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Squaryl Group Modified Phosphoglycolipid Analogs as Potential Modulators of GPR55

Received 00th January 20xx, Accepted 00th January 20xx DOI: 10.1039/x0xx00000x

Feiqing Ding,^{a,†} Adam T. Guy,^{b,†} Peter Greimel,^b Yoshio Hirabayashi,^c Hiroyuki Kamiguchi,^{*b} and Yukishige Ito^{*a}

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Lysophosphatidyl glucoside (LPGIc) is a structurally unique glycolipid that acts as a guidance cue for extending axons during central nervous system development by activating the class A G protein coupled receptor (GPR) 55 of spinal cord sensory axons. GPR55 not only plays an important role during development, but is also implicated in many disease states, rendering molecules that target GPR55 of widespread interest. In this study, we developed synthetic access to a novel class of LPGIc analogues featuring a squaryl diamide group as surrogate for the phosphodiester. We report the facile synthesis of a series of LPGIc analogues, their GPR dependent biological activity and a systematic analysis of the structure-activity relationship in regards to GPR55 modulation. The lead compound featuring identical configuration at all stereocenters compared to natural LPGIc exhibits an activity to repel axons of dorsal root ganglion (DGR) nociceptive neurons.

Phosphatidyl-β-D-glucoside (PtdGlc) is a structurally unique glycolipid (Scheme 1) which was identified in mammalian cells in 2001.¹ The proposed structure² was rigorously confirmed by chemical synthesis.³ Mammalian PtdGlc exhibits a strikingly homogeneous fatty acid composition and is enriched in lipid rafts.⁴ More recently, we revealed that PtdGlc is locally converted to lyso-phosphatidyl-β-D-glucoside (LPGlc, Scheme 1), released from radial glia in the developing spinal cord and regulates sensory axon wiring by specifically repelling nerve growth factor (NGF)-responsive dorsal root ganglion (DRG) axons.⁵ Extensive screening of G-protein coupled receptors identified GPR55 as a receptor of LPGlc.

GPR55 is a class A G-protein coupled receptor first identified in 1999.⁶ It is expressed in various organs⁷ including central and peripheral nervous systems, gastrointestinal tract, bone, and immune systems. Initially it was considered as a

Email: yukito@riken.jp (YI) and hiroyuki.kamiguchi@riken.jp (HK) †These authors contributed equally. cannabinoid receptor⁸ but more recently lysophosphatidyl inositol⁹ and LPGIc⁵ have been reported as its endogenous ligands. It has been shown that GPR55 plays important roles in development^{5,6} and is implicated in many disease states such as neuropathic pain,¹⁰ cancer,¹¹ and inflammation.¹² To-date a number of GPR55 agonists and antagonists have been developed,^{13,14} but the need for new specific GPR55 ligands remains,¹⁵ for both clinical and research needs.



Scheme 1 Structures of PtdGlc, LPGlc, and a squaryl analogue 1A.

To develop a new class of GPR55 modulators, we sought an approach based on the recently identified native ligand structure. LPGIc degradation is expected to proceed either via the phosphodiester (PDE) or the fatty acid (FA) ester group. The PDE links the carbohydrate head group with the sn-3 position of the glycerol backbone, while the FA ester attached to the sn-1 position. Consequently, we envisaged to replace them with appropriate bioisosteres.¹⁶ As a suitable PDE surrogate, we identified the squaryl (SQ) group, which has been exploited as a bidirectional electrophile suitable for glycan conjugation¹⁷ and protein labeling.¹⁸ The migration prone FA ester was planned to be replaced by a thioether to not only increase the stability but also to satisfy the demand for ease and economy of production. In this context, LPGIc FA ester to ether exchange did not abrogate the activity of LPGIc, while the amide analogue was completely inactive (Guy, Greimel, et al., unpublished). Additionally, the mannose and galactose analogues of LPGIc are also inactive,⁵ strongly suggesting to preserve the equatorial configuration of the C-2 and -4 hydroxy group of the glucose residue. Based on the above rationale, compound 1A (Scheme 1) was envisaged as our lead compound. Sekine and co-workers reported that

^{a.} Synthetic Cellular Chemistry Laboratory, RIKEN, 2-1 Hirosawa, Wako, Saitama 351-0198, Japan.

