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Research Article

Identification of Physiologically Active Substances as Novel Ligands for MRGPRD

Makiko Uno,¹ Satoko Nishimura,² Keisuke Fukuchi,² Yasuyuki Kaneta,² Yoko Oda,² Hironobu Komori,³ Shigeki Takeda,⁴ Tatsuya Haga,⁵ Toshinori Agatsuma,² and Futoshi Nara¹

Correspondence should be addressed to Makiko Uno, uno.makiko.mv@daiichisankyo.co.jp

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Mas-related G-protein coupled receptor member D (MRGPRD) is a G protein-coupled receptor (GPCR) which belongs to the Mas-related GPCRs expressed in the dorsal root ganglia (DRG). In this study, we investigated two novel ligands in addition to beta-alanine: (1) beta-aminoisobutyric acid, a physiologically active substance, with which possible relation to tumors has been seen together with beta-alanine; (2) diethylstilbestrol, a synthetic estrogen hormone. In addition to the novel ligands, we found that transfection of MRGPRD leads fibroblast cells to form spheroids, which would be related to oncogenicity. To understand the MRGPRD novel character, oncogenicity, a large chemical library was screened in order to obtain MRGPRD antagonists to utilize in exploring the character. The antagonist in turn inhibited the spheroid proliferation that is dependent on MRGPRD signaling as well as MRGPRD signals activated by beta-alanine. The antagonist, a small-molecule compound we found in this study, is a potential anticancer agent.

1. Introduction

G protein-coupled receptors (GPCRs) compose the largest family of membrane-bound receptors and many members still have unrevealed functions. Efforts have been made to find ligands for GPCRs, and a broad range of ligands, including small organic compounds [1, 2], eicosanoids [3], peptides [4], and proteins [5], have been identified. Members of the Mas-related GPCR (Mrgpr) family are known to be expressed mainly in the subpopulations of sensory neurons [6]. This has resulted in their having another name sensory neuron specific receptors (SNSRs) [7]. Mas-related G-protein coupled receptor member D (MRGPRD), formerly called Mas-related gene D (MRGD) and also referred to as hGPCR45 [8] or TGR7 [9], belongs to the Mrgpr family and

is known to be mainly expressed in the dorsal root ganglia (DRG) [6, 9–12]. When phylogenic trees of the hMRGPR family and the mouse MRGPR family are compared, variation of the members found in those phylogenic trees are not parallel [6]. Members of the human MRGPR family may cover other functions from the ones revealed by studies with the mouse MRGPR family. We hypothesized MRGPRD could possess functions besides the ones in the DRG and found a spheroid forming activity. At this time, for MRGPRD, beta-alanine (β Ala) is the only known ligand that would be physiological [9]. When we screened a physiologically active substance library and a known small compound library, we found two novel ligands for MRGPRD, that is, beta-aminoisobutyric acid (β AIBA) and diethylstilbestrol (DES). As β Ala promoted spheroid growth of MRGPRD stably

¹ Exploratory Research Laboratories II, Daiichi Sankyo Co., Ltd., Tokyo 134-8630, Japan

² Biologics Research Laboratories, Daiichi Sankyo Co., Ltd., Tokyo 134-8630, Japan

³ Oncology Research Laboratories, Daiichi Sankyo Co., Ltd., 1-16-13 Kitakasai, Edogawa-ku, Tokyo 134-8630, Japan

⁴ Department of Chemical Biology, Gunma University Graduate School of Engineering, 1-5-1 Tenjin-cho, Kiryu, Gunma 376-8515, Japan

⁵ Institute for Biomolecular Science, Faculty of Science, Gakushuin University, 1-5-1 Mejiro, Tokyo 171-8588, Japan

expressing cells, it would be useful to obtain antagonists in order to analyze the tumorigenic character of MRGPRD. We obtained MU-6840 as an antagonist of MRGPRD by high-throughput-screening at a large small chemical library. Since it specifically inhibits the growth of spheroids formed with cells stably expressing MRGPRD, MU-6840 is a potential anticancer agent. In this paper, we will discuss the novel ligand, the antagonist, and the novel character of MRGPRD.

