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Identification of a GPER/GPR30 antagonist with improved estrogen receptor counterselectivity

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ABSTRACT

GPER/GPR30 is a seven-transmembrane G protein-coupled estrogen receptor that regulates many aspects of mammalian biology and physiology. We have previously described both a GPER-selective agonist G-1 and antagonist G15 based on a tetrahydro-3H-cyclopenta[c]quinoline scaffold. The antagonist lacks an ethanone moiety that likely forms important hydrogen bonds involved in receptor activation. Computational docking studies suggested that the lack of the ethanone substituent in G15 could minimize key steric conflicts, present in G-1, that limit binding within the ER α ligand binding pocket. In this report, we identify low-affinity cross-reactivity of the GPER antagonist G15 to the classical estrogen receptor ER α . To generate an antagonist with enhanced selectivity, we therefore synthesized an isosteric G-1 derivative, G36, containing an isopropyl moiety in place of the ethanone moiety. We demonstrate that G36 shows decreased binding and activation of $ER\alpha$, while maintaining its antagonist profile towards GPER. G36 selectively inhibits estrogen-mediated activation of PI3K by GPER but not $ER\alpha$. It also inhibits estrogenand G-1-mediated calcium mobilization as well as ERK1/2 activation, with no effect on EGF-mediated ERK1/2 activation. Similar to G15, G36 inhibits estrogen- and G-1-stimulated proliferation of uterine epithelial cells in vivo. The identification of G36 as a GPER antagonist with improved ER counterselectivity represents a significant step towards the development of new highly selective therapeutics for cancer and other diseases.

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1. Introduction

Estrogens mediate a range of physiological processes, including roles in reproduction, the immune, nervous and cardiovascular systems [1]. Additionally, estrogen signaling plays a role in breast, ovarian and other types of cancer [2]. Estrogen signaling is mediated through at least three receptors, including the soluble nuclear receptors $ER\alpha$ and $ER\beta$, and GPER, a seven transmembranespanning G protein-coupled receptor (GPCR) [3]. All three of these receptors can mediate gene transcription events, either directly, as

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in the case of ER α and ER β [1], or indirectly, as in the case of GPER [4,5]. ER α , ER β and GPER can also mediate rapid signaling events through activation of MAPK, PI3K, Src kinase and related pathways [2,3]. *In vivo*, these receptors vary in their tissue distribution and the estrogen responsiveness of a given tissue is determined both by receptor expression, co-regulator expression and by signaling interplay between receptors in response to estrogen [1,3].

Estrogen receptors overlap in some physiological functions as well as in their ligand specificity. The triphenylethylene derivative tamoxifen is representative of the selective estrogen receptor modulator (SERM) anti-estrogen class of therapeutics that inhibit the binding of the endogenous ligand, 17 β -estradiol (E2, 1), to ER α and ER β , while fulvestrant (faslodex, ICI182,780) represents the "pure" antiestrogen class of therapeutics that initiates ER α and ER β receptor down-regulation. Paradoxically, both of these compounds act as GPER agonists [6,7], and the resulting agonist/antagonist prop-

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erties of these compounds vary by receptor status and/or tissue [8]. Thus, compounds with selectivity towards a single receptor are of great use for probing receptor function in complex systems where multiple estrogen receptors are expressed. We have previously developed a GPER-selective agonist, G-12[9], and a GPER-selective antagonist, G15 3 [10], which have been used to elucidate GPER function in a variety of systems, both in vitro as well as in vivo. For example, G-1 has been used to define a role for GPER in proliferation [4,11], protein kinase C activation [12], PI3K activation [9] and calcium mobilization in a range of cells [9,13,14]. Additionally, G-1 has been used in in vivo systems to investigate the roles of GPER in uterine epithelial proliferation [10], thymic atrophy [15], immune regulation in experimental autoimmune encephalomyelitis (a murine model of multiple sclerosis) [15,16] and vascular regulation [17]. The antagonist G15 has now been used by several groups to establish the role of GPER in estrogen-mediated events, including vasodilation [18], zebrafish oocyte maturation [19], neuroprotection [20] and inhibition of chondrogenesis [21]. These results indicate that selective ligands for GPER have a wide range of functional applications and can contribute to our understanding of the contributions of GPER signaling in complex systems expressing ER α and/or ER β in addition to GPER.

