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Identification of the PGRMC1 protein complex as the putative sigma-2 receptor binding site

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Abstract

The sigma-2 receptor, whose gene remains to be cloned, has been validated as a biomarker for tumor cell proliferation. Here we report the use of a novel photoaffinity probe, **WC-21**, to identify the sigma-2 receptor binding site. **WC-21**, a sigma-2 ligand containing both a photoactive moiety azide and a fluorescein isothiocyanate group, irreversibly labels sigma-2 receptors in rat liver; the membrane-bound protein was then identified as PGRMC1 (progesterone receptor membrane

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Author contributions: J.X., C.Z., K.T.W., and R.H.M. conceived and coordinated the experiments and wrote the manuscript. W.C. and D.Zhou designed and synthesized **WC-21**, **WC-26** and synthesized siramesine under the supervision of R.H.M.; S.V., Z.T. and D.Zeng synthesized compounds **SV119**, **SW120** and [¹²⁵I]**RHM-4** under the supervision of R.H.M.. J.X. designed and carried out the protein purification and photoaffinity labeling study. F.P., J.M.R., and F.Z. performed the cell culture and western blot analysis under the supervision of C.Z.. K.C.C. designed and performed the caspase-3 flow cytometry study under the supervision of R.S.H.. F.J. and D.S. designed and performed the PGRMC1 knockdown experiments with HEK 293T cells under the supervision of W.G.H..

component-1). Immunocytochemistry reveals that both PGRMC1 and **SW120**, a fluorescent sigma-2 receptor ligand, colocalizes with molecular markers of the endoplasmic reticulum and mitochondria in HeLa cells. Overexpression and knockdown of the PGRMC1 protein results in an increase and a decrease in binding of a sigma-2 selective radioligand, respectively. The identification of the putative sigma-2 receptor binding site as PGRMC1 should stimulate the development of unique imaging agents and cancer therapeutics that target the sigma-2 receptor/PGRMC1 complex.

Introduction

Sigma-1 and sigma-2 receptors were initially characterized in the mid-1970's using radioligand binding studies¹. Radiolabeled benzomorphans such as [³H](+)-pentazocine bind potently and selectively to sigma-1 versus sigma-2 receptors. Characterization of the sigma-2 receptor has relied on radioligand binding studies using [³H]DTG in the presence of (+)-pentazocine or dextrallorphan to block binding to sigma-1 receptors¹. The sigma-1 receptor was purified, sequenced and cloned from guinea pig brain in 1996, and shows no sequence homology with any mammalian proteins cloned to date². Ligands binding to the sigma-1 receptor modulate the release of neurotransmitters; several of these compounds have shown promise as antipsychotics, antidepressants, and drugs blocking the effects of psychostimulants^{3,4}. Recent studies indicate that the sigma-1 receptor is expressed in postsynaptic sites, regulates ion channels, and may function as a protein chaperone⁵⁻⁸. To date, the sigma-2 receptor has not been purified, sequenced or cloned.

Although its structure is not known, the sigma-2 receptor has been validated as a biomarker for tumor cell proliferation both *in vitro* and *in vivo*, and as a target for chemotherapy. Several studies have shown that: the sigma-2 receptor density in proliferative breast cancer cells is about 10-fold higher than in quiescent breast cancer cells⁹; sigma-2 receptor expression is upregulated during the transition from quiescence to proliferation and down-regulated during the transition from proliferation to quiescence¹⁰; sigma-2 receptor density can be used to determine the proliferative status of solid tumors¹¹; sigma-2 receptor ligands can serve as chemotherapeutics for treating solid tumors¹²⁻¹⁴; and, sigma-2 receptor radioligands can be used to image tumors by positron emission tomography (PET) or single photon emission computed tomography (SPECT)¹⁵⁻¹⁹. However, since the sigma-2 receptor gene has not been cloned, the exact role this receptor plays in tumor and normal cell proliferation is currently unknown. Our group has developed sigma-2 selective ligands, including [³H]RHM-1, [¹²⁵I]RHM-4, and **SW120** to directly identify, locate, and quantify this protein using a variety of receptor binding and molecular imaging techniques^{13-15, 18, 20-22}.