2. Materials and Methods

2.1. Materials. Dulbecco's Modified Eagle Medium (DMEM), Minimum Essential Medium Alpha without nucleic acids (α -MEM), RPMI1640, Opti-MEM I reduced serum media (Opti-MEM), Grace's Insect Medium, Penicillin-Streptomycin, Hanks' Balanced Salt Solution (HBSS), and LipofectAMINE 2000 were purchased from Invitrogen Corporation, USA. DMEM inositol-free was purchased from MP Biomedicals, USA. pSV2-dhfr was purchased from ATCC, USA. Fetal bovine serum (FBS) was purchased from Sanko Junyaku, Japan. PathDetect NFAT cis-Reporting System and PathDetect Elk1 trans-Reporting System were purchased from Agilent Technologies, USA. pRL-TK was purchased from Promega, USA. FBS dialyzed was purchased from JHR Biosciences, USA. Guanosine diphosphate (GDP), guanosine 5'-O-(3-thiotriphosphate) (GTPyS), saponin, LiCl, formic acid, and bovine serum albumin (BSA) were purchased from Sigma-Aldrich Inc., USA. Complete EDTA-free was purchased from Roche Diagnostics. LEADseeker beads, RNA binding YSi, myo-[2-3H]inositol and GTPyS [35S] were purchased from GE Healthcare, UK.

2.2. Cell Culture. HEK293 cells (DS Pharma Biomedical Corporation, Japan) were maintained in DMEM with 10% FBS and Penicillin-Streptomycin at 37°C under 5% CO₂. CHO/dhfr⁻ cells (DS Pharma Biomedical Corporation, Japan) expressing hMRGPRD (described in this paper) were maintained in α-MEM with 10% FBS dialyzed and Penicillin-Streptomycin at 37°C under 5% CO₂. NIH/3T3 cells (ATCC, USA) were maintained in RPMI1640 with 10% FBS and Penicillin-Streptomycin at 37°C under 5% CO₂.

2.3. Endogenous Signal Assay. Human adrenergic alpha 1A receptor (ADRA1A, AY389505.1) gene and human MRG-PRD (hMRGPRD, NM_198923.2) gene were cloned into pcDNA3.1 (Invitrogen, USA) (hereafter referred to as pcDNA-ADRA1A and pcDNA-hMRGPRD, resp.). Using LipofectAMINE 2000, 20000 HEK293 cells in each well of a 96-well multiplate were transfected with a set of 3 plasmids, consisting of one from each category as follows: (1) pNFAT-Luc plasmid (PathDetect NFAT cis-Reporting System), or pFR-Luc plasmid with pFA2-Elk1 plasmid (PathDetect Elk1 trans-Reporting System); (2) pRL-TK; (3) pcDNA3.1, pcDNA-ADRA1A, or pcDNA-hMRGPRD. Twenty-four hours after the transfection, the cells were lysed and transferred to OptiPlate-96. Samples were prepared in accordance with the manufacturer's instructions for the

Dual-Luciferase Reporter Assay System (Promega) and measured with the ARVO SX 1420 Multilabel Counter (Wallac). The signal intensity of each sample was calculated by dividing the value of firefly luciferase luminescent by the value of Renilla luciferase luminescent within the same sample. The cell culture media used in this assay was DMEM with 10% FBS, which may contain β Ala less than 0.5 μ M, at least 40 times lower than the EC50 of β Ala to MRGPRD.

