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Sphingosine-1-phosphate receptors : tools and molecular pharmacology

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**SPHINGOSINE-1-PHOSPHATE RECEPTORS:
TOOLS AND MOLECULAR PHARMACOLOGY**

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Cover: a red figured kylix, showing Oedipus sitting before the Sphinx of Thebes c. 470 B.C, inv. no. 16541, Museo Gregoriano Etrusco. Photo made by M.Sarri was used with permission of the Vatican Museum.

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**SPHINGOSINE-1-PHOSPHATE RECEPTORS:
TOOLS AND MOLECULAR PHARMACOLOGY**

ACADEMISCH PROEFSCHRIFT

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aan de Universiteit van Amsterdam
op gezag van de Rector Magnificus
Prof.dr. D.C. van den Boom
ten overstaan van een door het college
voor promoties ingestelde commissie,
in het openbaar te verdedigen
in de Agnietenkapel

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Faculteit der Geneeskunde

*“Traveler, there is no path.
Paths are made by walking”*
(Antonio Machade, Spanish poet, 1875-1939)



Abbreviations

aa	amino acid
AA	arachidonic acid
AC	adenylyl cyclase
ADP	adenosine diphosphate
ATP	adenosine triphosphate
$[Ca^{2+}]_i$	intracellular calcium
cAMP	cyclic adenosine monophosphate
CFP	cyan fluorescent protein
CHO	Chinese hamster ovary
DAG	diacyl glycerol
DNA	desoxyribonucleic acid
EC	endothelial cell
ECL	electrogenerated chemi-luminescence
EDG	endothelial differentiation gene
EDTA	ethylenediaminetetraacetic acid
EF-1	elongation factor-1
eNOS	endothelial nitric oxide synthase
ERK	extracellular signal-regulated kinase
FCS	foetal calf serum
GDP	guanine diphosphate
GEF	guanine nucleotide exchange factor
GFP	green fluorescent protein
GIRK	G-protein inwardly rectifying K^+
GPCR	G-protein coupled receptor
GRK	G-protein coupled receptor kinase
GTP	guanine triphosphate
HBSS	Hank's balanced salt solution
HEK	human embryonic kidney
HRP	horse radish peroxidase
HUVEC	human umbilical vein endothelial cell
IBMX	3-isobutyl-1-methylxanthine
IGF	insulin-like growth factor
IP_3	inositol 1,4,5-trisphosphate

Jnk	c-Jun NH ₂ terminal kinase
LDL	low density lipoprotein
LPA	lysophosphatidic acid
MAPK	mitogen-activated protein kinase
MEK	mitogen extracellular kinase
NO	nitric oxide
PBS	phosphate buffered saline
PCR	polymerase chain reaction
PhS1P	phytosphingosine 1-phosphate
PI3K	phosphatidylinositol 3-kinase
PIP2	phosphatidylinositol biphosphate
PKC	protein kinase C
PLC	phospholipase C β
PTX	pertussis toxin
RAEC	rat aortic endothelial cell
RFU	relative fluorescence units
RGS	regulators of G-protein signaling
Rho	ras homology
RNA	ribonucleic acid
ROCK	rho-associated kinase
S1P	sphingosine-1-phosphate
SM	sphingomyelin
TFMPP	m-trifluoromethylphenylpiperazine
UTP	uridine 5'-triphosphate
VSMC	vascular smooth muscle cell





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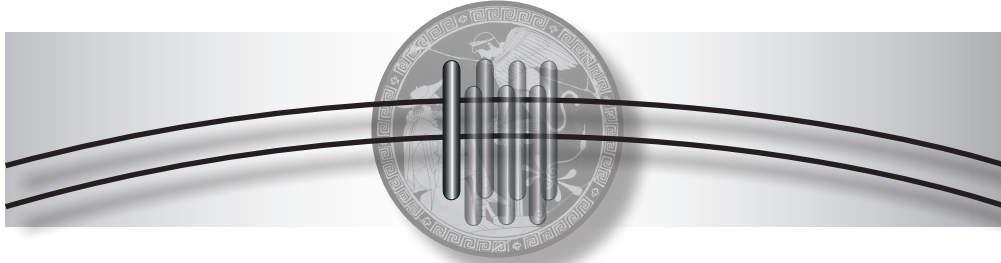
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General introduction

*“What animal is that which
in the morning goes on four feet,
at noon on two,
and in the evening upon three?”*



Introduction to sphingolipids

In 1884, the physician and chemist Johann Ludwig Wilhelm Thudichum (1829-1901) discovered two lipids in the brain characterized by a common aliphatic amino alcohol, for which he used the term “sphingo” (sphinx-like) (Thudichum, 1884) (Figure 1).

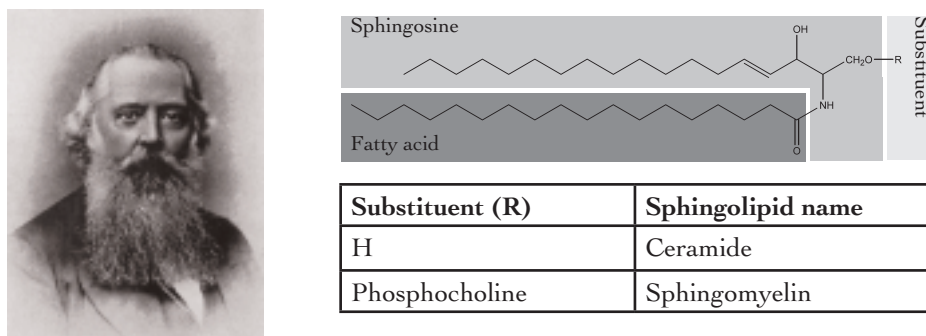


Figure 1. Johann Ludwig Wilhelm Thudichum (1829–1901) and the general sphingolipid structure.

In Greek mythology, the sphinx is a demon of death, destruction and bad luck, depicted as a female with the breast, paws and claws of a lion, sometimes with a snake-tail and eagle or gryphon wings. Sitting on a high rock near Thebes, she would post a riddle to all who came before her: “What animal is that which in the morning goes on four feet, at noon on two, and in the evening upon three?” Anyone who could not correctly answer the riddle, would be strangled by her (“sphingo” in Greek means “to strangle”). Until one day Oedipus came along and answered: “Man, who in childhood creeps on hands and knees, in manhood walks erect, and in old age with the aid of a staff.” Frustrated that her riddle was solved, the sphinx threw herself off the rock and perished. The Thebans were so grateful to Oedipus for ridding them of the monstrous sphinx, that they made him their king.

Sphingosine-backbone containing lipids (later named sphingolipids by Herb Carter (Carter et al., 1947)) were to Thudichum a part of the dangerous riddle of the brain, similar to the riddle posed by the sphinx. It was already clear at that time that sphingolipids had many different effects, but it still took several decades to solve just parts of their riddle.

Metabolism of sphingolipids

Sphingolipids are a class of lipids representing more than 1000 structurally related compounds in total. At first, sphingolipids like sphingomyelin (SM) and glycosphingolipids were regarded as metabolically inactive building blocks of membranes. In 1989, this idea was challenged when upon extracellular stimulation, sphingomyelinase was able to produce ceramide and phosphorylcholine from sphingomyelin (Okazaki et al., 1989). This was quickly supported by many groups which led to the general belief that this pathway, designated the “SM cycle” or “SM-ceramide pathway”, resulted in the release of ceramide as a second messenger. Ceramide could play a role in lipid-mediated signal transduction and influence important processes like apoptosis and cell-cycle arrest (Igarashi, 1997). Nowadays it is known that in this pathway (Figure 2), ceramide can be enzymatically transformed into sphingosine, which is adenosine triphosphate (ATP)-dependently phosphorylated by sphingosine kinase 1 or 2 (SphK1 or SphK2) into sphingosine-1-phosphate (S1P) (Le Stunff et al., 2002; Spiegel & Milstien, 2003; Tani et al., 2007). S1P can be irreversibly degraded to hexadecenal and phosphoethanolamine by S1P lyase (SPL) (Le Stunff et al., 2002; Spiegel & Milstien, 2003; Tani et al., 2007). In addition, S1P can be dephosphorylated either by S1P phosphohydrolase (SPP) or by membrane-associated lipid phosphate phosphohydrolases of which three forms exist (Le Stunff et al., 2002; Spiegel & Milstien, 2003; Tani et al., 2007).

Discovery of sphingolipid receptors

In 1990, Hla and Maciag published the discovery of a transcript which was induced in differentiating human endothelial cells (Hla & Maciag, 1990). Based upon sequence homology with other genes it became evident that this transcript encoded for a protein that belonged to the family of G-protein coupled receptors (GPCRs). Since the ligand for this receptor was unknown, they termed this orphan receptor endothelial differentiation gene 1 (EDG)-1 (Hla & Maciag, 1990). In the following years, several other genes were identified based upon homology to EDG-1 (Okazaki et al., 1993; Masana et al., 1995; Yamaguchi et al., 1996; An et al., 1998; Gräler et al., 1998; Glickman et al., 1999; Im et al., 2000(a); Im et al., 2000(b)), making up a family of in total eight orphan GPCRs, named EDG-1 through EDG-8. Research intensified and resulted eight years after the discovery of EDG-1 in the identification by two independent groups simultaneously of S1P as the ligand for the EDG-1 receptor (Lee et al., 1998; Zondag et al., 1998).

Four additional receptors of the eight-member EDG receptor group turned out to be potently activated by SIP (namely EDG-3,5,6 and 8), while the other three (EDG-2,4 and 7) were activated by a different phospholipid: lyso-phosphatidic acid (LPA). Subsequently, consistent with IUPHAR guidelines, the EDG receptors with high affinity for SIP were renamed to SIP₁₋₅ receptors in chronological order of their discovery, the ones being activated by LPA to LPA₁₋₃ receptors (Chun et al., 2002). Beside the first group of EDG receptors, three more GPCRs have been reported to have high affinity for SIP: GPR3, GPR6 and GPR12 (Uhlenbrock et al., 2002), but it remains to be confirmed whether they indeed are SIP receptors (Alewijns & Michel, 2006). Recently, also two more LPA receptors have been identified, named LPA₄ (previously GPR23) (Noguchi et al., 2003) and LPA₅ (previously GPR92) (Lee et al., 2006).

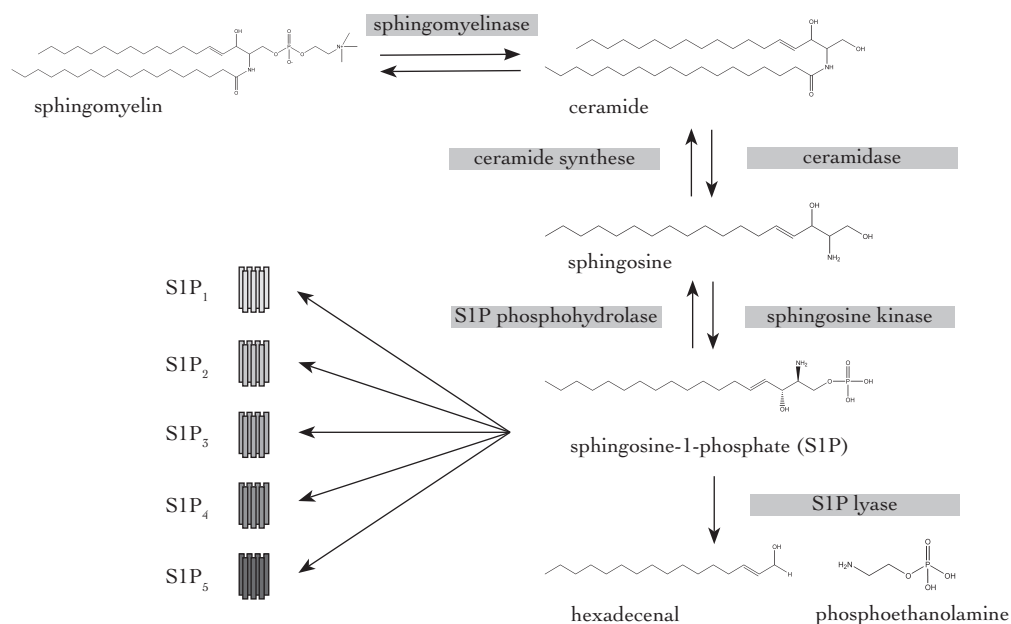


Figure 2. Sphingolipid metabolism.

G-protein coupled receptors and their signaling

As mentioned before, the SIP receptors belong to the GPCR family, which is the largest family of membrane receptors. These receptors couple to intracellular effector systems via the activation of G-proteins thereby transducing signals from the outside to inside of the cell (for a review on GPCRs, see Kristiansen, 2004). GPCRs generally consist of an extracellular N-terminus, an intracellular C-terminus

and seven hydrophobic transmembrane domains connected by hydrophilic loops (Figure 3). Activation of GPCRs accounts for approximately 80% of the signals induced by hormones or neurotransmitters (Birnbaumer et al., 1990). GPCRs are divided into seven classes: Class A (Rhodopsin like), Class B (Secretin like), Class C (Metabotropic glutamate/pheromone), Class D (Fungal pheromone), Class E (cAMP receptors), Ocular albinism proteins and the Frizzled/Smoothed family. SIP receptors belong to the first class, the class A GPCRs.

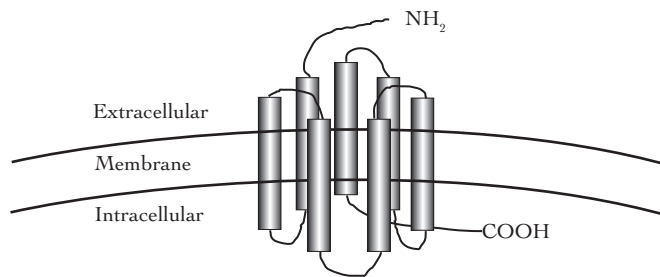


Figure 3. Schematic structure of a GPCR.

As the name already indicates, activation of GPCRs subsequently results in coupling to and activation of heterotrimeric G-proteins which consist of an α -subunit bound to a $\beta\gamma$ -dimer (for a review on heterotrimeric G-proteins see Carbrera-Vera et al., 2003). To date, 28 α , 6 β and 12 γ -subunits have been described (Hermans, 2003; Carbrera-Vera et al., 2003) opening the possibility of a multitude of different combinations. G-proteins are divided into 4 subfamilies, based on the sequence similarity of the α -subunit: G_s (G_s and G_{olf}), G_i (G_{ir} , G_{ic} , G_g , G_{i1-3} , G_o and G_z), G_q (G_q , G_{11} , G_{14} and $G_{15/16}$) and G_{12} (G_{12} and G_{13}). In general, the G_s -subfamily activates adenylyl cyclase (AC) subtype 1-9 (Pierce et al., 2002), resulting in increased levels of cAMP whereas the G_i -subfamily does the opposite by inhibiting activation of AC1, AC5 and AC6 thereby decreasing the formation of cAMP (Kamenetsky et al., 2006). The $\beta\gamma$ -subunits of the G_i -subfamily can, dependent upon the subtype of AC, either activate (AC2, AC4 and AC7) or inhibit (AC1) adenylyl cyclase. In addition, $\beta\gamma$ -subunits can activate many more effectors, amongst others phosphatidylinositol 3-kinase (PI3K) (Pierce et al., 2002), G-protein coupled receptor kinase (GRK)1,2,4, ATP-sensitive K⁺-channels, Ca²⁺-channels, small GTP binding proteins (rho and rac).

G_o , which belongs to the same subfamily as G_i is involved in the inhibition of Ca^{2+} -channels and the activation of G-protein inwardly rectifying K^+ (GIRK)-channels. The G_q -subfamily of G-proteins activates phospholipase C (PLC) β_{1-4} (Pierce et al., 2002), which leads to the hydrolysis of phosphatidylinositol biphosphate (PIP₂) resulting in the formation of diacylglycerol (DAG) and inositol 1,4,5-trisphosphate (IP₃). DAG subsequently activates protein kinase C (PKC). IP₃ regulates increases in intracellular calcium ($[Ca^{2+}]_i$). The G_{12} -subfamily finally regulates ras homology (rho) guanine nucleotide exchange factors (rhoGEFs) (Jiang et al., 1998, Pierce et al., 2002).

GPCR signaling can generally be initiated by molecules, also referred to as agonists, which induce a receptor conformation that increases the affinity of the receptor for the G-protein. In the inactive state, the α -subunit of the heterotrimeric G-protein is bound to guanine diphosphate (GDP), but upon binding by a guanine nucleotide exchange factors (GEFs) such as a agonist-activated receptor, the GDP is rapidly exchanged for guanine triphosphate GTP (Figure 4). This will lead to a reduced affinity of the α -subunit for the $\beta\gamma$ -dimer and cause dissociation of the α -subunit from the $\beta\gamma$ -dimer.

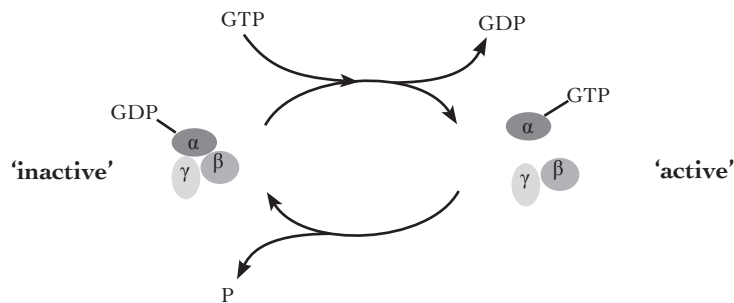


Figure 4. The G-protein activation/inactivation cycle.

For G-proteins belonging to the G_i -subfamily, dissociation of the trimer is blocked by pertussis toxin (PTX), a toxin produced by the bacterium *Bordetella pertussis*. It is the A-subunit of this toxin that is functional in inactivating the G_i -protein by catalysing adenosine diphosphate (ADP) ribosylation of the GDP bound α -subunit. Because of this ability to inactivate G_i (Wong & Rosoff, 1996), PTX has become a widely used pharmacological tool to investigate GPCR signaling (Wong & Rosoff, 1996). After dissociation, both α and $\beta\gamma$ -subunits will diffuse through the membrane

to activate or inhibit a large variety of effector proteins as already described above thereby affecting many cellular effects. GPCR signaling is terminated when the GTP bound to the α -subunit is hydrolyzed to GDP by the intrinsic GTPase activity of the α -subunit. The resulting α -GDP will dissociate from the effector and re-unite with the $\beta\gamma$ -complex to complete the G-protein activation/inactivation cycle. The intrinsic GTPase activity of the α -subunit increases upon coupling to an effector protein but it can also be increased by GTPase-activating proteins such as regulators of G-protein signaling (RGS) proteins (for a review on RGS proteins see Willars, 2006). The RGS proteins thus on one hand accelerate the termination of GPCR signaling while these proteins on the other hand increase the amount of G-proteins available for signaling.

As can be concluded from the description above, compounds that do not change the affinity of the receptor for the G-protein will not initiate signaling. Some of these compounds are also referred to as antagonists if they can antagonize the effects of agonists by occupying the binding-site on the receptor. In the mid-1990s it became clear that GPCRs can exhibit constitutive activity meaning that they can spontaneously adopt conformations that are able to bind to and activate G-proteins (for review see Bond & IJzerman, 2006). This phenomenon led to a general reclassification of ligands as some of these ligands were shown to decrease the constitutive receptor activity thereby acting as inverse agonists, whereas others, that did not affect the constitutive activity, were classified as neutral antagonists.

Originally GPCRs were considered to selectively activate only one specific family of G-proteins. It has now been shown that GPCRs can couple to and activate different G-protein subtypes simultaneously (Woodside, 2002). Momentarily, it is still not fully understood what determines the G-protein specificity of GPCRs. Mutagenesis studies have revealed certain critical domains in the intracellular loops and/or the C-terminus of a GPCR to be important for coupling to and activation of specific G-proteins (Moro et al., 1993; Parent et al., 1996; Havlickova et al., 2002; Gilchrist et al., 2002; Lawson & Wheatley, 2004; Kostenis et al., 2005). However, this has not yet revealed one conserved sequence which determines binding to G_s , G_i , and/or $G_{q/11}$ -proteins specifically. In this context, one domain which has been investigated by many groups is the highly conserved D/ERY motif, situated at the border of the 3rd transmembrane domain and the 2nd intracellular loop (Figure 5). This region contains a triplet of amino acids, Asp/Glu-Arg-Tyr. Conservation of these residues is 86 % for the Asp/Glu residue, 97 % for the Arg residue and 67 % for the Tyr

	TM 3	IL 2	TM 4
Serotonin _{1A} receptor	GQVTCDFIALDVLCCCTSSILHLCAIAL	DRY WAI TDPI DYPRRAA	AALISL TWLIGFLISIPPM
Serotonin _{1B} receptor	GQVVCDFWLSDDITCCTASILHLCVIAL	DRY WAI TDVAEYPKRAA	AVMIALVWVFSISISLPPFF
Serotonin _{1D} receptor	GQILCDIWLSSDITCCTASILHLCVIAL	DRY WAI TDVALEYAGHAA	TMIATVWVAISICISIPPLF
Serotonin _{1E} receptor	GYFLCEVWLSVDMTCCTCSILHLCVIAL	DRY WAI TNAEYAKRAAL	MLT VWTISIFISMPPLE
Serotonin _{1F} receptor	GQVVCDIWLSVDITCCTCSILHLSAIAL	DRY RAITDAVEYPKHAG	IMITIVWISVVFISMPPLE
Serotonin _{2A} receptor	PSKLCAVWIYLDVLFSTASIMHLCAISL	DRY VAI QNPIHSRTKAF	LKIIAVWTVISVIGSMPIPV
Serotonin _{2B} receptor	PLVLCPAWLFLLDVLSTASIMHLCAISV	DRY IAIKKPKIQSRATA	FIKITVWVLLISIGIAIPVPI
Serotonin _{2C} receptor	PRYLCPVWISLDVLFSTASIMHLCAISL	DRY VAIRNPIESRTKAIM	KIAIVWVAISIGVSVPIV
Serotonin ₄ receptor	GEVFCLVRTSLDVLTTASIFHLCCISL	DRY YAICCP LVYPLRI	ALMLGGCWWIPTFISFLPIM
Serotonin _{5A} receptor	GRRLCQLWIACDVLCCCTASINWVTAIAL	DRY WSITRTRLRRKCVSN	VMIALTWALSAVISLAPLL
Serotonin ₆ receptor	ARGLCCLWTA FDMCCSASILNLCISL	DRY LLLSPLRYPLRAL	ALVLGAWSLAALASFLPLL
Serotonin ₇ receptor	GHEFCNVFIAMDVMCCCTASIMTLCVISI	DRY LGITRPLTYGKCM	AKMILSVWLLSASITLPPLE
SIP ₁ receptor	TPAQWFLREGSMFVALSASVFSLLAIAI	ERY ITMLKHLNNFRFL	LLISACWVISLILGGLPIM
SIP ₂ receptor	TPVQWFAREGSFAFITLSASVFSLLAIAI	ERH VAI AKKLYKSCRM	LLLIGASWLSLVLGGPLIL
SIP ₃ receptor	SPTVWFLREGSMFVALGASTCSLLAIAI	ERH LTMIKPYDKRRHR	VFLIGMCWLI AFTL GALPIL
SIP ₄ receptor	APAQWFLREGLLFTALAASTFSLFTAG	ERF ATMVRPVAKTSRV	YGGFGLCWLLAALLGMLPIL
SIP ₅ receptor	SPALWFAREGGVFVALTASVLSLLAIAI	ERS LTMARPVSSRGR	T LAMAAAANGVSLLLGLPAL
LPA ₁ receptor	TVSTWLLRQGLIDTSLTASVANLLAIAI	ERH ITVFRQLHTNRRV	VVVVVIWMTAIVMGAIPSV
LPA ₂ receptor	SLEGWFLRQGLLDTSLTASVATLLAIAV	ERH RSVMAQLHSRGRV	VMLIVGVVVAALGLGLLPAH
LPA ₃ receptor	TVNRWFLRQGLDSSLTASLTNLLVIAV	ERH MSIMRNLTKKR	V TLLILLVVAIAIFMGAVPTL
α_{1B} adrenoceptor	GRIFCDIWAADVLCCTASILSILCAISI	DRY IGVRYPLVTRRKA	LLALLSVVWVLSVISIGPLL
α_{1A} adrenoceptor	GRVFCNIWAADVLCCTASIMGLCIISI	DRY IGVSYPLRYQRR	GLMALLCVWALSLSVISIGPLF
α_{1D} adrenoceptor	GRAFCDVWAADVLCCTASILSILCTISV	DRY VGVRHSLKYERKA	AAIALLWVVALVSVVGPPL

Figure 5. Alignment of the TM3 and IL2 of a selection of Class A GPCRs. The bold letters represent the highly conserved D/ERY motif. Borders of the transmembrane domains (TM) and intracellular loops (IL) are indicated on top.

residue respectively for class A GPCRs (Mirzadegan et al., 2003). Several studies have been performed to investigate the involvement of this domain on G-protein coupling (Ohyama et al., 2002; Capra et al., 2004; Favre et al., 2005; Rovati et al., 2007). Although, this motif is generally believed to play an important role in G-protein coupling and/or activation, the effects of mutations in this motif are not consistent and seem to differ among GPCRs, even within a certain class. For class A GPCRs this has already led to the idea that the receptors belonging to this group can probably be divided into two groups that make use of this motif in a different way (Capra et al., 2004). For one group including e.g. the α_{1B} -adrenergic receptor (Scheer et al., 2000), the β_2 -adrenergic receptor (Rasmussen et al., 1999), the histamine H₂ receptor (Alewijns et al., 2000) and the vasopressin type II receptor (Morin et al., 1998), this motif is involved in keeping the receptor in an inactive ground state. For this group of receptors, mutations in this motif generally retain G-protein binding and increase or induce constitutive receptor activity which is in some cases accompanied by an increase in agonist affinity (Rovati et al., 2007). In the second group, including e.g. the α_{2A} -adrenergic receptor, the M₁ muscarinic acetylcholine receptor and the thromboxane A_{2a} receptor, mutations in the D/ERY motif result in changes in agonist-induced responses, without inducing constitutive receptor activity. So far, many regions in the GPCR structure have been found to be involved in specificity of G-protein coupling, but no overall domain or region has been identified yet.

Since GPCRs can couple to and activate different subtypes of G-proteins simultaneously, activation of a receptor by an agonist may result in the activation of more than one signal transduction pathway. In classical pharmacology, it was generally believed that activation of a receptor by an agonist always produces the same effect, regardless of which agonist is used. The effect can however change in intensity, measured by efficacy or potency. It is now becoming clear that this idea is out of date. This means that a ligand can be a full agonist on one pathway but only a partial agonist or even an antagonist for another pathway (Hermans, 2003) (Figure 6).

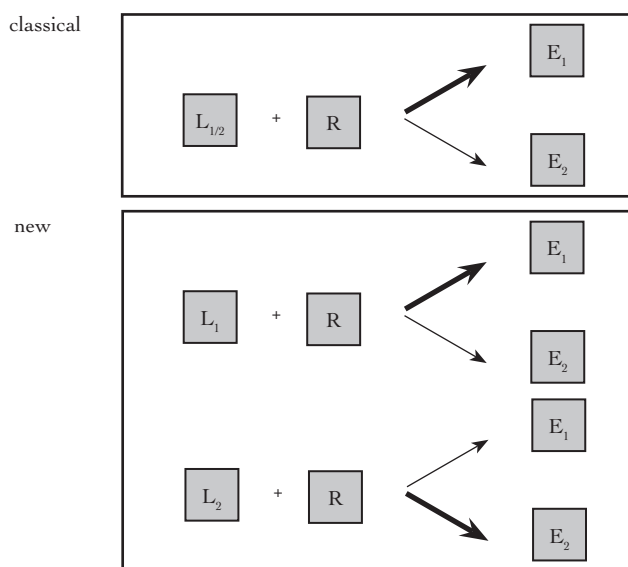


Figure 6. Schematic view of classical pharmacology and ligand-directed signaling (L = Ligand, R = Receptor, E = Effect).

A very illustrative example of ligand-directed signalling comes from the serotonin field. Here it has been shown that *M*-trifluoromethylphenylpiperazine (TFMPP) and bufotenin, both believed to be full agonists for the 5-HT₂ serotonin receptor, indeed had differential effects on different pathways (Stout et al., 2002). TFMPP was a full agonist at IP accumulation but had only partial intrinsic activity at arachidonic acid (AA) release. In contrast, bufotenin was a full agonist at AA release and only a partial agonist at IP accumulation (Stout et al., 2002). Many names for this concept are used currently, for example agonist-specific trafficking, signaling-selective agonism, functional selectivity and ligand-directed signaling (Perez & Karnik, 2005; Urban et al., 2007). Since there is no consensus yet, we will name this concept ligand-directed signaling.

Desensitization of GPCR signaling

When a GPCR is subject to long-term ligand exposure, the risk of over-stimulation exists. This is prevented by a process called desensitization, resulting in a lower response to the same level of agonist exposure and/or a need to increase the concentration of agonist to maintain the same response (for a review, see Pierce et al., 2002). Desensitization can involve many processes and mechanisms, some of which directly affect receptor responsiveness and can be either homologous or heterologous. In the first case, desensitization is a result of activation of the receptor itself, while heterologous desensitization is caused by activation of other receptors or stimuli. One mechanism of desensitization starts with the recruitment of GRKs to an activated receptor. The GRKs subsequently phosphorylate the receptor, thereby enabling the binding of arrestins. To date, six GRK isoforms are identified (GRK₁₋₆) which are all activated by protein kinases, like PKC or c-Src (Pronin & Benovic, 1997; Sarnago et al., 1999). Beside GRKs, other kinases like PKC or mitogen-activated protein kinase (MAPK) can also phosphorylate the receptor (Diviani et al., 1997; Smith et al., 1998). Phosphorylation of the receptor and binding of arrestins to the receptor will uncouple the receptor from the G-protein thereby terminating signaling. A second mechanism important in desensitization of GPCR signaling involves internalization of the receptor, which can be arrestin-dependent or independent (Bünemann & Hosey, 1999). The GPCR is transported from the cell membrane into clathrin-coated pits from where they can be either recycled to the membrane or be degraded, in which case the pits are called lysosomes. The described mechanisms are all rapid desensitization, while there is also long-term desensitization. This involves receptor down-regulation via increased degradation and/or reduced de novo synthesis.

S1P receptor signaling

The signaling pathways of S1P receptors have been extensively investigated and summarized in several excellent reviews (Pyne & Pyne, 2001; Kluk & Hla, 2002; Sanchez & Hla, 2004; Anliker & Chun, 2004 (a&b); Ishii et al., 2004; Young & Van Brocklyn, 2006; Meyer zu Heringdorf & Jakobs, 2007). An overview of the signaling of S1P receptors is described here and schematically depicted in figure 7.

- The SIP_1 receptor is ubiquitously expressed throughout the body with the highest overall expression in the brain (brain > lung \approx spleen > heart/vasculature > kidney) (Chae et al., 2004(b)). The SIP_1 receptor is a unique member of the SIP receptor family as it is the only receptor which selectively couples to a single G-protein family, i.e. G_i -proteins (Lee et al., 1996; Ancellin & Hla, 1999; Windh et al., 1999), to inhibit AC and subsequently decrease cAMP formation (Zondag et al., 1998; Van Brocklyn et al., 1998; Okamoto et al., 1998; Kon et al., 1999).