Particularly in systems expressing multiple estrogen receptors, the utility of these compounds derives from and is limited by their selectivity for GPER vs. ER α and ER β . These small molecules are based on a common tetrahydro-3H-cyclopenta[c]quinoline scaffold, with the key difference between agonist G-1 and antagonist G15 being the presence of an ethanone moiety on G-1 [9,10]. Since G15 lacks this bulky substituent group and, as we demonstrate here, exhibits increased ER α/β activity compared to G-1, we synthesized G36, a G-1 analog that contains an isopropyl moiety substituted for the ethanone in G-1. We postulated that the increased bulk present in G-1 and G36 would increase steric clashes within the binding pocket of ER α and ER β compared to G15, thus limiting the binding and downstream signaling activity observed when high doses of G15 are used.

In this report, we describe the identification and characterization of G36 4, a GPER-selective antagonist with enhanced selectivity for GPER and decreased activity towards ER α and ER β compared to the previously described antagonist, G15. We show that high doses of G15 induce low levels of transcription via ER α and that G36 minimizes this off-target effect. Additionally, G36 maintains equal efficacy as an antagonist of GPER compared to G15 in a range of functional assays, both *in vitro* and *in vivo*.

2. Materials and methods

2.1. Molecular docking

The crystal structure of the human estrogen receptor (ER) ligand-binding domain in complex with 17β-estradiol from the RCSB Protein Data Bank (PDB ID: 1ERE) was used [22]. The receptor was prepared for docking using the standard protocol implemented in fred_receptor, a wizard like graphical utility that prepares an active site for docking with FRED (version 2.2.5, OpenEye Scientific Software, Inc., Santa Fe, NM, USA, www.eyesopen.com, 2010). Three-dimensional conformations for the chosen ligands (E2, G-1, G15, and G36) were generated using OMEGA (version 2.3.2, OpenEye Scientific Software, Inc., Santa Fe, NM, USA, www.eyesopen.com, 2010) and the ligands were docked in the binding site of the receptor with FRED, using the default docking parameters. The Chemgauss3 scoring function, which uses Gaussian smoothed potentials to measure the complementarity of ligand poses within the active site, was used to evaluate and compare how well the different ligands fit in the ligand binding site.

2.2. Chemical synthesis and characterization of G36 (4-(6-bromo-benzo[1,3]dioxol-5-yl)-8-isopropyl-3a,4,5,9b-tetrahydro-3H-cyclopenta[c]quinoline)

A catalytic amount of Sc(OTf)₃ (0.049 g, 0.1 mmol, 10 mol%) in anhydrous acetonitrile (1 mL) was added to a mixture of 6bromopiperonal (0.229 g, 1.00 mmol), 4-isopropylaniline (0.135 g, 1.0 mmol) and cyclopentadiene (0.33 g, 5.0 mmol) in acetonitrile (3 mL). The reaction was stirred at ambient temperature ($\sim 23 \circ \text{C}$) for 2h with monitoring of product formation by thin layer chromatography using 20% ethyl acetate/hexanes eluent (R_f 0.5). The product crystallized from acetonitrile, and was filtered and washed with additional acetonitrile (5 mL) to provide the pure endo isomer as colorless solid (390 mg, 94%). Mp 135-136 °C; FT-IR (KBr): 3435, 2955, 1613, 1504, 1040 cm⁻¹; ¹H NMR (300 MHz, CDCl₃) δ 7.18 (s, 1H), 7.02 (s, 1H), 6.92 (d, *J*=2.0 Hz, 1H), 6.87 (d, *J*=8.0, 2.0 Hz, 1H), 6.57 (d /= 8.0 Hz, 1H), 5.99 (d, /= 1.3 Hz, 1H), 5.97 (d, /= 1.3 Hz, 1H), 5.89–5.85 (m, 1H), 5.68–5.64 (m, 1H), 4.88 (d, /=3.5 Hz, 1H), 4.10 (d, J=8.2 Hz, 1H), 3.45 (bs, 1H), 3.22-3.11 (m, 1H), 2.84-2.73 (m, 1H), 2.64–2.55 (m, 1H), 1.83–1.75 (m, 1H), 1.22 (d, /=1.5, 3H), 1.20 (d, J = 1.5 Hz, 3H); ¹³C NMR (75 MHz, CDCl₃) δ 147.4, 147.2, 143.1, 139.9, 134.7, 134.0, 130.2, 126.8, 125.9, 124.2, 116.0, 113.0, 112.8, 108.1, 101.7, 56.8, 46.2, 42.2, 33.3, 31.3, 24.3, 24.1; HPLC-MS: Electrospray positive ion (cone voltage 62 V, capillary voltage 3 kV). G36 (1 mg/mL CH₃CN, 20 μ L) was injected onto a Symmetry[®] C18 $(5 \,\mu m, 3.0 \,mm \times 150 \,mm, Waters)$ column and eluted with 60–90% acetonitrile (gradient 1.5% min⁻¹) in water showed a single peak at 22.42 min (Fig. 4). UV-vis λ_{max} 294 nm (Fig. 5). ESI-MS *m*/*z* (ES+) calcd for C₂₂H₂₂BrNO₂ (M+H)⁺ 412.08; found 412.11; HRMS: calcd [M+H]⁺ for C₂₂H₂₂BrNO₂ 412.0905, found 412.0912.