2.4. Guanosine 5'-O-(3-Thiotriphosphate) Binding Assay. pFast-Bac-hMRGPRD-Giα, in which bovine Giα (NM-174324) was fused to hMRGPRD in frame and cloned into pFast-Bac (Invitrogen Corporation), was served to obtain baculovirus possessing hMRGPRD-Giα, using the Bac-to-Bac Baculovirus Expression System with pFast-Bac1 in accordance with the manufacturer's instructions. Sf9 cells in suspension culture were infected with the baculovirus possessing hMRGPRD-Gi α at MOI = 2 to 5 and maintained for 72 hours at 27°C. Cells were harvested and fractured using the general method for the N₂ cavitation system. Briefly, cells were suspended in a membrane preparation buffer (20 mM) Hepes-KOH, 1 mM EDTA, 2 mM MgCl₂, Complete EDTAfree, pH7.4) before being put in the N₂ cavitation apparatus (Fike Metal Products), and then pressurized at 600 psi for 30 minutes on ice. Burst cells were collected and separated from the nuclear fraction by centrifugal separation (1,800 xvg for 10 minutes). The supernatant was centrifuged at 100,000 x g for 30 minutes to obtain a membrane fraction. The pellet was then resuspended in a membrane stock buffer (20 mM Tris-HCl, 10% glycerol, Complete EDTA-free, pH7.4) and kept as a hMRGPRD-Gi α expressing membrane at -80° C. One microgram of the hMRGPRD-Gi α expressing membrane, 0.44 nM GTPyS [35 S], 5 μ M GDP, the compound of interest, and 3.65 mg/mL LEADseeker beads were prepared with an assay buffer (20 mM Hepes, pH7.3, 100 mM NaCl, 10 mM MgCl₂, 1 mM EDTA, 100 μg/mL saponin) in standard white 384 well assay microplates (Corning Inc.) at final volume of $60 \,\mu\text{L}$. After incubating the microplates at room temperature for 1 hour, $10\,\mu\text{L}$ of $100\,\mu\text{M}$ GTPyS was added to each well. The microplates were measured using LEADseeker (GE Healthcare). The assay buffer used in this assay did not contain any β Ala.

2.5. Calcium Flux Assay. CHO/dhfr⁻ cells were transfected with pSV2-dhfr and pSR-hMRGPRD, where hMRGPRD was cloned into an original vector containing SRα promoter, using LipofectAMINE2000 to establish hMRGPRD/CHO/dhfr⁻ cells, stable CHO/dhfr⁻ cell lines which overexpress hMRGPRD. Three thousand six hundred hMRGPRD/CHO/dhfr⁻ cells were plated onto each well of black 384 well cell culture microplates (Corning Inc.) and cultured for 2 days. In accordance with the manufacturer's instructions for the FLIPR Calcium 3 Assay Kit (Molecular Devices), the loading buffer, in which Calcium 3 Assay Solution was diluted with a HH buffer (HBSS, 25 mM Hepes, pH7.4, 2.5 mM probenecid), was added to each well after media disposal and kept at 37°C for 1 hour under 5% CO₂. For the agonist screening, the HH buffer containing a test compound was added to

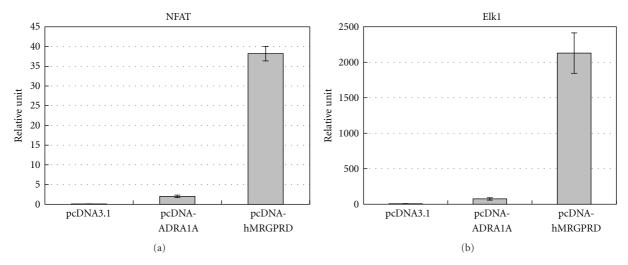


FIGURE 1: Signaling pathways of MRGPRD. Representative graphs of relative luciferase intensity of (a) NFAT and (b) Elk1. The results are the average of three independent samples.

each well and then the fluorescence transition caused by 488 nm excitation light was detected using FLIPR (Molecular Devices). For the antagonist screening, the HH buffer containing the test compound was added and incubated for 10 minutes and then the fluorescence transition was detected as the HH buffer containing β Ala was added to each well. The assay buffer used in this assay did not contain any β Ala.

2.6. IP Assay. Ten thousand hMRGPRD/CHO/dhfr- cells were plated onto each well of 96-well cell culture microplates and cultured for 6 hours, which were then labeled using labeling media (DMEM inositol-free, 10% FBS dialyzed, 1 nCi/mL myo-[2-3H]inositol) and cultured overnight. Labeling media was replaced with an IP assay buffer (200 mM HBSS, 25 mM Hepes, pH7.0, 0.1% BSA, 20 mM LiCl) that contained test compounds and was incubated for 1 hour for the agonists screening and incubated for 15 minutes for the antagonists screening. For the antagonist screening, in addition to the test compounds, β Ala was added and incubated for 1 hour. After removing the assay buffer, 20 mM formic acid was added and incubated for 3 hours at 4°C. Formic acid lysate was added to RNA binding YSi and shaken vigorously in OptiPlate-96 (Perkin Elmer). The microplates were incubated at room temperature overnight and examined for radioactivity using TopCount (Perkin Elmer). The assay buffer used in this assay did not contain any β Ala.