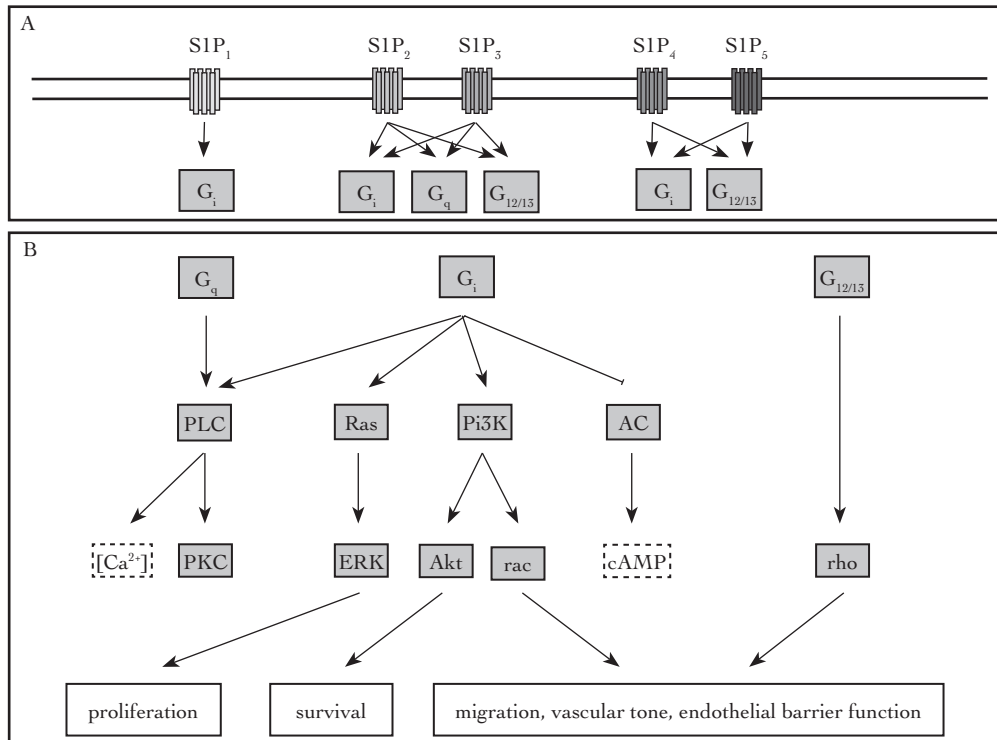


Figure 7. G-protein coupling of SIP receptors (A) and the subsequent signaling pathways of the specific G-proteins (B). (PLC: phospholipase C; Pi3K: phosphatidylinositol 3-kinase; AC: adenylyl cyclase; PKC: protein kinase C; ERK: extracellular signal-regulated kinase).

The SIP_1 receptor has also been shown to activate ERK-1/-2 in a PTX-dependent manner in various systems (Lee et al., 1996; Lee et al., 1998; Zondag et al., 1998; Okamoto et al., 1998; Okamoto et al., 1999). Small, G_i -dependent increases in intracellular calcium ($[Ca^{2+}]_i$) levels have also been observed, most likely via the activation of phospholipase C β (PLC) by the $\beta\gamma$ -subunit of G_i -proteins, since the SIP_1 receptor does not couple to G_q -proteins (Zondag et al., 1998, Okamoto et al., 1998; Van Brocklyn et al., 1998; Kon et al., 1999, Ancellin & Hla, 1999; Lee et al., 1999; Okamoto et al., 1999, An et al., 1999). Support for the activation of

PLC via the $\beta\gamma$ -subunit of G_i is the finding that preincubation with U73122 (a PLC inhibitor) or overexpression of $G_{\alpha t}$ (a $\beta\gamma$ -subunit scavenger) can indeed block the SIP_1 -mediated $[Ca^{2+}]_i$ increases (Okamoto et al., 1998). However, conflicting results were reported with regard to whether these $[Ca^{2+}]_i$ increases are indeed PLC mediated (Kon et al., 1999; Okamoto et al., 1999). Recently, experiments in smooth muscle cells using silencing of SIP_1 or specific activation of the SIP_1 receptor with the agonist SEW2871 showed that the SIP_1 receptor does not activate PLC β (Hu et al., 2006 (b)). With regard to the small GTPases rho and rac, the SIP_1 receptor can activate rac, but the data on rho are inconclusive (Kluk & Hla, 2002). In human umbilical vein endothelial cells (HUVECs), activation of rho has been shown (Lee et al., 1998; Paik et al., 2001) whereas in smooth muscle cells the SIP_1 -mediated activation of rho was not confirmed (Hu et al., 2006(b)). In addition, Chinese hamster ovary (CHO) cells either showed no effect (Okamoto et al., 2000) or an activation of rho by the SIP_1 receptor (Paik et al., 2001).

- Expression of the SIP_2 receptor is like the SIP_1 receptor also seen in many different cell types where it can activate G_i , G_q and $G_{12/13}$ -proteins (Ancellin & Hla, 1999; Windh et al., 1999; An et al., 1999). Via activation of G_q -proteins, stimulation of the SIP_2 receptor leads to increases in $[Ca^{2+}]_i$ via activation of PLC and subsequent IP_3 formation (Kon et al., 1999; Ancellin & Hla., 1999 An et al., 1999; Gonda et al., 1999; Kupperman et al., 1999). The SIP_2 receptor can also activate ERK-1/-2, c-Jun NH₂-terminal kinase (Jnk) and p38 MAPK (An et al., 1997; Okamoto et al., 1999; Gonda et al., 1999; Kupperman et al., 2000; An et al., 2000). It has recently been shown that the combination of SIP_2 and SIP_3 receptor activation is required for activation of Akt (Means et al., 2007). The SIP_2 receptor can activate one of the small GTPases, namely rho (Okamoto et al., 2000; Gonda et al., 1999), but it can inhibit the activation of rac and lower its basal activity (Okamoto et al., 2000). This most likely occurs via the activation of phosphatase and tensin homolog (PTEN), which antagonizes the PI3-Akt pathways, resulting in a down-regulation of rac and Akt activity (Sanchez et al., 2007). Although the SIP_2 receptor through G_i can regulate AC (Kon et al., 1999; Gonda et al., 1999) it seems to increase rather than decrease cAMP levels in CHO cells. Interestingly, these studies suggest that this is not caused via the activation of G_s (Windh et al., 1999) and also not via $[Ca^{2+}]_i$ (Kon et al., 1999). Probably, SIP_2 receptor activation leads to an inhibition of phosphodiesterase activity, the enzymes which normally break down cAMP, consequently resulting in an increase in cAMP levels.

- The $S1P_3$ receptor is expressed in the same tissues as mentioned for the $S1P_1$ and $S1P_2$ receptor with minor differences in expression level. Like the $S1P_2$ receptor, it couples to G_r , G_q and $G_{12/13}$ -proteins (Ancellin & Hla, 1999; Windh et al., 1999). Activation of the $S1P_3$ receptor regulates AC. However, this effect has been shown to be either an inhibition in a PTX-dependent manner or a stimulation of AC (Kon et al., 1999; Okamoto et al., 1999). The $S1P_3$ receptor is also linked to the PLC/IP₃/[Ca²⁺]_i pathway (Kon et al., 1999; Okamoto et al., 1999; An et al., 1999; Ishii et al., 2001), suggested to be even more efficiently than the $S1P_2$ receptor (Ancellin et al., 1999). The increase in [Ca²⁺]_i has been shown to be affected both by G_i and G_q -coupled pathways (Sato et al., 1999). Activation of the MAPK cascade has been shown via detection of ERK-1/-2 phosphorylation and c-Jun mRNA and protein expression (An et al., 1997; Okamoto et al., 1999; An et al., 2000). $S1P_3$ activation also results in activation of rho and rac (Okamoto et al., 2000; Liu et al., 2000; Paik et al., 2001). In mouse embryo fibroblasts however, no activation of rac was measured upon S1P stimulation (Ishii et al., 2001).

- The $S1P_4$ receptor shows a very restricted expression pattern, being found only in the lung (Jolly et al., 2002) and lymphoid tissues (Gräler et al., 1998). The affinity of S1P for the $S1P_4$ receptor is lower than for the other four receptors, whereas the affinity for phyto-S1P is higher. It couples to G_i and $G_{12/13}$ -proteins, but not to G_s or G_q -proteins (Gräler et al., 2003). S1P-stimulation of the $S1P_4$ receptor results in activation of ERK-1/-2, PLC and elevations of [Ca²⁺]_i (Van Brocklyn et al., 2000; Yamazaki et al., 2000). PLC is activated through the G_i -pathway since it is blocked by PTX treatment (Gräler et al., 2003). The $S1P_4$ receptor can activate rho, but the effect on rac still has to be determined (Gräler et al., 2003).

- The $S1P_5$ receptor expression is limited to the white matter of the brain (Im et al., 2000(b); Terai et al., 2003), skin and spleen (Im et al., 2000(b); Ishii et al., 2001). The $S1P_5$ receptor activates G_i and $G_{12/13}$, but not G_s and G_q -proteins (Im et al., 2000(b); Malek et al., 2001). Activation of the $S1P_5$ receptor inhibits AC, which is PTX-sensitive (Im et al., 2000(b); Malek et al., 2001). Interestingly, the inhibition of AC is already evident in the absence of S1P, suggesting constitutive activity of this receptor subtype (Niedernberg et al., 2003). In this way, inhibition of ERK activity, but not rho occurs (Niedernberg et al., 2003). $S1P_5$ can also activate Jnk and inhibit ERK-1/-2 activation, but not via G_i -activation (Malek et al., 2001). It has been shown that via G_i , $S1P_5$ can activate PI3K and it can activate rho kinase via $G_{12/13}$ (Li et al., 2003; Novgorodov et al., 2007). $S1P_5$ can also activate Akt (Jaillard et al., 2005)

Important structures in S1P receptors

Among GPCRs, entire motifs, single amino acids or general properties of amino acids have been conserved (Figure 5). This indicates their importance in specific receptor functions, e.g. ligand binding or G-protein coupling. 3D structures of many GPCRs, including the S1P receptors are currently only computational predictions from the DNA sequence, based upon the crystal structure of the bovine rhodopsin receptor provided by Palczewski et al. in 2000. These predicted structures can provide information about important amino acids for binding the ligand, and important domains for coupling to G-proteins.

For the S1P receptors, most research has been done on the S1P₁ receptor, which revealed some residues that are critical for interaction with its ligand, S1P (Fujiwara et al., 2007). Based upon these predictions, mutant receptors can be made and studied for verification the models. These studies indeed confirmed some important amino acids for interaction with S1P. Arg¹²⁰ and Arg²⁹² form two ion pairs with the anionic amine group of S1P, while Glu¹²¹ forms one ion pair with the amine group of S1P (Parrill et al., 2000). These sites are also predicted to be important for interaction with SEW2871, which strengthens the prediction (Fujiwara et al., 2007). Another important region for S1P binding are 2 tyrosine residues in the N-terminal tail of the S1P₁ receptor. It has been shown that O-sulfation of these two amino acids is required for high-affinity S1P binding (Fieger et al., 2005). This seems to be a S1P₁ receptor specific feature, since the other S1P receptors do not contain a tyrosine with the required neighboring aspartic acid and only the S1P₂ receptor contains a tyrosine flanked by a glutamic acid, but this is not a site for sulfation. For the S1P₄ receptor, Trp²⁹¹, Glu²⁸⁴ and Thr¹²⁷ are important for S1P binding according to computational predictions (Vaidehi et al., 2002). This has been opposed by a model by Holdsworth et al. in 2004 since all mentioned residues are directed towards extracellular loops, and are thus unlikely to interact with S1P. They show an important function of Glu¹²² which once mutated to Gln switches the S1P₄ receptor from S1P to LPA responsiveness (Holdsworth et al., 2004).

A very interesting motif in S1P receptors which has not yet been studied is the D/ERY motif. A striking difference can be found when comparing this motif to the G-protein coupling of these receptors. The S1P₁ receptor, which only couples to G_{i/o}-proteins, contains an ERY motif whereas the other S1P receptor subtypes contain an ERH/S/F motif and couple to G_q, G_{12/13}-proteins (S1P_{4,5}) and to G_i-proteins (S1P_{2,3}) (Figure 7).

Desensitization of S1P receptors

Desensitization of S1P receptors has not been that intensively studied yet, even though it seems to be the functional mechanism of a new drug, FTY720 (Matloubian et al., 2004). Shortly after S1P was found to be the ligand for the EDG-1 receptor, a paper appeared which showed S1P-induced trafficking of this receptor, C-terminally tagged with GFP (Liu et al., 1999). S1P was able to internalize the EDG-1-GFP receptor and after S1P wash out, the receptor was recycled to the membrane. Internalization of a C-terminal GFP-tagged S1P₁ receptor was also seen after insulin-like growth factor (IGF)-1 and IGF-2 stimulation, which could be blocked by dimethylsphingosine, showing the involvement of sphingosine kinase (El-Shewy et al., 2006). A loss of sensitivity to S1P has been shown after 10 minutes of S1P or FTY720-P (the phosphorylated, active form of FTY720) stimulation of renal mesangial cells (Xin et al., 2004). This loss of sensitivity could also be induced heterologously by stimulation with ATP, UTP or adenosine. When ATP or UTP were used, increased cAMP levels, PKC activation and another staurosporine-sensitive protein kinase were involved. The S1P₁ receptor internalized to endocytic vesicles and lysosomes from which it could be recycled to the membrane (Liu et al., 1999). Internalization of the receptor required GRK2 activation and subsequently two different pathways have been shown for two different ligands (Oo et al., 2007). The S1P₁ receptor will be recycled to the membrane after S1P-induced internalization, but will be broken down after FTY720-P-induced internalization (Oo et al., 2007). This has been ascribed to rapid polyubiquitinylation of the S1P₁ receptor by FTY720-P, but not by S1P (Oo et al., 2007). In contrast to the S1P₁ receptor, the S1P₃ receptor was shown to internalize without being phosphorylated (Rutherford et al., 2005).

S1P signaling in the vasculature

The two major cell types in the vasculature are endothelial cells (ECs) and vascular smooth muscle cells (VSMCs). The S1P_{1,3} receptors are differentially expressed on each cell type (Michel et al., 2007) (Table 1). Interestingly, since both cell types express all enzymes necessary to produce S1P (Yatomi et al., 2001), S1P can be formed locally in the vessel wall by endothelial as well as vascular smooth muscle cells (Yatomi et al., 2001; Alewijnse et al., 2004). However, S1P is also a major constituent in blood and plasma in concentrations of 200-900 nM (Murata et al., 2000). It can be released from activated platelets (Yatomi et al., 2000) and is a

major constituent of erythrocytes (Pappu et al., 2007), so it may also reach vascular cells via the bloodstream. SIP is a constituent of HDL (Murata et al., 2000), but it is unknown if it can activate its receptors in the vessel wall in this form. Most of the plasma SIP is bound to proteins and hence unlikely to be available for receptor interaction. In blood, the free SIP concentration is unknown but most likely lower than in plasma. The effect of SIP on both ECs and VSMCs and the total vessel is described below and depicted in figure 8.

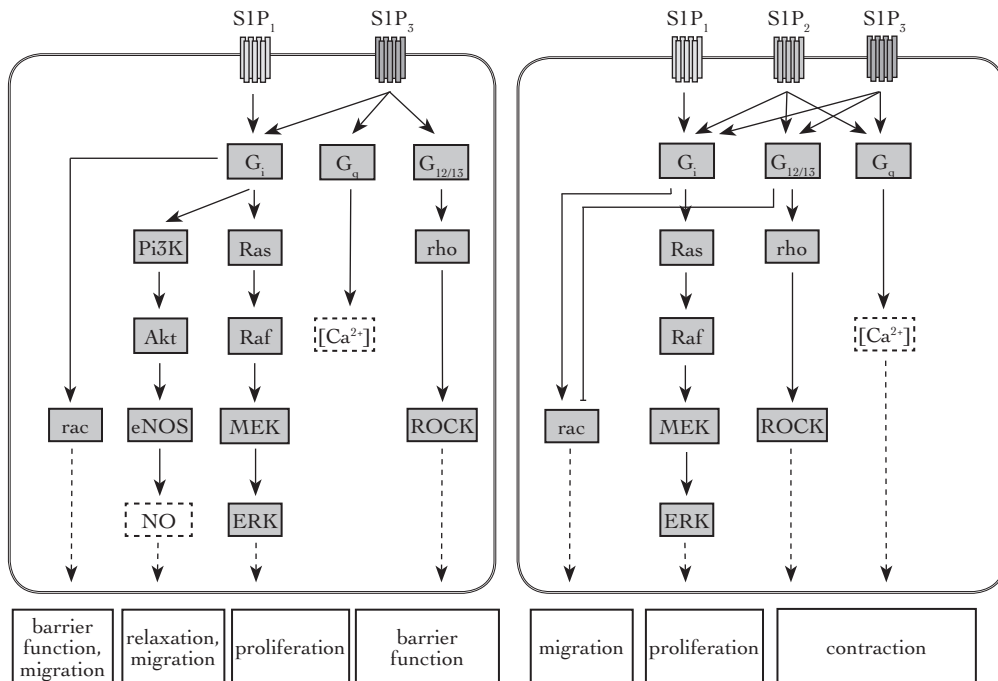


Figure 8. S1P receptor signaling in endothelial cells (left panel) and vascular smooth muscle cells (right panel) (Pi3K: phosphatidylinositol 3-kinase; eNOS: endothelial nitric oxide synthase; NO: nitric oxide; ERK: extracellular signal-regulated kinase; ROCK: rho-associated kinase; MEK: mitogen extracellular kinase).

Endothelial cells

The S1P₁ and S1P₃ receptors are most abundantly expressed in the endothelium (Panetti, 2002) (Table 1). On ECs, one of the effects of S1P is maintaining barrier function. Although the underlying mechanism is not clear yet, it involves activation of the small GTPase rac (McVerry & Garcia, 2004). Barrier function requires reorganization of actin which occurs via Tiam1/rac1 activation via S1P₁ (Singleton et al., 2005). The S1P₃ receptor seems to have the opposite effect, thus decreasing rather than increasing barrier function (Singleton et al., 2006). Besides

the effect on barrier function, S1P also causes migration of ECs via rearrangement of the cytoskeleton through activation of rac1. Activation of rho is also involved in migration of ECs. Inhibition of PI3-K decreases migration induced by S1P (Tamama & Okajima, 2002), but this does not affect S1P-mediated rac1 activation (Tamama & Okajima, 2002; Gonzalez et al., 2006). S1P-induced activation of PI3-K can also lead to the activation of endothelial nitric oxide synthase (eNOS) through Akt, resulting in the production of the vasodilator NO and subsequent relaxation of a vessel. A final effect of S1P on ECs is proliferation in a G_i -sensitive way via both S1P₁ and S1P₃ receptors. Rho activation plays a key role in this process, whereas ERK activation is not involved (Tamama & Okajima, 2002; Panetti, 2002).

Vascular smooth muscle cells

VSMCs generally express high levels of the S1P₂ receptor compared to the S1P₁ and S1P₃ receptor (Table 1). S1P affects migration of VSMCs, which is stimulated via S1P₁ and S1P₃ receptor activation, but inhibited via S1P₂ receptor activation (Waeber et al., 2004). Migration occurs via the activation or inhibition of rac1. The effect of S1P on VSMC migration is thus depending on the S1P receptor expression profile. S1P can also induce proliferation of VSMCs via all three receptors. In case of S1P₁ activation, the PI3-K pathway is involved. Through S1P-mediated activation of G_q , $[Ca^{2+}]_i$ will rise via the PLC pathway, which results in contraction of the vessel.

Vascular tone

The effect of S1P on vascular tone has been investigated by several groups, but conflicting results have been found, possibly due to differences in species, methods, vascular bed, vessel function and S1P concentrations used. S1P-induced constriction has mainly been found in smaller vessels (Bischoff et al., 2000(a); Bischoff et al., 2000(b); Sugiyama et al., 2000; Bischoff et al., 2001; Tosaka et al., 2001; Coussin et al., 2002; Ohmori et al., 2003; Salomone et al., 2003; Hedemann et al., 2004; Czyborra et al., 2006), while relaxation was observed in both smaller and larger vessels (Mogami et al., 1999; Dantas et al., 2003; Tölle et al., 2005; Roviezzo et al., 2006).

The effect of S1P on vascular tone is most likely the result of a balance between the effects on ECs and VSMCs. This balance may amongst others depend on the relative expression of the S1P_{1,3} receptors on both cell types.

Table 1. Vascular SIP receptor expression in the various cell types present in these vessels.

Whole artery			
	Rat	SIP ₁ , SIP ₂ , SIP ₃ , SIP ₅	Salomone et al., 2003
	Gerbil	SIP ₁ , SIP ₂ , SIP ₃	Scherer et al., 2006
	Human	SIP ₁ ≈ SIP ₃ > SIP ₂	Mazurais et al., 2002
Arterial Smooth Muscle			
	Mouse	SIP ₁	Chae et al., 2004b
	Rat	SIP ₂ > SIP ₃ > SIP ₁ ; SIP ₄ absent	Tamama et al., 2001
		SIP ₂ > SIP ₃ ; SIP ₁ , SIP ₄ and SIP ₅ absent	Ryu et al., 2002
		SIP ₂ ≈ SIP ₃ > SIP ₁	Coussin et al., 2002
		SIP ₃ , SIP ₄ and SIP ₅ absent	Yamaguchi et al., 2003
	Human	SIP ₂ > SIP ₃ > SIP ₁	Tamama et al., 2001
		SIP ₃ ; SIP ₂ in aorta but not coronary; SIP ₁ absent	Mazurais et al., 2002
		SIP ₂ > SIP ₃ ; SIP ₁ , SIP ₄ and SIP ₅ absent	Ryu et al., 2002
		SIP ₂ > SIP ₁ > SIP ₃ > SIP ₅ ; SIP ₄ absent	Damirin et al., 2005
Arterail endothelium			
	Mouse	SIP ₁	Chae et al., 2004
	Rat	SIP ₁ > SIP ₃ ; SIP ₂ absent	Tamama et al., 2001
	Bovine	SIP ₁ > SIP ₃ > SIP ₂	Wang et al., 1999
	Human	SIP ₁ > SIP ₃ > SIP ₂	Tamama et al., 2001
		SIP ₁ and SIP ₃ ; SIP ₂ absent	Kimura et al., 2000
		SIP ₁ ; SIP ₃ absent	Mazurais et al., 2002
Venous smooth muscle			
	Human	SIP ₁	Hla & Maciag, 1990
Venous endothelium			
	Human	SIP ₁	Hla & Maciag, 1990
		SIP ₁ > SIP ₃ ; SIP ₂ absent	Lee et al., 1999
		SIP ₁ > SIP ₃ > SIP ₂	Wang et al., 1999
		SIP ₁ > SIP ₃ ; SIP ₂ , SIP ₄ and SIP ₅ absent	Ryu et al., 2002

S1P receptors in vascular disease

Receptors that are important in development and/or functioning of any tissue are often also involved in diseases. This also holds true for the S1P receptors. Even though S1P can mediate wound healing for instance in the vessel by stimulating angiogenesis, it can become atherogenic and thrombogenic and influence cardiovascular diseases (Gardell et al., 2006). S1P is not only involved in vessel formation during the embryonic state, as was found in studies using S1P₁ receptor knockout mice, but also after birth. Therefore, S1P will have an effect on diseases in which angiogenesis (new vessel formation) plays a role, such as cancer, psoriasis and rheumatoid arthritis (Pyne & Pyne, 2000).

S1P also influences proliferation, which plays a role in atherosclerosis. In this disease, thrombus formation is induced by S1P released from thrombin-stimulated platelets (Gardell et al., 2006). Deposition of Ox-LDL in the thrombus results in activation of sphingomyelinase and production of S1P (Augé et al., 1999). S1P causes a decrease of the endothelial barrier permeability via G_i-protein activation through the S1P₁ and S1P₃ receptors. (Siess, 2002; McVerry & Garcia, 2004). S1P also stimulates migration and proliferation of smooth muscle cells into the atherosclerotic plaques, resulting in the occlusion of the vessel. More effects of S1P through its receptors are an acute and transient decrease in heart rate through activation of GIRK channels (Guo et al., 1999; Forrest et al., 2004; Karliner, 2004), decreases coronary artery blood flow (Ohmori et al., 2003; Karliner, 2004) and an initial increase in blood pressure in whole mice through increases of [Ca²⁺]_i due to release of Ca²⁺ from internal stores, followed by cytoskeletal re-arrangement (Forrest et al., 2004). The opposing effects of S1P on blood pressure make it hard to make a clear statement whether S1P is beneficial or has a negative effect on blood pressure.

Tools to study S1P receptors and their signaling

Knockout models

There are many ways to study the physiological role of newly discovered receptors. One of them is to generate receptor-null mice, also called knockout mice. The first report on S1P receptor knockout mice came in 2000, where Liu et al. described the essence of the S1P₁ receptor in vascular maturation (Liu et al., 2000). Disruption of the corresponding gene resulted in lethality at embryonic day E12.5 to E14.5. The cause was found to be a defective vascular maturation, since the VSMCs failed

to surround the nascent endothelial tubes even though they were in the vicinity. Conditional SIP_1 receptor knockouts in EC or VSMC showed that vessel coverage by VSMCs is controlled by the SIP_1 receptor in ECs (Allende et al., 2003).

Double or triple SIP receptor knockout mice indicated that besides the SIP_1 receptor, also the SIP_2 and SIP_3 receptor play an important role in vascular development (Kono et al., 2004). Even though the SIP_2 or SIP_3 single knockout mice were viable, and the SIP_1/SIP_3 double knockout did not alter the phenotype of the SIP_1 single null embryo, knocking out the SIP_2 receptor together with the SIP_1 receptor increased the severity of the phenotype of the single SIP_1 receptor knockout (Kono et al., 2004). These embryos died two days earlier in their embryonic phase than the SIP_1 single knockout embryos due to bleeding. The triple receptor knockout showed a similar or slightly more severe result than the double knockout (Kono et al., 2004). This indicated that during embryonic angiogenesis, these receptors function coordinately. The SIP_2 receptor single knockout mice had spontaneous seizures during 3-7 weeks of age, which occasionally led to death (Yang et al., 2002). The SIP_3 receptor single knockout mice had no obvious phenotype (Yang et al., 2002).

Besides vascular defects, SIP_1 knockout embryos also show abnormal limb development, which was caused by a loss of SIP_1 in endothelial cells (Chae et al., 2004 (a)). Although SIP_2 knockout mice do not have an obvious phenotype, they appear to have lower mesenteric and renal resistance (Lorenz et al., 2007). Another effect of knocking out the SIP_2 receptor was recently shown by Kono et al. in 2007, who found that these mice were deaf by one month of age, possibly due to vascular defects in the stria vascularis.

The SIP_3 null mice were successfully used to show that the observed slower heart rate (bradycardia) after administration of FTY720 was mediated via the SIP_3 receptor. Additionally, SIP receptor agonist-induced lymphopenia (inhibition of lymphocyte recirculation) was not altered in these mice, indicating that the SIP_3 receptor is most likely not involved in this process (Sanna et al., 2004).

siRNA

Short interfering (si) RNA is a technique in which a piece of a double stranded (ds) RNA fragment is brought into the cell via for example a lentiviral system. Inside the cell, the dsRNA is cleaved by a dicer resulting in siRNA pieces of 21-23 nucleotides. A single strand is incorporated into the RNA-induced silencing complex which can detect and destroy the target mRNA, thereby interrupting protein synthesis.

In case of SIP receptors, this relatively new method has been applied to study receptor function by specific knockdown of a receptor. In endothelial cells the importance of the SIP₁ receptor for actin reorganization was shown and also that the SIP₁ receptor may protect these cells from necrosis (Krump-Konvalinkova et al., 2005). Another study showed that together with the SIP₃ receptor, the SIP₁ receptor mediates the inflammatory responses in these cells (Lin et al., 2007). In vascular smooth muscle cells, siRNA proved that contraction is mediated exclusively via SIP₂ receptors (Hu et al., 2006(a)). For the SIP₅ receptor, usage of this method showed that in oligo-dendrocyte precursors cells migration was inhibited via activation of rho and ROCK through G_{α12/13} (Novgorodov et al., 2007).

Tagged receptors

When binding studies are unavailable, receptors can be identified by using a molecularly modified receptor, e.g. in receptor regulation studies. It is possible to add a tag to the receptor, e.g. HA-tag or green fluorescent protein (GFP), which is easily detectable by fluorescence microscopy or Western blot for example. As described before using tagged receptors, SIP has been shown to internalize the SIP₁ and SIP₃ receptor. When different ligands became available, it was also shown that receptor fate is dependent on which ligand is used. SIP₁-GFP internalized after stimulation with the SIP₁ receptor selective agonist SEW2871 and SIP will be recycled to the membrane, where FTY720-P stimulation leads to internalization followed by degradation (Jo et al., 2005). The same group showed later that this degradation was caused by poly-ubiquitination of the receptor, induced by one agonist, but not by other (Gonzalez-Cabrera et al., 2007).

It is important to realize that adding large tags to receptors can influence signaling and/or regulation of the receptor. This is illustrated in a study on the SIP₄ receptor where several different constructs were made. Addition of a C-terminal GFP tag resulted in no membrane localization of the receptor in contrast to their SIP₁-GFP receptor. The same group also constructed a N-terminally FLAG-tagged SIP₄

receptor, which again did not show membrane localization. Interestingly, a SIP_4 receptor with both tags was expressed at the cell surface. The SIP_4 receptor with a C-terminal FLAG tag did come to the cells surface, in contrast to the N-terminally FLAG tagged receptor. The same discrepancy was seen when a HA tag (N or C-terminal) was used instead of the FLAG tag. Subsequent experiments showed that SIP induces cell migration through the membrane expressed SIP_4 receptor (Kohno et al., 2003).

Specific antibodies

Antibodies directed against a specific receptor or against a ligand may be used to investigate receptor localization, visualize endogenous receptors or even to study regulation of receptors and/or receptor signaling. A SIP_1 receptor monoclonal antibody has been described which binds to the human and/or mouse SIP_1 receptor in membranes (Goetzl et al., 2004). Binding to the other SIP receptor subtypes has not been studied. It does not inhibit binding of SIP to the receptor, but it inhibits T cell chemotactic responses and suppression of T cell chemotaxis to chemokines, both induced by SIP. Interestingly, another monoclonal antibody, against SIP itself has been shown to reduce tumor progression or even eliminate tumors (Visentin et al., 2006). It also blocks EC migration, which is involved in blood vessel formation and was able to stop the protective effect of SIP against apoptosis of tumor cells. This could well indicate that SIP itself is a potential target in cancer therapy.

On the receptor side, not many antibodies exist and most of them that do exist are not very specific.

Transfection studies

An often used method to investigate receptor signaling are stable cell lines overexpressing receptors. Since they overexpress one specific receptor, signaling differences between the original cells and the stable cells are then assigned to the overexpressed receptor. A better control than using the original cells is using mock transfected cells. Mock transfected cells are made the same way as the stable cells, but are transfected with only the empty vector (plasmid) and not the plasmid containing the target gene DNA. This then rules out possible effects of the plasmid like antibiotic resistance.

A disadvantage of overexpression systems is the possibility that receptors will activate G-proteins they would normally not activate in native cells. Another thing

to keep in mind when interpreting the results obtained with stable cell lines is endogenous (native) receptors. Even when overexpressing the receptor of interest, the endogenous receptors are still present and functional in the cells. They can thus still cause part of the observed effect(s). Therefore it is preferable to create stable cell lines from cells that have no or as little as possible receptors expressed that have affinity for the used ligand.

