



Discovery of GPR183 Agonists Based on an Antagonist Scaffold

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The G protein-coupled receptor GPR183/EBI2, which is activated by oxysterols, is a therapeutic target for inflammatory and metabolic diseases where both antagonists and agonists are of potential interest. Using the piperazine diamide core of the known GPR183 antagonist (E)-3-(4-bromophenyl)-1-(4-(4-methoxybenzoyl)piperazin-1-yl)prop-2-en-1-one (NIBR189) as starting point, we identified and sourced 79 structurally related compounds that were commercially available. In vitro screening of this compound collection using a Ca²⁺ mobilization assay resulted in the identification of 10 compounds with agonist properties. To enable establishment of initial structure-activity relationship trends, these were supplemented with five inhouse compounds, two of which were also shown to be GPR183 agonists. Taken together, our findings suggest that the agonist activity of this compound series is dictated by the substitution pattern of one of the two distal phenyl rings, which functions as a molecular efficacy-switch.

The G protein-coupled receptor GPR183, also known as Epstein-Barr virus (EBV) induced gene 2 (EBI2), was identified in 1993 as one of the most up-regulated genes in EBV-infected cells. Before its deorphanization, the constitutive activity of GPR183 facilitated the discovery of $G\alpha_i$ coupling and ERK activation as

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downstream signaling events. [2] These two pathways along with β -arrestin coupling were subsequently shown to also be ligand-induced. [3] In 2011, GPR183 was deorphanized as two simultaneous papers revealed hydroxycholesterols (oxysterols) to be agonists of GPR183 with 7α ,25-dihydroxycholesterol (7α ,25-OHC, 1) (Figure 1A) as the most potent endogenous ligand. [4] The same two groups identified GPR183 as a chemotactic receptor with 1 acting as a potent chemoattractant, a finding that correlates well with the high expression of GPR183 in leukocytes [2a] and the observation that GPR183 differential expression is important for correct B cell positioning within lymphoid organs. [5]

The pharmacological interest in GPR183 stems from the implication of the receptor and its endogenous ligands in a variety of diseases like B cell malignancies, inflammatory/ autoimmune diseases and metabolic diseases. [6] Moreover, it was recently shown to regulate interferons and bacterial growth during *Mycobacterium tuberculosis* infection. [7]

Efforts to identify synthetic GPR183 ligands have resulted in a handful of small molecules, including agonists as well as

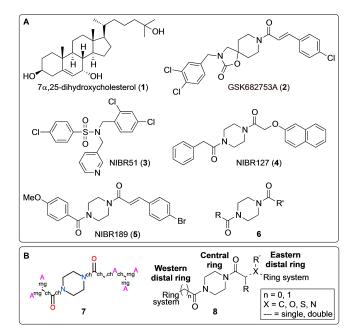


Figure 1. Known GPR183 ligands and general ligand structures. A) the endogenous agonist 1, the reported synthetic ligands 2–5, and the antagonist scaffold 6, B) the Markush formula 7 used as search query (ch: chain bond, rng: ring bond, A: any atom) and the general structure 8 for the 79 screened compounds.



Scheme 1. Synthesis of target compounds 88–92. Reagents and conditions: i) PhCH=CHCOCI or Ph(CH₂)₂COCI, TEA, DCM, rt, 24 h, 84–90 %; ii) 2-phenoxyacetic acid or 2-(4-chloro-2-methylphenoxy)acetic acid or 3-methoxy-4-methylbenzoic acid, EDC HCI, DIPEA, DMF, 0 °C to rt, 24 h, 92–97 %; iii) 2-chloroacetyl chloride, TEA, DCM, rt, 24 h, 99 %; iv) naphthalen-1-ol, Cs₂CO₃, KI, acetone, 35 °C, 24 h, 82 %; v) TFA:DCM, 1:1, rt, 1 h, 74 %.