2.7. Spheroid Assay. The Retro-X System (Clontech Laboratories, Mountain View, CA) was used to generate NIH/3T3-hMRGPRD cells, which are NIH/3T3 cells stably expressing hMRGPRD and NIH/3T3-RasV12 cells, which are NIH/3T3 cells stably expressing RasV12. Briefly, cDNA encoding hMRGPRD or RasV12 was cloned into a pLNCX vector. The viral solution obtained from 293-10A1 cells (established in our laboratory), which were transformed with either pLNCX-hMRGPRD or pLNCX-RasV12, was used to

establish those overexpressing cells. Five thousand NIH/3T3-hMRGPRD cells or NIH/3T3-RasV12 cells were plated onto each well of a CellTightSpheroid (Sumitomo Bakelite Co., Ltd.) and cultured overnight in RPMI1640 medium supplemented with 10% FBS, which may contain β Ala less than 0.5 μ M, at least 40 times lower than the EC₅₀ of β Ala to MRGPRD. Test compounds were added to each well and incubated for 72 hours. No agonists were used in this assay. In accordance with the manufacturer's instructions for the CellTiter-Glo Luminescent Cell Viability Assay (Promega), an assay solution was added to each well, and the cell lysate solution was then transferred to OptiPlate-96 to be measured with the ARVO SX 1420 Multilabel Counter (Wallac).

3. Results

3.1. Signaling Pathways and Constitutive Activity of MRGPRD. MRGPRD is a GPCR that induces intracellular calcium ion influx and inhibits adenylyl cyclase activity by G protein signaling and cAMP production in response to β Ala [6, 9, 13, 14]. To confirm these signaling pathways, we performed reporter systems using NFAT as a downstream signal of the Gqα subunit and Elk1 as a downstream signal of the Giα subunit. Together with either of those reporters, hMRGPRD or ADRA1A, which is known to couple to both the $Gq\alpha$ and $Gi\alpha$ subunits and used as a positive control, were transfected to the cells to measure their signals. The plasmid which constitutively expresses Renilla luciferase by thymidine kinase promoter was also transfected simultaneously as an internal control of the assay to normalize transfection efficacy and cell growth of each sample. The endogenous signaling activities of the GPCRs were observed as relative units which were the quotients of either NFAT or Elk1 signal divided by constitutive signal. As shown in Figures 1(a) and 1(b), both ADRA1A and hMRGPRD showed signals depending on the $Gq\alpha$ subunit as well as the $Gi\alpha$ subunit. No signal was seen for either pathways when no GPCR

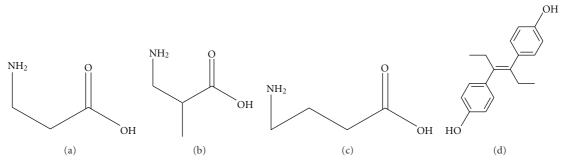


FIGURE 2: Structures of physiologically active substances which can activate MRGPRD: (a) beta-alanine (β Ala), (b) beta-aminoisobutyric acid (β AIBA), (c) gamma-aminobutyric acid (GABA), and (d) diethylstilbestrol (DES).

was added (Figures 1(a) and 1(b)). We found very high signals of hMRGPRD for both NFAT and Elk1 compared to those of ADRA1A: 19 times for NFAT and 28 times for Elk1, respectively (Figures 1(a) and 1(b)). The difference can be explained by assuming the "two-state model," where GPCRs exist in equilibrium between an inactive (R) state and an active (R^*) state [15]. As we used the same molar concentration of plasmid for hMRGPRD and ADRA1A and no ligand was added to this assay, these results indicate that the equilibrium of hMRGPRD is much more shifted to the R^* state than that of ADRA1A, in other words, hMRGPRD exhibits strong constitutive activity. These results indicate that expressing hMRGPRD in the cells can reinforce downstream signals without any ligand.