Progesterone receptor membrane component 1 (PGRMC1) is overexpressed in a variety of tumors in comparison with the corresponding normal cells, and thus represents an important biomarker for cancer progression and a potential target for anticancer drugs²³⁻²⁵. PGRMC-1, which exists in phosphorylated and dephosphorylated states during cancer cell proliferation²⁶, regulates cell growth and proliferation through interactions between its Cytochrome b5 binding domain and other potential binding partners which include Insig-1,

PAIR-BP1, and P450 proteins^{23, 27-31}. Recent studies suggest that PGRMC1 is a biomarker of cell proliferation and an excellent therapeutic target for inhibiting tumorigenesis^{24, 32, 33}. Interestingly, proteomic studies also showed that PGRMC1 is expressed in high levels in the proliferative cells of human endometrium³⁴.

The purpose of the current study was to identify the protein or protein complex that contains the sigma-2 receptor ligand-binding site as the first step in determining the role of this receptor in cell proliferation, and defining a new target for the development of novel tumor imaging agents and cancer therapeutics. Here we describe a novel strategy developed for the identification of the sigma-2 receptor. The sigma-2 receptor photoaffinity probe, **WC-21**, an analogue of **RHM-1**, contains an azide moiety for the photo-affinity tagging of the sigma-2 receptor, and a fluorescein isothiocyanate group (FITC) for protein visualization. **WC-21** was utilized to directly label the sigma-2 receptor binding site in rat hepatic membrane homogenates. Proteomic studies and consequent studies were carried out to determine the molecular identify of the sigma-2 receptor binding site and if it corresponded to any previously identified proteins. PGRMC1 was identified as the putative sigma-2 receptor binding site. The results of our studies provide strong experimental evidence supporting the localization of the putative sigma-2 receptor binding site within the PGRMC1 protein complex.

Results

Photoaffinity labeling and protein identification

The synthesis of sigma-2 receptor photoaffinity probe **WC-21**, which contains an azide moiety for the photoaffinity tagging of the protein and a FITC group for protein visualization, is described in detail in the supporting online material (Supplementary Figs. S1- S3). **WC-21** has high binding affinity for sigma-2 receptors ($K_i = 8.7$ nM) and relatively low binding affinity for sigma-1 receptors ($K_i > 4,000$ nM) (Fig. 1a). **WC-21** was incubated with rat liver membrane homogenates, and the **WC-21**-protein complex photo-crosslinked. The protein supernatant was enriched and separated by gel electrophoresis (Fig. 1b). Western blot analysis revealed a dominant protein band at ~24 kD that was labeled by FITC conjugated probe **WC-21** (Fig. 1b). Labeling of this protein band with **WC-21** could be blocked by well-characterized sigma-2 receptor ligands (Supplementary Fig. S4). Proteomic studies of the protein in the ~24 kD band identified two proteins, the 22 kD putative progesterone binding protein and the 25 kD 25-Dx protein, that share the same genetic name, *progesterone receptor membrane component 1* (PGRMC1)²³. Fig. 1c shows the protein sequence of the two peptides, **gdqpgasgdndddeppplpr** and **dftpaelr**, detected by mass spectrometry and the Cytochrome b5 domain (underlined) of PGRMC1. The gene for PGRMC1 is located on q22-q24 of the X chromosome.

Three other proteins, glutathione-S-transferase alpha type 2, glutathione S-transferase mu 1 (with 2 matched peptides), and NADH dehydrogenase (ubiquinone) flavoprotein 2 (with 1 matched peptide), were identified from the proteomics study. Based on the localization of sigma-2 receptors within the plasma membrane^{9, 35, 36}, endoplasmic reticulum and mitochondria^{21, 22}, these proteins were excluded as potential candidates since they are all intracellular enzymes which happen to overlap with PGRMC1 protein band.

The pharmacological profile of [¹²⁵I]RHM-4 in HeLa cells

[¹²⁵I]RHM-4 (Fig. 2a) is a useful radioligand for measuring sigma-2 receptor density in tumors and normal tissues¹⁹. AG-205, a reported PGRMC1 ligand^{33,37}, and the known sigma-2 receptor ligands, DTG, siramesine, **SV119**, and **WC-26**, readily displaced [¹²⁵I]RHM-4 binding in HeLa cell membrane homogenates (Fig. 2a). Similar results were also observed in mouse mammary 66 tumor cell membrane homogenates (Supplementary Fig. S5).