2.3. Reagents

 17β -estradiol was from Sigma (St. Louis, MO). G-1 and G15 were synthesized as previously described [9,10,23]. pERK antibodies were from Cell Signaling (Danvers, MA). Goat–anti-rabbit HRP was from GE-Amersham (Piscataway, NJ). DMEM and RPMI 1640 were obtained from Fisher Scientific (Pittsburgh, PA). 32% paraformaldehyde (PFA) was obtained from Electron Microscopy Sciences (Hatfield, PA).

2.4. Cell culture and transfection

COS7 cells were maintained in DMEM containing 10% fetal bovine serum, 2 mM L-glutamine, 100 units/mL penicillin and 100 μ g/mL streptomycin. SKBr3 cells were maintained in RPMI containing 10% fetal bovine serum, 2 mM L-glutamine, 100 units/mL penicillin and 100 μ g/mL streptomycin. Cells were grown as a monolayer at 37 °C, in a humidified atmosphere of 5% CO₂ and 95% air. For microscopy experiments, cells were seeded onto 12 mm glass coverslips and allowed to adhere for at least 12 h prior to transfection. Twenty-four hours prior to PI3K or pERK experiments, medium was replaced with serum-free, phenol red free RPMI 1640. For experiments requiring transient transfection of PH-mRFP, ER α -GFP, ER β -GFP or GPER-GFP, Lipofectamine2000 (Invitrogen) was used according to manufacturers instructions. PH-mRFP was transfected at 1/4 the recommended amount to achieve appropriate expression levels.

2.5. ERE activation assays

ERE activity was determined using MCF-7 cells stably transfected with an ERE-GFP reporter construct [24]. Briefly, cells were deprived of estrogen for 4 days (with one intermediate medium change) in phenol red free DMEM/F12 supplemented with 10% charcoal-stripped FBS. In 24 well plates, ~80,000 cells were seeded, and 24h later treated with the indicated compounds (dissolved in DMSO, final DMSO 0.1% for each compound added) for 24h in triplicate, trypsinized, washed and analyzed for green fluorescence by flow cytometry. Mean fluorescence intensities were determined and normalized to E2 values following subtraction of vehicle control values.

2.6. Ligand binding assays

Binding assays for ER α and ER β were performed as previously described [7]. Briefly, COS7 cells were transiently transfected with either ER α -GFP or ER β -GFP. Following serum starvation for 24 h, cells ($\sim 5 \times 10^4$) were incubated with G36 for 20 min in a final volume of 10 μ L prior to addition of 10 μ L 20 nM E2-Alexa633 in saponin-based permeabilization buffer. Following 10 min at RT, cells were washed once with 200 μ L PBS/2%BSA, resuspended in 20 μ L and 2 μ L samples were analyzed on a DAKO Cyan flow cytometer using HyperCytTM as described [25].