3.2. Screening a Physiologically Active Substance Library. Currently, β Ala and gamma-aminobutyric acid (GABA) are commonly regarded as physiological ligands of MRGD and both molecules are deeply related to neurotransmission [9]. When phylogenic trees of the hMRGPR family and the mouse MRGPR family are compared, variation of the members found in those phylogenic trees are not parallel [6]. This difference may cause the hMRGPR family members to have functions besides the ones revealed by the studies with mice. Therefore, we hypothesized hMRGPRD may have other functions besides neurotransmission. Since finding physiological ligands may help us deduce possible functions, we executed GTPγS binding assay using Giα fused hMRG-PRD to screen a physiologically active substance library. For GTPyS binding assay, as the membrane fraction of hMRGPRD-Gi α expressing Sf9 cells was used and no β Ala was contained in the assay buffer (20 mM Hepes, pH7.3, 100 mM NaCl, 10 mM MgCl₂, 1 mM EDTA, 100 μg/mL saponin) (see Section 2.4), the signals we detected were completely dependent on the compounds added to be tested. Compounds considered as hits were tested with a calcium flux assay. For the calcium flux assay, only transient signals were detected and compounds were added with a buffer (HBSS, 25 mM Hepes, pH7.4, 2.5 mM probenecid) that did not contain β Ala (see Section 2.5); therefore, the signals detected were again completely dependent on the compounds added for testing. We found 3 compounds that were positive, each of which in turn underwent an IP

TABLE 1: Agonistic potency of physiologically active substances to induce signals of hMRGPRD.

EC ₅₀ (μM)	GTPγS	Calcium flux	IP
β Ala	219.8 ± 115.7	22.6 ± 11.4	353.3 ± 217.0
β AIBA	821.7 ± 120.9	53.4 ± 27.3	378.3 ± 290.5
GABA	Not determined	4385 ± 871.1	>1000
DES	50.7 ± 18.1	10.4 ± 3.0	not available

EC₅₀ (sample concentration producing 50% of the maximal response) was calculated from dose-response curves and shown in μ M. Multiple assays were executed to calculate EC₅₀s: calcium flux assay (Calcium flux), GTP γ S binding assay (GTP γ S), and IP assay (IP).

assay. For the IP accumulation assay, DMEM inositol-free media with 10% FBS dialyzed was used to label myo-[2, 3] inositol, then the labeling media was replaced with an IP assay buffer (200 mM HBSS, 25 mM Hepes, pH7.0, 0.1% BSA, 20 mM LiCl), that did not contain any β Ala (see Section 2.6); therefore, accumulated IPs were attributed to the compound added for testing. The 3 compounds which we considered as hits were β Ala (Figure 2(a)), β AIBA (Figure 2(b)), and DES (Figure 2(d)) whose EC₅₀s are shown in Table 1. Representative dose-response curves of these compounds in each assay are shown in Figure 3. In all assays, the activity value of hMRGPRD with the saturated concentration of β Ala was set as 100% for each assay system, and activity values are shown in percentages. For the IP assay, since the cells were kept in the assay buffer containing compound for an hour, a much higher DES concentration than the cells could tolerate was needed to calculate EC_{50} ; therefore, we could not obtain EC₅₀ of DES in the IP assay. β AIBA was a newly found agonist, which was as potent as β Ala, and had a structure similar to β Ala (Figure 2). We also tested gamma-aminobutyric acid (GABA), which shares the same moiety with β Ala and β Ala (Figure 2) and had showed some activity for hMRGPRD in previous study [9]. Although with much weaker activity, GABA also worked as an MRGPRD agonist (Table 1, Figure 3).