Binding activity in PGRMC1 siRNA and cDNA treated HeLa cells

Because PGRMC1 is a known protein, a number of molecular biology tools are available to study the relationship between the expression of this protein and the binding of sigma-2 selective ligands in cancer cells. Knockdown studies using a PGRMC1-specific siRNA (Fig. 2b) reduced the binding of [¹²⁵I]RHM-4 to HeLa cells (Fig. 2c), indicating that the PGRMC1 complex has binding properties similar to the sigma-2 receptor. Similar results were observed in Human Embryonic Kidney (HEK) 293T cells using the PGRMC1 specific siRNA and the sigma-2 selective fluorescent probe, **SW120** (Supplementary Fig. S6). Cells treated with nontargeting siRNA were used as controls. Overexpression studies were also conducted in HeLa cells using a cDNA for the PGRMC1. HeLa cells transfected with the PGRMC1 cDNA displayed an increase in protein levels (Fig 2d) and a 60% increase in binding of [¹²⁵I]RHM-4 (Fig 2e) over cells transfected with vector alone. Sigma-1 receptor binding activities, determined using radioligand [³H](+)-pentazocine binding were found to be not changed in the nontargeting siRNA and PGRMC1 siRNA treated cells, or in cells transfected with vector and PGRMC1 cDNA (Supplementary Fig. S7).

Sigma-2 receptor and PGRMC1 in mammary 66 and 67 cells

A good correlation between sigma-2 receptor binding of [¹²⁵I]RHM-4 and the PGRMC1:actin ratio was observed in mouse mammary adenocarcinoma cells lines 66 and 67 after 6 (proliferative cells), 12 (early quiescent phase) and 18 (late quiescent phase) days in culture (Supplementary Fig. S8). These data are consistent with the expression of sigma-2 receptors in 66 and 67 cells⁹.

Caspase-3 activation and PGRMC1 upregulation by WC-26

Studies have shown that sigma-2 receptor ligands can induce caspase-3-dependent cell death^{14,38}. In the current study, functional assays were conducted to examine the ability of the PGRMC1 to regulate caspase-3 activation by the sigma-2 receptor ligand, **WC-26** (Fig. 3a). In HeLa cells, knocking down the PGRMC1 resulted in a blunting of the effect of **WC-26** to induce caspase-3 activation relative to HeLa cells treated with the non-targeting siRNA (Fig. 3b). Also, previous studies have shown that AG-205, a PGRMC1 ligand, induced an upregulation of PGRMC1 in A549 lung cancer cells^{33,37}. The sigma-2 receptor ligand **WC-26** induced an upregulation of PGRMC1 protein in a dose-dependent manner in nontargeting siRNA-treated cells (Fig. 3c). Only a minimal increase in PGRMC1 expression was observed in the PGRMC1 knockdown cells.

Intracellular localization of PGRMC1 and sigma-2 receptors

The intracellular localization of PGRMC1 and sigma-2 receptors was compared using confocal microscopy. PGRMC1 proteins were visualized with an anti-PGRMC1 antibody, whereas sigma-2 receptors were visualized using the fluorescent probe, **SW120**²¹ (Fig. 4a). Both the sigma-2 receptor probe (Fig. 4b, d) and anti-PGRMC1 antibody (Fig. 4c, e) colocalized with markers of the mitochondria (Fig. 4b, c) and the endoplasmic reticulum (Fig. 4d, e). An additional set of high magnification images are shown in Supplementary Fig. S9.

Discussion

In recent years, considerable effort has been focused on the identification of cancer biomarkers which could be used in early detection, patient prognosis, and prediction of response to different types of chemo- and radiotherapy. In many regards, the sigma-2 receptor possesses a unique ability to serve as both a diagnostic and therapeutic biomarker of solid tumors. The 10-fold higher density of sigma-2 receptors in proliferating breast cancer cells versus quiescent breast cells *in vitro* and *in vivo*^{9, 11} suggest that measuring the sigma-2 receptor status of solid tumors should provide an assessment of the proliferating:quiescent cell ratio in solid tumors, which could be used by an oncologist to select between certain types of chemotherapy (e.g., cell cycle specific versus cell cycle nonspecific) and radiation therapy (hyperfractionated versus conventional radiation therapy). Furthermore, preclinical studies of sigma-2 receptor agonists have shown promise as chemotherapeutic agents since they are capable of killing tumor cells via a variety of mechanisms including caspase- and non-caspase-mediated apoptosis. However, the widespread acceptance of the sigma-2 receptor as a cancer biomarker has been hampered by the fact that the molecular identity of this protein was not known.