2.7. Intracellular calcium mobilization

SKBr3 cells (1 × 10⁷ cells/mL) were incubated in Hanks balanced salt solution (HBSS, Gibco) containing 3 μ M indo1-AM (Invitrogen) and 0.05% pluronic acid for 1 h at RT. Cells were then washed twice with HBSS, incubated at RT for 20 min, washed again with HBSS, resuspended in HBSS at a density of 10⁸ cells/mL and kept on ice until assay, performed at a density of 2 × 10⁶ cells/mL. Ca⁺⁺ mobilization was determined ratiometrically using λ_{ex} 340 nm and λ_{em} 400/490 nm at 37 °C in a spectrofluorometer (QM-2000-2, Photon Technology International) equipped with a magnetic stirrer.

2.8. PI3K activation

The PIP3 binding domain of Akt fused to mRFP1 (PH-mRFP1) was used to localize cellular PIP3. COS7 cells (cotransfected with GPER-GFP or ER α -GFP and PH-mRFP1) cells were plated on coverslips and serum starved for 24 h followed by stimulation with ligands as indicated. The cells were fixed with 2% PFA in PBS, washed, mounted in Vectashield and analyzed by confocal microscopy using a Zeiss LSM510 confocal fluorescent microscope. Cells were preincubated with 1 μ M G36 as indicated for 20 min prior to stimulation and G36 was present during stimulation.

2.9. Western blotting

Cells were serum starved in phenol red-free RPMI 1640 for 24 h prior to treatment for pERK Western blots. For activation of pERK, cells were treated as indicated, washed once with ice-cold PBS and lysed using NP-40 buffer. Twenty µg protein was loaded per lane. Cells were preincubated with G15 or G36 as indicated prior to stimulation and inhibitors were present during stimulation. Band quantitation was carried out using ImageJ [26].

2.10. Mouse uterine estrogenicity assay

C57Bl6 female mice (Harlan) were ovariectomized at 10 weeks of age. E2 and G36 were dissolved in absolute ethanol at 1 mg/mL and further diluted in ethanol. For treatment, $10 \,\mu$ L of the appropriate dilution (of single or of combined compounds) was added to 90 μ L aqueous vehicle (0.9% NaCl with 0.1% albumin and 0.1% Tween-20). Ethanol alone (10 μ L) was added to 90 μ L aqueous vehicle as control (sham). At 12 days post-ovariectomy, mice were injected subcutaneously with compound. Eighteen hr after injection, mice were sacrificed and uteri were dissected, fixed in 4% paraformaldehyde, and embedded in paraffin. Five-micron sections were placed on slides, and proliferation in uterine epithelia was quantitated by immunofluorescence using anti-Ki-67 antibody (LabVision) followed by goat anti-mouse IgG conjugated to Alexa488 (Invitrogen). Nuclei were counterstained with 4',6diamidino-2-phenylindole (DAPI). At least 4 animals per treatment were analyzed, and the Ki-67 immunodetection was repeated three times per mouse.

3. Results and discussion

3.1. G-1/G15 ERE-mediated activity and design & synthesis of G36

In our original identification of the GPER antagonist G15 [10], we speculated that the ethanone moiety of G-1 might be critical for GPER activation through the formation of hydrogen bonds with the receptor and that loss of this bonding would result in a compound that could still bind but not activate GPER, thus acting as a competitive antagonist. In our initial characterization of G15, we detected weak binding of G15 to ER α and ER β at high concentrations ($\geq 10 \,\mu$ M) of G15; however, there were no effects of G15 on estrogen receptor-mediated signaling observed in multiple functional assays [10]. These results supported the use of G15 as a selective GPER antagonist for in vitro and in vivo investigations. In order to assess the structural basis of the observed low-affinity binding, we performed docking studies with the 3-dimensional Xray crystal structure of the estrogen-bound ligand binding domain of ER α (PDB ID 1ERE) [22]. E2, as expected, displayed a high binding score (-99.6) for the receptor site, the docking pose being almost identical to the ligand in the X-ray structure (RMSD = 0.53 Å). Docking of G-1 resulted in a significantly worse score (-52.8), mostly due to a steric clash with the Arg394 residue in the binding site (Fig. 1), which could explain the high level of GPER selectivity shown by this ligand. However, the docking score of G15(-76.2) indicates that this ligand has a lower steric clash than G-1 within the binding site and might therefore exhibit binding to the receptor, but with a much lower affinity than E2.