The most used cells in S1P receptor research are CHO, human embryonic kidney (HEK) 293 cells and HUVECs. However, endogenous S1P receptor expression of all of these cells is not very well investigated yet. The few reports that do exist show that in CHO-K1 cells, endogenous mRNA expression of the S1P₁, S1P₂ and S1P₄ receptor was detected (Holdsworth et al., 2005). In HEK293 cells, no endogenous expression of the S1P₁ receptor was found (Lee et al., 1998). The primary cell line HUVECs, like other ECs (see above), most likely express the S1P₁ and S1P₃ receptor, but this is based on functional data, not on direct measurements of receptor presence itself (Butler et al., 2004).

Ligands

Pharmacological S1P receptor studies have been hampered due to a scarce availability of ligands (Figure 9). The few that are available are described below.

Agonists: The novel immunosuppressive drug FTY720, a chemical modification of myriocin (ISP-1) from the fungus *Isaria sinclairii*, already was described in 1996 (Suzuki et al., 1996) but it was only in 2002 when it was found to exert its effect via S1P receptors (Brinkmann et al., 2002). They showed that only the phosphorylated form (FTY720-P), which resembles the structure of S1P, activates S1P_{1,3-5} receptors. Recently, clinical trials using FTY720 revealed an effect on heart rate. Administration of this compound induced bradycardia in healthy subjects (Schmouder et al., 2006). This FTY720-induced reduction in heart rate was also shown in mice, but was absent in S1P₃ receptor knockout mice (Forrest, 2004). VPC23153 and VPC24191 are synthetic agonists for the S1P₁ and S1P₃ receptor. Pytosphingosine 1-phosphate (PhS1P) was found as an agonist to S1P₄, with much higher affinity than for the S1P₅ receptor (Candelore et al., 2002). The first report on the S1P₁ selective agonist SEW2871 appeared in 2005 (Bolick et al., 2005).

Antagonists: The first S1P receptor antagonist was suramin, an antagonist for the S1P₃ receptor, but suramin also blocks other receptor ligand interactions (Ancellin & Hla, 1999) and has receptor-independent effects on G-protein function

(Freissmuth et al., 1996). SB649146, a S1P₁ receptor inverse agonist attenuated the S1P-induced activation of p42/p44 MAPK in HEK293 cells (Waters et al., 2006). There was no effect of this compound on GTPγS binding in membranes expressing the S1P₂, S1P₃ or LPA₁ receptor. VPC23019 is a synthetic antagonist for the S1P₁ and S1P₃ receptor. A specific S1P₂ receptor antagonist, JTE-013 was described in 2002 (Osada et al., 2002). A new S1P₃ antagonist, BML-241, was found upon a search for compounds that inhibit S1P-induced rises in [Ca²⁺]_i for 37% at 10 μM (Koide et al., 2002). Recently, a specific S1P₁ receptor antagonist, named W146, was described (Sanna et al., 2004).

A common problem with newly found or developed ligands is that they are usually poorly characterized. It is therefore important that compounds are investigated thoroughly before usage.

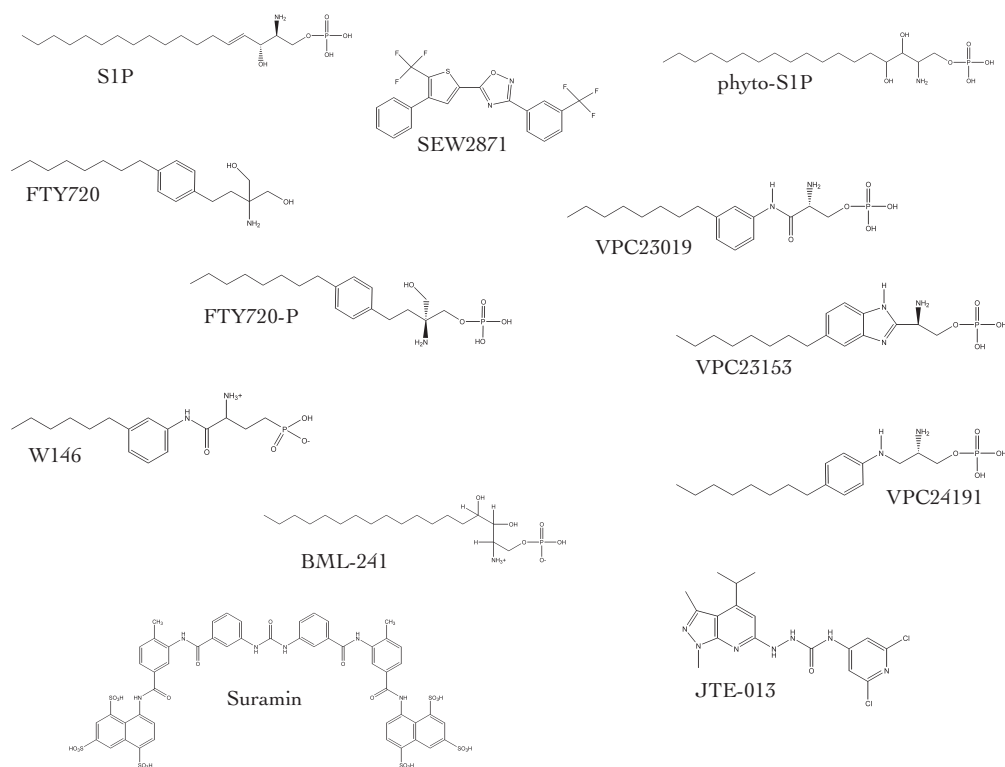


Figure 9. Chemical structures of several S1P receptor ligands.

Aim of the project

When one aims to study GPCRs, many different tools are needed. In the relatively new field of S1P receptors, these tools are not overly present. The few that do exist are usually poorly characterized due to their short existence. Therefore, the first part of this thesis was aimed at characterizing and/or improving existing tools and to develop new ones. This part focussed on characterizing one specific new antagonist by studying its effect on various signaling pathways of S1P receptors and other receptors (chapter 2), and on setting up a new method to study the internalization mechanisms of the S1P receptors in more detail (chapter 3).

The second part of this project focused on the application of the available or newly developed tools to characterize the S1P receptors. Our newly developed immunocytochemical method was applied to compare the internalization of these receptors (chapter 5) and to discover if the D/ERY motif was involved in this process. The latter by also looking at the effect of mutations in this motif on G_i or G_q -coupled pathways for the S1P₁₋₃ receptors (chapter 6). The many new ligands that are being developed open a pharmacological way to study certain effects induced via S1P receptors. They were used to assign effects of S1P receptor ligands in the rat aorta to S1P receptor subtypes expressed in endothelial or smooth muscle cells (chapter 4). These ligands also need to be thoroughly tested, which we did by investigating their effects on the multiple signaling pathways activated via the S1P₃ receptor (chapter 7).

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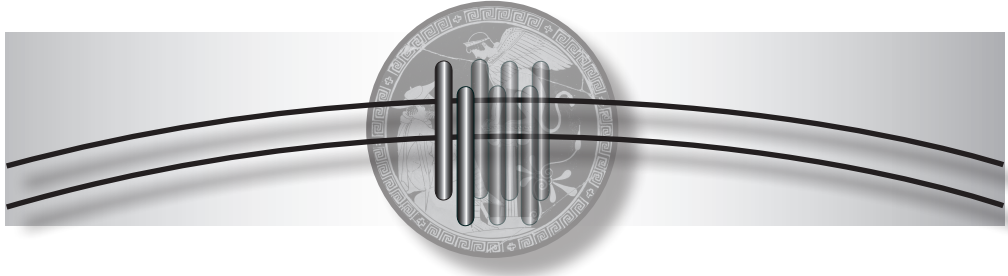
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*BML-241,
a selective SIP_5 receptor antagonist?*



Chapter 2

BML-241, a selective SIP_3 receptor antagonist?

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Abstract

Based upon a screening of $[Ca^{2+}]_i$ responses in HeLa cells overexpressing SIP receptors, it was suggested that BML-241 is a lead compound to develop a selective SIP_3 receptor antagonist. We have studied the proposed selective antagonistic properties of BML-241 for the SIP_3 receptor in more detail.

Chinese hamster ovary (CHO) cells stably transfected with the SIP_3 , SIP_2 or α_{1A} -adrenoceptor were used to investigate the effect of BML-241 on $[Ca^{2+}]_i$ elevations mediated via different receptors. CHO-K1 cells were used to study ATP-induced $[Ca^{2+}]_i$ elevations. Also the effect on SIP_3 -mediated inhibition of forskolin-induced cAMP accumulation and on binding to the α_{1A} -adrenoceptor were investigated. In addition, the effect of BML-241 on the inhibition of phenylephrine-induced contraction of rat mesenteric artery was studied in an organ bath.

High concentrations of BML-241 (10 μ M) inhibited $[Ca^{2+}]_i$ elevations induced by SIP_3 and SIP_2 receptor stimulation, whereas lower concentrations were ineffective. The high concentration of BML-241 also inhibited $[Ca^{2+}]_i$ increases via purinergic and α_{1A} -adrenoceptor stimulation. Moreover, BML-241 (10 μ M) inhibited α_{1A} -adrenoceptor-mediated contraction of rat mesenteric artery. However, it did not displace $[^3H]$ -prazosin from the α_{1A} -adrenoceptors in concentrations up to 100 μ M. BML-241 (10 μ M) did not affect the SIP_3 -mediated decrease of forskolin-induced cAMP accumulation. We conclude that BML-241 is a low potency inhibitor of $[Ca^{2+}]_i$ responses, rather than a specific antagonist of the SIP_3 receptor.

Introduction

Sphingosine-1-phosphate (SIP) is a bioactive sphingolipid found in high concentrations in blood (Yatomi et al., 1997), it is produced and stored in platelets, from which it is released upon stimulation. SIP can also be produced by many other cell types from membrane phospholipids to act in an auto- and/or paracrine manner.

SIP regulates many different biological functions like cell growth, proliferation, apoptosis, lymphocyte recirculation and angiogenesis via specific receptors. The first SIP receptor was discovered as an abundantly expressed gene in differentiating endothelial cells (Hla et al., 1990). Though orphan at first, the product of the endothelial differentiation gene 1 (EDG-1) turned out to have high affinity for SIP and was, therefore, renamed SIP₁ (Lee et al., 1998). The family of SIP receptors currently consists of 5 members with close homology. The SIP₁, SIP₂ and SIP₃ receptors are ubiquitously expressed, whereas the SIP₄ and SIP₅ receptor show a more restricted expression pattern (Alewijjnse et al., 2004). The SIP₁ receptor predominantly activates G_i-proteins, the SIP₂ and SIP₃ receptor activate G_p, G_{q/11} and G_{12/13}-proteins and the SIP₄ and SIP₅ receptors activate G_i and G_{12/13}-proteins (Spiegel et al., 2003).

The specific role of SIP receptor subtypes has predominantly been studied using genetic approaches because of the limited availability of selective agonists and antagonists. Research with knockout mice has for instance shown a crucial role for the SIP₁ receptor in the formation of blood vessels (Liu et al., 2000). However, pharmacological tools are still needed.

In 2002, a new compound, BML-241, was reported from a screening of [Ca²⁺] responses in HeLa cells overexpressing SIP receptors and was proposed to be a lead compound to develop a selective antagonist of the SIP₃ receptor (Koide et al., 2002). However, conclusions were based on measurements of one BML-241 concentration and on the comparison of only the SIP₁ and SIP₃ receptor in one assay. Therefore, we have investigated the proposed selectivity of the SIP₃ receptor antagonistic properties of BML-241 in more detail.

Materials and Methods

Materials

pcDNA3.1 containing the entire coding region of the human SIP₂ or SIP₃ receptor was purchased from UMR cDNA Resource Center (Rolla, MO, USA). Cell culture media, hygromycin B, Lipofectamine™ 2000, Optimem, Hank's Balanced Salt Solution (HBSS), pOG44, pcDNA3.1/HisA/C, pcDNA5/FRT/TO and CHO-FlpIn cells, were obtained from Invitrogen (Breda, The Netherlands). Pluronic acid and Fluo-4 AM were obtained from Molecular Probes (via Invitrogen). Foetal calf serum (FCS), enzyme free cell dissociation buffer and penicillin/streptomycin were obtained from Gibco (via Invitrogen). LANCE™ cAMP 384 kit, white 384 well optiplates and Microscint-O were obtained from Perkin Elmer (Zaventem, Belgium). Restriction enzymes (BamHI, ApaI, XhoI and HindIII) were obtained from Fermentas Life Sciences (St. Leon-Rot, Germany). SIP was obtained from Avanti-Polar Lipids, Inc (via InstruChemie B.V., Delfzijl, The Netherlands). Probenecid, Hepes, EGTA, Triton X-100, 3-isobutyl-1-methylxanthine, bovine serum albumin (BSA) (fatty acid free) and activated charcoal were obtained from Sigma Aldrich (Zwijndrecht, The Netherlands). Clear bottom black 96 well plates were obtained from Greiner Bio One (Alphen aan den Rijn, The Netherlands). BML-241 (2(R,S)-undecylthiazolidine-4(R)-carboxylic acid) was obtained from Biomol (via Tebu-Bio, Heerhugowaard, The Netherlands). CHO-K1 cells were purchased from ECACC (Zwijndrecht, The Netherlands). [³H]-prazosin (specific activity 80 Ci/mmol) was obtained from Amersham (Diegem, Belgium). CHO cells stably expressing the human α_{1A}-adrenoceptor at a density of approximately 2 pmol/mg protein were generated as previously reported (Keffel et al., 2000).

Molecular cloning

An N-terminal HisG-tag was added to the SIP₂ or SIP₃ receptor via cloning into pcDNA3.1/HisC or pcDNA3.1/HisA using BamHI & ApaI or BamHI & XhoI digestion, respectively. A second cloning step was done using HindIII & ApaI or HindIII & XhoI to clone the HisG-tagged SIP₂ or SIP₃ receptor into the expression vector pcDNA5/FRT/TO using standard cloning techniques.

Transfection e³ cell culture

Transfection of plasmids into CHO-FlpIn cells was done according to the manufacturer's protocol with Lipofectamine™ 2000. 600 µg/ml hygromycin B was used to select for positive clones, which were subsequently separately grown to confluency. Expression of the SIP_2 or SIP_3 receptor was confirmed by immunofluorescence against the HisG-tag. CHO-FlpIn cells stably expressing the SIP_2 or SIP_3 receptor were passaged 1:5 every two or three days in F-12 Nutrient Mixture (Ham) with L-glutamine, supplemented with 100 U/ml penicillin, 100 µg/ml streptomycin, 313 µg/ml hygromycin B and 10 % charcoal-stripped FCS. CHO-K1 cells were passaged 1:10 every two or three days in F-12 Nutrient Mixture (Ham) with L-glutamine, supplemented with 100 U/ml penicillin, 100 µg/ml streptomycin and 10 % FCS. CHO-K1 cells stably expressing α_{1A} -adrenoceptor were cultured as described previously (Keffel et al., 2000). All cell lines were cultured at 37 °C in humidified air containing 5 % CO_2 .

Intracellular calcium measurement

Cells were plated in a black, clear bottom 96 wells plate at 40,000 cells/well. After growth overnight, cells were loaded for 1 hour with basic buffer (HBSS containing 20 mM Hepes and 2.5 mM probenecid) containing 4 µM Fluo-4 AM ester and 0.42 % v/v pluronic acid and incubated at 37 °C. After loading, cells were washed twice with basic buffer and incubated at 37 °C with basic buffer in the presence or absence of BML-241 for 60 min. Fluorescence was measured at 37 °C using an excitation filter at 485 nm and emission filter at 520 nm on a NOVOstar (BMG Labtech, via Isogen, IJsselstein, The Netherlands). After measuring the basal level for 10 s, the ligand was added at 10 % v/v basic buffer. After 50 sec 10 % v/v Triton X-100 in basic buffer was added at 10 % v/v to determine the maximal signal (F_{max}). After 20 sec 0.1 M EGTA in basic buffer was added at 10 % v/v to determine the minimal signal (F_{min}). The increase in $[Ca^{2+}]_i$ upon ligand stimulation was calculated as the difference between the $[Ca^{2+}]_i$ for the basal level and after adding a ligand. The $[Ca^{2+}]_i$ was calculated via the equation: $[Ca^{2+}]_i = Kd * ((F - F_{min}) / (F_{max} - F))$. Kd is the dissociation constant of the binding of Fluo-4 to calcium (345 nM).

cAMP assay

The LANCE™ cAMP 384 kit was used to determine cAMP concentrations according to the manufacturer's protocol. CHO-FlpIn cells stably expressing the SIP₃ receptor were incubated for 1 hour with or without 10 μM BML-241 in F-12 Nutrient Mixture (Ham) with L-glutamine and detached from the surface using enzyme free cell dissociation buffer. Cells were washed once with HBSS and subsequently resuspended in stimulation buffer, containing HBSS with 0.5 mM 3-isobutyl-1-methylxanthine, 0.05% BSA and 5 mM Hepes.

Stimulation mixtures consisted of stimulation buffer with 3 μM forskolin, 10 μM BML-241 or vehicle and the concentration response curve of SIP ranging from 10 pM to 1 μM.

Cells were added to the stimulation mixtures 1:1 in a 384 well optiplate at 2500 cells/well and stimulated for 60 minutes. Detection of the cAMP formed during stimulation was done according to the manufacturer's protocol. Measurements were done on a platereader (Victor 2, Wallac, Perkin Elmer) 3 hours after adding detection buffer and antibody mixture.

Radioligand binding studies

Confluent 20 cm dishes with CHO-FlpIn cells stably expressing α_{1A}-adrenoceptor (Keffel et al., 2000) were washed once in PBS. Cells were scraped in ice-cold PBS, and spun down for 10 min at 200xg, 4 °C. The pellet was washed once in 50 mM Tris, 0.5 mM EDTA and then homogenised in 50 mM Tris, 0.5 mM EDTA on ice using a Turrax. The suspension was spun down for 20 min at 50,000xg at 4 °C. The pellet was resuspended in 50 mM Tris, 0.5 mM EDTA and homogenised again. Protein concentration was determined according to Bradford (Bio-Rad, Veenendaal, The Netherlands). Competition binding studies were performed as previously described (Keffel et al., 2000). Briefly, incubations were performed for 45 min at 25 °C in a total volume of 1 ml containing [³H]-prazosin and 50 mM Tris, 0.5 mM EDTA, pH 7.5. Non-specific binding was defined in the presence of 10 μM phen-tolamine. The incubations were terminated by rapid vacuum filtration over Whatman GF/C filters, which were washed with ice-cold binding buffer. After addition of Microscint-O, radioactivity of bound fraction on the filter was counted by a TopCount scintillation counter (Perkin Elmer, Zaventem, Belgium).

Vascular contraction studies

The experiments followed a protocol approved by the institutional Animal Ethical Committee. Adult male Wistar rats (280 – 320 g, Charles Rivers, Maastricht, The Netherlands) were anaesthetized by i.p. injection of 75 mg/kg pentobarbitone (O.B.G., Utrecht, The Netherlands). 500 I.U. heparin (Leo Pharma B.V., Weesp, The Netherlands) was administered i.p. to prevent blood coagulation. The mesenteric vascular bed was exteriorized 10 cm below the pylorus and immediately placed in Krebs-Henseleit buffer (118.0 mM NaCl, 4.7 mM KCl, 25.0 mM NaHCO₃, 1.2 mM MgSO₄, 2.5 mM CaCl₂, 1.1 mM KH₂PO₄ and 5.6 mM glucose) at room temperature, aerated with 5 % CO₂ / 95 % O₂, pH 7.4. Mesenteric vessels from the third order branches, with a length of about 3 mm, were isolated from fat and surrounding tissue, and a 40 µm diameter stainless-steel wire was inserted into the lumen of the vessel. The vessel was dissected and the segments were then transferred into the organ baths of a 4-channel wire myograph (Danish Myo Technology, Aarhus, Denmark) and subjected to a normalization procedure according to Mulvany & Halpern (Mulvany & Halpern, 1977). In short, the individual circumference was adjusted to 90 % of the value that the particular vessel would have had at a transmural pressure of 100 mmHg. Afterwards, the arteries were equilibrated for an additional 45 min and the buffer was refreshed every 20 min. The preparations were contracted twice for 10 min with a depolarizing high K⁺ Krebs-Henseleit solution (100 mM NaCl was replaced by 120 mM KCl) at intervals of 15 min. Subsequently, the vessels were precontracted with the α_1 -adrenoceptor agonist phenylephrine (0.3 µM). After reaching a steady level, one concentration of the endothelium-dependent vasodilator methacholine (10 µM) was added to assess the endothelial integrity. Vessels were excluded when relaxation was less than 80 %. After washing, again 100 mM KCl was added to the vessel segments to obtain the maximal contractile response. After washing and an additional 20 min equilibration, 10 µM BML-241 was added to the bathing solution. After 30 min, a cumulative concentration response curve (30 nM – 30 µM) for phenylephrine was constructed. Isometric force of contraction was measured continuously and all data presented are normalized to the contractile response obtained by the third 100 mM KCl-induced contraction.

Data analysis

Concentration response curves were analysed by fitting sigmoidal functions to the experimental data using Prism 4 (Graphpad Software, Inc., San Diego, CA, USA). Data are expressed as means \pm s.e.m. Differences between two groups were compared using a paired Student's t-test. One-way ANOVA with a Dunnett's correction was used where appropriate. $p < 0.05$ was considered significant.

Results

Effect of BML-241 on SIP-induced $[Ca^{2+}]_i$ elevation in CHO cells stably expressing the SIP_3 receptor

SIP concentration-dependently increased the $[Ca^{2+}]_i$ in CHO-FlpIn cells stably expressing the SIP_3 receptor. The calcium tracing shows a rapid ligand-induced increase followed by a decrease (Figure 1A), the former being characterized by a pEC_{50} of 8.1 ± 0.2 and an E_{max} of 2837 ± 419 nM ($n = 9$, Figure 2A). A 60 min pre-incubation with $10 \mu M$ BML-241 caused a rightward shift of the concentration-response curve without affecting the maximum responses ($pEC_{50} = 7.2 \pm 0.2$, $E_{max} = 2697 \pm 403$ nM, $n = 8$, $p < 0.05$, Figure 2A). Lower concentrations of BML-241 (1 or $3 \mu M$) did not significantly influence the $[Ca^{2+}]_i$ response (data not shown).

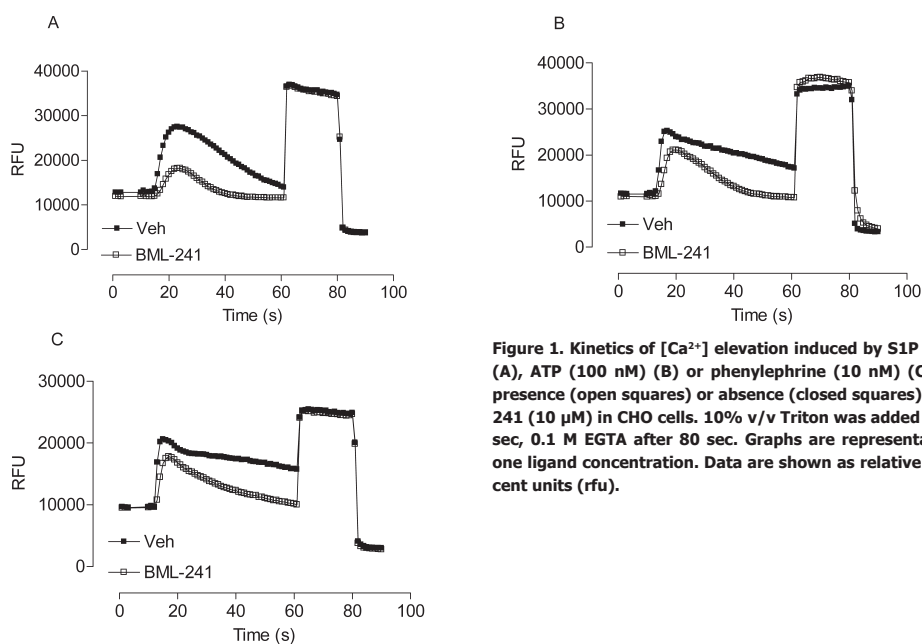


Figure 1. Kinetics of $[Ca^{2+}]_i$ elevation induced by SIP (10 nM) (A), ATP (100 nM) (B) or phenylephrine (10 nM) (C) in the presence (open squares) or absence (closed squares) of BML-241 ($10 \mu M$) in CHO cells. 10% v/v Triton was added after 60 sec, 0.1 M EGTA after 80 sec. Graphs are representative for one ligand concentration. Data are shown as relative fluorescent units (rfu).

Effect of BML-241 on SIP -induced $[Ca^{2+}]_i$ elevation in CHO cells stably expressing the SIP_2 receptor

SIP also concentration-dependently increased the $[Ca^{2+}]_i$ in CHO-FlpIn cells stably expressing the SIP_2 receptor. The concentration response curve was significantly shifted to the right after a pre-incubation with 10 μM BML-241, whereas the maximal response was not affected ($pEC_{50} = 7.7 \pm 0.1$ vs. 7.3 ± 0.1 , $p < 0.05$, $E_{max} = 2698 \pm 257$ vs. 2309 ± 359 nM, $n = 6-7$, Figure 2B). However, lower concentrations of BML-241 (1 or 3 μM) did not show such inhibition (data not shown).

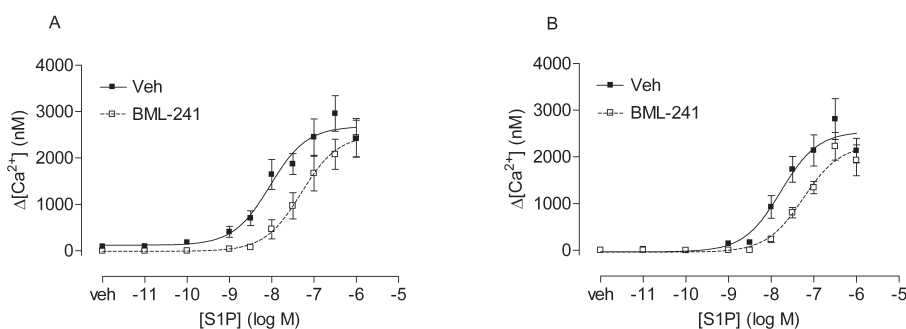


Figure 2. Concentration response curve of SIP -induced $[Ca^{2+}]_i$ elevation in CHO-FlpIn cells stably expressing the SIP_3 -receptor ($n=9$) (A) or the SIP_2 -receptor (B) in the presence (open squares) or absence (closed squares) of BML-241 (10 μM) ($n=6-7$).

Effect of BML-241 on $[Ca^{2+}]_i$ elevation in CHO cells via the purinergic receptors and α_{1A} -adrenoceptor

In CHO-cells expressing the purinergic receptor or α_{1A} -adrenoceptor ATP and phenylephrine also increased $[Ca^{2+}]_i$. However, the fluorescent signal remained elevated longer compared to SIP -mediated stimulation (Figure 1A,B,C). Therefore, we measured the effect of BML-241 on peak responses ('early') as well as after 60 s ('late'). Pre-incubation with BML-241 (10 μM) caused a significant rightward shift of the ATP-induced $[Ca^{2+}]_i$ elevation in the early phase ($pEC_{50} = 6.9 \pm 0.1$ vs. 6.4 ± 0.1 , $n=5$, $p < 0.05$, Figure 3A), but had no significant effect on the late response ($pEC_{50} = 6.3 \pm 0.1$ vs. 6.1 ± 0.1 , $n=5$). On the other hand, this concentration of BML-241 did not significantly alter the E_{max} of the early response ($E_{max} = 1463 \pm 72$ vs. 1536 ± 152 nM, $n=5$), but significantly decreased the E_{max} of the late response ($E_{max} = 972 \pm 43$ vs. 697 ± 62 nM, $n=5$, $p < 0.05$). Similarly, BML-241 (10 μM) significantly shifted the phenylephrine-induced $[Ca^{2+}]_i$ elevation to the right for the early

phase ($pEC_{50} = 8.4 \pm 0.1$ vs. 8.1 ± 0.1 , $n=5$, $p<0.05$, Figure 3B), as well as for the late phase ($pEC_{50} = 7.9 \pm 0.1$ vs. 7.7 ± 0.1 , $n=5$, $p<0.05$). BML-241 ($10 \mu M$) did not significantly alter the E_{max} of the early response ($E_{max} = 1719 \pm 98$ vs. 1693 ± 148 nM, $n=5$), but significantly decreased the late response ($E_{max} = 1241 \pm 58$ vs. 889 ± 43 nM, $n=5$, $p<0.05$).

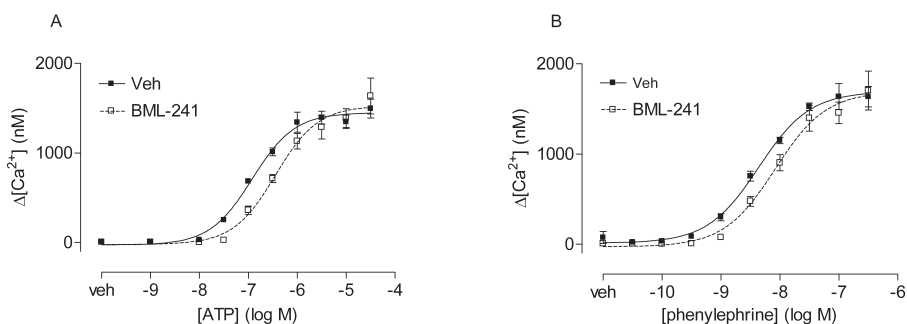


Figure 3. Concentration response curve of the early phase of ATP-induced (A) and phenylephrine-induced (B) $[Ca^{2+}]$ elevation in CHO-K1 cells in the presence (open squares) or absence (closed squares) of BML-241 ($10 \mu M$) ($n=5$).

Effect of BML-241 on phenylephrine-induced contraction of the mesenteric artery

To investigate non-selective effects of BML-241 on a tissue level, an organ bath setup was used. In mesenteric artery preparations phenylephrine causes contraction via the α_1 -adrenoceptor. Addition of BML-241 ($10 \mu M$) caused a significant rightward shift of the concentration response curve for phenylephrine ($pEC_{50} = 5.9 \pm 0.1$ vs. 5.4 ± 0.1 , $n=6-7$, $p<0.05$, Figure 4) and significantly decreased the E_{max} ($E_{max} = 160 \pm 6$ vs. 132 ± 9 %, $n=6-7$, $p<0.05$, Figure 4). BML-241 had no significant influence on KCl-induced constriction (data not shown).

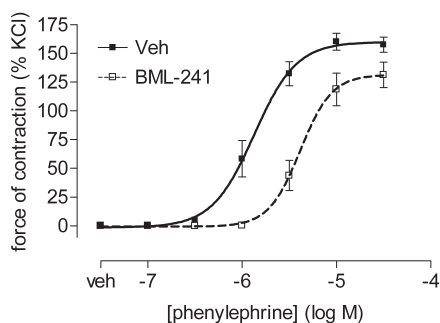


Figure 4. Phenylephrine-induced contraction of rat mesenteric artery in the presence (open squares) or absence (closed squares) of BML-241 ($10 \mu M$) ($n=6-7$). Data are expressed as percentage of the third KCl (120 mM)-induced contraction.

Effect of BML-241 on binding of [3H]-prazosin to the α_{1A} -adrenoceptor

In competition binding experiments, BML-241 in concentrations up to 100 μM did not cause detectable inhibition of radioligand binding to α_{1A} -adrenoceptors, whereas phenylephrine as the positive control exhibited a concentration-dependent displacement of the radioligand (Figure 5).

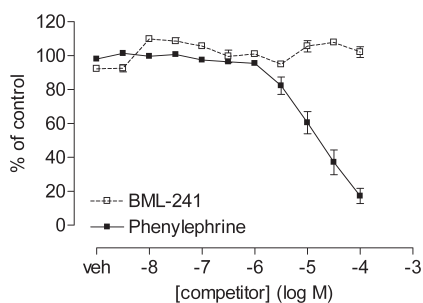


Figure 5. Displacement of [3H]-prazosin on membranes of CHO cells stably expressing the human α_{1A} -adrenoceptor by increasing concentrations of phenylephrine or BML-241 (n=3). Data are expressed as % of control.