The goal of the current study was to complete a series of steps aimed at determining the molecular identity of the sigma-2 receptor. The first step involved the development of a sigma-2 selective photoaffinity probe which could irreversibly tag sigma-2 receptors in rat liver membrane homogenates, the tissue source used in sigma-2 receptor binding assays. The photoaffinity labeling and mass spectrometry sequence analysis indicated that a strong candidate protein for the sigma-2 receptor was PGRMC1. A review of the literature revealed a number of similarities between the expression of PGRMC1 in cancer cells and normal tissues and the density of sigma-2 receptors in the same cancer cells and tissues. The availability of siRNA, cDNA, and monoclonal antibodies for PGRMC1 provided the tools needed to study the relationship between the expression of this protein and the binding of sigma-2 selective ligands in cancer cells. The intracellular localization of PGRMC1 and sigma-2 receptors in mitochondria and endoplasmic reticulum, reduction in sigma-2 receptor binding and caspase-3 mediated cell death in PGRMC1 siRNA treated HeLa cells, and the increase in sigma-2 receptor binding upon overexpressing the PGRMC1 clearly demonstrate that sigma-2 receptor ligands bind to the PGRMC1 protein complex. Therefore, this study is the first to report that the putative sigma-2 receptor ligand binding site is located within the PGRMC1 protein complex. In addition, the observations that AG-205, a reported PGRMC1 ligand, displaces [¹²⁵I]RHM-4 from HeLa cell membrane homogenates, and that both

AG-205 and the sigma-2 receptor ligand **WC-26** induce an upregulation of the PGRMC1 protein in lung cancer cells^{33, 37} and HeLa cells further support our conclusions.

The identification that the putative sigma-2 receptor binding site resides within the PGRMC1 protein complex is significant for a variety of reasons. First, it provides a link between receptor binding studies using sigma-2 radioligands and a known protein, the PGRMC1, which has been validated as a biomarker of cancer and cell proliferation. Second, it provides a gene whose expression can be examined across a wide panel of tumor cells. The genetic mutations and single-nucleotide polymorphism (SNP) of PGRMC1 can also be readily examined in assessing the possible roles of this gene in normal cell function and in malignant transformation. Finally, the results reported here provide an important scientific bridge between the sigma-2 receptor and the PGRMC1, and make available a variety of radioligands, fluorescent probes, and potential small molecule ligands to study the sigma-2 receptor/PGRMC1 complex in solid tumors and cancer cells using a variety of experimental techniques.

Methods

Chemical synthesis of the compounds and radiotracers

The tritiated compound, [³H]**RHM-1**, was made by American Radiolabeled Chemicals, Inc. (St. Louis, MO, USA) via *O*-alkylation of the corresponding phenol precursor^{17, 20}; the specific activity of the radioligand was 80 Ci/mmol. [³H](+)-pentazocine was purchased from Perkin-Elmer (Boston, MA, USA). [¹²⁵I]**RHM-4** was prepared by an iododestannylation reaction of the corresponding tributyltin precursor^{13, 19}. **SV119**, **SW120** and **WC-26**, and siramesine were synthesized by our group according to published methods^{12, 21, 39, 40}. The synthesis of the novel FITC conjugated probe, **WC-21** (Supplementary Figures S1-S3), is described in the supplementary online methods. AG-205 was purchased from TimTec (Newark, DE, USA). DTG and (+)-pentazocine were purchased from Sigma Aldrich (Milwaukee, WI, USA). Haloperidol was purchased from Tocris Bioscience (St. Louis, MO, USA). PGRMC1 recombinant protein was purchased from Assay Designs, Inc. (Ann Arbor, MI, USA).

Cell membrane and hepatic membrane homogenates

Human HeLa and HEK 293T cells were grown in Minimum Essential Media (MEM) containing 10% fetal bovine serum, 2 mM L-glutamine, 1% nonessential amino acids, and 1 × penicillin/streptomycin at 37 °C in a humidified 5% CO₂/95% air atmosphere. The mouse mammary tumor 66 cells were grown as previously described⁹. The cells were harvested from T75 flasks after 6, 12, or 18 days in culture. For binding assays, cells were mechanically scraped, rinsed with ice-cold PBS, transferred to a 15 mL conical centrifuge tube and centrifuged for 10 min at 200 × g. The cell pellet from each T75 flask was resuspended in 1 mL of 50 mM Tris-HCl at pH 8.0, and homogenized using an Ultra-Turrax T8 polythion homogenizer (IKA Works, Inc, Wilmington, NC, USA) at speed 3 for 20 seconds. Hepatic cells membrane homogenates were prepared from the liver of male Sprague-Dawley rats as previously described²⁰.

Photoaffinity labeling and protein identification

For photoaffinity labeling, 100 nM of **WC-21** was incubated in a 96 well plate with rat liver membrane homogenates (~ 300 µg of protein in a volume of 150 µL) for 1 h, then the reactions were harvested and filtered into 96 well filter plate after being washed three times. The washed filters were irradiated with 254 nm ultraviolet light for 2 min at a distance of 1 cm; the protein-loaded filters were then punched out, pooled and crushed by homogenization in 2% sodium dodecyl sulfate (SDS) solution in a 15 mL centrifuge tube, following a centrifugation of 10 minutes at speed of $800 \times g$, the supernatant was mixed with protein gel sample buffer and boiled for 2 min.