To test whether G15 exhibits any activity towards ER α , we utilized a highly-sensitive estrogen response element (ERE) transcriptional assay, in which estrogenic activity through either ER α (or potentially ER β) results in activation of a stably transfected ERE-GFP reporter construct in MCF-7 cells yielding a readout of GFP



Fig. 1. Docking poses of selected ligands in the ER binding site: E2 green, G-1 magenta, G15 yellow. Arg394, which displays steric clashes with G-1 is depicted in orange in the lower left corner. The Pro399-Leu402 loop was hidden in order to better present the ligands. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of the article.)



Fig. 2. High concentrations of G15 mediate weak activation of ERE. MCF-7 cells stably transfected with an ERE-GFP reporter were treated for 24h with the indicated concentration of E2, G-1 or G15. Whereas E2 shows half maximal activation at approximately 100 pM, G-1 shows no activation up to 10 μ M. G15 exhibits limited activation (~15–20%) at concentrations ranging from 1 to 10 μ M.

expression (Fig. 2) [24]. In this assay, high doses ($\geq 1 \mu$ M) of G15 (concentrations $10^5 - 10^6$ greater than that required for E2) resulted in weak activation of the ERE reporter ($\leq 20\%$ of E2 response), consistent with the initial results that G15 binds weakly to ER α and ER β at high concentrations. To establish that the expression of GFP is mediated by ER α (or potentially ER β), we coincubated E2 and G15 with ICI182,780 (a full ER antagonist that results in ER downregulation), which yielded complete ablation of GFP induction by both E2 and G15 (data not shown). Together, these results demonstrate that at high concentrations, G15 is capable of mediating limited ER-dependent transcriptional activity.

With these results in mind, we revisited the structural similarity of the GPER-selective agonist G-1 and G15. Both GPER-selective compounds are based on the same cyclopenta[c]quinoline scaffold,

with the difference being the presence of an ethanone moiety on G-1 which is lacking in G15 (Fig. 3A). Based on the docking study, we postulated that the presence of this side chain on G-1 resulted in additional steric conflicts and reduced binding ability of G-1 for ER α . Since we speculated that the inability of G15 to form hydrogen bonds through the ethanone group is responsible for its antagonist activity, we further hypothesized that replacing the reactive ethanone group with a hydrophobic isopropyl group might generate a GPER antagonist with increased selectivity for GPER over $ER\alpha$ and $ER\beta$ compared to G15. Thus, we identified the isopropyl group as a potential isosteric replacement of the ethanone group that would lack both H-bond donor and acceptor capacity. The target compound G36 was synthesized using a three-component Povarov cyclization of 4-isopropylaniline, 6-bromopiperonal and cyclopentadiene catalyzed by Sc(OTf)₃ in acetonitrile (Fig. 3B). The product crystallized from acetonitrile to afford G36 in excellent yield as a racemic mixture of the analytically pure syn/endo diastereomer.

3.2. ER docking, binding and activation of G36

Docking analysis with G36 yielded a score (-52.3) very similar to that of G-1 with a similar steric clash of the isopropyl group with Arg 394 (Supplemental Fig. 1). To examine the binding selectivity of G36 towards ER α and ER β , we compared the binding of G36 to that of G15 and E2 using a competitive fluorescent estrogen ligand binding assay. Results from dose-response curves confirm that G15 displays weak but significant binding to ER α and ER β only at the highest concentration, 10 μ M, with G36 lacking detectable binding activity to both ER α and ER β over the entire range of concentrations tested (Fig. 4). It should be noted that the binding of G15 to ER α and ER β is nevertheless weak, between 10⁴ and 10⁵ fold less than that of E2. These results suggested that G36 may exhibit reduced acti-



Fig. 3. (A) Structures of estrogen (17β-estradiol) 1, G-1 2, G15 3 and G36 4. (B) Synthetic route for G36.



