6'-methylenebis(2-(tert-butyl)-4-methylphenol)
1817-74-9	4,4'-dinitrodiphenylmethane	bis(4-nitrophenyl)methane
42240-73-3	Bis(4-amino-2,3-dichlorophenyl)methane	4,4'-methylenebis(2,3-dichloroaniline)
3236-63-3	2,2'-methylenebis(4-methylphenol)	2,2'-methylenebis(4-methylphenol)
2467-03-0	2,4'-dihydroxydiphenylmethane	2-(4-hydroxybenzyl)phenol
457-68-1	4,4'-difluorodiphenylmethane	bis(4-fluorophenyl)methane
101-68-8	4,4'-diphenylmethane diisocyanate, (4,4'-Methylenebis(phenyl Isocyanate))	bis(4-isocyanatophenyl)methane
139-25-3	4,4'-diisocyanato-3,3'-dimethyldiphenylmethane	bis(4-isocyanato-3-methylphenyl)methane
19471-12-6	3,3'-diaminodiphenylmethane	3,3'-methylenedianiline
2467-02-9	2,2'-dihydroxydiphenylmethane	2,2'-methylenediphenol
1844-01-5	4,4'-dihydroxytetraphenylmethane	4,4'-(diphenylmethylene)diphenol
174462-43-2	2,3,4,4'-tetrahydroxydiphenylmethane	4-(4-hydroxybenzyl)benzene-1,2,3-triol

Table S2.

PDB IDs of ER α utilized for calculating of the volumes of each ligand-binding pocket. PDB IDs are listed in alphabetical order.

PDB ID	ligand name	pocket volume (Å ³)
1A52	estradiol	389.6
1ERE	17beta-estradiol	461.4
1GWQ	raloxifene	425.4
1QKU	estradiol	407.1
1X7E	WAY-244	510.8
1X7R	genistein	457.9
2P15	ortho-trifluoromethylphenylvinyl estradiol	518.0
2QA6	4-(6-hydroxy-1 <i>H</i> -indazol-3-yl)benzene-1,3-diol	318.1
2QGT	(9beta,11alpha,13alpha,14beta,17alpha)-11-(methoxymethyl)estra-1(10),2,4-triene-3,17-diol	417.8
2QSE	4-OH-PhIP	545.0
2QXM	PhIP	529.1
3Q95	estriol	422.0
3UU7	bisphenol A	428.0
3UUA	bisphenol AF	623.0
3UUC	bisphenol C	316.8
4MGA	4-tert-octylphenol	153.3
4MGC	benzophenone-2	314.0
4MGD	HPTE	395.1
4PP6	resveratrol	340.4
4PPP	fluoro-resveratrol	388.5
4PPS	(1 <i>S</i> ,3alpha <i>R</i> ,5 <i>S</i> ,7alpha <i>S</i>)-5-(4-hydroxyphenyl)-7alpha-methyloctahydro-1 <i>H</i> -inden-1-ol	466.4
4TV1	propylparaben	317.0
4ZN7	diethylstilbestrol	422.6
5DI7	(1 <i>S</i> ,3alpha <i>R</i> ,5 <i>S</i> ,7alpha <i>S</i>)-5-(4-hydroxy-2-methylphenyl)-7alpha-methyloctahydro-1 <i>H</i> -inden-1-ol	485.8
5DID	(1 <i>S</i> ,3alpha <i>R</i> ,5 <i>S</i> ,7alpha <i>S</i>)-5-(2,3-difluoro-4-hydroxyphenyl)-7alpha-methyloctahydro-1 <i>H</i> -inden-1-ol	460.4
5DIE	(1 <i>S</i> ,3alpha <i>R</i> ,5 <i>S</i> ,7alpha <i>S</i>)-7alpha-methyl-5-(2,3,5-trifluoro-4-hydroxyphenyl)octahydro-1 <i>H</i> -inden-1-ol	468.5
5DIG	(1 <i>S</i> ,3alpha <i>R</i> ,5 <i>S</i> ,7alpha <i>S</i>)-5-[4-hydroxy-2-(trifluoromethyl)phenyl]-7alpha-methyloctahydro-1 <i>H</i> -inden-1-ol	497.6
5EGV	3-chloranyl-4-[4-(2-chloranyl-4-oxidanyl-phenyl)furan-3-yl]phenol	395.0
5EI1	2-(4-hydroxyphenyl)-3-iodanyl-imidazo[1,2- α]pyridin-6-ol	389.9
5EIT	2-(4-hydroxyphenyl)-3-(trifluoromethyl)imidazo[1,2- α]pyridin-6-ol	381.4
5JMM	biochanin A	617.0
5KR9	coumestrol	350.9
5KRM	(1 <i>S</i> ,7alpha <i>S</i>)-5-(2,5-difluoro-4-hydroxyphenyl)-7alpha-methyl-2,3,3alpha,4,7,7alpha-hexahydro-1 <i>H</i> -inden-1-ol	475.8
5KRO	(8 <i>R</i> ,9 <i>S</i> ,13 <i>S</i> ,14 <i>S</i> ,17 <i>S</i>)-13-methyl-17-(methyl(phenyl)amino)-7,8,9,11,12,13,14,15,16,17-decahydro-6 <i>H</i> -cyclopenta[α]phenanthren-3-ol	378.5
5TLL	(<i>E</i>)-2-chloro-4'-hydroxy-4-((hydroxyiminio)methyl)-[1,1'-biphenyl]-3-olate	359.5
5TLU	(14beta,17alpha)-21-(4-aminophenyl)-19-norpregna-1(10),2,4-trien-20-yne-3,17-diol	416.8
5TMZ	(8 <i>S</i> ,9 <i>S</i> ,13 <i>S</i> ,14 <i>S</i> ,17 <i>S</i>)-16-(3-methoxybenzyl)-13-methyl-7,8,9,11,12,13,14,15,16,17-decahydro-6 <i>H</i> -cyclopenta[α]phenanthrene-3,17-diol	436.8
5TN1	(8 <i>S</i> ,9 <i>S</i> ,13 <i>S</i> ,14 <i>S</i> , <i>E</i>)-17-((4-isopropylphenyl)imino)-13-methyl-7,8,9,11,12,13,14,15,16,17-decahydro-6 <i>H</i> -cyclopenta[α]phenanthren-3-ol	425.9