For western blot analysis of the **WC-21**-labeled protein band, lysates containing 35 µg of protein were run on a 12% acrylamide gel and transferred to a polyvinylidene fluoride (PVDF) membrane (Bio-Rad Laboratories, Hercules, CA, USA). The position of the predominant band of the ligand-labeled protein in the gel was determined using a horseradish peroxidase-conjugated goat anti-FITC antibody (Bioscience Resource Project, Saco, ME, USA) at a 1:1,000 dilution (Supplementary Fig. S3).

Western blot analysis with a horseradish peroxidase (HRP) conjugated anti-FITC antibody revealed the predominant protein band at ~24 kD was labeled by **WC-21** (Fig. 1b). This protein band labeling could be blocked by sigma-2 receptor ligands, such as DTG, haloperidol and **RHM-1** (Supplementary Fig. 3). The 24 kD band on the silver stained gel was excised and trypsinized; the trypsinized protein solution was further confirmed as tagged with **WC-21** an antibody specific for FITC. Proteomic study of the protein solution identified two proteins, the 22 kDa putative progesterone binding protein and the 25 kDa 25-Dx protein, which share the same genetic name, progesterone receptor membrane component 1 (PGRMC1)

For sequencing studies, proteins were enriched by SDS polyacrylamide gel electrophoresis (SDS-PAGE) on 4-15% gradient gels using ~100 µg of protein per loading well. Ten gel pieces containing proteins ranging from 20-26 kD molecular weight were excised and subjected to electro-elution with Midi GeBAflex-tube (Gerard Biotech, Oxford, OH, USA). The eluted protein samples were dialyzed with 7,000 MWCO SnakeSkin pleated dialysis tubing (Pierce Biotechnology, Inc. Rockford, IL, USA) and lyophilized; the enriched protein sample was further separated with two 12% gels. One gel was silver stained for protein band visualization using a mass spectrometry compatible kit from Bio-Rad Laboratories (Hercules, CA, USA) and the other gel was transferred to a PVDF membrane for western blot analysis with a horseradish peroxidase conjugated anti-FITC antibody (Fig. 1B). The FITC positive protein band was found at molecular weight of 24 kD, which was determined using the protein standards. The 24 kD band was excised and trypsinized. MALDI-mass spectrometry analysis of the trypsinized proteins was performed by the Proteomics & Mass Spectrometry facility of the Donald Danforth Plant Science Center (St. Louis, MO, USA) to identify the proteins.

PGRMC1 siRNA treatment

HeLa cells were plated on 6-well plates at a cell density of 1.5×10^5 cells/well. 24 h after plating, the cells were treated with either nontargeting siRNA (catalogue number D-001210-02-20, Thermo Scientific Dharmacon, Inc., Lafayette, CO, USA) or human PGRMC1 siRNAs (catalogue number L-010642-00-0020, Thermo Scientific Dharmacon, Inc., Lafayette, CO, USA) mixed with DharmaFECT 2 Transfection Reagent (Thermo Scientific Dharmacon, Inc., Lafayette, CO, USA) to yield a final concentration of 50 nM. siRNA transfection was performed according to the siRNA transfection protocol outlined by Dharmacon. 48 h after siRNA treatment, cells were collected for the sigma-2 receptor binding assay. The silencing of PGRMC1 was confirmed by western blot analysis.

Overexpression of PGRMC1

HeLa cells were plated in 100 mm dishes at 1.5×10^6 /dish. 24 h after plating, the cells were transfected with the pCMV6-XL4 vector only or human PGRMC1 cDNA (OriGene Technologies, Inc., Rockville, MD) using a Fugene 6 transfection reagent (Roche Applied Science, Indianapolis, IN). Transfection was carried out according to the guidelines of the manufacturer. For transfection of HeLa cells, 18 μ L of Fugene 6 and 10 μ g human PGRMC1 cDNA were used. After 24 h, the cells were collected for the sigma-2 receptor binding assays. The overexpression of PGRMC1 was confirmed by western blot analysis.