Fig. 4. G36 exhibits improved binding selectivity towards ER α and ER β compared to G15. Dose response profile of E2, G15 and G36 for competition of E2-Alexa633 binding to ER α -GFP (A) or ER β -GFP (B). G36 shows decreased binding to ER α and ER β at the highest concentration compared to G15.

vation of ERE in comparison to G15. In addition to being a highly sensitive functional assay, one benefit of the ERE assay is that it can be used to investigate both agonist and antagonist properties of compounds. We took advantage of this property to probe more fully the activities of G-1, G15 and G36 against $ER\alpha/\beta$ (Fig. 5). We confirmed that G-1, when administered to MCF7 cells stably expressing the ERE-GFP reporter, displayed no agonist activity at doses up to 10 µM; however, when G-1 is administered at high doses it does show weak antagonism of the 1 nM E2-mediated ERE response. This finding is noteworthy, as it suggests that G-1 should optimally be used at sub-micromolar concentrations when investigating GPER function in complex systems where transcriptional effects may be relevant, thus limiting possibly confounding results from potential inhibition of estrogen-mediated transcriptional effects via $\text{ER}\alpha/\beta$. Most published work has employed G-1 at sub-micromolar doses (typically 10–100 nM, occasionally $\sim 1 \,\mu$ M), so the finding that G-1 antagonizes $ER\alpha/\beta$ function at high doses does not detract from such studies, particularly since most studies are carried out over short (i.e. non-genomic) time frames and in the absence of added estrogen.

In contrast to G-1, G15 demonstrates limited agonism of the ERE reporter (Fig. 5) with ~15% activity at 1 μ M and ~25% activity at 10 μ M compared to the maximal E2 response (10 nM, see Fig. 2). Additionally, G15 inhibits the E2-induced ERE response by ~30% when administered at 10 μ M with no significant effect at 1 μ M. These results suggest that G15 likely functions as a partial agonist of transcription mediated by ERs. Our prior studies however demonstrated that G15 does not mediate rapid signaling events



Fig. 5. G36 exhibits reduced activity towards ERE activation and inhibition compared to G15. (A) Activation of ERE-GFP response in MCF7 cells by increasing concentrations of G-1, G15 and G36. (B) Inhibition of ERE response induced by 1 nM E2 as a function of increasing concentrations of G-1, G15 and G36. *p < 0.05 vs. DMSO alone (A) or 1 nM E2 (B).

via either the MAPK or PI3K pathways in ER α - or ER β -expressing cells [10], suggesting that the partial agonist activity seen here applies only to transcriptional signaling through the classical ERs and not to ER-mediated rapid signaling events. This separation of rapid and genomic signaling events has previously been observed with androgen receptor ligands, with some promoting genomic signaling and others preferentially activating non-genomic pathways [27,28]. In contrast to G15, G36 shows much lower activity in the ERE activation assay, with no significant ERE response elicited by G36 concentrations up to 1 μ M and only \sim 5% activation at 10 μ M compared to \sim 25% for G15 at 10 μ M. G36 also shows no ability to antagonize E2-mediated ERE response at concentrations up to 10 µM. It should be noted that the inhibitory effects of G-1 and G15 on estrogen-induced transcription were absent when E2 was used at 10 nM (data not shown), suggesting that the inhibitory effects are only present at low estrogen concentration. Nevertheless, these results, along with the ER α and ER β binding data, suggest that G36 has decreased activity via ER α and ER β compared to the previously described GPER antagonist, G15.

3.3. G36 selectively inhibits estrogen-mediated PI3K activity through GPER

We next investigated whether G36 maintains the inhibitory activity towards GPER displayed by G15 [10]. First, to examine PI3K activation of endogenous receptors, COS7 cells were transfected with PH-RFP, a fluorescent reporter of PIP3 localization whose nuclear translocation is used to assess PI3K activation [7]. Neither E2 nor G-1 nor G36 at a concentration of 10 μ M demonstrated any activation of PI3K, indicating that COS7 cells do not exhibit either receptor-mediated or non-specific responses to any of these ligands (Fig. 6A). Next, COS7 cells were transiently transfected with either ER α -GFP or GPER-GFP and PH-RFP. As expected, due to its minimal activity in ERE assays, G36 did not inhibit E2-mediated PI3K activation in cells expressing ER α -GFP (Fig. 6B). Conversely, in cells expressing GPER-GFP, G36 effectively inhibited PI3K activation induced by E2 (Fig. 6C), suggesting that the modifications made to the initial chemical scaffold had maintained the GPER antagonist activity while decreasing the ER α activity of G15.