^{b.} RIKEN Center for Brain Science.

^{c.} Cellular Informatics Laboratory, RIKEN

Electronic Supplementary Information (ESI) available: See DOI: 10.1039/x0xx00000x

parent compound phosphatidylglucoside.²⁰ Interestingly, all potential energy minima of **2** exhibit either gauch⁺ or gauch⁻ conformation of the second inner dihedral of the PDE (Supporting information Figure S4). In case of the squaryl analogue, a considerable number of potential energy minima is observed, with no obvious rotational barrier present in the anomeric linkage between the pyranose analogue and the squaryl group. The conformational space of 3 was ranked against representative conformations of each of the identified potential energy minima in 2, based on their ability to reproduce one of the carbonyl oxygen and the methyl group positions in 2. This revealed that 3 can mimic each of the three identified potential energy minima conformations of 2 adequately without the need to attain energetically unfavourable conformations. Superimposition of the lowest energy conformer of 2 associated with potential energy minima I and an equivalent conformation of 3, less than 3 kcal/mol above its respective lowest energy, reveals their high degree of similarity (Figure 1b). Equally matching conformations of 3 with conformations of 2 associated with potential energy minima II and III exhibited similar energy penalties (Supporting information Figure S5). This suggests that the envisaged glucose residue in combination with the squaryl aglycon is able to attain highly similar conformations compared to LPGIc and thus exhibits a high potential to fit into



Scheme 3 Syntheses of (a) chiral building blocks and (b) LPGIc analogues. the GPR55 binding pocket.

Our synthetic strategy for the envisaged LPGIc analogues is outlined in Scheme 2a. It is based on three fragments, namely glycosylamine (G-NH₂), squaryl diester (DES), and lipid

2 | J. Name., 2012, 00, 1-3

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1.180

+90

2/3

Figure 1 (a) Energy landscape and conformational similarity of equatorial phosphate 2

with equatorial squarate 3 on a pyranose ring, based on QM free energy calculations.

(b) The lowest energy conformation of compound 2 is superimposed on the most closest matching conformation of compound 3. Carbon, cyan or iceblue; oxygen, red or

squaryl diamides are polarizable and mimic PDE linkage in DNA

I, II and III) in case of an anomeric PDE and a clear rotational

barrier of the anomeric bond between $-45^{\circ} < \phi < 45^{\circ}$. This is in good agreement with reported NMR results of the LPGIc

To better understand the conformational characteristics of the squaryl group compared to an anomeric phosphate we performed quantum mechanics calculation of model compounds 2 and 3 (Figure 1a). The analysis revealed the presence of three potential energy minima (Figure 1a, labeled

magenta; hydrogen, white or grey; nitrogen, blue; phosphorous, tan.

+/-180

(a)

+90

(b)

oligomers.¹⁹

(a)

(b) 1/6/8 Α

в

с

D

G-NH₂

L-NH₂

(CH2)16CH

(CH₂)₁₆CH

-(CH2)16CH3

(CH2)16CH (CH₂)₁₆CH (CH₂)₁₆CH

Published on 09 July 2018. Downloaded on 7/10/2018 3:13:22 AM

DOI: 10.1039/C8CC04467H

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component (L-NH₂). As the sequential coupling of these components can be achieved without additional protectiondeprotection, a facile synthesis of various analogues is possible. This strategy also enables access to an array of molecules differing in chain length, stereochemistry, and functional groups. In addition, further diversification is possible by oxidation of the thioether to sulfoxide or sulfone, to allow a broader characterization of the GPR55 binding pocket.