Caspase-3 activation assay

WC-26 induced caspase-3 activation in PGRMC1 siRNA treated HeLa cells was quantified by flow cytometry using a FACScan (BD Biosciences, San Jose, CA, USA) equipped with an air-cooled argon laser. HeLa cells were fixed in 70% ethanol, washed twice with phosphate buffered saline (PBS), and incubated with rabbit anti cleaved-caspase-3 antibody (Cell Signaling Technology, Danvers, MA, USA) at a 1:50 dilution in BD Perm/WashTM buffer (BD Biosciences, San Jose, CA, USA) for 2 h at 37 °C. The cells were washed three times with PBS, and then incubated with phycoerythrin (PE) conjugated donkey anti rabbit IgG (BD Biosciences Pharmingen,) (1:300 dilution) for 1 h at room temperature. After washing with PBS, 5 μ L of 7-amino-actinomycin D (7-AAD, BD Biosciences, San Jose, CA, USA) was added at a concentration of 0.25 μ g of 7-AAD/ 1×10^6 cells. The PE-conjugated antibody was excited at 488 nm, and the emission collected using a 570 nm filter. The 7-AAD was excited at 488 nm, and the emission collected using a 650 nm long-pass filter.

Immunocytochemistry of sigma-2 receptors and PGRMC1

MitoTracker Red CMXRos and ER-TrackerTM Red dye were purchased from Invitrogen Corporation (Carlsbad, CA, USA). HeLa cells were plated in 35 mm diameter glass-bottom dishes at 2×10^5 cells/dish for 24 h. The live cells were incubated with 30 nM SW120 and either the mitochondria tracker, MitoTracker Red CMXRos (20 nM), or the endoplasmic reticulum tracker, ER-TrackerTM Red (500 nM), for 2 h at 37 °C. The labeled cells were then coverslipped and imaged by confocal microscopy as previously described^{21, 22}. The images were acquired using a confocal laser scanning microscope (LSM5 Pascal, Carl Zeiss, Thirwood, NY, USA). SW120 was excited using the 488 nm line from an argon laser, and

the emission collected through a 505-530 nm band-pass filter. MitoTracker and ER-Tracker were excited using the 543 nm line from a helium–neon laser, and the emission collected through a 560 nm long-pass filter.

For imaging PRGMC1, fixed HeLa cells were incubated with goat anti-PGRMC1 antibody at a 1:100 dilutions in the blocking serum overnight at 4 °C. After washing, the cells were incubated with FITC-conjugated donkey anti-goat IgG secondary antibody for 1 h. Rabbit anti-COX IV antibody (Cell signaling, Danvers, MA) and rabbit anti-GRP78 BiP antibody (Abcam, Cambridge, MA) were used for imaging Mito-marker and ER-marker, respectively. The secondary antibody used for imaging Mito-marker and ER-marker is TRITC-conjugated donkey anti-rabbit IgG. The FITC and TRITC antibody labeled cells were then coverslipped and imaged by confocal microscopy.

Supplementary Methods for receptor binding assay, western blot analysis of PGRMC1, real time PCR for PGRMC1 mRNA expression and synthesis of the novel photoaffinity probe **WC-21** and the ¹H NMR and ¹³C NMR spectra for compounds **2, 3, 4, 5** and **WC-21** (Supplementary Figures S10-S19) can be found in the Supplementary Information.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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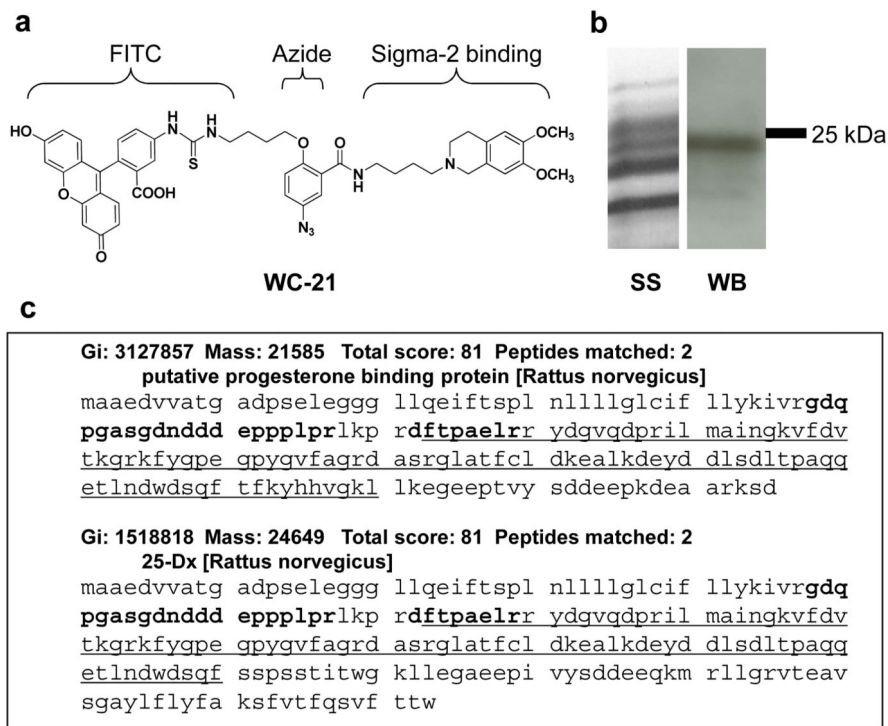