3.4. G36 inhibits estrogen- and G-1-mediated calcium mobilization

In order to determine the potency of G36 compared to G15, we assayed the ability of G36 to inhibit both E2- and G-1-mediated calcium mobilization in SKBr3 cells, which endogenously express only GPER and neither ER α nor ER β . Both E2 and G-1 mediated similar levels of calcium mobilization at 200 nM (Fig. 7A). To determine the potency of G36, cells were stimulated with a constant dose of E2 or G-1 and increasing concentrations of G36 were used to generate a dose-response curve (Fig. 7B). IC₅₀ values for G36 inhibition of E2 and G-1 mediated calcium mobilization were similar, with G36 having an IC₅₀ of 112 nM for E2 and 165 nM for G-1. These values compare favorably with those found for G15 in the same

assay [10], with G36 showing slightly higher potency for inhibiting E2-mediated (G36 IC₅₀ = 112 nM, G15 IC₅₀ = 190 nM) calcium mobilization and essentially identical potency for G-1-mediated (G36 IC₅₀ = 165 nM, G15 IC₅₀ = 185 nM) calcium mobilization. To assess whether G36 might act non-specifically to inhibit calcium mobilization, we tested G36 at 10 μ M for its effect on calcium mobilization by an unrelated GPCR, specifically endogenous purinergic receptors (Fig. 7C). G36 exhibited no inhibition of calcium mobilization by 1 μ M ATP, indicating high selectivity for GPER. Furthermore, G36 itself did not alter the basal calcium level in cells (Fig. 7C, first arrow). Thus, G36 affords enhanced selectivity as an antagonist of GPER with similar potency in the inhibition of GPER function when compared to G15.

3.5. G36 inhibits estrogen- and G-1-mediated but not EGF-mediated ERK activation

Since activation of the ERK/MAPK signaling pathway was one of the functions first attributed to GPER [6], we verified the activity of G36 as an antagonist of GPER function in ERK activation (Fig. 8). SKBr3 cells, which express only GPER and neither ER α nor ER β , were stimulated with E2 and the GPER-selective agonist G-1. With



Fig. 6. G36 inhibits E2-induced PI3K activation in cells expressing GPER, but not in cells expressing ERα. (A) COS7 cells (which lack endogenous ERα, ERβ and GPER) transiently transfected with only PH-RFP show no activation of PI3K, as evidenced by the lack of nuclear translocation of PH-RFP, in response to E2, G-1 or G36 (all at 10 µM). (B) COS7 cells transiently transfected with ERα-GFP and PH-RFP activate PI3K, as evidenced by nuclear translocation of PH-RFP in response to E2 and this response is not inhibited by the presence of G36. (C) In COS7 cells transiently transfected with GPER-GFP and PH-RFP, G36 inhibits the PI3K activation induced by E2.



Fig. 6. (Continued).

both stimuli, we detected a 5–6-fold increase in the level of pERK. Upon stimulation in the presence of either G15 or G36, we observed a significant decrease in the E2- and G-1-induced ERK activation. To test for possible non-specific effects of G36 or G15, we examined EGF-induced ERK activation, which was unchanged by the presence of either G15 or G36, illustrating the selectivity of each of these molecules for GPER.

3.6. G36 inhibits estrogen- and G-1-mediated uterine proliferation

In order to extend our findings *in vivo*, we evaluated the effect of G36 in the proliferative response of the uterine epithelium (Fig. 9). Upon administration of E2, uterine epithelia exhibit increased proliferation, as evidenced by the increase in the percent of cells



Fig. 7. G36 inhibits E2 and G-1 mediated calcium mobilization in SKBr3 cells. (A) SKBr3 cells, which endogenously express only GPER but neither ER α nor ER β , were monitored for calcium mobilization induced by either E2 (200 nM) or G-1 (200 nM). DMSO was added at the first arrow as a control and either E2 or G-1, as indicated at the second arrow. (B) E2- and G-1-mediated calcium mobilization are inhibited by increasing concentrations of G36 (pre-incubated for ~2 min, following the scheme in A). (C) Purinergic receptor activation (by 1 μ M ATP, second arrow) was not inhibited by pretreatment with 10 μ M G36 (first arrow). Data is normalized to E2 (200 nM) or G-1 (200 nM) as shown in A.