Effect of BML-241 on SIP_3 -induced inhibition of cAMP accumulation

In CHO-FlpIn cells stably expressing the SIP_3 receptor, SIP_3 inhibited forskolin-induced cAMP accumulation in a concentration-dependent manner ($pEC_{50} = 7.8 \pm 0.4$, n=4). Pre-incubation with BML-241 (10 μM) did not have a significant influence on the SIP_3 -induced inhibition of cAMP accumulation ($pEC_{50} = 8.1 \pm 0.5$, n=4). In mock transfected CHO cells, SIP_3 did not inhibit forskolin-induced cAMP accumulation (Figure 6).

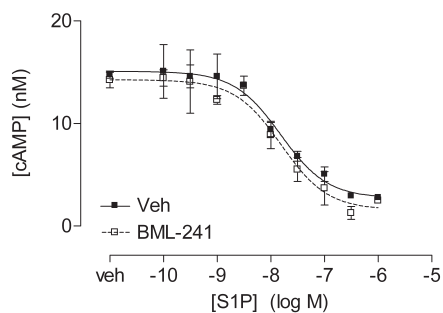


Figure 6. Concentration response curve of SIP_3 -induced inhibition of forskolin-induced cAMP accumulation in the presence (open squares) or absence (closed squares) of BML-241 (10 μM) in CHO-FlpIn cells stably expressing the SIP_3 -receptor (n=4).

Discussion

Research on SIP receptors has been hampered by a lack of specific ligands, particularly reagents which allow investigating the role of endogenous SIP. The SIP₃ receptor can stimulate phospholipase C, activate rho and regulate ERK1/2 activation (Watterson et al., 2005). Specific and particularly subtype-selective antagonists could help considerably to better understand the physiology of SIP responses as well as the mechanisms of drugs targeted at this system; moreover such antagonists have the potential to become useful drugs by themselves.

In 2002, it was reported from a screening of HeLa cells that a single high concentration of BML-241 (10 μM) inhibited the SIP₃-induced [Ca²⁺]_i response by 37 %, whereas SIP₁-induced responses were not affected (Koide et al., 2002). Based upon these data, BML-241 was proposed to be a lead compound to develop a selective antagonist of the SIP₃ receptor. Our study confirms that a high concentration of BML-241 indeed inhibits the SIP-induced [Ca²⁺]_i elevation in cells overexpressing the SIP₃ receptor. However, this high concentration (10 μM), was not selective for the SIP₃ receptor, but also affected the SIP₂-mediated [Ca²⁺]_i elevation.

To determine the specificity of BML-241 for SIP receptors, effects on purinergic receptors and α₁-adrenoceptors were studied using ATP and phenylephrine, respectively. Interestingly, a difference in kinetics of the [Ca²⁺]_i signal was observed compared to SIP stimulated cells stably expressing the SIP₂ or SIP₃ receptor. The purinergic receptor agonist ATP and the α_{1A}-adrenoceptor agonist phenylephrine also rapidly increased [Ca²⁺]_i, but the fluorescent signal remained elevated longer than after SIP receptor stimulation. Therefore, the effect of BML-241 was investigated in the 'early' and 'late' phases separately. BML-241 (10 μM) caused a significant rightward shift of the EC₅₀ in the early phase, but not in the late phase for the ATP-induced [Ca²⁺]_i response. However, for the phenylephrine-induced [Ca²⁺]_i response, BML-241 caused a significant rightward shift of the EC₅₀ of the early and the late phase. The maximal response for both the ATP-induced and the phenylephrine-induced [Ca²⁺]_i elevation was significantly decreased by BML-241 (10 μM) in the late phase but not in the early phase.

The relevance of the lack of selectivity for SIP receptors was confirmed at the tissue level by the inhibitory effect of BML-241 on the phenylephrine-induced contraction in a rat mesenteric artery preparation. The combination of BML-241 being

a low-potency and a non-selective antagonist of $[Ca^{2+}]_i$ responses could be due to low affinity binding to multiple receptors. However, BML-241 did not displace $[^3H]$ -prazosin from the α_{1A} -adrenoceptor, even at very high concentrations (100 μM), indicating that BML-241 did not act at the receptor level. Based upon these results, we conclude that BML-241 is a low-potency functional antagonist of $[Ca^{2+}]_i$ elevation rather than a specific S1P₃ receptor antagonist. In line with this conclusion, BML-241 showed a lack of antagonism on the G_i-mediated inhibitory effect of the S1P₃ receptor on forskolin-induced cAMP production. Recently, expression of S1P₁, S1P₂ and S1P₄ receptor mRNA was shown in CHO-K1 cells (Holdsworth et al., 2005). Hence, the S1P-induced inhibition of forskolin-induced cAMP accumulation could be caused via these endogenous receptors instead of the stably transfected S1P₃ receptor. However, in our mock transfected CHO-FlpIn cells, there is no effect of S1P on the forskolin-induced cAMP accumulation. Therefore, the effect of S1P can only be caused via the stably transfected S1P₃ receptor and not by endogenous S1P receptors. Due to the complexity of calcium signaling, identification of the low-affinity molecular target of BML-241 remains unclear and is beyond the scope of this paper. However, these results show that possible new S1P receptor antagonists, like BML-241, but also the new compound VPC23019 (Davis et al., 2005), a S1P₁/S1P₃ antagonist, for example, should be investigated thoroughly.

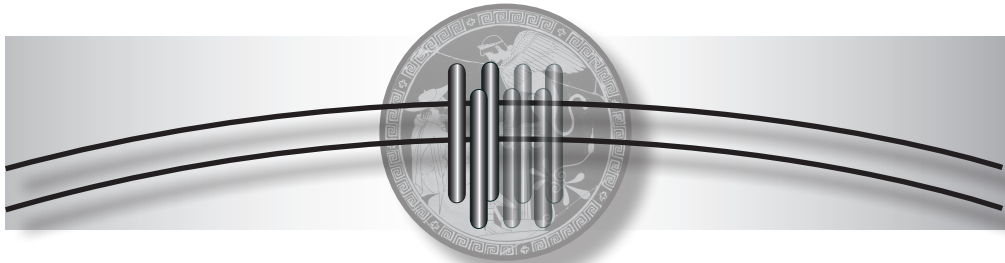
Acknowledgements

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*Validation of a rapid, non-radioactive method
to quantify internalization
of G-protein coupled receptors*



Chapter 3

Validation of a rapid, non-radioactive method to quantify internalization of G-protein coupled receptors

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Abstract

Agonist exposure can cause internalization of G-protein coupled receptors (GPCRs), which may be a part of desensitization but also of cellular signaling. Previous methods to study internalization have been tedious or only poorly quantitative. Therefore, we have developed and validated a quantitative method using a sphingosine-1-phosphate (S1P) receptor as a model. Due to a lack of suitable binding studies, it has been difficult to study S1P receptor internalization. Using a N-terminal HisG-tag, S1P₁ receptors on the cell membrane can be visualized via immunocytochemistry with a specific anti-HisG antibody. S1P-induced internalization was concentration dependent and was quantified using a microplate reader, detecting either absorbance, a fluorescent or luminescent signal, depending on the antibodies used. Among those, the fluorescence detection method was the most convenient to use. The relative ease of this method makes it suitable to measure a large number of data points, e.g. to compare the potency and efficacy of receptor ligands.

Introduction

When chronically exposed to agonist, G-protein coupled receptors (GPCRs) can undergo desensitization, i.e. the response to a given level of agonist exposure declines and/or more agonist is required to maintain the same response. Various mechanisms can be involved in desensitization, some of which directly affect receptor responsiveness. One of these mechanisms involves phosphorylation of activated receptors by GPCR kinases (GRKs). This enables binding of arrestins to the receptor, leading to uncoupling from the G-protein and hence impaired signaling. Another, not necessarily mutually exclusive mechanism of desensitization is internalization of the receptor, which can be either homologous (due to activation of the receptor) or heterologous (due to activation of other receptors or to other stimuli). Receptor internalization can be arrestin-dependent or independent (Bünemann & Hosey, 1999) and involves translocation of the receptor from the cell membrane to the inside of the cell. There it is entrapped inside clathrin-coated pits and can undergo either degradation or resensitization.

Several approaches have been used in the past to detect and quantify GPCR internalization. One classical method involves homogenization of tissue or cells, differential centrifugation to yield a plasma membrane and a microsomal fraction followed by electrophoresis and autoradiographical analyses of the receptor of interest in both fractions (Simpson et al., 1984). Another classical method involves labeling of all receptors in intact cells with a lipophilic radioligand and competition by a hydrophilic ligand, which will replace the radioligand from sites at the cell surface but not from those entrapped intracellularly (Staehein, 1983).

More recently, genetic approaches have been used. An example is the use of a C-terminal green fluorescent protein (GFP) (Barak et al., 1997; Kallal et al., 1998) to monitor receptor localization after ligand stimulation, sometimes combined with an algorithm to discriminate between internalized and membrane bound receptors (Conway et al., 1999; Schlag et al., 2004; Hirasawa et al., 2005; Fukunaga et al., 2006). This latter method uses a cell-based fluorescence imaging system ArrayScanII, where the amount of internalized receptor/GFP spots are counted and normalized against the number of nuclei stained with Hoechst. However, a GFP-, or other fluorescent tag is relatively large (200-300 amino acids) and can influence receptor function (McLean & Milligan, 2000). Other methods use a combination of techniques, like quantitative flow cytometry analysis of the cell surface receptors and immunocytochemical confocal microscopy analysis to visualize the subcellular

localization of the receptor of interest (Hirasawa et al., 1998).

Detailed quantitative studies of internalization are of interest as this process can be the mechanism of underlying physiological and pharmacological effects. For example, receptor internalization is thought to underlie the modulation of immune function by sphingosine-1-phosphate (S1P) (Matloubian et al., 2004). This also appears to be the mechanism of action of the immunosuppressive drug FTY720 (fingolimod) (Chiba et al., 2006; LaMontagne et al., 2006), which is undergoing clinical trials for the treatment of multiple sclerosis and previously also for renal graft rejection (Budde et al., 2002; Kappos et al., 2006).

In our search to quantify internalization of membrane receptors in a fast and easy way, we have setup a method, which uses a N-terminal HisG (6xHis+Gly)-tagged S1P₁ receptor. Since the HisG-tag is N-terminal, it is extracellular. Therefore, when immunocytochemistry is performed without permeabilization, only membrane-bound, tagged receptors will be detected. This way, internalization is quantified as a difference in membrane-fluorescence between unstimulated and agonist-stimulated cells.

The use of small, N-terminal tags has been described before (Von Zastrow et al., 1993; Wozniak & Limbird, 1996; Daunt et al., 1997; Vicentic et al., 2002; Miller, 2004). Of these, Daunt et al., 1997 described a method where cells were plated onto 24-well tissue culture dishes. After stimulations with an agonist, incubations with 1st and 2nd antibodies followed. Measurements were done after adding alkaline phosphatase substrate, resulting in a colorimetric reading. In this paper, we now describe the validation of a similar method by testing various antibody combinations and detection methods to quantify internalization of N-terminal HisG-tagged receptors. This resulted in immunocytochemical method using a specific anti-HisG 1st antibody combined with a fluorescent 2nd antibody (AlexaFluor® 488) as our preferred choice. The signal can either be visualized under the fluorescence microscope or measured in a fluorescence microplate reader. The latter results in quantitative measurement of the decrease in signal, caused by receptor internalization, in a direct, sensitive and fast way.

Materials & Methods

Materials

pcDNA3.1 containing the entire coding region of the human $S1P_1$ receptor was purchased from UMR cDNA Resource Center (Rolla, MO, USA).

Cell culture media, hygromycin B, Lipofectamine™2000, pcDNA3.1/His C, pOG44, CHO-FlpIn cells, Alexa Fluor® 488 goat anti-mouse (IgG), anti-HisG (mouse monoclonal IgG2a), anti-HisG-horse radish peroxidase (HRP) (mouse monoclonal IgG2a), pcDNA5/FRT/TO, NuPAGE® System, NuPAGE® Novex 4-12% Bis-Tris gel, MES running buffer, NuPAGE® transfer buffer and Invitrolon™ PVDF blotting membranes were obtained from Invitrogen (Breda, The Netherlands). BCA™ Protein Assay Kit was obtained from Pierce (Etten-Leur, The Netherlands). Formaldehyde solution was obtained from Sigma (Zwijndrecht, The Netherlands). Restriction enzymes (BamHI, XhoI, EcoRI and HindIII) were obtained from Fermentas Life Sciences (St. Leon-Rot, Germany). Black, clear bottom 96 well plates were obtained from Greiner Bio One (Alphen aan den Rijn, The Netherlands). $S1P$ was obtained from Avanti-Polar Lipids, Inc (via Instruchemie B.V., Delfzijl, The Netherlands). Chamber slides were obtained from Nunc (Amsterdam, The Netherlands). BM Chemiluminescence Blotting Substrate [POD] and ABTS (2,2'-azino-bis(3-ethylbenzthiazoline-6-sulphonic acid)) were obtained from Roche (Mannheim, Germany). Hyperfilm ECL was obtained from Amersham Biosciences (Diegem, Belgium). SEW2871 (5-(4-Phenyl-5-trifluoromethylthiophen-2-yl)-3-(3-trifluoromethylphenyl)-1,2,4-oxadiazole) was obtained from Calbiochem (via VWR, Amsterdam, The Netherlands). Goat anti-mouse-HRP was obtained from Jackson ImmunoResearch Europe Ltd. (via Sanbio B.V., Uden, The Netherlands). RIPA buffer was obtained from Boston bio products (Worcester, UK). Halt protease inhibitor cocktail was obtained from Pierce (via Perbio Science Nederland B.V. Etten-Leur, The Netherlands).

Molecular Cloning & Transfection

An N-terminal HisG-tag was added to the $S1P_1$ receptor via cloning into pcDNA3.1/HisC using BamHI&XhoI. A second cloning step was done using HindIII&XhoI to clone the HisG-tagged $S1P_1$ receptor into the expression vector pcDNA5/FRT/TO. Transfection of plasmids into CHO-FlpIn cells and cell culture was done as described previously (Chapter 2).

Western blot

Cells were washed and scraped in PBS, spun down, and lysed in RIPA buffer with protease inhibitor. Protein concentration was determined using the BCA kit according to the manufacturer's protocol. Ten μg of protein were loaded onto a 4-12 % NuPAGE® Bis-Tris Gel. Electrophoresis was carried out at 200 V for 40 min in NuPAGE® MES running buffer. Protein was transferred to Invitrolon™ PVDF blotting membranes at 30 V for 1 hour. The membrane was blocked for 1 hour at room-temperature in 50 mg/ml non-fat dry milk in PBS, followed by overnight incubation at 4 °C with a dilution of 1:5000 for mouse anti-HisG. After washing with PBS-Tween (0.1 %), the membrane was incubated with a 1:20,000 dilution of goat anti-mouse-HRP IgG at room-temperature. Detection was done using BM Chemiluminescence Blotting Substrate and Hyperfilm ECL.

Immunocytochemistry

Cells (40,000 cells/well) were plated on day one in 96 wells black, clear bottom plates or 8-well chamber slides. The next day, medium was changed to serum free medium. After incubation overnight, cells were stimulated with the indicated ligand in serum free medium for 30 minutes at 37 °C, unless otherwise indicated. The stimulation solutions were removed and cells were fixed by applying 4 % v/v formaldehyde in PBS for 15 minutes at room-temperature. After washing 3 times with PBS, cells were incubated with a 1st antibody for 1 hour at room-temperature. Cells were washed again 3 times with PBS and if necessary incubated with a 2nd antibody for 1 hour at room-temperature. Cells were washed 3 times with PBS followed by measurements on a platereader (Victor 2, Wallac, Perkin Elmer). The following antibody combinations were used: mouse anti-HisG 1:200 with AlexaFluor® 488 goat anti-mouse 1:500, mouse anti-HisG-HRP 1:200 and mouse anti-HisG 1:200 with goat anti-mouse-HRP 1:500. In case an HRP-bound antibody was used, either ABTS solution (50 $\mu\text{g}/\text{ml}$) or BM Chemiluminescence Blotting Substrate was added to generate an absorbance or luminescence signal respectively. For the absorbance signal a 405 nm filter was used and for the fluorescent signal a 490 nm excitation, and a 535 nm emission filter were used.

Data analysis

Concentration response curves were analyzed by fitting sigmoidal functions to the experimental data using Prism 4 (Graphpad Software, Inc., San Diego, CA, USA). Data are expressed as means \pm s.e.m. Differences between groups were compared using a one way ANOVA with a Dunnett's correction or a Student's t-test where appropriate. $P < 0.05$ was considered significant.

Results

Characterization of the HisG-tagged SIP_1 receptor

The presence of HisG- SIP_1 receptor in stably transfected CHO-FlpIn cells was confirmed in Western blot analysis as a band of approximately 50 kDa (Figure 1). The HisG- SIP_1 receptor was also visualized in the cell membrane by fluorescence microscopy using AlexaFluor® 488 goat anti-mouse as a secondary antibody (Figure 2). CHO-FlpIn cells stably expressing the untagged SIP_1 receptor were used as a negative control for both detection methods (Figures 1 and 2). Addition of the N-terminal HisG-tag to the SIP_1 receptor did not influence signaling via this receptor as it did not significantly alter the potency of SIP_1 to inhibit forskolin-stimulated cAMP accumulation (pEC_{50} SIP_1 9.0 ± 0.1 vs HisG- SIP_1 8.9 ± 0.1 ; $n=6$), measured as described previously (Chapter 2).

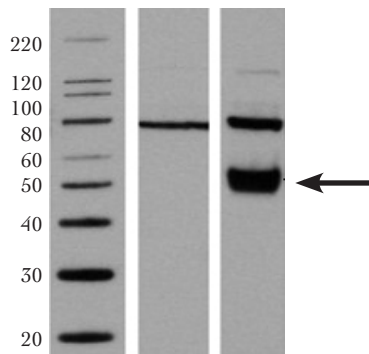


Figure 1. Western blot of CHO-FlpIn cells stably expressing the SIP_1 receptor (lane 1) and HisG- SIP_1 receptor (lane 2) using a mouse anti-HisG 1st antibody and a goat anti-mouse HRP 2nd antibody. HisG- SIP_1 is detected at approximately 50 kDa as indicated by the arrow. The band at 80 kDa is nonspecific. The blot shown is from a typical experiment out of 4.

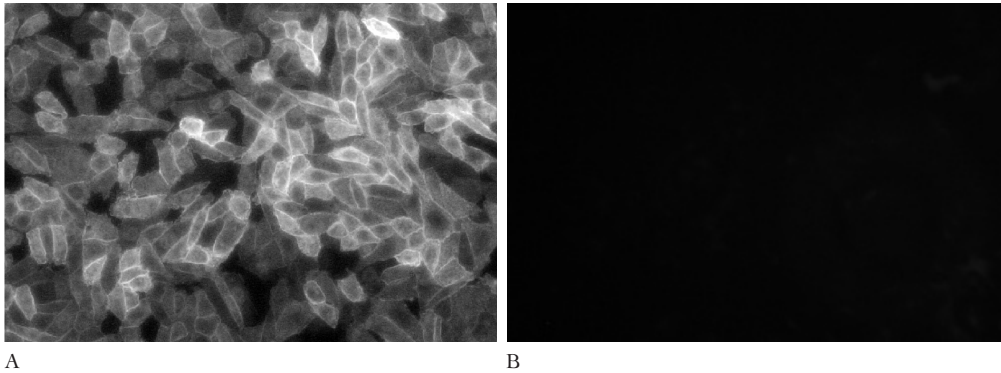


Figure 2. Fluorescence microscope pictures of CHO-FlpIn cells stably expressing the HisG-S1P₁ receptor(A) and the S1P₁ receptor (B) using a mouse anti-HisG 1st antibody and AlexaFluor® 488 goat anti-mouse 2nd antibody. Images shown are from one typical experiment out of 9.

Internalization of the S1P₁ receptor

A 30 min incubation with 1 μ M S1P or 10 μ M of the selective S1P₁ agonist SEW2871 (Sanna et al., 2004) at 37 °C decreased fluorescence (Figure 3A,C,E). No such decrease of fluorescence was seen when cells were incubated with either agonist at 4 °C (Figure 3B,D,F). S1P concentration-dependently internalized the HisG-S1P₁ receptor as visualized with a fluorescence microscope (Figure 4). Such decrease in fluorescence was also measured using a fluorescence microplate reader. The fluorescent signal of cells stably expressing the HisG-S1P₁ receptor was significantly higher than for cells stably expressing the untagged S1P₁ receptor. Stimulation with S1P (1 μ M) for 30 min significantly decreased this signal (Figure 5).

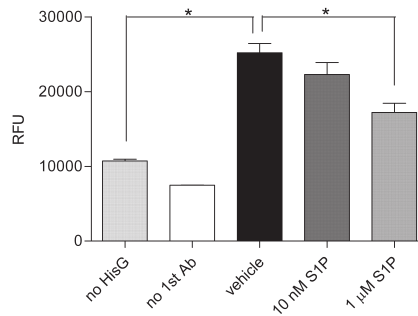


Figure 5. Quantitative measurements of cells stably expressing HisG-S1P₁ using a mouse anti-HisG 1st antibody and AlexaFluor® 488 goat anti-mouse 2nd antibody. Cells were stimulated at 37 °C for 30 min at the indicated S1P concentrations. The first bar represents cells stably expressing the S1P₁ receptor without a HisG-tag. The second bar represents cells stably expressing HisG-S1P₁ incubated without the 1st antibody. Statistical significance was shown between 'vehicle' and '1 μ M S1P' as well as between 'no HisG' and 'vehicle'. Data are shown as relative fluorescent units (RFU) \pm s.e.m. (n=3).

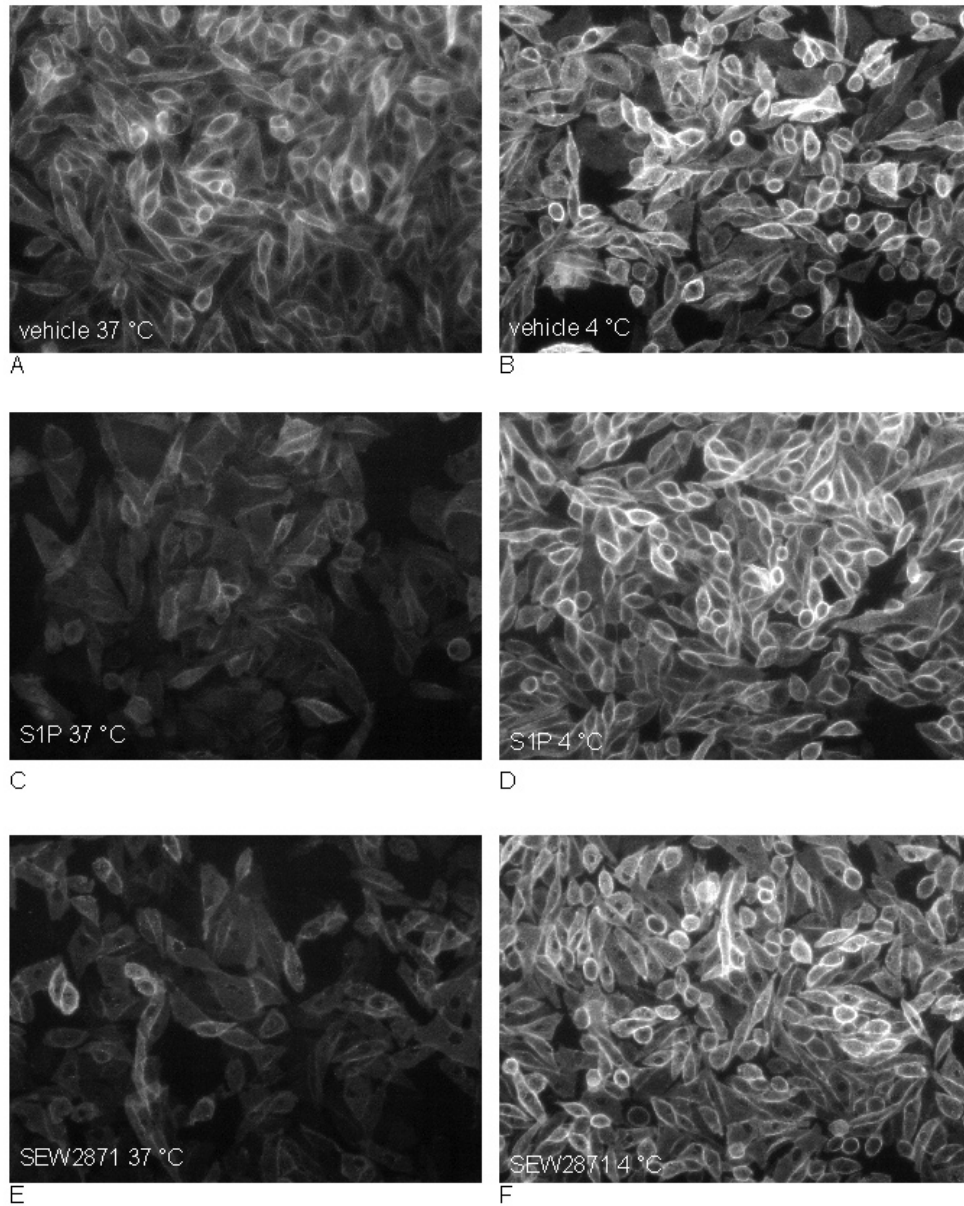


Figure 3. Fluorescence microscope pictures of CHO-FlpIn cells stably expressing the HisG-S1P₁ using a mouse anti-HisG 1st antibody and AlexaFluor® 488 goat anti-mouse 2nd antibody. Cells were stimulated for 30 min with 1 μM S1P (C,D) or 10 μM SEW2871 (E,F) or vehicle (A,B) at either 37 °C (left panel) or 4 °C (right panel). Images shown are from one typical experiment out of 4.

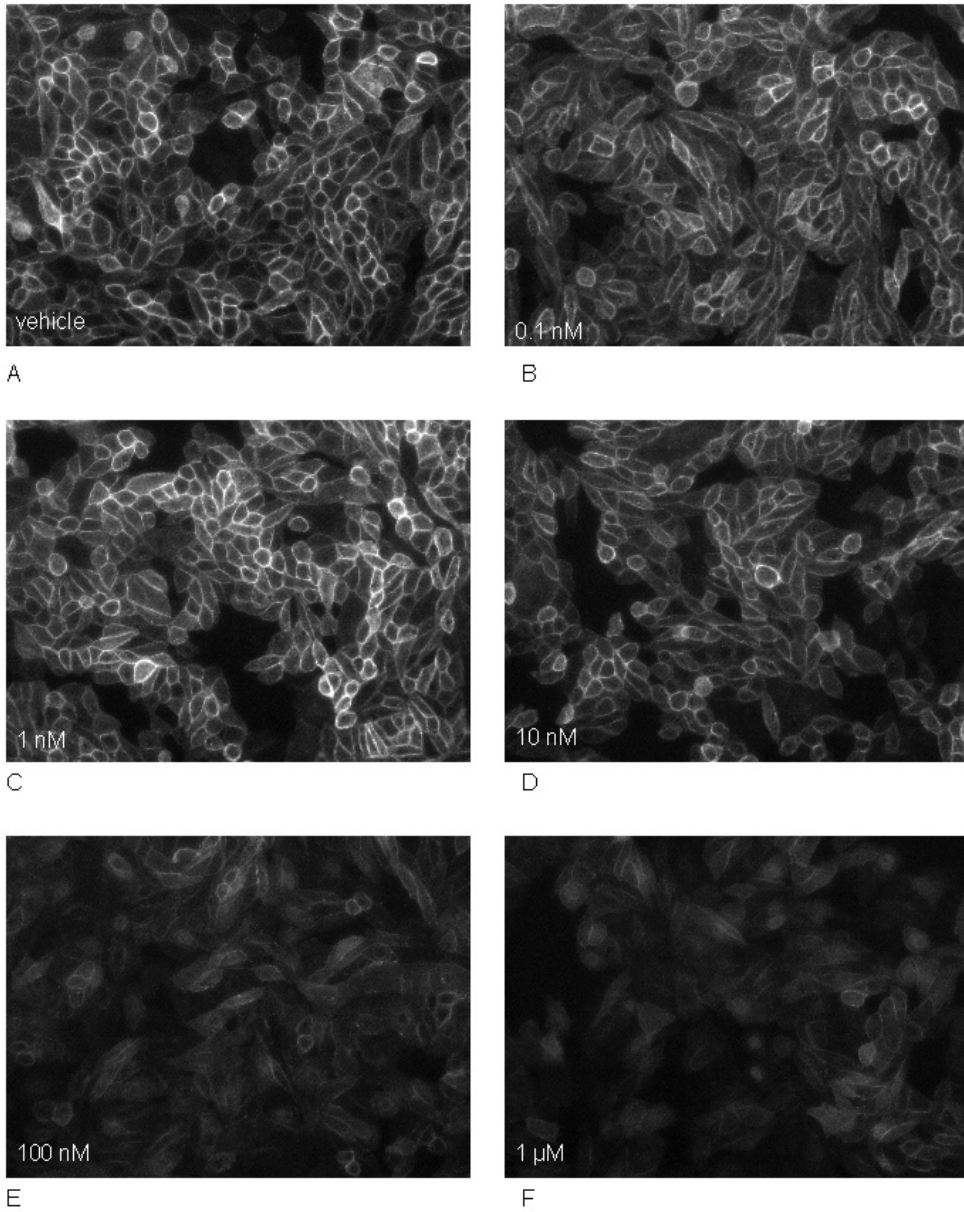


Figure 4. Fluorescence microscope pictures of CHO-FipIn cells stably expressing the HisG-S1P₁ receptor using a mouse anti-HisG 1st antibody and AlexaFluor® 488 goat anti-mouse 2nd antibody. Cells were stimulated with S1P at the indicated concentrations (vehicle (A), 0.1 nM (B), 1 nM (C), 10 nM (D), 100 nM (E) or 1 μM (F)) at 37 °C for 30 min. Images shown are from one typical experiment out of 4.