3.3. Antagonists Screening. One of the nearest neighbors of hMRGPRD is MAS1 oncogene, which shows focus formation when transfected into NIH-3T3 cells [6, 16]. Also, there is information that β Ala, β AIBA, and GABA are related to some

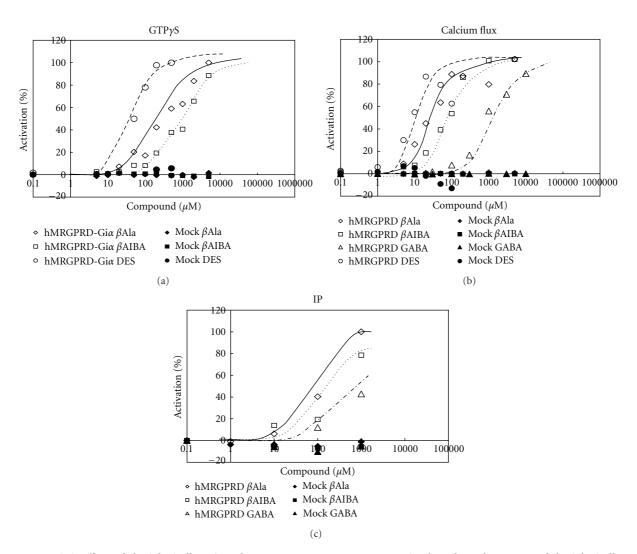


FIGURE 3: Agonistic effects of physiologically active substances on MRGPRD. Representative dose-dependent curves of physiologically active substances in (a) GTP γ S binding assay, (b) calcium flux assay, and (c) IP assay are shown. Each value expresses the percentage of control with maximum efficacy in each assay. β Ala on MRGPRD (open diamond), β AIBA on MRGPRD (open square), GABA on MRGPRD (open triangle). DES on MRGPRD (open circle), β Ala on mock (closed diamond), β AIBA on mock (closed square), GABA on mock (closed triangle), DES on mock (closed circle).

kinds of tumors [17-22]. We took notice of oncogenicity and found that hMRGPRD-transformed fibroblast cells could form spheroids. The details of this feature will be reported elsewhere. We next screened a large chemical library to obtain MRGPRD antagonists which would help us examine the novel character of MRGPRD, the spheroid forming activity. Using β Ala as a ligand for MRGPRD, a highthroughput screening (HTS) for antagonists was executed by the calcium flux method utilizing FLIPR, and MU-6840 (Figure 4) was selected as a hit. MU-6840 did not inhibit signals of the following GPCRs: CCR4, CCR5, CCR7, CCR8, CXCR3, S1PR1, P2Y12R, OGR1, OXER1, GPR84, GPR14, or GHS-R (data not shown). As shown in Table 2, MU-6840 inhibited β Ala signals through hMRGPRD dosedependently in all methods tested here, namely, calcium flux (Figure 5(a)), GTPyS binding (Figure 5(b)), IP accumulation (Figure 5(c)), and spheroid proliferation (Figure 5(d)).

We did not use the endogenous signaling assay mentioned above (Section 2.3) because it is not suitable for detecting the effects of compounds due to the following reasons: (a) the transient transfection assay requires 24 hours to garner enough luciferase expression to be measured quantitatively; (b) cells cannot tolerate compound addition at the same time as transfection, as this would cause cell damage; (c) if the compound is added after suitable luciferase expression has been reached, the translated luciferase prior to compound addition remains and only the luciferase being transcribed/translated during compound addition would be inhibited, which would make the basal signal high enough to mask the inhibitory effect of the compound.

In Figures 5(a), 5(b), and 5(c), values are shown in percentages: the value at adding only β Ala with no antagonist is set as 0%; the value at adding neither ligand nor antagonist is set as 100%. For the calcium flux assay (Figure 5(a)), as

FIGURE 4: Structure of hMRGPRD antagonist, MU-6840.

Table 2: Antagonistic activity of MU-6840 for hMRGPRD.

	Calcium flux	GTPyS	IP	Spheroid
IC ₅₀ (nM)	691.7 ± 120.9	1703.3±353.9	473.8 ± 160.3	655.9 ± 204.3

 IC_{50} (sample concentration producing 50% of the maximal inhibition) was calculated from dose-response curves and shown in nM. Multiple assays were used to calculate IC_{50} s: calcium flux assay (Calcium flux), GTP γ S binding assay (GTP γ S), IP assay (IP), and spheroids growth assay (Spheroid).

no endogenous activity could be measured, the maximum inhibition never exceeded 100%. On the other hand, for the GTP γ S binding assay (Figure 5(b)) and IP accumulation assay (Figure 5(c)), the maximum inhibitions exceeded 100% as these assays are able to detect hMRGPRD's strong constitutive activity. These excesses indicate that MU-6840 can act as an inverse agonist.