staining positive for the proliferation marker Ki-67. As we have previously observed [10], G-1 can also induce significant proliferation of uterine epithelia, albeit to a substantially lesser extent than E2. This result is most likely due to E2 activating the full cohort of estrogen receptors in the mouse (ER α , ER β and GPER), whereas G-1 only activates GPER, which alone cannot fully recapitulate the response of ER α and ER β . G36 administered alone had no effect on proliferation, whereas G36 administered with G-1 completely abrogated the proliferative response to G-1, as expected of a GPER antagonist. Combination treatment with E2 and G36 resulted in a significant decrease in proliferation compared to E2 treatment, consistent with the inhibition previously observed with G15. These *in vivo* results confirm our *in vitro* findings that G36 is a GPER antag-



Fig. 8. G36 inhibits ERK activation induced by E2 and G-1 via GPER. SKBr3 cells were stimulated with E2 (10 nM), G-1 (10 nM) or EGF (10 ng/mL) in the presence of vehicle control (DMSO), 1 μ M G15 or G36 (pre-incubated for 15 min) and analyzed for pERK and total ERK levels by Western blot. Quantitation of pERK activation was from three independent experiments. *p < 0.05 vs. DMSO alone.



Fig. 9. G36 inhibits GPER-mediated proliferation *in vivo*. Epithelial uterine cell proliferation was assessed in the presence of E2, G-1, G36, G-1+G36 or E2+G36. Amounts injected were as follows: E2: $10 \mu g/kg$; G-1: $10 \mu g/kg$; G36 (alone or in combination): $50 \mu g/kg$. Ki-67 positivity of the uterine epithelium was determined by immunofluorescence microscopy. *p < 0.05 vs. sham; **p < 0.05 vs. E2.

onist and demonstrate the utility of GPER-selective agonists and antagonists in a complex model system.

4. Conclusions

In summary, we have developed a second generation GPER antagonist G36 with improved performance and selectivity that exhibits significantly decreased off-target effects on ER α and ER β compared to our previously described antagonist, G15. It is however important to note the context in which both G-1 and G15/G36 are likely to be used. G-1 has typically been used to determine the contribution of GPER activation to a cellular or physiological response in the absence of estrogen, either by its omission in culture medium or following ovariectomy in mice. Thus, although G-1 may

exhibit slight inhibitory activity on estrogen-mediated ERE activation, its complete lack of stimulatory activity represents the more important consideration. For G-1, it is not clear that the inhibitory activity is a result of ER binding, as we were previously unable to detect significant competition even at 10 µM [9]. It is therefore possible that this inhibitory effect is indirect as a result of rapid signaling initiated by GPER. Interestingly, very high doses of G-1 have recently been shown to inhibit estrogen-mediated uterine epithelial cell proliferation in vivo through inhibition of stromal ERK1/2 activation and ER α phosphorylation [29]. Alternatively, G15 demonstrates both binding at high concentrations as well as stimulatory activity in the absence of estrogen and inhibitory activity in the presence of estrogen. The observation of binding suggests that these functional effects may be direct via $ER\alpha$ binding. However, G15 would typically not be used alone (except as a negative control) and thus the stimulatory activity would likely not be of concern, unless potentially used in combination with G-1, where ERE activation could confound results. Furthermore, any effects would be limited to "long-term" assays where ERE activation could be involved and not rapid or acute events (e.g. <1 h).

The improved selectivity of G36 in terms of <~5% activation and inhibition of ERE-mediated transcription at 10 μ M will make it highly useful for probing the functions of GPER in a wide range of complex assays and model systems that express one or both of the classical ERs in addition to GPER. Nevertheless, as with any pharmacological agent, conclusions regarding the involvement of the presumed target should be confirmed with the use of siRNA or knockout mice where possible. Building from the numerous documented successful applications using G15 to examine the contributions of GPER to estrogen physiology, we believe that the development of G36 will give researchers a more selective tool to investigate GPER function.

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.jsbmb.2011.07.002.

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