Choice of detection method

Instead of the antibody combination mouse anti-HisG and AlexaFluor® 488 goat anti-mouse (Figure 6A), other combinations were tested to optimize the method. The use of a 1st antibody mouse anti-HisG-HRP (Figure 6B), or a 1st antibody mouse anti-HisG with a 2nd antibody goat anti-mouse-HRP (Figure 6C) combined with the BM Chemiluminescence Blotting Substrate (POD) resulted in a luminescence signal which proved very sensitive. However, the signal was very unstable over time, which required addition of the substrate by the microplate reader. The combination of a 1st antibody mouse anti-HisG-HRP (Figure 6D), or a 1st antibody mouse anti-HisG with a 2nd antibody goat anti-mouse-HRP (Figure 6E) combined with ABTS resulted in an absorbance signal. This signal however, had less

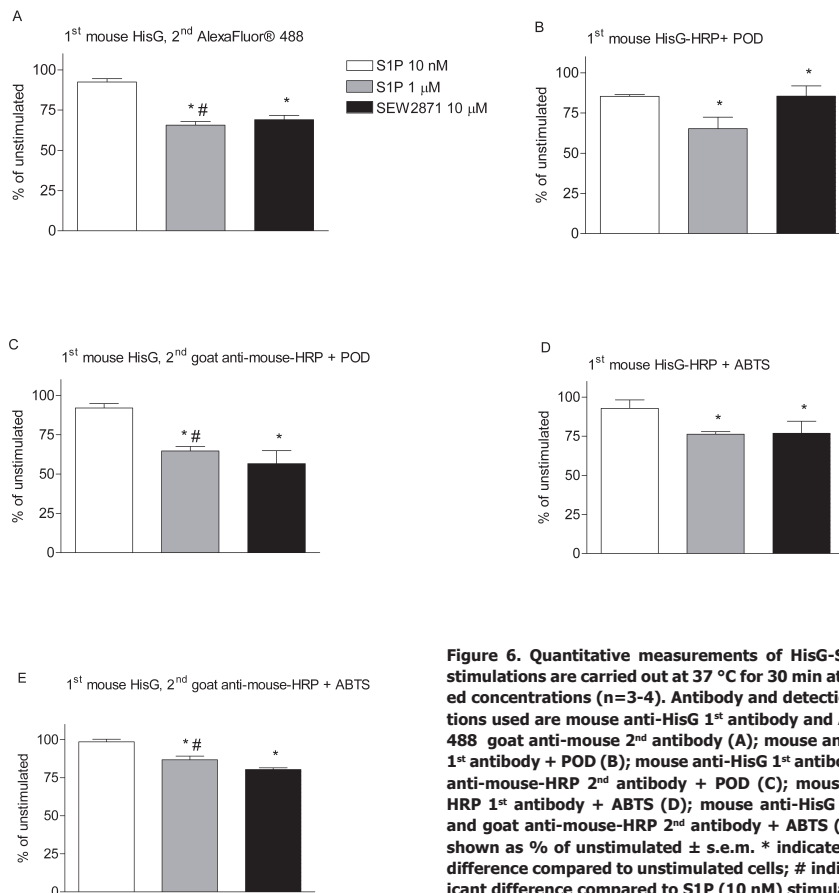


Figure 6. Quantitative measurements of HisG-S1P. Ligand stimulations are carried out at 37 °C for 30 min at the indicated concentrations (n=3-4). Antibody and detection combinations used are mouse anti-HisG 1st antibody and AlexaFluor® 488 goat anti-mouse 2nd antibody (A); mouse anti-HisG-HRP 1st antibody + POD (B); mouse anti-HisG 1st antibody and goat anti-mouse-HRP 2nd antibody + POD (C); mouse anti-HisG-HRP 1st antibody + ABTS (D); mouse anti-HisG 1st antibody and goat anti-mouse-HRP 2nd antibody + ABTS (E). Data are shown as % of unstimulated ± s.e.m. * indicates significant difference compared to unstimulated cells; # indicates significant difference compared to S1P (10 nM) stimulated cells.

sensitivity compared to the other two detection methods. The use of a 1st antibody mouse anti-HisG-HRP resulted in a loss of sensitivity compared to the combination of a 1st antibody mouse anti-HisG with a 2nd antibody goat anti-mouse-HRP.

When the combination of a 1st antibody mouse anti-HisG and a 2nd antibody AlexaFluor® 488 goat anti-mouse was used, we measured a concentration-dependent internalization of HisG-S1P₁ after stimulation with either S1P or SEW2871 (pEC₅₀ S1P = 7.4 ± 0.1 (n=8) and pEC₅₀ SEW2871 = 5.7 ± 0.2 (n=3) respectively, Figure 7).

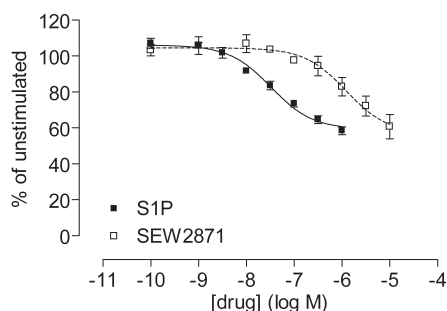


Figure 7. Effect of increasing concentrations of S1P (closed squares) or SEW2871 (open squares) on CHO-FipIn cells stably expressing the HisG-S1P₁ receptor, using a mouse anti-HisG 1st antibody and AlexaFluor® 488 goat anti-mouse 2nd antibody. Stimulations were carried out at 37 °C for 30 min. The data shown represent the mean ± s.e.m. of 6 independent experiments performed in triplicate.

Discussion

Previously reported methods to study the internalization of GPCRs were very labor-intensive, required specific equipment, added bulky tags and/or were poorly quantitative. We have developed and validated a method which uses stably transfected cells expressing a receptor with a minor modification only, i.e. an N-terminal HisG-tag, which did not influence receptor signaling. Several antibody combinations and detection methods were tested. The luminescent approach was very sensitive, but highly dependent on strict adherence to a specific time interval between substrate addition and actual signal measurement. The absorbant signal obtained using another detection substrate resulted in a loss of sensitivity. The antibody combination of a 1st HisG specific antibody and a 2nd AlexaFluor® 488 fluorescent antibody proved the most convenient, resulting in a sensitive and stable signal. Compared to previously published methods using a similar approach (Von Zastrow et al., 1993; Wozniak & Limbird, 1996; Daunt et al., 1997; Vicentic et al., 2002),

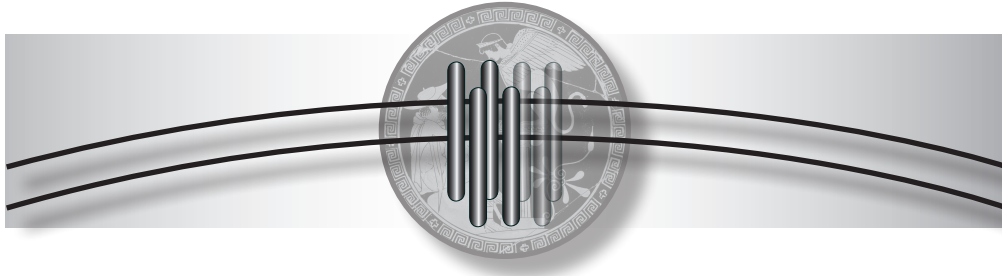
this immunocytochemical detection of the N-terminal HisG-tag is suitable to assess a large number of samples e.g. to construct detailed concentration response curves and/or compare multiple ligands. In this regard, we showed that SIP was approximately 50 times more potent than SEW2871 in internalising the HisG-SIP₁ receptor. This is in good agreement with the previously reported potency difference between the two agonists for activating the receptor (Sanna et al., 2004). Since comparable methods have been described for adrenergic receptors as mentioned above (Daunt et al., 1997; Vicentic et al., 2002; Wozniak & Limbird, 1996; Von Zastrow et al., 1993), it is likely that the method we describe here is also suitable for receptors other than our model receptor, the SIP₁ receptor.

While our technique does not allow following internalization in real time, the ability to process a large number of samples compensates for that by studying multiple time points. The only limitation to our method is the inability to measure internalization of endogenous receptors. However, this would be possible when specific receptor antibodies are available, which is not yet the case for our model receptor. In conclusion, we have validated a method to quantitatively measure receptor internalization in which we use an N-terminal HisG-tag combined with immunocytochemistry. It proves to be sensitive enough to discriminate between different receptor ligands in a fast and non-radioactive way.

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*Pharmacological and biochemical characterization
of the vasoactive properties
of sphingosine-1-phosphate receptors in rat aorta*



Chapter 4

Pharmacological and biochemical characterization of the vasoactive properties of sphingosine-1-phosphate receptors in rat aorta

Manuscript submitted

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Abstract

The bioactive sphingolipid sphingosine-1-phosphate (S1P) exerts diverse cellular effects via a family of five specific G-protein coupled receptors, named S1P₁₋₅. In the cardiovascular system, the S1P₁, S1P₂ and S1P₃ receptors are differentially expressed in both vascular smooth muscle cells (VSMCs) and endothelial cells (ECs). The vasoactive effects of S1P are, therefore, a result of the balance between expression of these receptors on either cell type. We have explored this balance in the rat aorta using new synthetic S1P receptor ligands combined with biochemical methods. In intact isolated rat aortic rings, agonists at the S1P₁ and/or S1P₃ receptor (FTY720, FTY720-P, SEW2871 and VPC23153) did not induce vasoconstriction, but rather endothelium dependent vasorelaxation of pre-contracted preparations. In rat aortic ECs only the G_i-coupled S1P₁ receptor was highly expressed. In these cells, S1P, FTY720, FTY720-P and SEW2871 inhibited cAMP accumulation via S1P₁. Additionally, VPC23019 competitively antagonized the FTY720-P-induced inhibition of the cAMP accumulation. VSMCs showed low mRNA expression of all three S1P receptors. S1P-induced elevation of [Ca²⁺]_i in these cells was observed via predominantly the S1P₂ receptor, since JTE-013 partially inhibited this response. In addition, minor S1P₁ and/or S1P₃-mediated effects on cAMP accumulation were observed in VSMCs. In conclusion, by using newly available pharmacological tools, we have shown vasodilation-promoting S1P₁ receptor-mediated effects in ECs and vasoconstricting-promoting S1P₂ receptor-mediated effects in VSMCs, which for most compounds results in vasorelaxation in rat aorta.

Introduction

Sphingosine-1-phosphate (S1P) is an important lipid mediator generated from membrane lipids upon cell activation. S1P can be synthesized from sphingomyelin, a major constituent of most cellular membranes, via the action of several enzymes (for a review see Hannun et al., 2001). Besides local production in the vessel wall (Mulders et al., 2006), S1P may reach smooth muscle or endothelial cells via the bloodstream, since S1P is a constituent of high-density lipoproteins (Murata et al., 2000), can be released from activated platelets (Yatomi et al., 2000) and is a major constituent of erythrocytes (Pappu et al., 2007). S1P has profound effects on vascular cells as it shows anti-apoptotic and growth-inducing properties on both smooth muscle and endothelial cells, but also has vasoactive properties (Peters & Alewijnse, 2007). Although intracellular targets have been suggested (Payne et al., 2002), most of the aforementioned effects are mediated via specific S1P receptors present on the cell membrane. Until now, five different S1P receptor subtypes have been described, named S1P₁₋₅ of which in the cardiovascular system the S1P₁, S1P₂ and S1P₃ receptors are the predominant S1P receptors (Kluk & Hla, 2001).

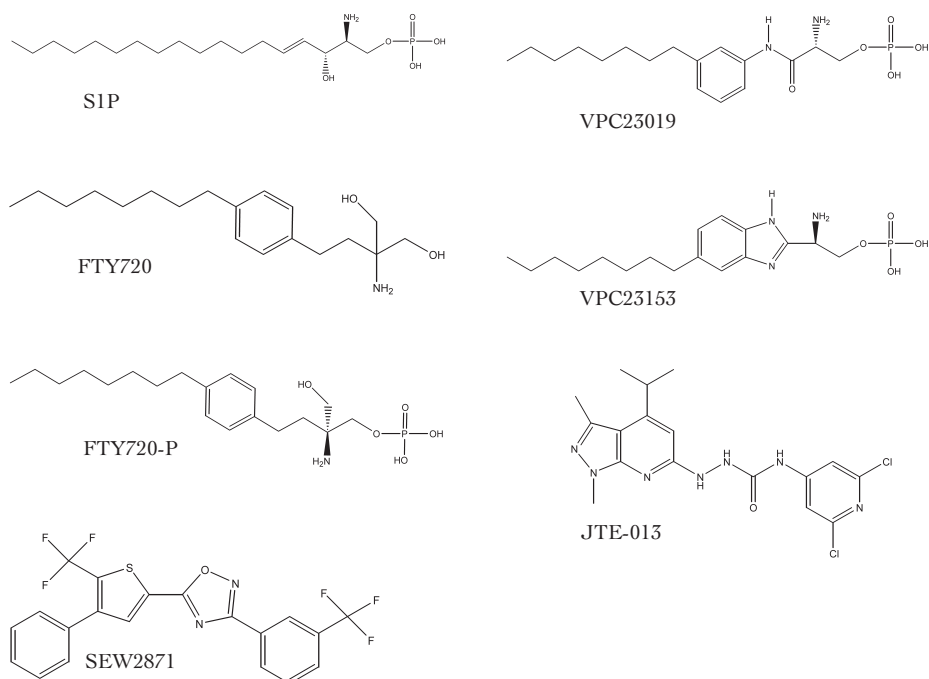


Figure 1. Chemical structure of ligands for the S1P₁, S1P₂ and S1P₃ receptors.

S1P increases intracellular calcium in both vascular smooth muscle and endothelial cells via the activation of S1P receptors. Increases in intracellular calcium in vascular smooth muscle cells are linked to increased constriction. In contrast, a rise in intracellular calcium in the endothelial cell will induce NO production via activation of endothelial NO synthase (eNOS) resulting in vasodilation. Furthermore, activation of endothelial S1P₁ and S1P₃ receptors may lead to enhanced NO production via the activation of the PI3Kinase/Akt pathway and subsequent eNOS phosphorylation (Morales-Ruiz et al., 2001; Nofer et al., 2004; Mulders et al., 2006). The net effect of S1P on the vessel, therefore, depends on the relative expression of S1P receptor subtypes in smooth muscle and endothelial cells. Indeed, S1P has been shown to induce both, vasodilation and vasoconstriction depending, amongst others, on species, applied techniques, and vessel type used (Waeber et al., 2004; Hemmings, 2006). However, research up till now on the role of S1P and its receptors in the vascular system has been hampered by the sparse availability of specific receptor ligands.

Table 1. Overview of properties and pharmacological characteristics of ligands for the S1P₁, S1P₂ and S1P₃ receptors.
¹ Young and van Brocklyn, 2006; ² Mandala et al., 2002 IC₅₀ (binding); ³ Sanna et al., 2004: EC₅₀ (GTPγS); ^{4,1} Davis et al., 2005: pK_b (GTPγS); ^{4,2} Davis et al., 2005: pK_i (binding); ⁵ Ohmori et al., 2003: IC₅₀.

Ligand	Properties	Pharmacological characteristics		
		S1P ₁ (nM)	S1P ₂ (nM)	S1P ₃ (nM)
S1P	S1P _{1,5} agonist	24.7 ¹ 0.47 ²	18.4 ¹ 0.31 ²	19.7 ¹ 0.17 ²
FTY720	-	300 ²	>10000 ²	>10000 ²
FTY720-P	S1P _{1,3,4,5} agonist	0.21 ²	>10000 ²	5.0 ²
SEW2871	S1P ₁ agonist	13 ³	>10000 ³	>10000 ²
VPC23153	S1P _{1,3} agonist	no values available		
VPC23019	S1P _{1,3} antagonist	32 ^{4,1}	no effect	1047 ^{4,1} 1175 ^{4,2}
JTE-013	S1P ₂ antagonist	13.8 ^{4,2}	17 ⁵	no effect

Recently several agonists and antagonists of the S1P₁, S1P₂ and S1P₃ receptors have become available (Table 1, Figure 1) which makes it possible for the first time to pharmacologically characterize vascular S1P receptor function. This study was designed to use the newly available S1P receptor ligands to investigate vascular S1P receptor signaling in vitro by combining contractile effects in isolated rat aortas and designate these effects to specific S1P receptors in either smooth muscle or endothelial cells.

Materials and Methods

Materials

Medium 199, Hanks' Balanced Salt Solution (HBSS), DNaseI Amp grade and Trizol were obtained from Invitrogen (Breda, The Netherlands). Phosphate buffered saline (PBS), foetal calf serum (FCS), enzyme-free cell dissociation buffer and penicillin/streptomycin were obtained from Gibco (via Invitrogen). LANCE cAMP 384 kit and white 384-well optiplates were obtained from Perkin-Elmer (Zaventem, Belgium). SIP, (R)-Phosphoric acid mono-[2-amino-2-(3-octylphenylcarbonyl)-ethyl] ester (VPC23019), (R)-Phosphoric acid mono-[2-amino-2-(6-octyl-1H-benzimidazol-2-yl) ethyl] ester (VPC23153) and were obtained from Avanti-Polar Lipids Inc. (via InstruChemie B.V., Delfzijl, The Netherlands). Probenecid, HEPES, ethyleneglycoltetraacetate, Triton X-100, 3-isobutyl-1-methylxanthine, bovine serum albumin (regular and fatty-acid free), collagenase, elastase, gelatin, RNAlater, forskolin, sodium citrate, ATP, bovine serum albumine (BSA) and trypsin inhibitor were obtained from Sigma-Aldrich (Zwijndrecht, The Netherlands). Clear bottom black 96-well plates were obtained from Greiner Bio One (Alphen aan den Rijn, The Netherlands). Pluronic acid and Fluo-4 AM were obtained from Molecular Probes (via Invitrogen). qPCR primers were obtained from Biogio (Nijmegen, The Netherlands). N-(2,6-dichloro-4-pyridinyl)-2-[1,3-dimethyl-4-(1-methylethyl)-1H-pyrazolo[3,4-b]pyridin-6-yl]-hydrazinecarboxamide (JTE-013) was obtained from Tocris (Bristol, UK). Isopropanol, ethanol, and chloroform were obtained from Merck (Nottingham, UK). SYBR green supermix was obtained from Biorad Laboratories BV (Veenendaal, The Netherlands). Rat aortic endothelial cells (RAECs), attachment factor and rat endothelial cell growth medium were obtained from Cell Applications, Inc. (San Diego, CA). 2-Amino-2-[2-(4-octylphenyl)ethyl]-1,3-propanediol (FTY720) was obtained from ITK Diagnostics BV (Uithoorn, the Netherlands). 2-Amino-2-[2-(4-octylphenyl)ethyl]-1,3-propanediol mono(dihydrogen phosphate) ester (FTY720-P) was synthesized according to previously described methods (Albert et al., 2005). 5-[4-Phenyl-5-(trifluoromethyl)-2-thienyl]-3-[3-(trifluoromethyl)phenyl]-1,2,4-oxadiazole (SEW2871) was obtained from Cayman Chemical Company (Ann Arbor, MI)

Contraction and relaxation studies

Male Wistar rats (Charles River, Maastricht, The Netherlands) weighing 240-280 grams were anesthetized with pentobarbital (100 mg/kg i.p.) The thoracic aorta

was isolated and placed in an oxygenated Tyrode's solution (118.5 mM NaCl, 4.7 mM KCl, 1.2 mM MgSO₄, 0.025 mM Na₄EDTA, 1.8 mM CaCl₂, 1.2 mM KH₂PO₄, 5.5 mM glucose, 25 mM NaHCO₃) at room temperature and cleaned of superficial fat and loose connective tissue. Thereafter, the aorta was cut into rings of approximately 3 mm length that subsequently were mounted between two triangular stainless steel hooks in water jacketed (37 °C) organ baths containing Tyrode's solution. This was continuously gassed with a mixture of 95% O₂ / 5% CO₂. Isometric force of contraction was recorded on a Powerlab data acquisition system (AD Instruments, Bella Vista, NSW, Australia). The resting tension was set to 10 mN and the segments were allowed to equilibrate for at least 45 min. After this equilibration period the preparations were exposed thrice to a depolarizing 40 mM K⁺ solution, with 30 min intervals. The presence of a functional endothelium was assessed by precontracting the segments with 1 μM phenylephrine, and a subsequent addition of 10 μM methacholine. Segments that had a relaxation less than 80% were excluded. For all relaxation studies, the vessel segments were precontracted with 1 μM phenylephrine.

Vascular smooth muscle cell isolation and cell culture

Thoracic rat aorta was cleaned from fat-tissue and blood clots and incubated while rotating for 30 min at 30 °C in digestive medium (medium 199 supplemented with 100 U/ml penicillin and 100 μg/ml streptomycin (basal medium), collagenase 1 mg/ml, elastase 42.5 μg/ml, BSA 1 mg/ml and trypsin inhibitor 0.5 mg/ml). The adventitia was removed, the aorta was washed in basal medium and resuspended in digestive medium. After 45 min incubation at 37 °C while rotating, the aorta was cut into small pieces, which were resuspended in digestive medium and incubated for 2.5 hours at 37 °C while rotating. Basal medium plus 20 % FCS was added and the mixture was spun down. The pellet was washed twice in basal medium plus 20 % FCS and transferred to a T25 tissue culture flask coated with 0.1 % gelatine in PBS. After reaching confluency, cells were passaged 1:1 to a T75 tissue culture flask coated with 0.1 % gelatine in PBS, in basal medium plus 10 % FCS. VSMCs were passaged 1:2 after reaching confluency in tissue culture flasks coated with 0.1 % gelatine in PBS, in basal medium plus 10 % FCS.

RAECs were cultured as described in the manufacturer's protocol.

Intracellular calcium measurements

Intracellular calcium measurements were performed as previously described (Chapter 2) with minor changes. In short, cells were plated in a black, clear bottom 96 wells plate at 40,000 cells/well for RAECs, or 10,000 cells/well for VSMCs. After one day growth, cells were serum starved overnight. Cells were then loaded for 1 hour with 4 μM Fluo-4 AM ester in HBSS containing 20 mM Hepes, 0.42 % v/v pluronic acid and 2.5 mM probenecid and incubated at 37 °C. After loading, cells were washed twice with HBSS containing 20 mM Hepes and 2.5 mM probenecid and incubated at 37 °C with HBSS containing 20 mM Hepes and 2.5 mM probenecid in the presence or absence of JTE-013 (1 μM or 10 μM) for 30 min. The fluorescence signal was measured at basal level, followed by ligand stimulation, Triton (5 % v/v) addition (resulting in F_{max}) and 250 mM EGTA addition (resulting in F_{min}). The $[\text{Ca}^{2+}]_i$ was calculated via the equation: $[\text{Ca}^{2+}]_i = Kd * ((F - F_{\text{min}}) / (F_{\text{max}} - F))$. Kd being the dissociation constant of the binding of Fluo-4 to calcium (345 nM). The increase in $[\text{Ca}^{2+}]_i$ upon ligand stimulation was calculated as the difference between the $[\text{Ca}^{2+}]_i$ for the basal level and after adding a ligand.

cAMP assay

The LANCE™ cAMP 384 kit was used to determine cAMP concentrations according to the manufacturer's protocol. RAECs or VSMCs, after overnight serum-starvation, were detached from the surface using trypsin. Cells were washed once with HBSS and subsequently resuspended in buffer, containing HBSS with 0.05 % BSA (fatty-acid free) and 5 mM Hepes. Stimulation mixtures consisted of stimulation buffer with 3 μM forskolin, 0.5 mM 3-isobutyl-1-methylxanthine, with or without VPC23019 and the concentration response curve of the indicated compounds. All compounds were two times concentrated. Cells were added to the stimulation mixtures 1:1 in a 384 well optiplate at 2500 cells/well and stimulated for 1 hour. Detection of the cAMP formed during stimulation was done according to the manufacturer's protocol in a total volume of 20 μl . Measurements were done using a platereader (Victor 2, Wallac, Perkin Elmer) 3 hours after adding detection buffer and antibody mixture.

RNA isolation and cDNA synthesis

Cultured cells were detached from the cell-culture dish using Trizol. RNA was isolated as described below.

Total RNA was isolated according to the manufacturer's protocol with minor changes, using a second chloroform extraction to remove traces of phenol in the aqueous phase, a high salt solution (0.8 M sodium citrate, 1.2 M NaCl) together with isopropanol to precipitate RNA and a second wash of the RNA pellet with 75 % ethanol. RNA purity was verified on the Experion (Biorad Laboratories) and the RNA concentration was determined by spectrophotometry at the Nanodrop (Isogen Life Science). To eliminate genomic DNA contamination 1 µg of total RNA was treated with 1 µl DNase I, Amp Grade. cDNA was synthesized by reverse transcription using the iScript cDNA Synthesis kit according to the manufacturer's protocol. A control for the presence of genomic DNA, in which no cDNA was synthesized, was made for each sample. The cDNA of 1 µg RNA was diluted 1:50 for use in real-time quantitative PCR.

Primer design:

Oligonucleotide primers with a fluorophore near the 3' end containing a hairpin (non-capital letters) to render the fluorescence quenching capability, were designed using the D-LUX designer software (Invitrogen), based on sequences from the GenBank database: S1P₁ receptor (138 bp product), forward TTCCTGGTTCTGGCTGTGCT, reverse actccTTGAATTTGCCAGCGGAGtC; S1P₂ receptor (95 bp product), forward CCAACGGAGGCACTGACTAAT, reverse ATGTCTAGCCCTAAACTCGAGCC; S1P₃ receptor (79 bp product), forward cacagGCATCTTCACAGCCATTCTGGtG, reverse CTGCGGCTGCTGGACTTGAC; elongation factor-1 (EF-1, 96 bp product), forward GCAAGCCCATGTGTGTTGAA, reverse TGATGACACCCACAGCAACTG; p0 ribosomal protein (91 bp product), forward cacagaAGGGTCCTGGCTTTGTCTGtG, reverse CGCAAATGCAGATGGATCG. Each primer pair was tested for selectivity, sensitivity and PCR efficiency. Constitutively expressed EF-1 and p0 ribosomal protein were selected as endogenous controls to correct for potential variation in RNA loading.

Real-time quantitative PCR

Relative quantification of mRNA was performed on a MyiQ Single-Color Real-Time PCR Detection System (Biorad Laboratories) following the thermal protocol:

95 °C for 3 minutes to denature, 45 cycles at 95 °C for 10 s followed by 60 °C for 45 s for annealing and extension. The final reaction mixture of 15 µl consisted of diluted cDNA, 1x iQ SYBR Green Supermix, 200 nM forward primer and 200 nM reverse primer. All reactions were performed in duplicate in 96-well plates. Controls for genomic DNA were included for each cDNA sample and also a negative control containing only both primers and the iQ SYBR Green Supermix. All data were normalized to the expression of elongation factor 1 and ribosomal protein p0.

Data analysis

Concentration response curves were analyzed by fitting sigmoidal functions to the experimental data using Prism 4 (Graphpad Software, Inc., San Diego, CA, USA). Data are expressed as means ± s.e.m. Differences between groups were compared using a one way ANOVA with a Dunnett's correction or a Student's t-test where appropriate. $P < 0.05$ was considered significant.

Results

Contraction and relaxation studies

None of the tested ligands affected the applied pretension of isolated rat aortic rings (data not shown), indicating that none of the compounds induces contraction in this preparation. Relaxant responses were measured in phenylephrine-precontracted preparations. The phenylephrine (1 µM)-induced pre-contraction amounted to 17.5 ± 0.2 mN in the presence of endothelium ($n=49$) or 19.3 ± 0.5 mN ($n=14$) in the absence of endothelium. All compounds, with the exception of S1P, induced a concentration-dependent relaxation in the phenylephrine-precontracted vascular preparations (Figure 1) with maximal effects of 39.4 ± 7.1 % (FTY720, $n=5$); 36.4 ± 4.0 % (FTY720-P, $n=4$); 18.0 ± 6.6 % (VPC23153, $n=5$) or 16.2 ± 4.6 % (SEW2871, $n=4$) ($p < 0.05$). FTY720-P and its non-phosphorylated parent compound FTY720, induced comparable vasorelaxation, however FTY720-P-induced relaxation was much faster compared to FTY720 (1-2 min versus 8 min). Endothelium removal completely diminished relaxation for all compounds (Figure 2).

S1P receptor-mediated signaling in RAECs

Both, S1P and FTY720-P, potentially inhibited the forskolin-induced cAMP accumulation ($pEC_{50} = 7.8 \pm 0.2$ ($n=10$) and 8.5 ± 0.3 ($n=5$), respectively) in RAECs (Figure 3A) ($p < 0.05$). FTY720 and SEW2871 also inhibited forskolin-induced cAMP accumula-

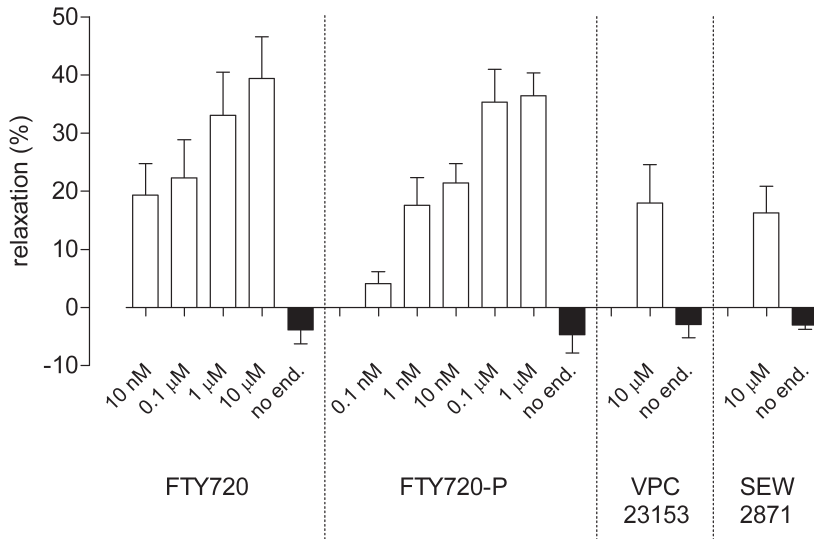


Figure 2. Ligand-induced relaxation of precontracted (1 μM phenylephrine) rat aortic rings with (white bars) and without (black bars) endothelium by FTY720, FTY720-P, VPC23153 and SEW2871. Data are shown as % relaxation of phenylephrine-induced contraction and are means ± s.e.m. of 2-5 experiments.

tion, but with a lower potency ($pEC_{50} = 6.2 \pm 0.9$ (n=5) and 5.9 ± 0.2 (n=8), respectively) than the two former compounds (figure 3A). Surprisingly, the $SIP_{1,3}$ agonist VPC23153 (Table 1, Figure 1) did not decrease but rather increased forskolin-induced cAMP accumulation ($pEC_{50} = 5.8 \pm 0.3$ (n=3)) (Figure 3B). The $SIP_{1,5}$ antagonist VPC23019 (Table 1, Figure 1) concentration-dependently shifted the FTY720-P curve to the right yielding a Schild plot with a slope close to unity and a pK_B of 7.4 (Figure 4). The SIP_2 receptor antagonist JTE-013 (Table 1, Figure 1)

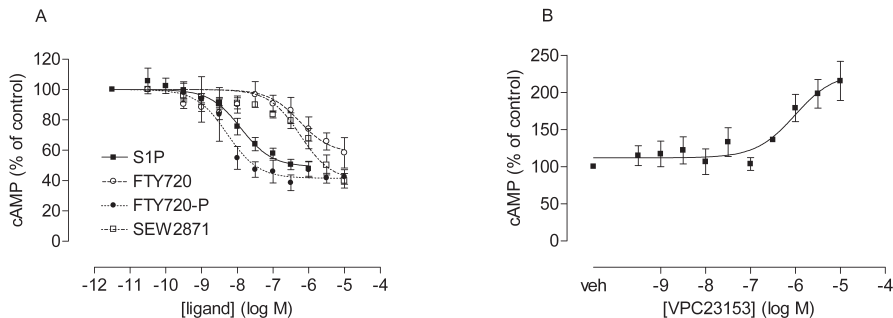


Figure 3. SIP_1 , FTY720 and FTY720-P and SEW2871-induced inhibition of forskolin (3 μM)-induced cAMP accumulation in RAECs (A). Data are expressed as % of forskolin-induced cAMP production of vehicle control (100% = 16.6 ± 3.7 nM (S1P, n=10), 19.0 ± 4.4 nM (SEW2871, n=8), 19.0 ± 6.9 nM (FTY720, n=5), 16.9 ± 5.9 nM (FTY720-P, n=5). Effect of VPC23153 on forskolin (3 μM)-induced cAMP accumulation in RAECs (B). (100% = 6.4 ± 2.1 nM (n=3)).

(10 μ M) did not affect concentration response curves of agonist-mediated inhibition of forskolin-induced cAMP levels (data not shown). Both S1P and FTY720-P slightly increased $[Ca^{2+}]_i$ in RAECs but only at high concentrations (1 μ M) (107 ± 17 nM and 96 ± 43 nM, respectively), whereas the other ligands failed to do so.