Figure 1. Identification of the putative sigma 2 receptor binding site as PGRMC1 using a novel photoaffinity probe

a, WC-21 contains a FITC group and a photoactive azide (N₃) moiety, and displays high sigma-2 receptor affinity and selectivity; $K_i = 8.7 \pm 1$ nM for sigma-2 receptor and $K_i > 4,000$ nM for sigma-1 receptor. **b**, Following UV cross-linking, ligand-labeled proteins were enriched and separated through SDS-PAGE; WC-21 cross-linked proteins can be visualized using an anti-FITC antibody; SS is silver staining and WB is western blot. **c**, Proteomic studies of the FITC positive band identified two proteins which shared the same genetic name: progesterone receptor membrane component 1 (PGRMC1). The protein structure shows the peptides (bold) detected by mass spectrometry as well as the Cytochrome b5 domain (underlined).

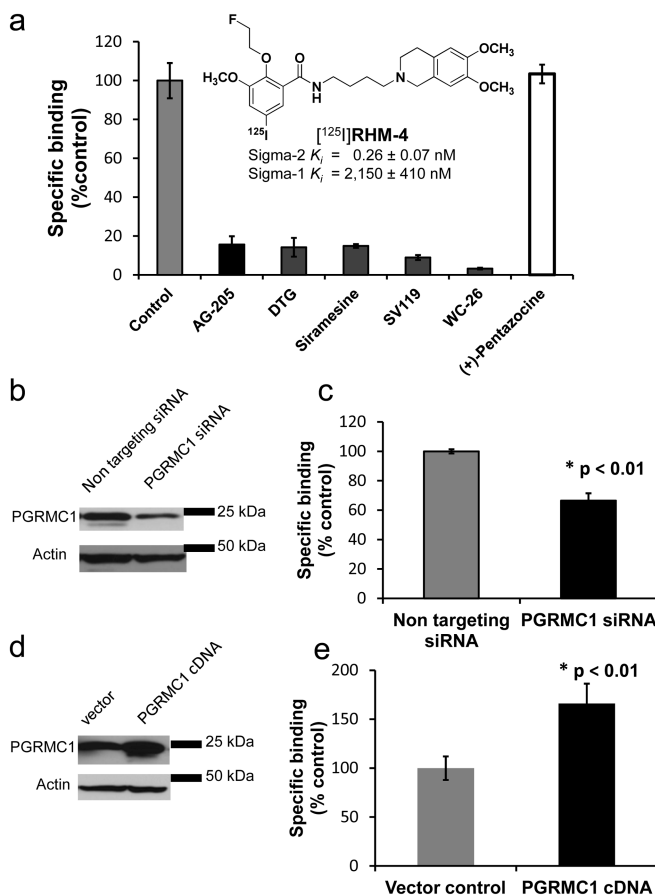


Figure 2. The pharmacological profile of $[^{125}\text{I}]\text{RHM-4}$ and sigma-2 receptor binding activity in PGRMC1 siRNA and cDNA treated HeLa cells

a, Competitive binding studies of $[^{125}\text{I}]\text{RHM-4}$ were carried out with the PGRMC1 ligand AG-205, sigma-2 receptor ligands DTG, WC-26, SV119, and siramesine or with the sigma-1 receptor ligand (+)-pentazocine. $[^{125}\text{I}]\text{RHM-4}$ binding was blocked by AG-205 and the sigma-2 ligands but not by (+)-pentazocine. $n = 2$, sample in triplicate. **b**, Typical western blot confirming PGRMC1 protein expression knockdown in PGRMC1 specific siRNA treated HeLa cells relative to nontargeting siRNA treated controls compared to actin (loading control). **c**, The bar graph shows reduced sigma-2 receptor binding activity in the PGRMC1 knockdown cells; * $p < 0.01$, Student's t-test, $n = 3$. **d**, Typical western blot confirming increased PGRMC1 protein expression in PGRMC1 transfected HeLa cells relative to vector transfected controls. **e**, Bar graph showing increased sigma-2 receptor binding activity in PGRMC1 transfected cells; * $p < 0.01$, Student's t-test, $n = 3$. Error bars in **a**, **c**, **d** represent SEM.