Owing to hMRGPRD's endogenous activity (Figures 1(a) and 1(b)), when hMRGPRD is stably expressed, NIH/3T3, which would not form spheroids by itself, can form spheroids with no ligand, such as β Ala. In this spheroid assay (see Section 2.7), the culture media, RPMI1640, does not contain β Ala. In addition, β Ala which would come from FBS, taking up 10% of the media, could be negligible for the following reasons: the β Ala concentration of normal serum is around $4\,\mu\mathrm{M}$ [23, 24] which makes $\beta\mathrm{Ala}$ concentration in this media less than $0.5 \,\mu\text{M}$; EC₅₀s of β Ala to hMRGPRD is higher than 20 μ M (Table 1). When β Ala was added to this assay system, spheroid growth of NIH/3T3-hMRGPRD was promoted, although no effect was seen with the spheroid growth of NIH/3T3-RasV12 cells, overexpressing oncogenic gene RasV12 (data not shown). It shows that the spheroid proliferation of NIH/3T3-hMRGPRD is dependent on MRGPRD signaling pathways. In order to determine the effect of the compounds, a spheroid proliferation assay is more suitable than a focus formation assay. For a spheroid assay, it is common to use RasV12 as a positive control [25, 26]. In the spheroid assay, MU-6840 inhibited NIH/3T3hMRGPRD spheroid proliferation, which was dependent on basal activity of MRGPRD, that is, independent of any agonist. This result indicates that MU-6840 exhibits inverse agonism. Furthermore, MU-6840 did not inhibit spheroid proliferation of NIH/3T3-RasV12 cells (Figure 5(d)). Taking

all things into account, MU-6840 shows inhibitory activity specific to MRGPRD.

4. Discussion

MRGPRD belongs to the Mas-related GPCR family, which has been shown to be expressed in the DRG [6, 9–12]. When phylogenic trees of the hMRGPR family and the mouse MRGPR family are compared, variation of the members found in those phylogenic trees are not parallel [6]. This difference may cause the hMRGPR family members to have functions besides the ones revealed by the studies with mice. Being one of the members of hMRGPR, we hypothesized that MRGPRD could have other functions besides the ones in DRG and found a spheroid forming activity.

In this study, we found 2 physiologically active substances as novel ligands for MRGPRD: β AIBA and DES (Figures 2(b) and 2(d)). β Ala has long been widely known to act as a ligand for glycine receptors and GABA receptors in the central nervous system [27-32]. Since MRGPRD is known to be expressed in the DRG [6, 9–12], it is reasonable that β Ala acts as a ligand of MRGPRD to have some roles in the nervous system. Besides the nervous system, MRGPRD may also have roles in renin-angiotensin system by releasing arachidonic acid in response to angiotensin (Ang) II metabolites, although no direct signals, that is, calcium flux, IP accumulation, and so forth, of MRGPRD in response to Ang II metabolites are shown [33]. On the other hand, besides being one of the nearest members of MAS1 oncogene [6, 16], there is some information that β Ala, as well as β AIBA, is related to some types of tumors. Rat or mouse mammary tumors contain high levels of taurine, β Ala, and GABA, which are enriched in neural tissue and which are not seen in normal rat or mouse mammary tissue [17]. β Ala concentrations in nerve-related tissue are shown to be about $50 \,\mu\text{M}$ in rat sciatic nerve and $60 \,\mu\text{M}$ in cat brain [24, 34], which are higher than EC₅₀ of β Ala to MRGPRD in a calcium flux assay (Table 1). Cancer patients were more prone to have increased urinary β AIBA excretion than patients with different diseases [18]. The percentage of high urinary β AIBA excreter was higher in urinary bladder carcinoma patients compared to patients with urinary bladder papilloma or cystitis, or healthy volunteers [19]. The massive amounts of β AIBA in the urine of Burkitt's lymphoma patients were measured, and the degree of elevation of β AIBA excretion seemed to be related to the amount of tumor mass present [20, 21]. β AIBA excretion in patients with chronic myeloid leukemia was directly correlated to the leukocyte count, the indicator of tumor cell mass [22]. The average β AIBA level in the urine of patients with Burkitt's lymphoma is about 170 times higher than that of normal subjects [21]. Since the serum concentrations of β AIBA in normal subjects are about $2.5 \,\mu\mathrm{M}$ [35], even if the difference of the serum $\beta\mathrm{AIBA}$ level is not as conspicuous as that of in urine, only 30 times difference is enough to exceed the EC₅₀ of β AIBA to MRGPRD in a calcium flux assay (Table 1). All together, these suggest that β Ala and β AIBA may be involved in tumorigenicity by enhancing MRGPRD oncogenic signals. However, it does not exclude the possibility that increasing concentrations of β Ala