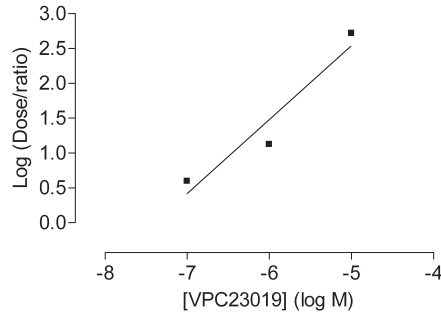


Figure 4. Schild plot of the effect of VPC23019 on the FTY720-P-induced inhibition of forskolin-induced cAMP accumulation in RAECs. Cells were pre-incubated for 30 min with the indicated concentrations of VPC23019. (n=6). Slope = 1.06, $pK_b = 7.4$.

S1P receptor-mediated signaling in VSMCs

In contrast to the endothelial cells, S1P, FTY720-P and SEW2871 did not affect forskolin-induced cAMP accumulation in rat aortic VSMCs to an extent that was reliably quantifiable. However, S1P induced an elevation of $[Ca^{2+}]_i$ in VSMCs, which was inhibited by JTE-013 (Figure 5) ($p < 0.05$). This antagonist did not significantly inhibit ATP-induced increases in $[Ca^{2+}]_i$ (vehicle: $pEC_{50} = 5.5 \pm 0.1$, $E_{max} = 1539 \pm 402$ nM (n=5); JTE-013 (1 μ M): $pEC_{50} = 5.4 \pm 0.2$, $E_{max} = 1404 \pm 403$ nM (n=4)). VPC23019, VPC23153, FTY720 and FTY720-P did not elevate $[Ca^{2+}]_i$.

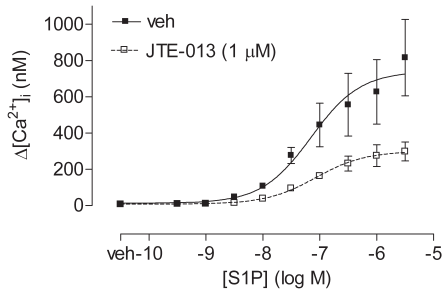


Figure 5. The effect of JTE-013 (1 μ M) on concentration-dependent S1P-induced elevation of $[Ca^{2+}]_i$ in VSMCs. $E_{max} = 816 \pm 211$ nM (veh), $298 \pm 52^*$ nM (1 μ M) and $180 \pm 40^*$ nM (10 μ M). * $p < 0.05$ (n=5).

S1P receptor mRNA expression

Quantitative PCR showed the presence of S1P₁, S1P₂ and S1P₃ receptor mRNA in VSMCs (C_t value ranging from 30-32 (n=3), corrected for P0 and EF-1). RAECs only showed S1P₁ receptor mRNA expression (approximately 46 times higher than in VSMCs) (C_t value around 25 (n=3), corrected for P0 and EF-1), whereas mRNA for the S1P₂ and S1P₃ receptors was not detected up to cycle 40.

Discussion

Besides regulators of growth, sphingolipids (including S1P) are vasoactive compounds. S1P has been shown to induce vasodilation and vasoconstriction, amongst others depending on vessel type, animal species and technique used (reviewed in Hemmings, 2006). These processes are most likely mediated by high affinity S1P receptors, although the existence of intracellular targets of S1P cannot be ruled out (Payne et al., 2002). Within the vasculature, mainly the S1P₁₋₃ receptor subtypes are expressed and the ultimate effect of S1P on a given vessel is, therefore, dependent on the differential expression of these receptor subtypes in the endothelium and smooth muscle cells. This study was performed to use agonists and antagonists of the S1P receptors that have become recently available to pharmacologically characterize S1P receptor-dependent signaling in intact isolated rat aortic rings and cultured rat aortic endothelial and vascular smooth muscle cells. The immunosuppressant drug candidate FTY720 (fingolimod, now being developed for the treatment of multiple sclerosis (Kappos et al., 2006)) can be phosphorylated by sphingosine kinase and in its phosphorylated form (FTY720-P) is a potent agonist for all, except the S1P₂ receptor subtype (Mandala et al., 2002). VPC23153 is described to be an agonist at the human S1P₁ and S1P₃ receptor, whereas VPC23019 has been reported to have antagonistic effects at these receptors. SEW2871 is a highly selective S1P₁ agonist and JTE013 displays antagonistic activity towards the S1P₂ receptor (Table 1, Figure 1).

Both endothelial and vascular smooth muscle cells express S1P receptors (Michel et al., 2007) and the intracellular signaling pathways these receptors activate are known to affect vascular tone (Alewijns et al., 2004; Young & Van Brocklyn, 2006; Hemmings, 2006). Indeed, all of the aforementioned compounds tested in this study, with the exception of S1P, modulated vascular tone in rat aortic rings as they induced a concentration-dependent vasorelaxation in pre-constricted preparations (Figure 1). Under non-preconstricted conditions all compounds were devoid of any

activity. The observed relaxation was abolished by endothelium removal. qPCR analysis in cultured RAECs showed high mRNA expression of the G_i -coupled $S1P_1$ receptor, whereas $S1P_2$ and $S1P_3$ receptor mRNA was not detected. Intracellular calcium levels in RAECs were only slightly increased by high concentrations (1 μ M) of S1P or FTY720-P (less than 20% compared to VSMC responses), which might be due to non-specific effects.

With the exception of VPC23153, all tested agonists (FTY720, FTY720-P, S1P and SEW2871) inhibited forskolin-induced cAMP accumulation in RAECs. The order of potency of these compounds in this assay matched with their $S1P_1$ receptor potency (Table 1, Figure 1). VPC23153 did not decrease, but rather increased the forskolin-induced cAMP accumulation, indicating the possibility of inverse agonism. Both, increase and decrease in cAMP production in ECs were associated with vasorelaxation, indicating that this effect is not crucial for the final vasorelaxant effect. Therefore, the endothelium-dependent vasorelaxant effect seen in aorta preparations is most likely mediated via $\beta\gamma$ -subunits of G_i , as was recently shown (Gonzalez et al., 2006).

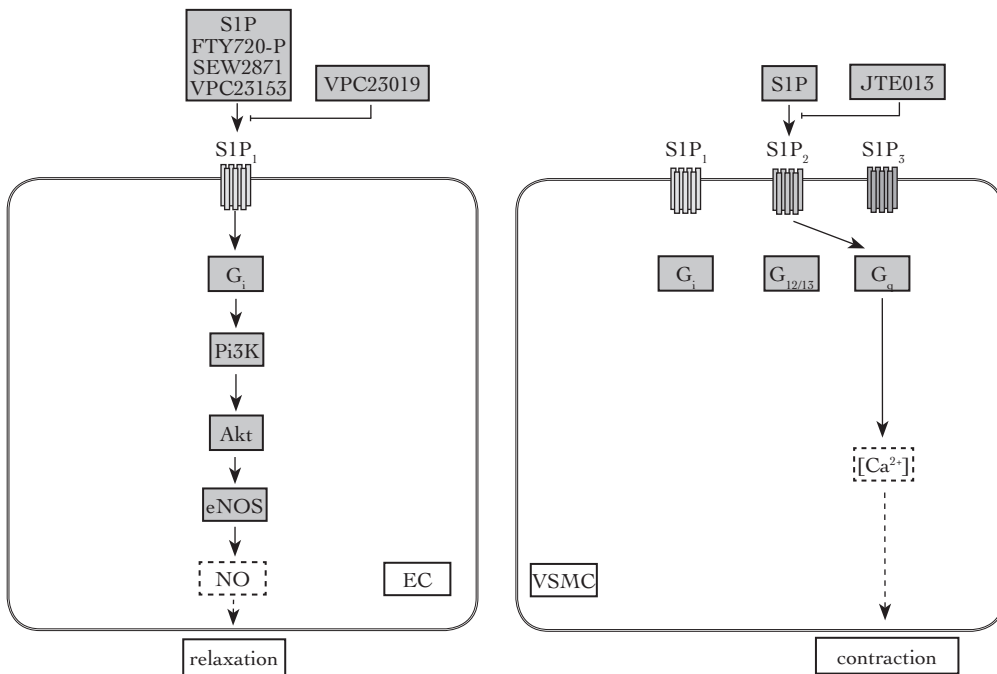


Figure 6. Schematic representation of the effect of S1P receptor ligands in ECs and VSMCs. $S1P_1$, $S1P_2$ and $S1P_3$ receptors activate signaling pathways leading to contraction or relaxation. Activation of the $S1P_1$ and probably also $S1P_3$ receptors in ECs lead to relaxation. In VSMCs, activation of $S1P_2$ receptors leads to contraction.

qPCR showed mRNA expression for the $S1P_1$, $S1P_2$ and $S1P_3$ receptors in cultured VSMCs. In line with these findings, FTY720, FTY720-P and SEW2871 inhibited forskolin-induced cAMP accumulation in these cells. However, this effect was relatively small, making reliable quantification not possible. Increases in $[Ca^{2+}]_i$ were only observed after stimulation with S1P, but not with any of the other ligands, which all lack $S1P_2$ agonism. Additionally, JTE-013 inhibited this S1P-induced elevation of $[Ca^{2+}]_i$. These findings indicate that in smooth muscle cells, S1P effects are predominantly mediated via the $S1P_2$ receptor.

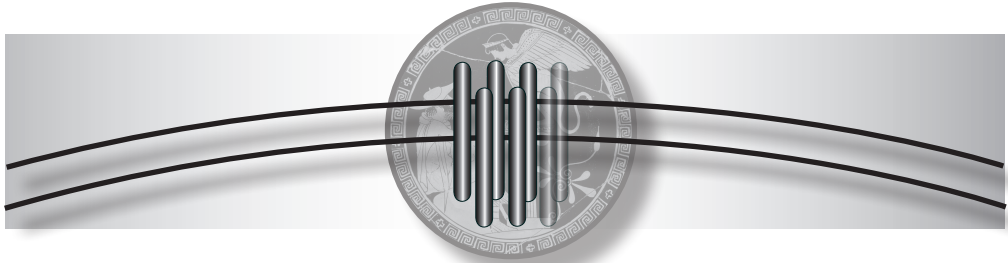
An overview of the effects of S1P and the new synthetic S1P receptor ligands on either EC and/or VSMC is drawn in figure 6. Our results show that in intact aortic rings, endothelium-dependent relaxation is induced by all synthetic agonists, most likely via the activation of the $S1P_1$ receptor on endothelial cells. However, the presence and influence of the $S1P_3$ receptor in these cells can not be ruled out. On VSMCs, only the ligand with $S1P_2$ agonistic activity, being S1P itself, increased $[Ca^{2+}]_i$, indicating that S1P signaling in these cells occurs predominantly via the $S1P_2$ receptor. Specific $S1P_2$ receptor activation in aortic rings can however not be investigated as these compounds do not act on $S1P_2$ receptors. Therefore, the net effect of these new ligands in the aorta shall be relaxation. In disease states that are characterized by vascular dysfunction, e.g. hypertension, diabetes or atherosclerosis, subtype selective S1P receptor agonist/antagonists may be used as modulators of vascular tone.

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*Differential agonist-induced internalization of
sphingosine-1-phosphate receptor
subtypes 1, 2 and 3*



Chapter 5

Differential agonist-induced internalization of sphingosine-1-phosphate receptor subtypes 1, 2 and 3

Manuscript in preparation

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Abstract

Regulation of sphingosine-1-phosphate (SIP) receptors is of major importance especially since it has been shown to be the mechanism of action for the immunosuppressive prodrug FTY720. Little is known about possible differential regulation of the different SIP receptor subtypes expressed in the cardiovascular system. This study was therefore set up to compare ligand-induced internalization of the SIP_{1,3} receptor subtypes. A recently developed immunocytochemical method was used to characterize internalization and resensitization of membrane expressed HisG-tagged SIP receptors. Inhibitors of several downstream signaling pathways were used to investigate the mechanism of receptor internalization. SIP₃ receptor internalization showed striking differences compared to SIP₁ and SIP₂ receptor internalization as it occurred much faster, required lower SIP concentrations and showed a higher reduction in the fraction of membrane receptors. The differences in signaling between the three receptor subtypes could only partly explain the differential internalization of these receptors. For the SIP₃ receptor activation of G_i-proteins is involved in SIP₃ receptor internalization, whereas elevations in [Ca²⁺]_i or activation of G_q-signaling are not important.

In conclusion, the SIP receptor subtypes expressed in the cardiovascular system differ in ligand-induced internalization, with G_i-activation being of influence. This difference could be important for the functioning of drugs targeted at SIP receptors, including FTY720.

Introduction

The bioactive lipid sphingosine-1-phosphate (S1P) is involved in the regulation of many important cellular processes such as migration and proliferation and lymphocyte trafficking. Most of the effects of S1P are mediated via five distinct G-protein coupled receptors (GPCRs), named S1P₁₋₅ (previously EDG-1,5,3,6 and 8, respectively) (Chun et al., 2002). The S1P₁, S1P₂ and S1P₃ receptors are ubiquitously expressed and are the most important sphingolipid receptors in the cardiovascular system (Spiegel & Milstien, 2000; Alewijnse et al., 2004), the S1P₄ receptor is mainly expressed in the lung and lymphoid tissues and the S1P₅ receptor in the white matter of the central nervous system (Anliker & Chun, 2004). S1P can be generated in many cell types by the concerted action of specific enzymes and its synthesis is tightly controlled through the regulation of activity of specific enzymes such as sphingosine kinase (Spiegel & Milstien, 2003).

In general, exposure of GPCRs to agonists initiates a series of regulatory processes, some of which directly affect receptor responsiveness. One of these mechanisms involves arrestin binding upon phosphorylation of the receptor by G-protein coupled receptor kinases. This uncouples the receptor from the G-protein and results in termination of signaling. Another mechanism is internalization of the receptor, either arrestin-dependent or independent (Bünemann & Hosey, 1999), and involves translocation of the receptor from the plasma membrane to clathrin-coated pits inside the cell from where it can either be recycled to the membrane or undergo degradation. It is nowadays well accepted that GPCR internalization not only regulates the functional activity and number of receptors present at the cell surface, but that it also mediates GPCR coupling to certain downstream effector pathways (von Zastrow, 2001).

As found for other GPCRs, S1P receptors are also subject to receptor regulatory processes upon ligand stimulation. High concentrations of S1P, 200-900 nM (Okajima, 2002), have been found in plasma and serum, suggesting that S1P is present in blood. The origin of S1P in blood, recently studied in mouse, was found to be mainly haematopoietic with erythrocytes being the most important contributor (Pappu et al., 2007). The concentrations of S1P in blood are sufficient to stimulate S1P receptors on cardiovascular cells such as endothelial cells and probably also smooth muscle cells, but it is not known whether this actually occurs, since the majority of S1P in blood is bound to lipoproteins, stored in platelets or bound to albumin (Okajima, 2002). Although it is still unclear if bound S1P is indeed func-

tionally active, regulatory processes can be speculated to be important in the cardiovascular system. Interestingly, SIP receptor internalization is suggested to be the mechanism of action of FTY720, an immunosuppressive prodrug that is now in clinical trials for the treatment of multiple sclerosis (Kappos et al., 2006). Although FTY720, once phosphorylated in vivo by sphingosine kinase-2, is a potent agonist for four of the five SIP receptors (SIP_{1,3,5}) (Mandala et al., 2002), the lymphopenia induced by this drug is suggested to be mainly due to internalization of the SIP₁ receptor expressed on lymphocytes (Matloubian et al., 2004). Besides the importance of SIP₁ receptor regulation in the immune system, this process may also be of major importance for the SIP receptors in the cardiovascular system. Except from some recent work on the SIP₁ and SIP₃ receptor (Liu et al., 1999; Spiegel & Milstien, 2000; Rutherford et al., 2005), relatively little is known about regulation of the SIP receptors. Therefore, we have investigated differences in regulation upon agonist treatment of the SIP receptor subtypes important in the cardiovascular system. For this purpose, we compared internalization of the SIP₁, SIP₂ and SIP₃ receptors using a recently described method that enables quantification of receptor internalization (Chapter 3). Our research revealed that these three SIP receptor subtypes differ strikingly in ligand-induced internalization and that these differences can partly be explained by differences in signaling properties between these receptor subtypes. For the SIP₃ receptor the internalization is partly dependent on G_i-activation whereas the activation of G_q-signaling pathways was not important.

Materials & Methods

Materials

pcDNA3.1 containing the entire coding region of human SIP_{1,3} receptors was purchased from UMR cDNA Resource Center (Rolla, MO, USA). Cell culture media, hygromycin B, LipofectamineTM2000, pOG44, CHO-FlpIn cells, Alexa Fluor[®] 488 goat anti-mouse (IgG), anti-HisG (mouse monoclonal IgG2a), anti-HisG-horse radish peroxidase (HRP) (mouse monoclonal IgG2a) and pcDNA5/FRT/TO were obtained from Invitrogen (Breda, The Netherlands). Restriction enzymes (ApaI, BamHI, XhoI, EcoRI and HindIII) were obtained from Fermentas Life Sciences (St. Leon-Rot, Germany). Black, clear bottom 96 well plates were obtained from Greiner Bio One (Alphen aan den Rijn, The Netherlands). SIP was obtained from Avanti-Polar Lipids, Inc (via Instruchemie B.V., Delfzijl, the Netherlands). Pertussis toxin, A23187, BAPTA-AM (1,2-Bis (2-aminophenoxy)ethane-N,N',N'-

tetraacetic acid, tetraacetoxymethyl ester), calphostine-C, U73122 (1-[6-[(17 β)-3-Methoxyestra-1,3,5[10]-trien-17-yl)amino]hexyl]-1H-pyrrole-2,5-dione) and Rp-cAMPS (Rp-Cyclic 3',5'-hydrogen phosphorothioate adenosine triethylammonium salt) were obtained from Sigma-Aldrich (Zwijndrecht, the Netherlands).

Molecular cloning

A N-terminal HisG-tag was added to the SIP_{1,3} receptors via cloning into pcDNA3.1/HisC using BamHI&XhoI (for SIP₁) or BamHI&ApaI (for SIP₂) or into pcDNA3.1/HisA using BamHI&XhoI (for SIP₃). A second cloning step was done using HindIII&XhoI (for SIP_{1,3}) or HindIII&ApaI (for SIP₂) to clone the HisG-tagged SIP receptors into the expression vector pcDNA5/FRT/TO. The cDNA of the SIP₃ receptor obtained from Guthrie contained a point mutation (G⁹⁶²A which results in R³²¹Q) which was removed by PCR using the following primers (forward: TGCCTGGTCAGGGGGCGGGGGGCCCG, reverse: CGGGCCCCCGCCCCCTGACCAGGCA). The SIP_{1,3} receptor coding DNA in the resulting plasmid as well as the stable cell line constructed with this plasmid have been confirmed by sequencing.

Transfection and cell culture

Generation of stable CHO-FlpIn cell lines expressing the indicated receptors and culturing of these cell lines was done as described previously (Chapter 2). The cell lines were checked for expression of the indicated HisG-SIP receptors by immunocytochemistry as described below. These experiments revealed no major differences in membrane fluorescence for the three receptor subtypes.

Calcium assay and cAMP assay

Intracellular calcium concentrations ($[Ca^{2+}]_i$) and cAMP levels were measured as described previously (Chapter 2).

Immunocytochemistry

Immunocytochemistry was performed as described previously (Chapter 3). In short, an N-terminal HisG-tag was added to the receptors which can be detected using a 1st anti-HisG antibody combined with a AlexaFluor® 488 goat-anti-mouse fluorescent 2nd antibody. The fluorescent signal can be measured on a microplate reader. Some adaptations were made for specific applications of this method. After overnight serum starvation, cells were stimulated with the indicated ligand in

serum free medium for 30 minutes at 37 °C (for concentration response curves), stimulated with 1 μ M S1P for the indicated time (for time course stimulations) or stimulated with 1 μ M S1P for 30 minutes at 37 °C followed by 3 times washing with serum free medium and incubation at 37 °C with serum free medium for the indicated time (for resensitization).

Data analysis

Concentration response curves were analyzed by fitting sigmoidal functions to the experimental data using Prism 4 (Graphpad Software, Inc., San Diego, CA, USA). Data are expressed as means \pm s.e.m. One-way ANOVA with a Dunnett's correction or Student's t-test was applied where appropriate. $P < 0.05$ was considered significant.

Results

Time course of internalization of the HisG-S1P₁, HisG-S1P₂ and HisG-S1P₃ receptor

Internalization of the different receptor subtypes was studied by determining the decrease of membrane receptors by an immunocytochemical method. For all receptors studied, incubation with 1 μ M S1P induced a time-dependent decrease in cell surface receptors. Internalization of the HisG-S1P₃ receptor was significantly faster ($t_{1/2} = 1.1 \pm 0.2$ min (n=4)) than of the HisG-S1P₁ ($t_{1/2} = 16.6 \pm 7.0$ min (n=4)) and HisG-S1P₂ receptors ($t_{1/2} = 19.1 \pm 5.4$ min (n=4)) ($p < 0.05$) (Figure 1). All further internalization experiments were done with 30 min agonist exposure.

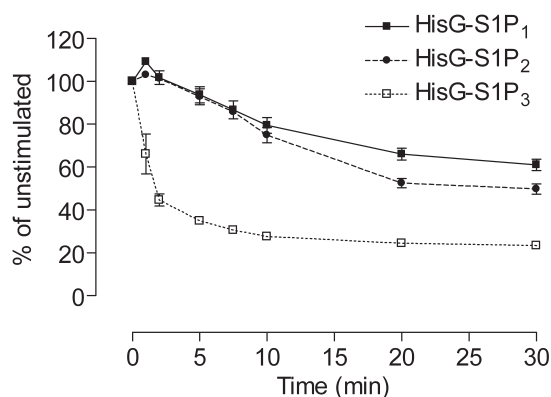


Figure 1. Time dependent S1P-induced internalization of HisG-S1P₁, HisG-S1P₂ or HisG-S1P₃ receptor subtypes. Stimulations with S1P (1 μ M) were carried out at 37 °C for the indicated time. Values are expressed as percentage of fluorescent signal measured for unstimulated cells ($t = 0$ min) which was set at 100% and are presented as means \pm s.e.m. (100% was for HisG-S1P₁: 54978 \pm 5892 RFU (n=5); HisG-S1P₂: 65091 \pm 4610 RFU (n=5); HisG-S1P₃: 62013 \pm 4941 RFU (n=5)).

Potency of internalization of the HisG-S1P₁, HisG-S1P₂ and HisG-S1P₃ receptor

The potency of S1P to induce internalization was similar for the HisG-S1P₁ and HisG-S1P₂ receptor ($pEC_{50} = 7.5 \pm 0.2$ (n=8) and 7.1 ± 0.2 (n=4)), while the HisG-S1P₃ receptor was more sensitive to S1P-induced internalization ($pEC_{50} = 8.0 \pm 0.2$ (n=3)) (Figure 2). Besides a difference in pEC_{50} between the subtypes in response to S1P, also the relative amount of internalization differed. Thus, the HisG-S1P₃ receptor internalized to a lower level than the HisG-S1P₁ or the HisG-S1P₂ receptor (HisG-S1P₃: 27 ± 3 % (n=3) vs HisG-S1P₁: 58 ± 2 % (n=8) and HisG-S1P₂: 42 ± 2 % (n=4), $p < 0.05$).

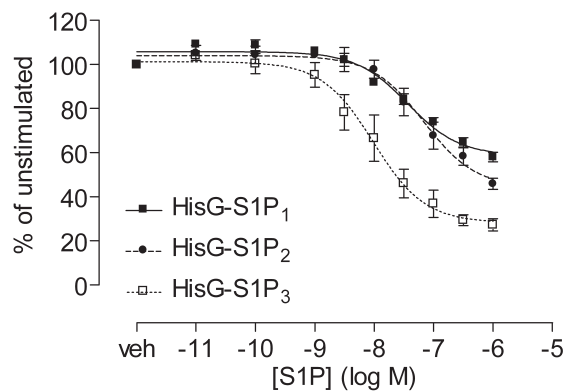


Figure 2. Concentration-dependent S1P-induced internalization of HisG-S1P₁, HisG-S1P₂ or HisG-S1P₃ receptor subtypes. Stimulations were carried out at 37 °C for 30 min. Values are expressed as percentage of fluorescent signal measured for unstimulated cells which was set at 100% and are presented as means \pm s.e.m. (100% was for HisG-S1P₁: 49374 ± 3620 RFU (n=8); HisG-S1P₂: 63161 ± 3235 RFU (n=4); HisG-S1P₃: 53837 ± 5899 RFU (n=3)).

Resensitization

Recycling of S1P receptors to the membrane was studied. The washing procedure that followed initial S1P or vehicle stimulation reduced the amount of membrane receptors for the HisG-S1P₃ receptor, but not for the HisG-S1P₁ or HisG-S1P₂ receptor. Therefore, we quantified the difference in membrane fluorescence between 30 min vehicle-treated and S1P-stimulated cells at different times after the initial stimulation. Significant differences remained in this regard for 1.5 h after washout of S1P for the HisG-S1P₂ receptor (Figure 3B) and for 2 h after washout of S1P for the HisG-S1P₁ receptor (Figure 3A). For the HisG-S1P₃ receptor the difference between the vehicle and S1P treated cells remained significant for at least 4 h after washout of S1P, but the fluorescence level was only restored to 80 % (yet still increasing) of the initial level (Figure 3C).

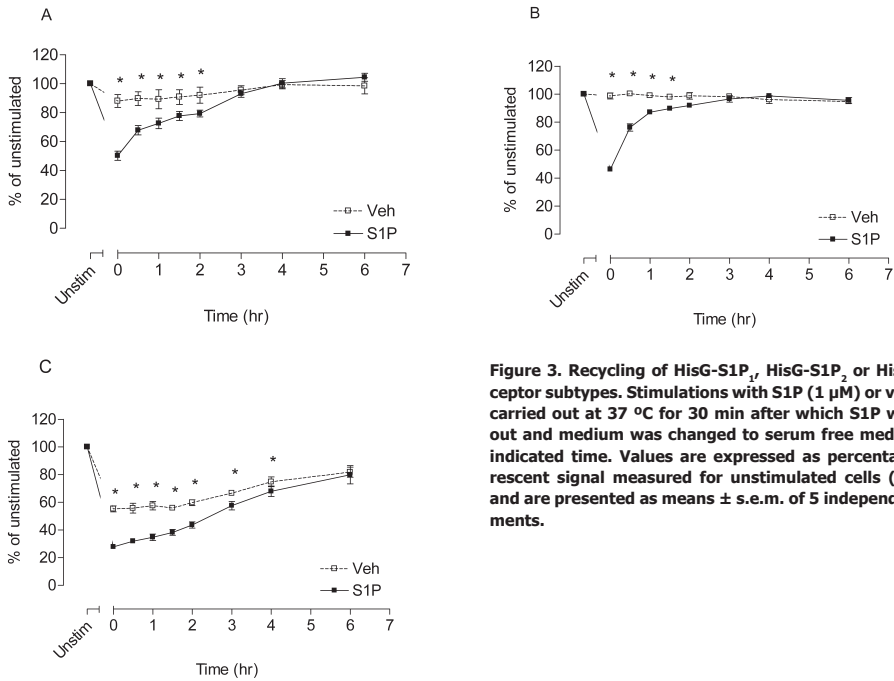


Figure 3. Recycling of HisG-S1P₁, HisG-S1P₂ or HisG-S1P₃ receptor subtypes. Stimulations with S1P (1 μM) or vehicle were carried out at 37 °C for 30 min after which S1P was washed out and medium was changed to serum free medium for the indicated time. Values are expressed as percentage of fluorescent signal measured for unstimulated cells (t = 0 min) and are presented as means ± s.e.m. of 5 independent experiments.

Signaling of the HisG-S1P₁, HisG-S1P₂ and HisG-S1P₃ receptor

Previously it was shown that the HisG tag did not affect signaling of the S1P₁ receptor as no difference in the potency of S1P to inhibit forskolin-induced cAMP accumulation in the cells expressing the tagged or untagged S1P₁ receptor was seen (Chapter 3). Also, no difference was observed after addition of the HisG tag on signaling of the S1P₂ and S1P₃ receptor (data not shown). There was no change in the

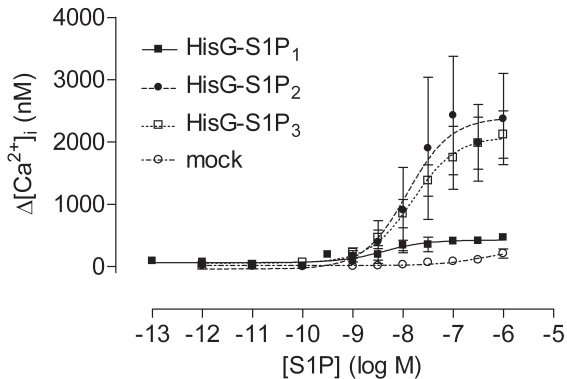


Figure 4. S1P-induced increases in [Ca²⁺]_i in mock transfected CHO-FlpIn or in CHO-FlpIn cells expressing the HisG-S1P₁, HisG-S1P₂ or HisG-S1P₃ receptor or in mock transfected cells. Data presented are means ± s.e.m. of 3 to 6 independent experiments.

potency of S1P to inhibit forskolin-induced cAMP accumulation in cells expressing the S1P₃ receptor and the HisG tag did not influence the potency of S1P to raise intracellular calcium levels in cells expressing the S1P₂ receptor (data not shown). S1P slightly but significantly increased elevations in [Ca²⁺]_i in cells expressing the HisG-S1P₁ receptor compared to mock transfected cells (HisG-S1P₁: E_{max} = 499 ± 29 nM (n=3) and mock: E_{max} = 205 ± 74 nM (n=3)). Cells expressing the HisG-S1P₂ receptor or the HisG-S1P₃ receptor showed significantly higher elevations in [Ca²⁺]_i upon S1P stimulation compared to mock transfected cells and cells expressing the HisG-S1P₁ receptor (HisG-S1P₂: pEC₅₀ = 7.6 ± 0.3; E_{max} = 2614 ± 813 nM (n=4) and HisG-S1P₃: pEC₅₀ = 7.9 ± 0.3; E_{max} = 2166 ± 438 nM (n=6)) (Figure 4).

S1P did not inhibit the forskolin-induced cAMP accumulation in mock transfected cells (data not shown) or in cells transfected with the HisG-S1P₂ receptor (Figure 5). However, S1P inhibited forskolin-induced cAMP accumulation in cells expressing the HisG-S1P₁ (pEC₅₀ = 9.5 ± 0.2; E_{max} = 22 ± 4 % (n=11)) or HisG-S1P₃ receptor (pEC₅₀ = 8.6 ± 0.2; E_{max} = 21 ± 1 % (n=9)) (Figure 5). Surprisingly, S1P increased the forskolin-induced cAMP accumulation via the HisG-S1P₂ receptor (pEC₅₀ = 7.0 ± 0.2; E_{max} = 170 ± 11 % (n=3)).