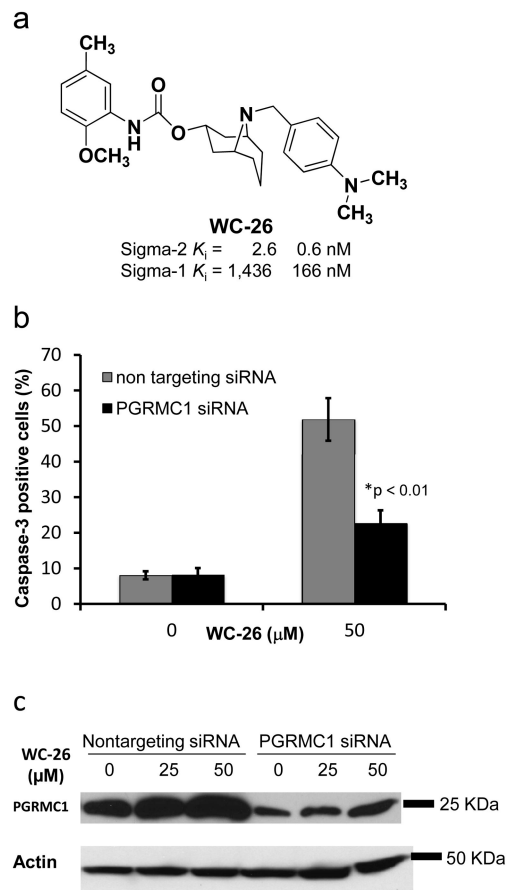


Figure 3. Caspase-3 activation and PGRMC1 upregulation induced by the sigma-2 receptor ligand, WC-26

a, Chemical structure of **WC-26**, a selective sigma-2 receptor ligand. **b**, Knockdown of PGRMC1 in HeLa cells resulted in decreased caspase-3 activation induced by **WC-26**. **c**, **WC-26** stimulated PGRMC1 expression in the same manner as AG-205³⁴, a PGRMC1 ligand; * $p < 0.01$, Student's t-test, $n=3$. Error bars in **b** represent SEM.

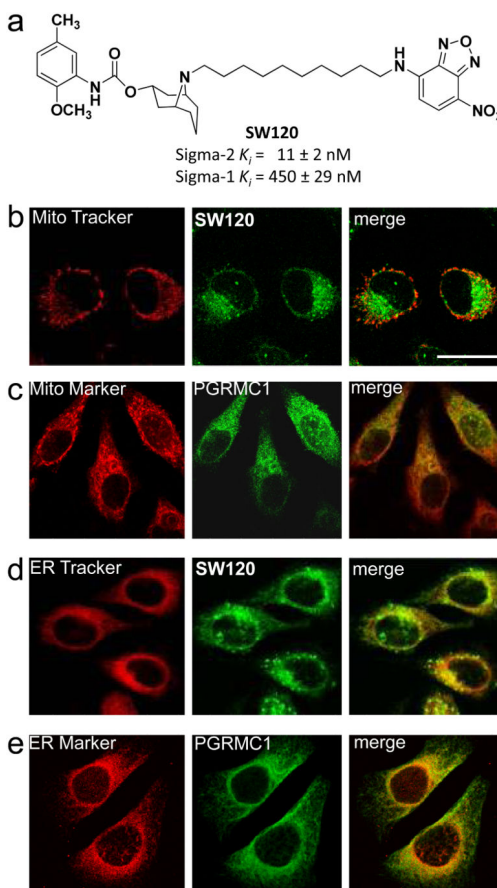


Figure 4. Intracellular co-localization of PGRMC1 and sigma-2 receptors in HeLa cells using confocal microscopy

a, Sigma-2 receptors labeled with **SW120** and PGRMC1 partially colocalized with mitochondria markers (**b, c**) and endoplasmic reticulum markers (**d, e**) in HeLa cells. For imaging sigma-2 receptors, live cells were incubated with **SW120** and either MitoTracker or ER-Tracker and then imaged by confocal microscopy. For imaging PRGMC1, cells were fixed and incubated with goat anti-PGRMC1 antibody followed by a FITC-conjugated secondary antibody (Mito-marker or ER-marker). Antibody stained HeLa cells were then coverslipped and imaged by confocal microscopy. Scale bar represents 20 μm .