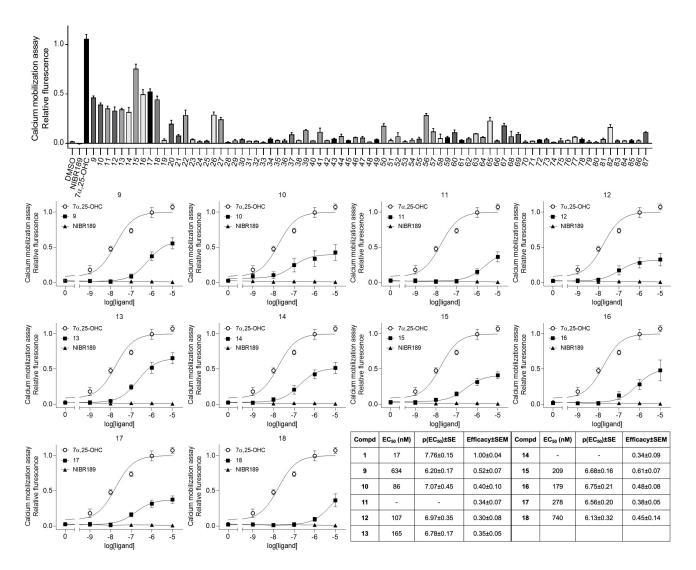


Figure 2. Top: Screening of compounds 9–87 in a calcium mobilization assay, with solvent (DMSO), the endogenous agonist 1, and the known antagonist 5 included as controls. Calcium response determined upon addition of 10 μ M compound and represented as relative fluorescence with 1 being the maximum response induced by 1. Middle: Calcium assay dose-response graphs of the 10 most active compounds (9–18) from the screen. Represented as relative fluorescence for each concentration used. Curves for compounds 1 and 5 added for reference. Bottom: Table showing potency (EC₅₀), p(EC₅₀) \pm SE and efficacy \pm SEM (determined as activity induced by 10 μ M ligand) for compounds 9–18. All data represent mean \pm SEM of 3 individual experiments performed in duplicates.



antagonists^[6a] (Figure 1A). Prior to the deorphanization of GPR183, a small molecule inverse agonist of GPR183 named GSK682753A (2) was identified; this compound was later shown to also inhibit 7 α ,25-OHC (1)-mediated GPR183 activity.^[2b,3] In 2014, screening of a library containing around 100 K compounds resulted in the discovery of the GPR183 agonist NIBR51 (3).^[8] This agonist was then used to rescreen the same compound library for GPR183 antagonists, leading to the identification of NIBR127 (4), which in turn was chemically optimized to give the more potent GPR183 antagonist NIBR189 (5).^[8] Using the molecular scaffold 6 (Figure 1A) of the known GPR183 antagonist 5 as starting point, we here report the discovery of a novel series of small molecule GPR183 agonists.

To identify commercially available compounds that were structurally related to the known GPR183 antagonist $5^{[8]}$ we first conducted a substructure screen of the Enamine Screening Collection (~2.7 M compounds) using the Markush formula 7 (Figure 1B) as search query. The results were further filtered for "drug-likeness" based on chemical criteria and subsequently clustered to ensure chemical diversity. A total of 79 compounds (9–87, supplementary Figure S1 and Table S1) were manually selected for experimental screening; these were all piperazine diamides of the general structure 8 (Figure 1B).

G protein activation induced by the 79 compounds was detected using CHO-K1 cells that were transiently transfected with GPR183 and a chimeric $G\alpha$ subunit Ggi4myr that is recognized as $G\alpha_i$ by $G\alpha_i$ -coupled receptors, but activates $G\alpha_n$ pathways, [9] consequently enabling Ca2+ release. Calcium release was measured for 100 seconds after ligand addition by utilizing a fluorescent indicator, and data was extracted as the change in fluorescence over time. Based on the G protein signaling efficacy induced by 10 μM of each compound (Figure 2, top), we selected 10 of the compounds (9-18) for further dose-response experiments (Figure 2, middle). Here, compounds 15 and 16 displayed the most favorable agonist properties, combining acceptable efficacy and potency (EC₅₀ of 209 nM and 179 nM, respectively). As previously shown, the antagonist 5 did not display any intrinsic activity, while the endogenous agonist 1 activated the receptor with a potency (EC₅₀) of 17 nM (Figure 2, shown as reference curves).

As all the screened compounds were built on a central piperazine diamide core, the structural variation was in the two distal ring systems and the spacers (Figure 1B). Of the 10 active compounds (Figure 3), four contained the same (*E*)-alkene spacer as the antagonist 5, two contained an ethylene spacer, and four contained the oxy-methylene spacer found in the known antagonist 4. However, the non-systematic structural variation in the distal ring systems made it difficult to identify clear structure-activity relationship (SAR) trends in this compound series.

We therefore designed and synthesized five additional compounds (Scheme 1): three reference compounds (88–90) that contained distal unsubstituted phenyl rings, as well as two crossover compounds (91 and 92) that combined the Western and Eastern ring systems of the top agonist hits 15 and 16.

The substituted piperazines **88** and **89** were synthesized in one step from commercially available phenyl(piperazin-1-yl)

Figure 3. Structures of the 10 compounds (**9–18**) that showed agonistic activity in the initial screening, grouped by type of two-atom spacer (C=C, C-C, C-O).