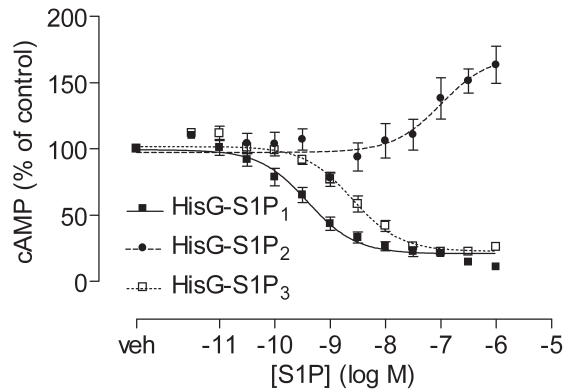


Figure 5. S1P-mediated effect on forskolin-induced cAMP accumulation in cells expressing the HisG-S1P₁, HisG-S1P₂ or HisG-S1P₃ receptor. Stimulations were carried out for 60 min at room temperature. The forskolin-induced cAMP accumulation in the absence of S1P was set at 100% and all values were expressed compared to this control (100 % was for HisG-S1P₁: 14.4 ± 1.9 nM (n=11); HisG-S1P₂: 4.3 ± 0.7 nM (n=3); HisG-S1P₃: 15.7 ± 1.3 nM (n=9)).

Influence of pertussis toxin (PTX) on internalization of the HisG-S1P₁, HisG-S1P₂ and HisG-S1P₃ receptor

The S1P-induced internalization of the HisG-S1P₁ and HisG-S1P₂ receptor were not significantly altered by overnight incubations with PTX (100 ng/ml (S1P-induced inhibition of cAMP accumulation was completely abolished under these conditions)) ($pEC_{50} = 7.2 \pm 0.1$ vs 6.9 ± 0.1 (n=4) for HisG-S1P₁; 6.8 ± 0.1 vs 6.8 ± 0.1 for HisG-S1P₂ (n=3)) (Figure 6 A,B). However, PTX slightly but significantly attenuated S1P-induced internalization of the HisG-S1P₃ receptor ($pEC_{50} = 8.1 \pm 0.1$ (without PTX) and 7.3 ± 0.1 (with PTX) (n=4)) (Figure 6C).

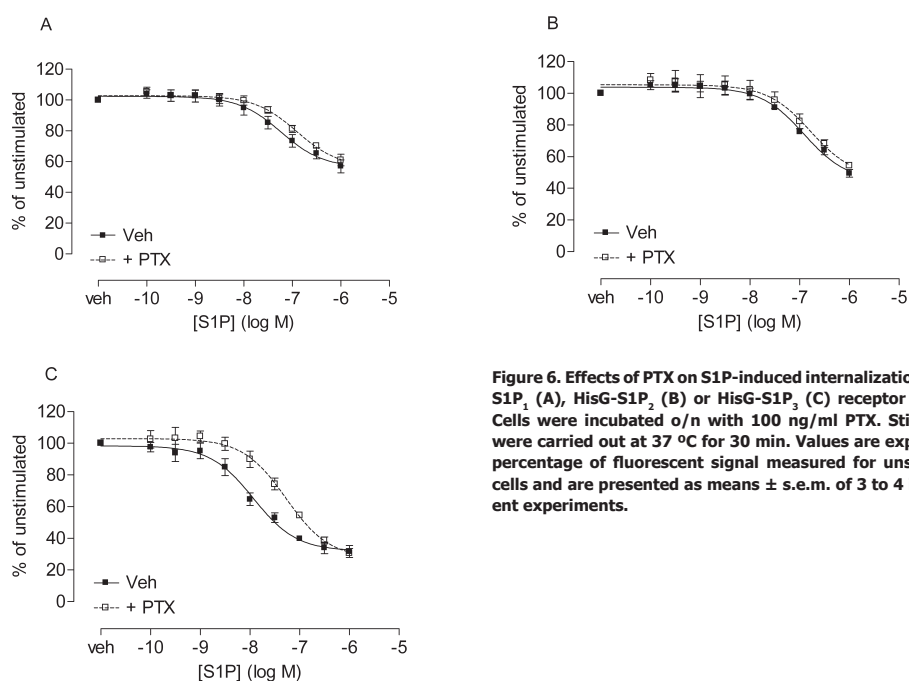


Figure 6. Effects of PTX on S1P-induced internalization of HisG-S1P₁ (A), HisG-S1P₂ (B) or HisG-S1P₃ (C) receptor subtypes. Cells were incubated o/n with 100 ng/ml PTX. Stimulations were carried out at 37 °C for 30 min. Values are expressed as percentage of fluorescent signal measured for unstimulated cells and are presented as means \pm s.e.m. of 3 to 4 independent experiments.

Signal transduction involved in internalization of HisG-S1P₁, HisG-S1P₂ and HisG-S1P₃ receptors

Pre-incubations with either the protein kinase A (PKA) inhibitor Rp-cAMPS (100 μ M), the phospholipase C (PLC) inhibitor U73122 (10 μ M) or the protein kinase C (PKC) inhibitor calphostine-C (1 μ M) did not alter S1P-induced HisG-S1P₃ receptor internalization (veh: $pEC_{50} = 7.8 \pm 0.2$ (n=5); Rp-cAMPS: $pEC_{50} = 7.9 \pm 0.2$ (n=4); U73122: $pEC_{50} = 8.0 \pm 0.1$ (n=3); calphostine-C: $pEC_{50} = 8.1 \pm 0.4$ (n=3)) (Table 1). The S1P-induced internalization of the other two subtypes was also not significantly altered by the different inhibitors (Table 1).

Table 1. Influence of Rp-cAMPS (100 μ M), calphostin-C (1 μ M) and U73122 (10 μ M) on SIP-induced internalization of the HisG-S1P₁, HisG-S1P₂ or HisG-S1P₃ receptor subtypes. Stimulations were carried out at 37 °C for 30 min. Values are expressed as percentage of fluorescent signal measured for unstimulated cells and are presented as means \pm s.e.m.

	veh	Rp-cAMPS	U73122	calphostine-C
HisG-S1P ₁	7.1 \pm 0.1 (5)	7.0 \pm 0.2 (4)	7.0 \pm 0.1 (2)	7.2 \pm 0.2 (3)
HisG-S1P ₂	6.8 \pm 0.0 (5)	6.8 \pm 0.1 (4)	6.5 \pm 0.2 (3)	6.7 \pm 0.2 (3)
HisG-S1P ₃	7.8 \pm 0.2 (5)	7.9 \pm 0.2 (4)	8.0 \pm 0.1 (3)	8.1 \pm 0.4 (3)

Addition of the calcium ionophore A23187 (10 μ M) did not influence HisG-S1P₁ receptor internalization ($pEC_{50} = 7.0 \pm 0.1$ vs 7.0 ± 0.2 ($n=3$) (vehicle vs A23187)) (Figure 7A). Addition of the calcium chelator BAPTA-AM (10 μ M) also had no significant influence on HisG-S1P₃ receptor internalization ($pEC_{50} = 7.6 \pm 0.2$ vs 8.2 ± 0.2 . ($n=3$) (vehicle vs BAPTA-AM)) (Figure 7B). This BAPTA-AM concentration fully suppressed increases in $[Ca^{2+}]_i$ in CHO cells expressing the HisG-S1P₃ receptor, while the used A23187 concentration induced a similar increase in $[Ca^{2+}]_i$ compared to SIP (data not shown).

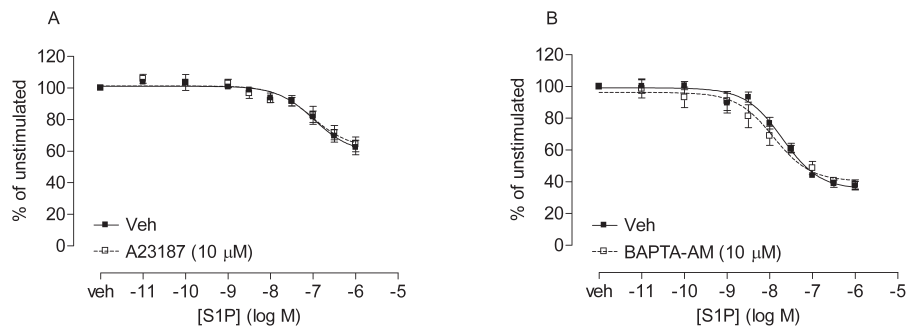


Figure 7. Influence of A23187 (10 μ M) and BAPTA-AM (10 μ M) on SIP-induced internalization of HisG-S1P₁ and HisG-S1P₃ receptor subtypes. Stimulations were carried out at 37 °C for 30 min. Values are expressed as percentage of fluorescent signal measured for unstimulated cells and are presented as means \pm s.e.m. of 3 independent experiments.

Discussion

Like most GPCRs, SIP receptors are susceptible to regulatory processes upon ligand stimulation. The interest for this mechanism has increased drastically since it has been the suggested mechanism of action for the immunosuppressive prodrug FTY720 (Gräler & Goetzl, 2004). We have used a previously reported, optimized method (Chapter 3) to compare SIP₁, SIP₂ and SIP₃ receptor internalization. Internalization of the SIP₁ receptor has been previously studied (Liu et al., 1999) using a C-terminal GFP-tagged SIP₁ receptor in HEK293 cells. Interestingly, the data obtained in that study and the present study for the SIP₁ receptor are in good agreement. The time to internalize half of the SIP₁ receptors ($t_{1/2}$) is comparable (~15 min). In addition, resensitization was also comparable, reaching about 80% of the pretreatment level of receptors at 2 h after SIP washout. This confirms that SIP receptor internalization can be validly studied using our newly developed method. Comparing internalization of the three SIP receptors ubiquitously expressed in the cardiovascular system, namely the SIP₁, SIP₂ and SIP₃ receptor (Alewijnse et al., 2004), revealed some interesting differences between the three receptor subtypes. While time-dependent SIP-induced internalization for the HisG-SIP₁ and HisG-SIP₂ receptor was comparable, the HisG-SIP₃ receptor was internalized much faster. In addition, the potency of SIP to induce internalization was significantly higher at the HisG-SIP₃ receptor compared to the other two subtypes. Furthermore, SIP-stimulation reduced membrane expression of HisG-SIP₃ receptor to lower levels than the HisG-SIP₂ receptor, which was in turn reduced to lower levels than the HisG-SIP₁ receptor. Finally, resensitization of the HisG-SIP₃ receptor to the membrane was slower than for the HisG-SIP₁ and HisG-SIP₂ receptor. Different SIP receptor subtypes activate different signaling pathways (Spiegel & Milstien, 2000). Could this be underlying the observed differences in internalization? When comparing signaling, the HisG-SIP₁ receptor inhibits cAMP accumulation, while the small SIP-induced increases in $[Ca^{2+}]_i$ seen in cells expressing the HisG-SIP₁ receptor are most mediated via $\beta\gamma$ -subunits of G_i , although an effect of endogenous SIP receptors is also possible (Holdsworth et al., 2005). The HisG-SIP₂ receptor induces elevations in $[Ca^{2+}]_i$ and surprisingly elevations of cAMP accumulation, even though the SIP₂ receptor is generally reported to activate G_i -proteins. This effect has been observed before in CHO cells overexpressing the SIP₂ receptor, but has not yet been fully understood (Windh et al., 1999). Stimulation of the HisG-SIP₃ receptor results in both inhibition of cAMP accumula-

tion and elevations of $[Ca^{2+}]_i$. Interestingly, pre-treatment with PTX, an inhibitor of G_i -activation, shifted the SIP-induced internalization curve of the HisG-SIP₃ receptor to the right, whereas it did not affect SIP-induced internalization of the HisG-SIP₁ and HisG-SIP₂ receptor. Thus, activation of the G protein influences internalization of the SIP₃ receptor, but not the SIP₁ or SIP₂ receptor.

We investigated internalization of these receptors in more detail by modulating both G_i and G_q -coupled signaling pathways. Incubations with Rp-cAMPs, an inhibitor of PKA, did not alter internalization. Thus, although inhibition of G_i -activation by PTX treatment showed an effect on internalization, inhibition of the downstream target PKA is not involved. Since there is no commercially available inhibitor of G_q -activation, we inhibited downstream signaling using inhibitors of the enzymes activated by the G_q -signaling cascade (U73122 for PLC and calphostin-C for PKC). Incubation with either inhibitor did not have any effect on SIP-induced internalization of any of the three SIP receptor subtypes. Thus, inhibiting the G_q -pathway at different levels has no influence on SIP-induced receptor internalization.

To study whether the $[Ca^{2+}]_i$, which is increased via SIP₃ receptor activation but not via SIP₁ receptor activation, influenced internalization, we applied the calcium ionophore A23187 to HisG-SIP₁ expressing cells to increase $[Ca^{2+}]_i$ and applied a calcium chelator BAPTA-AM to HisG-SIP₃ expressing cells to eliminate a possible effect on $[Ca^{2+}]_i$. Both approaches did not show any influence on SIP-induced receptor internalization of either subtypes. This means that changes in $[Ca^{2+}]_i$ are not involved in the differences seen between SIP-induced internalization of the SIP receptor subtypes.

We thus conclude that for the HisG-SIP₃ receptor, G_i -activation is of influence on SIP-induced internalization, but PKA, which is downstream in the G_i -pathway, does not have any effect. Influence of G_i -sensitive pathways in receptor internalization is also seen in for example α_{2A} and α_{2B} -adrenergic receptors, although partially (Olli-Lahdesmaki et al., 2004) or neuropeptide Y receptors (Pheng et al., 2003). Furthermore, although there are no available inhibitors of G_q -activation, none of the enzymes in this pathway, i.e. PLC and PKC seem to have any influence in HisG-SIP₃ receptor internalization. However, even though downstream signaling is not of influence, a direct effect of G_q -activation itself on SIP receptor internalization can not yet be ruled out. In addition, simulating the activation of both the G_q and G_i -pathways by increasing or decreasing $[Ca^{2+}]_i$ does not affect internalization of HisG-SIP₁ or HisG-SIP₃.

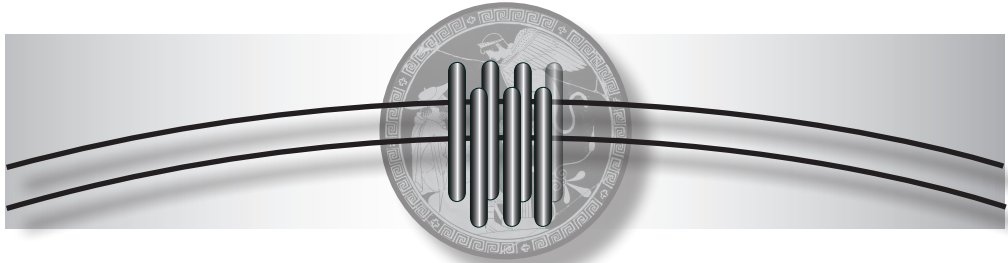
The data described in this article show a clear difference in the internalization of the HisG-SIP₁, HisG-SIP₂ and HisG-SIP₃ receptor subtypes. In part, these differences can be linked to a difference in signaling pathways activated by specific receptor stimulation, since G_i, but not G_q-activation seems to influence HisG-SIP₃ receptor internalization. However, this does not fully account for the observed differences. An additional explanation may be found in a recent investigation which showed that in contrast to SIP₁, the SIP₃ receptor internalization is independent of receptor phosphorylation (Rutherford et al., 2005). The observed differences in SIP receptor regulation could have implications for the functioning of drugs targeted at SIP receptors including FTY720.

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*Involvement of the third amino acid of the D/ERY motif
in G-protein selectivity of SIP receptors*



Chapter 6

Involvement of the third amino acid of the D/ERY motif of SIP receptors in G-protein selectivity

Manuscript in preparation

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Martin C. Michel, Stephan L.M. Peters & Astrid E. Alewijnse

Abstract

The highly conserved D/ERY motif is known to play an important role in G-protein coupling and/or constitutive activity of G-protein coupled receptors (GPCRs). However, the specific amino acid residues that are important in G-protein selectivity and which for instance restrict some GPCRs from coupling to multiple G-proteins are not exactly known. Among sphingosine-1-phosphate (SIP) receptors only the SIP₁ receptor, which couples to G_{i/o}-proteins only, contains an ERY motif whereas the other SIP receptor subtypes show some variance in the third amino acid of this motif. In this paper we investigated the role of the third amino acid residue of the D/ERY motif in regulating the preference for certain signaling responses, we compared wild-type and mutant SIP receptors. Mutating the ERY motif of the SIP₁ receptor to ERH, the motif present in the wild-type SIP₂ and SIP₃ receptor, or ERA reduced SIP-induced G_i-mediated signaling without inducing calcium elevations. The G_i-mediated responses of the SIP₁ specific agonist SEW2871 were also decreased at both mutant receptors, whereas the effects of FTY720-P at both mutant receptors were unchanged suggesting that these compounds induce different conformational states of the receptor. Mutating the ERH motif of the SIP₂ or SIP₃ receptor to an ERY or ERA motif did not induce major changes in signaling of these receptors, except for an SIP-induced increase in cAMP accumulation in the SIP₂ mutant receptor containing an ERY motif.

We conclude that the tyrosine residue of the D/ERY motif does not restrict the SIP₁ receptor from activating multiple G-proteins; however, mutating this residue revealed differences in response patterns induced by either SIP, SEW2871 or FTY720-P.

Introduction

The bioactive sphingolipid sphingosine-1-phosphate (SIP) regulates many different cellular functions including proliferation, migration and differentiation via a family of specific G-protein coupled receptors (GPCRs). This family currently consists of five subtypes (SIP₁₋₅), of which the SIP₁, SIP₂ and SIP₃ receptors are ubiquitously expressed, whereas the SIP₄ receptor is mainly found in lymphoid tissue and the SIP₅ receptor in the brain (Anliker & Chun, 2004).

GPCRs exert their signals primarily via activation of heterotrimeric G-proteins, of which there are four subfamilies, namely G_{i/o}, G_s, G_{q/11} and G_{12/13} (for review see Hermans, 2003). With 28 α subunits, six β and 12 γ -subunits being described (Hermans, 2003; Carbrea-Vera et al., 2003), many different combinations are possible, resulting in a diversity of pathways to activate.

Most GPCRs couple preferentially to one subfamily of G-proteins (89 % of the extensive group reviewed by Wong, 2003). However, a small fraction of GPCRs shows promiscuous coupling as they can activate multiple subfamilies of G-proteins. At present, it is not precisely known what restricts GPCRs from coupling to multiple G-proteins. It has been suggested that a specific amino acid region situated at the boundary of the third transmembrane domain and the second intracellular loop of the GPCR may play a role in G-protein selectivity. This region contains a triplicate of amino acids; D/ERY (Asp/Glu-Arg-Tyr), which is highly conserved among class A GPCRs (for 86 %, 97 % and 67 % respectively) (Flanagan, 2005). Indeed, several studies have found this D/ERY motif to be involved in constitutive receptor activity or G-protein coupling based upon site-directed mutagenesis within this motif (Capra et al., 2004; Favre et al., 2005; et al., 2006). Most of these studies have been done on the aspartate/glutamic acid and/or arginine residue while the tyrosine residue has been much less investigated, possibly due to the fact that it is the least conserved of the three. The effect of mutations in the D/ERY motif seems to differ between receptors or receptor families (reviewed in Rovati et al., 2007). It has been suggested that within the class A GPCRs, to which the SIP receptors belong, there may be two groups that differently use the D/ERY motif (Capra et al., 2004). In one group this motif is involved in keeping the receptor in an inactive ground state. In this group, which includes the α_{1B} -adrenoceptor and the histamine H₂ receptor, mutations in this motif do not affect ligand binding, but result in an increase or induction of constitutive receptor activity, sometimes accompanied by an increase in agonist affinity. In the second group of receptors, which includes

the α_{2A} -adrenoceptor and M_1 muscarinic acetylcholine receptor, mutations in the D/ERY motif result in changes in agonist-induced responses without inducing constitutive activity.

Unlike most GPCR, four out of five members of the SIP receptor family can couple to and activate multiple G-protein subfamilies. The SIP_1 receptor is the only receptor which exclusively activates G_i -protein coupled pathways (Ancellin & Hla, 1999). The SIP_2 and SIP_3 receptor activate G_r , G_q and $G_{12/13}$ -proteins (Ancellin & Hla, 1999) and the SIP_4 and SIP_5 receptor activate G_i and $G_{12/13}$ -proteins (Im et al., 2000; Malek et al., 2001; Gräler et al., 2003). Interestingly, the SIP_1 receptor is also the only subtype that contains a tyrosine residue, as found in most GPCRs at the third position of the D/ERY motif. The SIP_2 , SIP_3 , SIP_4 and SIP_5 receptors contain either a histidine (SIP_2 and SIP_3), or phenylalanine (SIP_4) or a serine (SIP_5) at this position. From these observations, it may thus be hypothesized that the tyrosine residue may contribute to restricting the SIP_1 receptor from promiscuous coupling.

In this study we, therefore, investigated whether this third amino acid is indeed important in G-protein selectivity by mutating the ERY motif of the SIP_1 receptor to an ERH (as present in the SIP_2 and SIP_3 receptor) or an ERA motif (generating a “neutral” mutation). Chinese hamster ovary (CHO)-FlpIn cells were transfected with either the wild-type (wt) or mutant receptors, and SIP-induced changes in $[Ca^{2+}]_i$ and cAMP accumulation (typical G_q and G_i -coupled pathways) in these cell lines were subsequently studied. Besides the natural ligand SIP, a selective SIP_1 agonist (SEW2871) and the phosphorylated form of the new immunosuppressive prodrug FTY720, an agonist for the $SIP_{1,3,5}$ receptor were used (Mandala et al., 2002).

Material & Methods

Materials

pcDNA3.1 containing the entire coding region of the human SIP_1 , SIP_2 or SIP_3 receptor was purchased from UMR cDNA Resource Center (Rolla, MO, USA). Cell culture media, hygromycine B, Lipofectamine™ 2000, Optimem, Hank's Balanced Salt Solution (HBSS), pOG44, pcDNA3.1/HisA/C, pcDNA5/FRT/TO and CHO-FlpIn cells, were obtained from Invitrogen (Breda, The Netherlands). Pluronic acid and Fluo-4 AM were obtained from Molecular Probes (via Invitrogen). Restriction enzymes (ApaI, BamHI, XhoI, EcoRI and HindIII) were obtained

from Fermentas Life Sciences (St. Leon-Rot, Germany). Foetal calf serum (FCS), enzyme free cell dissociation buffer and penicillin/streptomycin were obtained from Gibco (via Invitrogen). LANCE™ cAMP 384 kit and white 384 well optiplates were obtained from Perkin Elmer (Zaventem, Belgium). SIP was obtained from Avanti-Polar Lipids, Inc (via Instuchemie B.V., Delfzijl, The Netherlands). Probenecid, HEPES, EGTA, Triton X-100, 3-isobutyl-1-methylxanthine, bovine serum albumin (BSA) (fatty acid free) and activated charcoal were obtained from Sigma Aldrich (Zwijndrecht, The Netherlands). Black, clear bottom 96 well plates were obtained from Greiner Bio One (Alphen aan den Rijn, The Netherlands). 5-[4-Phenyl-5-(trifluoromethyl)-2-thienyl]-3-[3-(trifluoromethyl)phenyl]-1,2,4-oxadiazole (SEW2871) was obtained from Cayman Chemical Company (Ann Arbor, MI, USA). 2-Amino-2-[2-(4-octylphenyl)ethyl]-1,3-propanediol mono(dihydrogen phosphate) ester (FTY720-P) was synthesized according to previously described methods (Albert et al., 2005).

Molecular cloning and transfection

A N-terminal HisG-tag was added to the SIP_{1,3} receptors via cloning into pcDNA3.1/HisC using BamHI&XhoI (for SIP₁) or BamHI&ApaI (for SIP₂) or into pcDNA3.1/HisA using BamHI&XhoI (for SIP₃). A second cloning step was done using HindIII&XhoI (for SIP_{1,3}) or HindIII&ApaI (for SIP₂) to clone the HisG-tagged SIP receptors into the expression vector pcDNA5/FRT/TO.

Mutations in the ERY/H motif of the SIP_{1,3} receptor were made via site directed mutagenesis using the primers listed in table 1. All constructs were confirmed by sequencing of the plasmids and the stable cell lines.

Transfection of plasmids into CHO-FlpIn cells and clonal selection was done as described previously (Chapter 2). Membrane expression of the receptors was confirmed using an immunocytochemical method to detect membrane receptors by detecting the N-terminal HisG-tag as described previously (Chapter 3). The cDNA of the SIP₃ receptor obtained from Guthrie contained a point mutation (G⁹⁶²A which results in R³²¹Q) which was removed by PCR using the following primers (forward: TGCCTGGTCAGGGGGCGGGGGCCCG, reverse: CGGGCCCCCGCCCCCTGACCAGGCA). The SIP₃ receptor coding DNA in the resulting plasmid as well as the stable cell line constructed with this plasmid have been confirmed by sequencing.

Table 1. Primers used for mutations in the ERY/H motif of the SIP_{1,3} receptors via site directed mutagenesis.

Receptor + Mutation	Primer
SIP ₁ Y ¹⁴⁵ A	Forward: CATCGCCATTGAGCGCGCAATCACAATGCT
	Reverse: GCGCTCAATGGCGATGGCGAGGAGACTG
SIP ₁ Y ¹⁴⁵ H	Forward : CGCCATTGAGCGCCATATCACAATG
	Reverse : CATTGTGATATGGCGCTCAATGGCG
SIP ₂ H ¹⁵¹ A	Forward : ATCGCCATTGAGCGCGCAGTGGCCATTGCC
	Reverse : GCGCTCAATGGCGATGGCCAGGAGGCTGAA
SIP ₂ H ¹⁵¹ Y	Forward : CCATTGAGCGCTACGTGGCCATTGC
	Reverse : GCAATGGCCACGTAGCGCTCAATGG
SIP ₃ H ¹⁵⁷ A	Forward : GCCATCGAGCGGGCCTTGACAATGAT
	Reverse : ATCATTGTCAAGCCCCGCTCGATGGC
SIP ₃ H ¹⁵⁷ Y	Forward : GCCATCGAGCGGTACTTGACAATGA
	Reverse : TCATTGTCAAGTACCGCTCGATGGC

Cell culture

CHO-FlpIn cells stably expressing the SIP₁, SIP₂ or SIP₃ wt or mutant receptors were passaged 1:5 every two or three days in F-12 Nutrient Mixture (Ham) with L-glutamine, supplemented with 100 U/ml penicillin, 100 µg/ml streptomycin, 313 µg/ml hygromycin B and 10 % charcoal-stripped FCS. All cell lines were cultured at 37 °C in humidified air containing 5 % CO₂.

cAMP assay

The LANCE™ cAMP 384 kit was used to determine cAMP concentrations according to the manufacturer's protocol. CHO-FlpIn cells, after overnight serum-starvation, were detached from the surface using dissociation buffer. Cells were washed once with HBSS and subsequently resuspended in stimulation buffer, containing HBSS with 0.05% BSA (fatty-acid free) and 5 mM HEPES. Stimulation mixtures consisted of stimulation buffer with 1 mM (final concentration 0.5 mM) 3-isobutyl-1-methylxanthine, 6 µM (final concentration 3 µM) forskolin and the concentration response curve of the indicated compounds. Cells were added to the stimulation mixtures 1:1 in a 384 well optiplate at 2500 cells/well and stimulated for 60 minutes at room temperature with a total volume of 20 µl/well. Detection of the cAMP formed during stimulation was done according to the manufacturer's protocol. Measurements were done on a microplate reader (Victor 2, Wallac, Perkin Elmer) 16-24 hours after adding detection buffer and antibody mixture.

Intracellular calcium measurement

Intracellular calcium measurements were performed as previously described (Chapter 2) with minor changes. CHO-FlpIn cells were plated in a black, clear bottom 96 wells plate at 40,000 cells/well. After one day growth, cells were serum starved overnight. Cells were then loaded for 1 hour with basic buffer (HBSS containing 20 mM HEPES, 2.5 mM probenecid) containing 4 μ M Fluo-4 AM ester and 0.42 % v/v pluronic acid and incubated at 37 °C. After loading, cells were washed twice with basic buffer and incubated at 37 °C with basic buffer for 60 min. The fluorescence signal was measured at basal level, followed by ligand stimulation, Triton (5 % v/v) addition (resulting in F_{\max}) and 250 mM EGTA addition (resulting in F_{\min}). The $[Ca^{2+}]_i$ was calculated via the equation: $[Ca^{2+}]_i = K_d * ((F - F_{\min}) / (F_{\max} - F))$. K_d being the dissociation constant of the binding of Fluo-4 to calcium (345 nM). The increase in $[Ca^{2+}]_i$ upon ligand stimulation was calculated as the difference between the $[Ca^{2+}]_i$ for the basal level and after adding a ligand.

Data analysis

Concentration response curves were analyzed by fitting sigmoidal functions to the experimental data using Prism 4 (Graphpad Software, Inc., San Diego, CA, USA). Data are expressed as means \pm s.e.m. One-way ANOVA with a Dunnett's correction or Student's t-test was applied where appropriate. $P < 0.05$ was considered significant.

Results

Effect of mutations in the ERY motif of the SIP_1 receptor on signaling

In cells expressing the wt SIP_1 receptor SIP inhibited forskolin-induced cAMP accumulation. At both, the SIP_1 Y¹⁴³A and SIP_1 Y¹⁴³H mutant receptors, the potency of SIP to inhibit forskolin-induced cAMP accumulation was significantly reduced (wt $pEC_{50} = 9.7 \pm 0.1$ (n=8); Y¹⁴³A $pEC_{50} = 7.8 \pm 0.2$ (n=4); Y¹⁴³H $pEC_{50} = 8.9 \pm 0.3$ (n=3); $p < 0.05$) (Figure 1A). This difference was not seen upon stimulation with FTY720-P (wt $pEC_{50} = 10.2 \pm 0.1$ (n=9); Y¹⁴³A $pEC_{50} = 10.5 \pm 0.2$ (n=5); Y¹⁴³H $pEC_{50} = 9.8 \pm 0.1$ (n=4)) (Figure 1B). The potency of the SIP_1 specific agonist SEW2871 to inhibit forskolin-induced cAMP accumulation was significantly decreased at the SIP_1 Y¹⁴³A and the SIP_1 Y¹⁴³H mutant receptor compared to the wt SIP_1 receptor (wt $pEC_{50} = 8.2 \pm 0.1$ (n=12); Y¹⁴³A $pEC_{50} = 7.5 \pm 0.3$ (n=7); Y¹⁴³H $pEC_{50} = 7.7 \pm 0.1$ (n=4) $p < 0.05$) (Figure 1C).

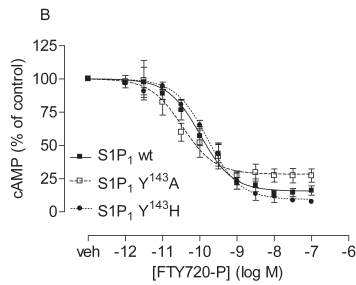
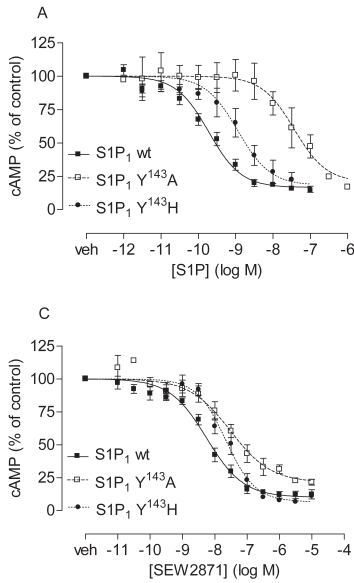


Figure 1. Concentration response curve of S1P(A), FTY720-P(B) or SEW2871(C)-induced inhibition of forskolin-induced cAMP accumulation for cells expressing the S1P₁ (wt, Y¹³⁷A or Y¹³⁷H) receptor. Ligand stimulations were carried out at room temperature for 60 min. Values are expressed as % of forskolin-induced cAMP production of vehicle control which was set at 100 % and are presented as means ± s.e.m. (100 % was for S1P S1P₁: 6.5 ± 1.0 nM (n=8); S1P₁ Y¹⁴³A: 11.1 ± 2.8 nM (n=4); S1P₁ Y¹⁴³H: 12.0 ± 2.1 (n=3); FTY720-P S1P₁: 6.3 ± 0.9 nM (n=9); S1P₁ Y¹⁴³A: 12.7 ± 3.6 nM (n=5); S1P₁ Y¹⁴³H: 12.8 ± 3.6 nM (n=4); SEW2871 S1P₁: 6.7 ± 0.9 nM (n=12); S1P₁ Y¹⁴³A: 11.1 ± 2.0 nM (n=7); S1P₁ Y¹⁴³H: 12.1 ± 1.7 nM (n=4)).