5TN3	(8 <i>S</i> ,9 <i>S</i> ,13 <i>S</i> ,14 <i>S</i>)-17-((4-isopropylphenyl)amino)-13-methyl-7,8,9,11,12,13,14,15,16,17-decahydro-6 <i>H</i> -cyclopenta[α]phenanthren-3-ol	475.9
5TN4	(<i>S</i>)-5-(4-hydroxy-3,5-dimethylphenyl)-2,3-dihydro-1 <i>H</i> -inden-1-ol	388.9
5TN5	(1 <i>S</i> ,3α <i>S</i> ,5 <i>S</i> ,7α <i>S</i>)-5-(4-hydroxyphenyl)-7α-methyloctahydro-1 <i>H</i> -inden-1-ol	419.0
5TN6	(1 <i>S</i> ,1' <i>S</i> ,3α <i>S</i> ,7α' <i>S</i>)-7α'-methyl-1',2,2',3,3',3α <i>S</i> ,4',6',7',7α' <i>S</i> -decahydro-1,5'-spirobi[indene]-1',5'-diol	448.0
5TN7	(<i>E</i>)-3'-fluoro-4'-hydroxy-3-((hydroxyiminio)methyl)-[1,1'-biphenyl]-4-olate	387.1
5TN8	(<i>E</i>)-4'-hydroxy-3-((hydroxyiminio)methyl)-[1,1'-biphenyl]-4-olate	355.3
5U2B	(8 <i>R</i> ,9 <i>S</i> ,13 <i>S</i> ,14 <i>S</i> ,17 <i>S</i>)-13-methyl-17-(phenylamino)-7,8,9,11,12,13,14,15,16,17-decahydro-6 <i>H</i> -cyclopenta[α]phenanthren-3-ol	515.8

Table S3.

PDB IDs of ER β agonist structures utilized for calculating of the volumes of each ligand binding pocket. PDB IDs are listed in alphabetical order.