17

18

methanone (93) via coupling with the corresponding acyl chlorides (Scheme 1). The same starting material (93) was reacted with the corresponding carboxylic acids in the presence of EDC to afford 90 and 91 in excellent yields. Coupling of Bocpiperazine (94) and chloroacetyl chloride yielded intermediate 95. This was further reacted with 1-naphthol and Boc-deprotected to give 96. Subsequent amide coupling employing 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide (EDC) led to compound 92 in excellent yield.

As $G\alpha_i$ is the direct G protein-signaling pathway elicited by GPR183, [2a] we switched to this pathway for the functional tests of the synthesized compounds **88–92**, and included **15** and **16** for comparison with the calcium release experiment in the initial screen. Hence, the GPR183 agonist activity of the five inhouse compounds was experimentally tested using a BRET assay to determine $G\alpha_i$ coupling at various concentrations of the compounds. CHO-K1 cells were transiently transfected with GPR183 and the CAMYEL (cAMP sensor using YFP-Epac-Rluc) BRET biosensor, which changes conformation in response to cAMP levels; consequently, activation of $G\alpha_i$ leads to a rise in BRET signal.

While the reference compounds **88–90** were devoid of agonist activity (Figure 4A–C), the crossover compounds **91** and **92** displayed agonist profiles similar to the initial agonist hits **16** and **15**, respectively, i.e. similar potency and partial agonist properties, meaning that the efficacy did not reach that of the full agonist **1** (Figure 4D–E). The potency of the reference endogenous agonist **1** obtained here (19 nM) (Figure 4H) was comparable to the value in the calcium assay, as were the

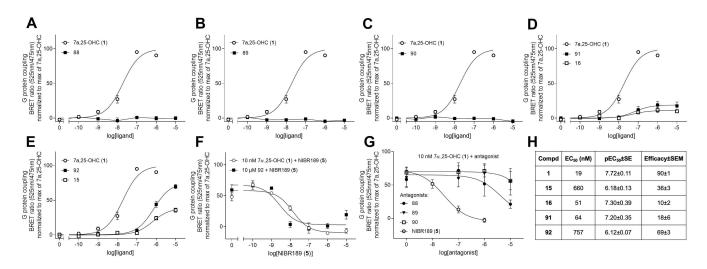


Figure 4. Dose-response curves representing GPR183 G protein-coupling upon addition of various compounds. Data is normalized to the maximum activity of the endogenous agonist 1 in each individual experiment. A–C) The reference compounds 88–90. D–E) The crossover compounds 91 and 92 with initial agonist hits 15–16. F) Activity of a fixed concentration of the endogenous agonist 1 and the synthesized agonist 92 with increasing amount of the antagonist 5. G) Activity induced by 10 nM of the endogenous agonist 1 after preincubation with the reference compounds 88–90 or the antagonist 5. H) Table showing potency (EC_{50}), $p(EC_{50})$ \pm SE and efficacy \pm SEM (determined as activity induced by 1 μ M for compound 1 and by 10 μ M for the other ligands) for the compounds that displayed agonist properties. All data is presented as mean \pm SEM of three individual experiments.

potencies for compounds **15** and **16**. To verify that the activity was mediated by GPR183, **92** was also tested in the presence of increasing concentrations of the antagonist **5**, which potently inhibited the activity induced by both **92** and the positive control **1** (pIC $_{50}\pm$ SE of 8.56 ± 0.27 and 7.86 ± 0.16 , respectively) (Figure 4F). Given the structural similarities with previously reported GPR183 antagonists, we also tested the antagonist properties of the reference compounds **88–90**. Compounds **89** and **90** did not inhibit the G protein activity induced by **1**, while **88** displayed weak antagonistic activity with 79% inhibition at the highest concentration tested (10 μ M). In contrast, the reference antagonist **5** resulted in almost 100% inhibition at a lower concentration (1 μ M), corresponding to an inhibitory potency of 28 nM (Figure 4G).

The lack of agonist activity for the unsubstituted reference compounds (inactive 88–90 vs. active 11/16) means that the substitution pattern on the distal rings is generally important for receptor activation. Notably, the data for the aryloxy-based compounds (inactive 90 vs. active 15/91) shows that the agonist activity is dictated by the nature of the Eastern ring system.

Comparison of the active and inactive compounds that contained the oxy-methylene spacer indicated a pattern: aside from the 1-naphthyl (15/92), the active compounds (16/17/18/91) are all para/ortho-disubstituted (Figure 3, Scheme 1). This is in contrast to the comparable inactive compounds 70–74 (supplementary Figure S1), which contain para- and/or meta-substituents. Thus, the lack of agonist activity for certain compounds seems to be linked to an inappropriate Eastern ring system, as also demonstrated by the thiophene-containing compounds 18 (active, Figure 3) and 69 (inactive, Figure S1). Taken together, the Western ring seems to be tolerant to modifications, as a simple benzene ring (15/91) is sufficient for

activity, along with other variations (16/92, 17, 18). On the other hand, the Eastern ring is sensitive to modifications, with minor variations in the substitution pattern having a dramatic impact on agonist activity.