To begin with, glucosylamine $\mathbf{4}^{21}$ readily available from Glc was reacted with diethyl squarate to give mono amide **5**



Figure 2 (a) LPGIc analogues used for axon turning assay. (b) Biological activity of squaryl group modified analogues assessed by axon turning assay. Compounds were tested at an in-pipette concentration of 10 μ M. Bars show mean turning angle ± SEM, numbers in parentheses indicate the number of axons tested. (c) Effect of GPR55 inhibitor on chemorepulsive response to compound 1A. Compound 1A was used at an intrapipette concentration of 20 μ M. Treating axons with bath-applied ML193, a specific inhibitor of GPR55, abolished 1A-induced chemorepulsion. Bars show mean turning angle ± SEM, numbers in parentheses indicate the number of axons tested. (**P<0.01, Kruskal-Wallis test. (d) Representative images of axon turning assays using compound 1A. Numbers in each panel show the minutes after onset of concentration gradient, arrows indicate the source of the gradient. Scale bar, 10 μ m. Assay conditions are further described in the supporting information.

(Scheme 2b). On the other hand, a small library of the lipid component (**6A-G**) was created (see Supporting Information) as exemplified for **6A** (Scheme 3a). To obtain **6A-F** in

homochiral forms, they were synthesized from (*S*)- or (*R*)glycidol as starting materials. Subsequent reaction with **5** gave **1A-G** (Scheme 3b). In a similar manner, D-xylose analogues **8A**-**E** were prepared from **7**.

To confirm the relevance of our design of LPGIc analogues, activity of 1A to initiate axon turning was evaluated in comparison with its diastereomer 1D and the xylose analogue 8A, along with sulfones 1B and 1E (Figure 2a). Assay was conducted as previously reported⁵ with some modifications. Exploratory experiments (Figure 2b) conducted at in-pipette concentration of 10 μ M (approximately 10 nM near the tip of axons) (Figure 2b) indicated that the compound 1A has the most favourable activity to repel axons of DRG nociceptive neurons. Higher chemorepulsive response was observed when the in-pipette concentration was increased to 20 μ M (Figure 2c,d), comparable to the native $LPGlc^{5}$. On the other hand, no significant activity was observed for other compounds including the diastereomer 1D and the xylose analogue 8A.^{22,23} These results indicate the importance of C-6 and C-2' hydroxy groups in order for LPGIc analogues to activate GPR55. The potent analogue 1A did not affect axon extension (Figure S-2), and was blocked by the specific GPR55 inhibitor ML193,14c indicating that it indeed acts on GPR55.

In conclusion, we developed a facile and modular strategy to create novel class of LPGIc analogues. Compound 1A, exhibiting identical stereoconfiguration of its hydroxy groups as native LPGIc, showed the expected LPGIc-like activity to repel DRG nociceptive axons. This confirmed our initial hypothesis that the squaryl group can act as a bioisostere of anomeric phosphodiesters. Failure of the xylose analogue 8A to stimulate GPR55, similar to the Gal and Man analogues reported previously, further augments the high preference of the GPR55 binding pocket towards glucose head groups. The inactivity of the sulfone analogues, such as 1B, suggests the presence of spatial constraints close to the FA ester position. Based on this strategy, systematic synthesis of LPGIc analogues as potential modulators of GPR55 is possible and a more comprehensive structure-activity relationship study along this line is underway.

This work was supported by Grants-in Aid for Scientific Research 16H0629 (YI), 17K14970 (ATG), and 16K08259 (PG) from Japan Society for the Promotion of Science, AMED-CREST Grant JP18gm0910006 (HK), Special Postdoctoral Program (FD) and Integrated Lipidology Program (PG, YH) of RIKEN. We thank Dr. Hiroyuki Koshino for NMR measurements, Dr. Takemichi Nakamura for mass spectroscopy measurements, and Akemi Takahashi for technical assistance. The computing resources were provided in part by the HOKUSAI-GreatWave system at RIKEN. We are indebted to Dr. Atsushi Miyawaki and Dr. Yasushi Sako for their support.

Conflicts of interest

There are no conflicts to declare.

Notes and references

Published on 09 July 2018. Downloaded on 7/10/2018 3:13:22 AM

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Feiqing Ding, ^a, † Adam T. Guy, ^b, † Peter Greimel,
b Yoshio Hirabayashi, c Hiroyuki Kamiguchi, *b and Yukishige Ito
*a

We report the facile synthesis of a series of LPGlc analogs, their GPR dependent biological activity and a systematic analysis of the structure-activity relationship in regards to GPR55 modulation.