PDB ID	ligand name	pocket volume (Å ³)
1QKN	raloxifene	382.3
1U3Q	4-(6-hydroxy-benzo[delta]isoxazol-3-yl)benzene-1,3-diol	344.9
1U3R	2-(5-hydroxy-naphthalen-1-yl)-1,3-benzoxazol-6-ol	364.3
1U3S	3-(6-hydroxy-naphthalen-2-yl)-benzo[delta]isoxazol-6-ol	407.9
1X76	5-hydroxy-2-(4-hydroxyphenyl)-1-benzofuran-7-carbonitrile	354.4
1X78	[5-hydroxy-2-(4-hydroxyphenyl)-1-benzofuran-7-yl]acetonitrile	349.4
1X7B	2-(3-fluoro-4-hydroxyphenyl)-7-vinyl-1,3-benzoxazol-5-ol	314.0
1X7J	genistein	375.9
1YY4	1-chloro-6-(4-hydroxyphenyl)-2-naphthol	408.5
1YYE	3-(3-fluoro-4-hydroxyphenyl)-7-hydroxy-1-naphthonitrile	402.6
1ZAF	3-bromo-6-hydroxy-2-(4-hydroxyphenyl)-1 <i>H</i> -inden-1-one	503.8
2J7X	estradiol	423.0
2J7Y	(16alpha,17alpha)-estra-1,3,5(10)-triene-3,16,17-triol	372.6
2NV7	4-(4-hydroxyphenyl)-1-naphthaldehyde oxime	340.6
2YJD	4-(2-propan-2-yloxybenzimidazol-1-yl)phenol	468.1
2YLY	<i>n</i> -cyclopropyl-4-oxidanyl- <i>n</i> -[(2 <i>R</i>)-2-oxidanyl-2-phenyl-propyl]benzenesulfonamide	226.3
3OLL	estradiol	361.9
3OLS	estradiol	404.9
3OMO	2-(trifluoroacetyl)-1,2,3,4-tetrahydroisoquinolin-6-ol	289.0
3OMP	2-(trifluoroacetyl)-1,2,3,4-tetrahydroisoquinolin-7-ol	321.3
3OMQ	2-[(trifluoromethyl)sulfonyl]-1,2,3,4-tetrahydroisoquinolin-6-ol	346.9
4J24	estradiol	324.9
4J26	estradiol	434.6
4ZI1	2-(4-hydroxyphenyl)-7-methyl-3-phenyl-1 <i>H</i> -inden-5-ol	350.4
5TOA	estradiol	361.0

Table S4.

PDB IDs of ER β LBDs utilized for SiteFinder calculations to analyze ligand-binding sites. PDB IDs are listed in alphabetical order.

PDB ID	rank of 1st site	rank of 2nd site	ligand	active or inactive	position of H12*
1HJ1	1	7	ICI164,384	inactive	free
1L2J	1	none	(<i>R,R</i>)-5,11-cis-diethyl-5,6,11,12-tetrahydrochrysene-2,8-diol	inactive	CBS
1NDE	1	none	4-(2-([4-([3-(4-chlorophenyl)propyl)sulfanyl]-6-(1-piperazinyl)-1,3,5-triazin-2-yl]amino)ethyl)phenol	inactive	CBS
1QKM	1	9	genistein	inactive	free
1QKN	1	10	raloxifene	inactive	free
1U3Q	1	none	4-(6-hydroxy-benzo[d]isoxazol-3-yl)benzene-1,3-diol	active	active position
1U3R	1	none	2-(5-hydroxy-naphthalen-1-yl)-1,3-benzoxazol-6-ol	active	active position
1U3S	1	none	3-(6-hydroxy-naphthalen-2-yl)-benzo[d]isoxazol-6-ol	active	active position
1U9E	1	3	2-(4-hydroxy-phenyl)benzofuran-5-ol	active	active position
1X76	1	3	5-hydroxy-2-(4-hydroxyphenyl)-1-benzofuran-7-carbonitrile	active	active position
1X78	1	none	[5-hydroxy-2-(4-hydroxyphenyl)-1-benzofuran-7-yl]acetonitrile	active	active position
1X7B	1	none	2-(3-fluoro-4-hydroxyphenyl)-7-vinyl-1,3-benzoxazol-5-ol	active	active position
1X7J	5	none	genistein	active	active position
1YY4	1	none	1-chloro-6-(4-hydroxyphenyl)-2-naphthol	active	active position
1YYE	1	17	3-(3-fluoro-4-hydroxyphenyl)-7-hydroxy-1-naphthonitrile	active	active position
1ZAF	1	3	3-bromo-6-hydroxy-2-(4-hydroxyphenyl)-1 <i>H</i> -inden-1-one	active	active position
2FSZ	5	11	4-hydroxytamoxifen	inactive	free
2GIU	1	none	(9 <i>aS</i>)-4-bromo-9 <i>a</i> -butyl-7-hydroxy-1,2,9,9 <i>a</i> -tetrahydro-3 <i>H</i> -fluoren-3-one	inactive	CBS
2I0G	1	none	(3 <i>aS</i> ,4 <i>R</i> ,9 <i>bR</i>)-4-(4-hydroxyphenyl)-1,2,3,3 <i>a</i> ,4,9 <i>b</i> -hexahydrocyclopenta[c]chromen-8-ol	inactive	CBS
2I0J	1	9	(3 <i>aS</i> ,4 <i>R</i> ,9 <i>bR</i>)-4-(4-hydroxyphenyl)-1,2,3,3 <i>a</i> ,4,9 <i>b</i> -hexahydrocyclopenta[c]chromen-8-ol	inactive	free
2J7X	1	none	estradiol	active	active position
2J7Y	1	none	(16 <i>alpha</i> ,17 <i>alpha</i>)-estra-1,3,5(10)-triene-3,16,17-triol	active	active position
2JJ3	1	none	(3 <i>aS</i> ,4 <i>R</i> ,9 <i>bR</i>)-4-(4-hydroxyphenyl)-6-(methoxymethyl)-1,2,3,3 <i>a</i> ,4,9 <i>b</i> -hexahydrocyclopenta[c]chromen-8-ol	inactive	CBS
2NV7	1	13	4-(4-hydroxyphenyl)-1-naphthaldehyde oxime	active	active position
2POG	1	8	(3 <i>aS</i> ,4 <i>R</i> ,9 <i>bR</i>)-4-(4-hydroxyphenyl)-1,2,3,3 <i>a</i> ,4,9 <i>b</i> -hexahydrocyclopenta[c]chromen-9-ol	inactive	free
2QTU	1	none	(3 <i>aS</i> ,4 <i>R</i> ,9 <i>bR</i>)-2,2-difluoro-4-(4-hydroxyphenyl)-6-(methoxymethyl)-1,2,3,3 <i>a</i> ,4,9 <i>b</i> -hexahydrocyclopenta[c]chromen-8-ol	inactive	CBS
2YJD	1	3	4-(2-propan-2-yloxybenzimidazol-1-yl)phenol	active	active position
2YLY	2	5	<i>N</i> -cyclopropyl-4-oxidanyl- <i>N</i> -[(2 <i>R</i>)-2-oxidanyl-2-phenylpropyl]benzenesulfonamide	active	active position
2Z4B	1	none	(3 <i>aS</i> ,4 <i>R</i> ,9 <i>bR</i>)-2,2-difluoro-4-(4-hydroxyphenyl)-1,2,3,3 <i>a</i> ,4,9 <i>b</i> -hexahydrocyclopenta[c]chromen-8-ol	inactive	CBS
3OLL	1	8	estradiol	active	active position
3OLS	1	none	estradiol	active	active position
3OMO	1	none	2-(trifluoroacetyl)-1,2,3,4-tetrahydroisoquinolin-6-ol	active	active position