To conclude, *in vitro* screening of a selection of commercially available compounds enabled us to identify novel synthetic agonists for GPR183; by supplementing these hits with a small series of in-house compounds we were able to establish initial SAR trends. Our findings provide a basis for formulation of hypotheses regarding the structural requirements for agonist activity and suggest several avenues for further optimization aimed at potent GPR183 agonists with improved pharmacodynamic and pharmacokinetic properties. Such studies are currently underway in our laboratories.

Conflict of Interest

The authors declare no conflict of interest.

Keywords: agonists \cdot antagonists \cdot drug discovery \cdot GPR183 \cdot receptors

- M. Birkenbach, K. Josefsen, R. Yalamanchili, G. Lenoir, E. Kieff, J. Virol. 1993, 67, 2209–2220.
- [2] a) M. M. Rosenkilde, T. Benned-Jensen, H. Andersen, P. J. Holst, T. N. Kledal, H. R. Luttichau, J. K. Larsen, J. P. Christensen, T. W. Schwartz, J. Biol. Chem. 2006, 281, 13199–13208; b) T. Benned-Jensen, C. Smethurst, P. J. Holst, K. R. Page, H. Sauls, B. Sivertsen, T. W. Schwartz, A. Blanchard, R. Jepras, M. M. Rosenkilde, J. Biol. Chem. 2011, 286, 29292–29302.
- [3] T. Benned-Jensen, C. M. Madsen, K. N. Arfelt, C. Smethurts, A. Blanchard, R. Jepras, M. M. Rosenkilde, FEBS Open Bio 2013, 3, 156–160.
- [4] a) C. Liu, X. V. Yang, J. Wu, C. Kuei, N. S. Mani, L. Zhang, J. Yu, S. W. Sutton, N. Qin, H. Banie, L. Karlsson, S. Sun, T. W. Lovenberg, *Nature* 2011,



- 475, 519–523; b) S. Hannedouche, J. Zhang, T. Yi, W. Shen, D. Nguyen, J. P. Pereira, D. Guerini, B. U. Baumgarten, S. Roggo, B. Wen, R. Knochenmuss, S. Noel, F. Gessier, L. M. Kelly, M. Vanek, S. Laurent, I. Preuss, C. Miault, I. Christen, R. Karuna, W. Li, D. I. Koo, T. Suply, C. Schmedt, E. C. Peters, R. Falchetto, A. Katopodis, C. Spanka, M. O. Roy, M. Detheux, Y. A. Chen, P. G. Schultz, C. Y. Cho, K. Seuwen, J. G. Cyster, A. W. Sailer, *Nature* 2011, 475, 524–527.
- [5] a) J. P. Pereira, L. M. Kelly, Y. Xu, J. G. Cyster, *Nature* 2009, 460, 1122–1126; b) D. Gatto, K. Wood, R. Brink, J. Immunol. 2011, 187, 4621–4628.
- [6] a) S. Sun, C. Liu, Front. Pharmacol. 2015, 6, 60; b) L. Barington, F. Wanke, K. Niss Arfelt, P. J. Holst, F. C. Kurschus, M. M. Rosenkilde, J. Leukocyte Biol. 2018, 104, 313–322.
- [7] a) J. Tang, Y. Shi, L. Zhan, C. Qin, Microb. Pathog. 2020, 145, 104234; b) S.
 Bartlett, A. T. Gemiarto, M. D. Ngo, H. Sajiir, S. Hailu, R. Sinha, C. X. Foo, L.
 Kleynhans, H. Tshivhula, T. Webber, H. Bielefeldt-Ohmann, N. P. West,
- A. M. Hiemstra, C. E. MacDonald, L. V. V. Christensen, L. S. Schlesinger, G. Walzl, M. M. Rosenkilde, T. Mandrup-Poulsen, K. Ronacher, *Front. Immunol.* **2020**, *11*, 601534.
- [8] F. Gessier, I. Preuss, H. Yin, M. M. Rosenkilde, S. Laurent, R. Endres, Y. A. Chen, T. H. Marsilje, K. Seuwen, D. G. Nguyen, A. W. Sailer, J. Med. Chem. 2014, 57, 3358–3368.
- [9] E. Kostenis, J. Recept. Signal Transduction Res. 2002, 22, 267–281.

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