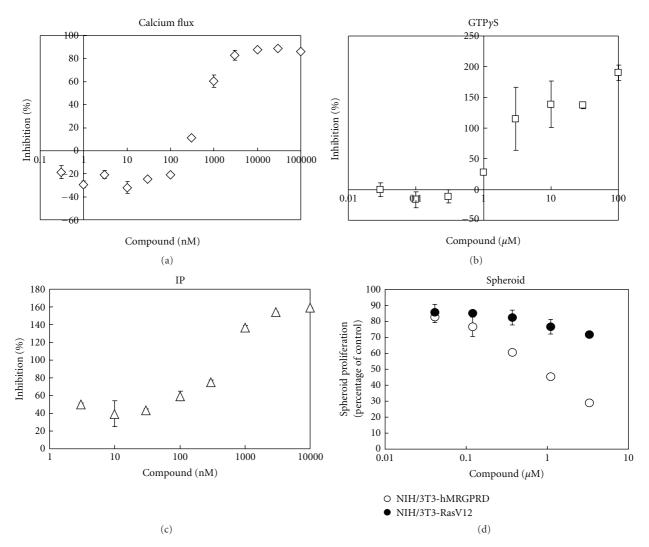


FIGURE 5: Inhibitory effects of MU-6840 on MRGPRD. Representative dose-dependent inhibition curves in (a) calcium flux assay, (b) GTP γ S binding assay, (c) IP assay, and (d) spheroid assay by MU-6840 are shown. Each value expresses the percentage of control with no compound present. The results are the average of three independent tests.

and β AIBA is just a sign of the existence of cells with high metabolic rates, such as cancer cells, as β Ala and β AIBA are degradation products of nucleic acids.

Besides β Ala, β AIBA, GABA, and DES, 5 synthetic agonists for MRGPRD are reported [14, 36]. At least two of them are specific to MRGPRD, in other words they do not activate other Mas-related GPCRs [36]. Considering EC₅₀s and the concentrations *in vivo*, we assume that β Ala and β AIBA are the ones which can contribute to the physiological functions of MRGPRD.

Another new aspect we would like to focus on is the high endogenous signaling activity of MRGPRD. There are publications that some GPCRs, such as human β_2 adrenoceptor, exhibit significant basal, agonist-independent G protein activation, which is associated with the number or ratio of the active (R^*) state receptors [15, 37, 38]. In the report of Ajit et al., they noted that "our studies indicate that MrgD RNA expression inversely correlated with larger

beta-alanine induced calcium transient functional responses as assessed by FLIPR responses" in the explanation of their selection of the cloned MRGPRD expressing cell for a calcium flux assay [14]. We also had a similar experience with this episode, and we assume this phenomenon clearly implies MRGPRD's high basal activity: if there are many MRGPRDs in the cells, the signal has already reached the maximum without any ligand, and no transient signal with the ligand could be detected, since FLIPR can only detect transient signal. Regarding MRGPRD's high endogenous signaling activity, just expressing a suitable amount of MRGPRD may add cells of a tumorigenic phenotype. High MRGPRD expression might attribute to some types of cancer, as we have detected a very high MRGPRD expression in some clinical cancer tissues (data not shown). MU-6840, a specific MRGPRD antagonist and possibly acting as an inverse agonist, has the potential ability to become an anticancer agent.

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