3OMP	1	10	2-(trifluoroacetyl)-1,2,3,4-tetrahydroisoquinolin-7-ol	active	active position
3OMQ	1	none	2-[(trifluoromethyl)sulfonyl]-1,2,3,4-tetrahydroisoquinolin-6-ol	active	active position
4J24	1	10	estradiol	active	active position
4J26	1	9	estradiol	active	active position
4ZI1	1	none	2-(4-hydroxyphenyl)-7-methyl-3-phenyl-1 <i>H</i> -inden-5-ol	active	active position
5TOA	1	7	estradiol	active	active position

* “CBS” means that H12 is located in an inactivated position on the ER β coactivator-binding site (CBS); 'free' means helix 12 is not visualized or is far outside of the LBD.

Table S5.

Compounds names, CAS RN, and ligand IDs of CDS-Core or Chemical IDs from the Protein Data Bank (PDB); 3D coordinates were utilized for docking simulation experiments. Chemical IDs from PDB are designated by three letters.

	common names	CAS RN®	Ligand ID
1	bisphenol C	14868-03-2	0D1 (PDB)
2	4,4'-(1,3-dimethylbutylidene)bisphenol	6807-17-6	ZUHRAX
3	2,2-bis(p-hydroxyphenyl)-1,1,1- trichloroethane (HPTE)	2971-36-0	-
4	bisphenol AF	1478-61-1	TIBVOQ
5	bisphenol Z	843-55-0	-
6	4,4'-(2-ethylhexylidene)bisphenol	74462-02-5	-
7	4,4'-(2-hydroxybenzylidene)-bis(2,3,6-trimethylphenol)	184355-68-8	-
8	bisphenol B	77-40-7	-
9	1,1-bis(4-hydroxy-3-methylphenyl)cyclohexane	2362-14-3	SIJHOJ
10	bisphenol M	13595-25-0	-
11	bisphenol AP	1571-75-1	-
12	α, α, α' -tris(4-hydroxyphenyl)-1-ethyl-4-isopropylbenzene	110726-28-8	-
13	2,2-bis(3-amino-4-hydroxyphenyl)hexafluoropropane	83558-87-6	-
14	9,9-Bis(4-hydroxyphenyl)fluorene	3236-71-3	ABUCOP
15	9,9-bis(4-hydroxy-3-methylphenyl)fluorene	15499-84-0	XOGJEI
16	bisphenol P	2167-51-3	-
17	2,2-bis[4-(4-aminophenoxy)phenyl]hexafluoropropane	69563-88-8	HOYZOL
18	2,2-bis(4-hydroxy-3-methylphenyl)propane	79-97-0	REGKOF
19	bisphenol A	80-05-7	2OH (PDB)
20	α, α' -bis(4-hydroxy-3,5-dimethylphenyl)-1,4-diisopropylbenzene	36395-57-0	ACAYIN