S1P, but not FTY720-P induced increases in $[Ca^{2+}]_i$ in mock transfected cells ($pEC_{50} = 7.0 \pm 0.3$; $E_{max} = 297 \pm 57$ nM (n=4)) (Figure 2). S1P and FTY720-P induced significantly higher increases in $[Ca^{2+}]_i$ in cells expressing the S1P₁ wt receptor (S1P: $pEC_{50} = 7.1 \pm 0.1$; $E_{max} = 857 \pm 122$ nM (n=6) $p < 0.05$; FTY720-P: $pEC_{50} = 6.2 \pm 0.2$; $E_{max} = 614 \pm 48$ nM (n=9)). These characteristics for S1P and FTY720-P were not significantly changed for both S1P₁ mutant receptors (S1P₁ Y¹⁴³A and S1P₁ Y¹⁴³H) from the S1P₁ wt receptor (S1P₁ Y¹⁴³A S1P: $pEC_{50} = 6.9 \pm 0.1$; $E_{max} = 456 \pm 32$ nM (n=3); FTY720-P: $pEC_{50} = 6.9 \pm 0.1$; $E_{max} = 344 \pm 56$ nM (n=3); S1P₁ Y¹⁴³H S1P: $pEC_{50} = 7.4 \pm 0.1$; $E_{max} = 824 \pm 67$ nM (n=7); FTY720-P: $pEC_{50} = 6.6 \pm 0.1$; $E_{max} = 705 \pm 92$ nM (n=9) (Figure 2)

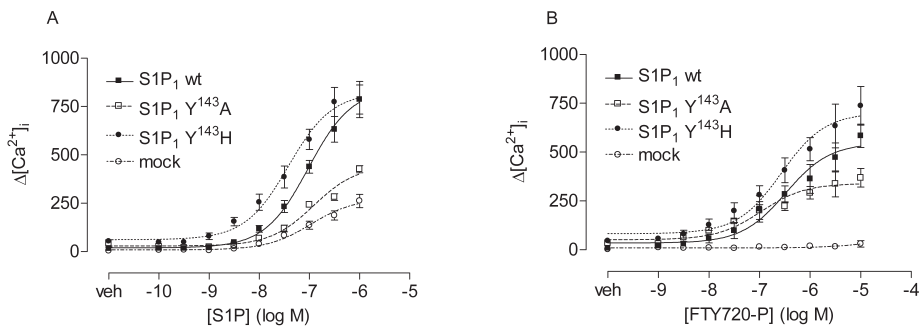


Figure 2. Concentration response curve of S1P(A) or FTY720-P(B)-induced elevations in $[Ca^{2+}]_i$ for cells expressing the S1P₁ (wt, Y¹⁴³A or Y¹⁴³H) receptor or mock transfected cells. Values are mean ± s.e.m. (n=3-9).

Effect of mutation in the ERH motif of the SIP₂ receptor on signaling

In CHO-FlpIn cells expressing the wild type *SIP*₂ receptor, *SIP* slightly increased forskolin-induced cAMP accumulation ($E_{\max} = 139 \pm 19$ % of basal (n=6)) (Figure 3A). A comparable *SIP* response was observed in cells expressing the *SIP*₂ H¹³¹A mutant receptor ($E_{\max} = 110 \pm 10$ % of basal (n=6)). Interestingly, the *SIP*-induced increase in forskolin-induced cAMP accumulation was significantly greater for the *SIP*₂ H¹³¹Y mutant receptor compared to the wt receptor ($E_{\max} = 258 \pm 28$ % of basal (n=6) (p<0.05)) (Figure 3A).

In addition to the effects on cAMP accumulation, stimulation of the *SIP*₂ wt cells with *SIP* also resulted in a concentration dependent increase in $[Ca^{2+}]_i$ ($pEC_{50} = 7.3 \pm 0.1$ (n=3); $E_{\max} = 1255 \pm 400$ nM (n=3)) (Figure 3B). This effect was not significantly altered in cells expressing the *SIP*₂ H¹³¹A or *SIP*₂ H¹³¹Y mutant receptor (Figure 3B). Because the *SIP*-induced elevations in $[Ca^{2+}]_i$ sometimes approach the effect observed for TritonX, which is used to determine the total concentration of calcium, in some cases (especially at high *SIP* concentrations) no reliable calculation of the change in $[Ca^{2+}]_i$ is possible.

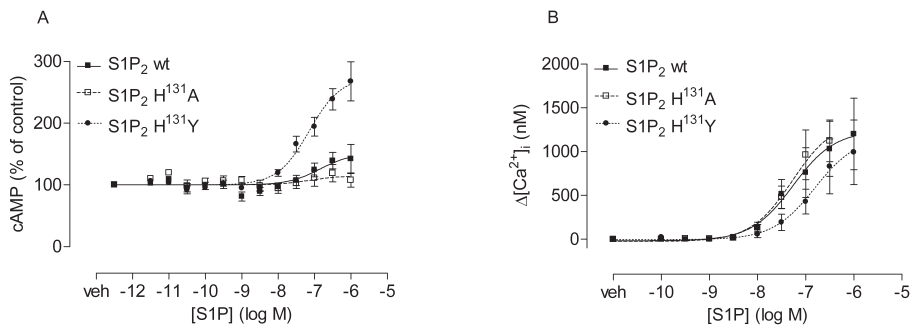


Figure 3. Concentration response curve of *SIP*-induced change in forskolin-induced cAMP accumulation for cells expressing the *SIP*₂ (wt, H¹³¹A or H¹³¹Y) receptor. Ligand stimulations were carried out at room temperature for 60 min. Values are expressed as % of forskolin-induced cAMP production of vehicle control which was set at 100% and are presented as means \pm s.e.m. (100% was for *SIP*₂ : 4.8 ± 0.6 nM (n=6); *SIP*₂ H¹³¹A: 5.4 ± 0.7 nM (n=6); *SIP*₂ H¹³¹Y: 6.1 ± 0.5 nM (n=6)) (A). Concentration response curve of *SIP*-induced elevations in $[Ca^{2+}]_i$ for cells expressing the *SIP*₂ (wt, H¹³¹A or H¹³¹Y) receptor. Values are mean \pm s.e.m. (n=3) (B).

Effect of mutation in the ERH motif of the SIP₃ receptor on signaling

In cells expressing the wt *SIP*₃ receptor *SIP* and FTY720-P inhibited forskolin-induced cAMP accumulation in a concentration dependent manner (Figure 4). The concentration response curve for *SIP* or FTY720-P was not significantly altered for the *SIP*₃ H¹³⁷A receptor or *SIP*₃ H¹³⁷Y mutant receptor compared to the *SIP*₃ wt receptor (Figure 4, Table 2).

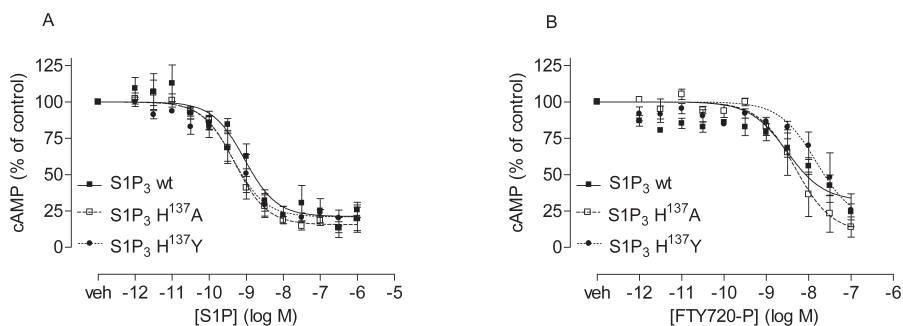


Figure 4. Concentration response curve of S1P(A) or FTY720-P(B)-induced inhibition of forskolin-induced cAMP accumulation for cells expressing the S1P₃ (wt, H¹³⁷A or H¹³⁷Y) receptor. Ligand stimulations were carried out at room temperature for 60 min. Values are expressed as % of forskolin-induced cAMP production of vehicle control which was set at 100% and are presented as means ± s.e.m. (100% was for S1P S1P₃: 13.5 ± 2.7 nM (n=5); S1P₃ H¹³⁷A: 17.6 ± 4.5 nM (n=5); S1P₃ H¹³⁷Y: 12.6 ± 3.4 (n=5); FTY720-P S1P₃: 15.3 ± 2.0 nM (n=3); S1P₃ H¹³⁷A: 23.0 ± 1.1 nM (n=3); S1P₃ H¹³⁷Y: 22.4 ± 4.5 (n=3).

Both, S1P and FTY720-P, potently induced elevations in [Ca²⁺]_i in cells expressing the wt S1P₃ receptor (Figure 5). This potency was only significantly increased for the S1P₃ H¹³⁷A receptor compared to the wt upon stimulation with FTY720-P (Figure 5). Also in these experiments, the data could in some cases not be reliably calculated as described for the S1P₂ receptors.

Table 2. pEC₅₀ and maximal inhibition values for the S1P₃ wt and mutant receptors in ligand-induced inhibition of forskolin-induced cAMP accumulation. * p<0.05 compared to wt.

ligand		wt	H ¹³⁷ A	H ¹³⁷ Y
S1P	pEC ₅₀	9.0 ± 0.2 (5)	9.5 ± 0.1 (5)	9.1 ± 0.3 (5)
	E _{max}	71 ± 8 %	80 ± 4 %	78 ± 3 %
FTY720-P	pEC ₅₀	8.2 ± 0.1 (3)	8.7 ± 0.1 (3)*	8.0 ± 0.1 (3)
	E _{max}	88 ± 4 %	94 ± 4 %	81 ± 2 %

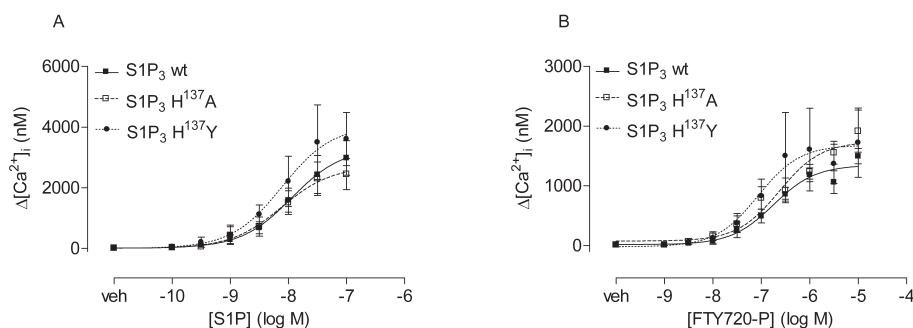


Figure 5. Concentration response curve of S1P(A) or FTY720-P(B)-induced elevations in [Ca²⁺]_i for cells expressing the S1P₃ (wt, H¹³⁷A or H¹³⁷Y) receptor. Values are mean ± s.e.m. (n=3-4).

Discussion

The specific molecular determinants of GPCRs controlling their coupling specificity to G-proteins have remained elusive. However, for some receptors, the D/ERY motif has been shown to be involved in G-protein coupling and activation (Favre et al., 2005; Capra et al., 2004; Rovati et al., 2007). For the family of S1P receptors, the third amino acid residue of this motif shows a remarkable variance, possibly correlating with the selectivity of the S1P receptor subtypes for specific G-proteins or the lack thereof. The S1P₁ receptor contains an ERY motif and couples exclusively to G_i-proteins. The other four subtypes show promiscuous coupling to G_i, G_q and G_{12/13}-proteins (S1P₂ and S1P₃) or G_i and G_{12/13} proteins (S1P₄ and S1P₅) and contain either an ERH (S1P₂ and S1P₃), ERF (S1P₄) or ERS (S1P₅) motif.

Mutating the ERY motif of the S1P₁ receptor to an ERH motif, which is present in the S1P₂ and S1P₃ receptor, showed an interesting change in signal transduction. Both, at the S1P₁ Y¹⁴⁵H and Y¹⁴⁵A mutant receptor the potency of S1P and SEW2871 to inhibit forskolin-induced cAMP accumulation was significantly decreased, which points in the direction of a reduced coupling to and activation of G_i-proteins. Similar effects were observed when mutating the tyrosine residue of the cannabinoid receptor CB₂, which is phylogenetically related to S1P receptors (Rhee et al., 2000; Alberich et al., 2004). In both studies, the DRY motif was mutated to a DRA motif and a loss of signaling to the G_i coupled pathway was observed.

Remarkably, the decrease in G_i-signaling of the S1P₁ Y¹⁴⁵H and Y¹⁴⁵A mutant receptor was not observed when using FTY720-P instead of S1P or SEW2871. Mutating the tyrosine residue of the ERY motif of the S1P₁ receptor thus seems to differentially affect the signaling of the ligands. Differential effects between S1P and SEW2871 on the one hand and FTY720-P on the other have also been observed when studying receptor fate after internalization. In that study S1P₁ receptors internalized upon stimulation with S1P or SEW2871 were shown to recycle back to the plasma membrane, while S1P₁ receptor internalized upon FTY720-P stimulation were broken down in lysosomes (Oo et al., 2007). The differences in receptor fate after internalization were assigned to polyubiquitinylation of the S1P₁ receptor by FTY720, but not by S1P (Oo et al., 2007). These results together with the results obtained in this study strongly suggest differences in the ligand-induced S1P₁ receptor conformation between S1P and SEW2871 versus FTY720-P.

The increases in [Ca²⁺]_i in CHO-FlpIn S1P₁ cells induced by both, S1P and FTY720-P were significantly different from the responses observed in mock-

transfected cells. Because the SIP and FTY720-P induced changes in $[Ca^{2+}]_i$ seem to be decreased for the $SIP_1 Y^{145}A$ mutant receptor, which also showed decreased G_i -signaling, the calcium effects are most likely G_i -mediated. However, definitive proof to confirm that the observed calcium effects are indeed G_i -mediated is still missing.

In contrast to the effects of mutations in the ERY motif observed for the SIP_1 receptor, the SIP_2 and SIP_3 receptors did not show many differences in G_i or G_q -activation when the ERH motif was mutated into ERY or ERA. No changes were seen for SIP and/or FTY720-P-induced inhibition of forskolin-induced cAMP accumulation or increases in $[Ca^{2+}]_i$ besides one remarkable observation. SIP increased forskolin-induced cAMP accumulation via the SIP_2 receptor rather than decreased, as one would expect based upon activation of G_i -proteins. Such an effect has been reported before in CHO cells (Windh et al., 1999), and was shown not to be induced via $[Ca^{2+}]_i$ (Kon et al., 1999). The authors suggest that it may be due to an inhibition of phosphodiesterase activity which results in less break-down of cAMP. This is, however, unlikely in our system, since the use of IBMX should inhibit all phosphodiesterase activity. The observed increase in cAMP was even more pronounced in the $SIP_2 H^{131}Y$ mutant receptor.

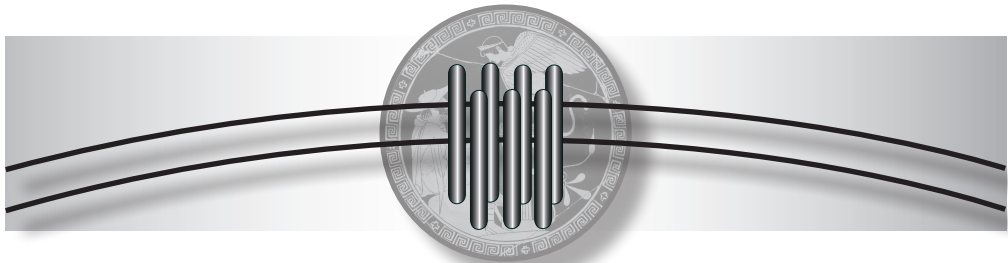
Overall, we conclude that mutations in the third amino acid of the D/ERY motif of the SIP receptor regulate G-protein mediated signaling, possibly via influencing the conformational state of the receptor. This could then clarify the observed differences in G_i -signaling in response to SIP and SEW2871 compared to FTY720-P. However, if the third amino acid is a tyrosine, it does not restrict the receptor from promiscuous coupling or influence constitutive receptor activity as seen for some other GPCRs.

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*Differential response patterns of several ligands
at the sphingosine-1-phosphate receptor
subtype 3*



Chapter 7

Differential response patterns of several ligands at the sphingosine-1-phosphate receptor subtype 3

Manuscript in preparation

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Abstract

Recently, several new ligands for the sphingosine-1-phosphate (S1P) receptor subtype 3 have become available (VPC23019, VPC23153, VPC24191 and the active metabolite of FTY720). The reported potencies and efficacies of these ligands were often based on one read-out system only, but the SIP_3 receptor is known to activate multiple signaling pathways. Therefore, we have compared the potencies and efficacies of these ligands in three different assays of SIP_3 receptor function (inhibition of cAMP accumulation, elevation of intracellular calcium and receptor internalization).

Our study indicates that the potency and/or efficacy of compounds can vary between assays and that for some compounds the potency can even vary within a particular assay at different stimulation times. Most strikingly, VPC23019 which has been described as an $SIP_{1,3}$ antagonist, behaved as a partial agonist in all three assays. In addition, the potency and efficacy in increasing $[Ca^{2+}]_i$ compared to inhibiting cAMP accumulation or inducing SIP_3 internalization was lower for most compounds. Furthermore beside activating G_i -coupled pathways S1P, in contrast to FTY720-P, seemed to additionally activate G_s -coupled pathways in high concentrations.

Based upon our findings we conclude that the SIP_3 receptor ligands tested in this study exhibited somewhat different cellular response patterns. An explanation for the observed differences could be that some functional responses require a higher efficacy than others. Alternatively, our observations could point into the direction of ligand-directed signaling for the SIP_3 receptor, but definitive proof of the latter awaits further studies.

Introduction

The sphingosine-1-phosphate (S1P) receptors, which bind the bioactive lipid S1P, represent a family of five proteins (named S1P₁ to S1P₅) which all belong to the family of G-protein coupled receptors (GPCRs) (Chun et al., 2002). One of the S1P receptor subtypes, the S1P₃ receptor, is known to play a critical role in the control of cardiac rhythm (Forrest et al., 2004; Sanna et al., 2004) and lung epithelial barrier function (Gon et al., 2005). Interestingly, the role of the S1P₃ receptor in cardiac rhythm was recognized due to the cardiac side-effects of FTY720 (Sanna et al., 2004), a S1P agonist which is momentarily in clinical trials for the treatment of multiple sclerosis. Studies using S1P₃ knock-out mice, subsequently, suggested that the negative chronotropic effect of FTY720 was predominantly related to the S1P₃ receptor expressed in atrial myocytes and on the sino-atrial node (Forrest et al., 2004; Sanna et al., 2004). Recently, it has also been shown that the cardioprotective effects of high density lipoproteins are mediated via S1P₃ receptors (Theilmeier, 2006). Furthermore, the anti-atherosclerotic action of FTY720 in ApoE^{-/-} is also believed to be mediated for a major part by S1P₃ receptors (Keul et al., 2007).

In lung, S1P₃ is exclusively expressed in pulmonary epithelium and activation of S1P₃ results in degradation of ZO-1 and claudin leading to a decreased epithelial tight junction formation (Gon et al., 2005). This will lead to the development of paracellular gaps and pulmonary edema. Based upon these data S1P₃ receptors may represent potential new targets in cardiac diseases and disease states associated with a disregulated epithelial barrier function as e.g. Adult Respiratory Distress Syndrome.

In recent years, several new S1P receptor ligands with some specificity for the S1P₃ receptor have become commercially available which could be useful tools to further elucidate the physiological role of S1P₃ receptors in e.g. the cardiovascular system (Davis et al., 2005). Of these ligands VPC23019, was characterized as a S1P_{1/3} receptor antagonist whereas VPC23153 and VPC24191 were described to have S1P_{1/3} agonistic activity. Although S1P₃ receptors are known to activate G_i as well as G_q and G_{12/13}-related signal transduction pathways (Young et al., 2006), the pharmacological characterization of these compounds was based on a single functional read-out system (³⁵S-GTPγS binding) only (Davis et al., 2005). However, it is generally known, and has e.g. extensively been studied for the S1P₅ receptor (Niedernerg et al., 2003), that the choice of the functional assay system can have implications for the observed efficacy and/or potency of a ligand. In this study

we, therefore, determined the pharmacological properties of the mentioned VPC ligands and the active metabolite of FTY720 on different SIP_3 -receptor mediated functional responses in transfected CHO-FlpIn cells expressing the human SIP_3 receptor.

Materials & Methods

Materials

pcDNA3.1 containing the entire coding region of the human SIP_3 receptor was purchased from UMR cDNA Resource Center (Rolla, MO, USA). Cell culture media, hygromycin B, Lipofectamine™ 2000, Optimem, Hank's Balanced Salt Solution (HBSS), pOG44, pcDNA3.1/HisA, pcDNA5/FRT/TO and CHO-FlpIn cells, were obtained from Invitrogen (Breda, The Netherlands). Restriction enzymes (BamHI, XhoI and HindIII) were obtained from Fermentas Life Sciences (St. Leon-Rot, Germany). Pluronic acid and Fluo-4 AM were obtained from Molecular Probes (via Invitrogen). Foetal calf serum (FCS), enzyme free cell dissociation buffer and penicillin/streptomycin were obtained from Gibco (via Invitrogen). LANCE™ cAMP 384 kit and white 384 well optiplates were obtained from Perkin Elmer (Zaventem, Belgium). SIP_3 (R)-Phosphoric acid mono-[2-amino-2-(6-octyl-1H-benzotriazol-2-yl) ethyl] ester (VPC23153), (R)-Phosphoric acid mono-[2-amino-2-(3-octyl-phenylcarbamoyl)-ethyl] ester (VPC23019) and (S)-Phosphoric acid mono-[2-amino-3-(4-octyl-phenylamino)-propyl] ester (VPC24191) were obtained from Avanti-Polar Lipids (via Instruchemie B.V., Delfzijl, The Netherlands). Probenecid, EGTA, Triton X-100, 3-isobutyl-1-methylxanthine, pertussis toxin (PTX), bovine serum albumin (BSA) (fatty acid free) and activated charcoal were obtained from Sigma Aldrich (Zwijndrecht, The Netherlands). Black, clear bottom 96 well plates were obtained from Greiner Bio One (Alphen aan den Rijn, The Netherlands). 2-Amino-2-[2-(4-octylphenyl)ethyl]-1,3-propanediol mono(dihydrogen phosphate) ester (FTY720-P) was synthesized according to previously described methods (Albert et al., 2005).

Molecular cloning, transfection & cell culture

A N-terminal HisG-tag was added to the SIP_3 receptor via cloning into pcDNA3.1/HisA using BamHI&XhoI. A second cloning step was done using HindIII&XhoI to clone the HisG-tagged SIP_3 receptor into the expression vector pcDNA5/FRT/TO. CHO-FlpIn cells stably expressing HisG-tagged

SIP₃ receptors were constructed and cultured as described before (Chapter 2). The HisG-tag has no influence on signal transduction induced via this receptor (data not shown). The cDNA of the SIP₃ receptor obtained from Guthrie contained a point mutation (G⁹⁶²A which results in R³²¹Q) which was removed by PCR using the following primers (forward: TGCCTGGTCAGGGGGCGGGGGCCCG, reverse: CGGGCCCCCGCCCCCTGACCAGGCA). The SIP₃ receptor coding DNA in the resulting plasmid as well as the stable cell line constructed with this plasmid have been confirmed by sequencing.

Calcium- and cAMP assay

Intracellular calcium concentrations ($[Ca^{2+}]_i$) and cAMP levels were measured as described previously (Chapter 2). Stimulations for cAMP measurements were done for 60 minutes unless otherwise indicated. In some cases, SIP elevated forskolin-induced cAMP levels after an initial decrease. For calculations of potency, the curve fit was based on the data points up to those SIP concentrations. To determine the efficacy, the E_{max} of SIP was set to 100 % and the E_{max} of the other compounds was expressed relative to this.

Immunocytochemistry

Immunocytochemistry was performed as described previously (Chapter 3). In short, after overnight serum starvation, cells were stimulated with the indicated ligand in serum free medium for 30 minutes at 37 °C. The N-terminal HisG-tag added to the SIP₃ receptor is detected using a 1st anti-HisG antibody combined with AlexaFluor® 488 goat-anti-mouse 2nd fluorescent antibody. The fluorescent signal measured on a microplate reader indicates the amount of receptors on the membrane.

Data analysis

Concentration response curves were analyzed by fitting sigmoidal functions to the experimental data using Prism 4 (Graphpad Software, Inc., San Diego, CA, USA). Data are expressed as means \pm s.e.m. One-way ANOVA with a Dunnett's correction or Student's t-test was applied where appropriate. $P < 0.05$ was considered significant.

Results

Ligand-induced inhibition of forskolin-induced cAMP accumulation

S1P inhibited forskolin-induced cAMP accumulation in CHO-FlpIn cells expressing the HisG- SIP_3 receptor ($pEC_{50} = 9.0 \pm 0.1$, $n=14$) (Figure 1A, Table 1). FTY720-P had a significantly lower potency to inhibit cAMP accumulation in these cells ($pEC_{50} = 8.6 \pm 0.1$, $n=11$) (Figure 1A, Table 1). VPC23019, VPC23153 and VPC24191 inhibited forskolin-induced cAMP accumulation, but also with a lower potency than S1P (Table 1). While the efficacy of FTY720-P was similar to S1P, all three VPC compounds showed a significantly lower efficacy compared to S1P and FTY720-P, i.e. behaved as partial agonists (Table 1).

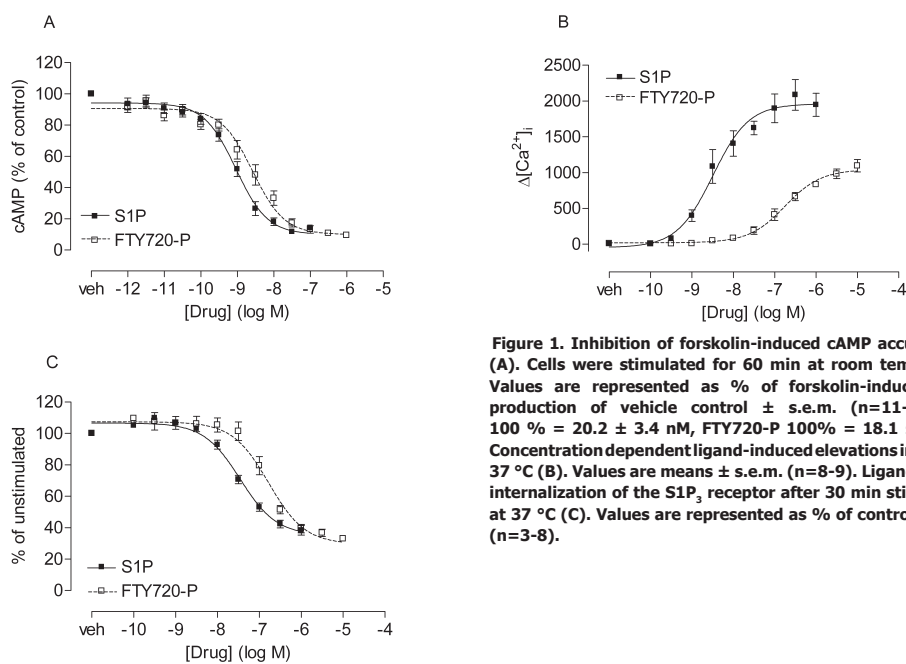


Figure 1. Inhibition of forskolin-induced cAMP accumulation (A). Cells were stimulated for 60 min at room temperature. Values are represented as % of forskolin-induced cAMP production of vehicle control \pm s.e.m. ($n=11-14$). S1P 100 % = 20.2 ± 3.4 nM, FTY720-P 100% = 18.1 ± 2.2 nM. Concentration dependent ligand-induced elevations in $[Ca^{2+}]_i$ at 37 °C (B). Values are means \pm s.e.m. ($n=8-9$). Ligand-induced internalization of the SIP_3 receptor after 30 min stimulations at 37 °C (C). Values are represented as % of control \pm s.e.m. ($n=3-8$).

Ligand-induced elevation of $[Ca^{2+}]_i$

S1P increased $[Ca^{2+}]_i$ in these cells ($pEC_{50} = 8.4 \pm 0.1$ ($n=9$)) (Figure 1B, Table 1). FTY720-P also increased $[Ca^{2+}]_i$, but with a significantly lower potency ($pEC_{50} = 6.8 \pm 0.1$ ($n=13$)) (Figure 1B, Table 1) and an efficacy of only 51 ± 4 % of that of S1P (Figure 1B, Table 1). VPC23019, VPC23153 and VPC24191 all raised $[Ca^{2+}]_i$ with significantly lower potencies and efficacies compared to S1P (Table 1).

Ligand-induced receptor internalization

All tested compounds induced internalization of the HisG- $S1P_3$ receptor after 30 min stimulations (for S1P and FTY720-P see figure 1C, for all compounds see Table 1). The potency of FTY720-P as well as all three VPC compounds to induce internalization was significantly lower than of S1P (Table 1). Only VPC23153 and VPC23019 had a lower efficacy compared to S1P (Table 1), whereas VPC24191 and FTY720-P had an efficacy comparable to that of S1P (Table 1).

Table 1. Potency and efficacy values for various ligands on the $S1P_3$ receptor for inhibition of forskolin-induced cAMP accumulation, induced elevations of $[Ca^{2+}]_i$, and induced internalization. The efficacy of S1P in each assay was arbitrarily set at 100 % (cAMP assay 100 % = 90 % inhibition of cAMP accumulation, calcium assay 100 % = 1947 nM and internalization assay 100 % = 60 % decrease compared to unstimulated cells. * $p < 0.05$ compared to S1P

Ligand	cAMP			$[Ca^{2+}]_i$			Internalization		
	pEC ₅₀	Efficacy (%)	n	pEC ₅₀	Efficacy (%)	n	pEC ₅₀	Efficacy (%)	n
S1P	9.0 ± 0.1	100 ± 2	14	8.4 ± 0.1	100 ± 8	9	7.5 ± 0.1	100 ± 4	8
FTY720-P	8.6 ± 0.1*	93 ± 2	11	6.8 ± 0.1*	51 ± 4*	13	6.8 ± 0.1*	109 ± 6	7
VPC23019	6.5 ± 0.1*	79 ± 6*	8	6.1 ± 0.1*	48 ± 4*	5	5.7 ± 0.2*	72 ± 10*	4
VPC23153	6.2 ± 0.1*	77 ± 7*	9	5.9 ± 0.1*	25 ± 2*	6	5.6 ± 0.4*	52 ± 18*	4
VPC24191	6.6 ± 0.1*	67 ± 8*	7	5.9 ± 0.1*	56 ± 10*	6	6.0 ± 0.3*	88 ± 5	4

Effects of various incubation times on forskolin-induced cAMP accumulation

To investigate whether the observed differences between ligand efficacies and potencies in both assays were a result of differences in stimulation times (60 min for cAMP and almost instantly for $[Ca^{2+}]_i$), shorter stimulation times were evaluated in the cAMP assay. The potency of S1P to inhibit forskolin-induced cAMP accumulation was increased when instead of 60 min, shorter stimulation times were

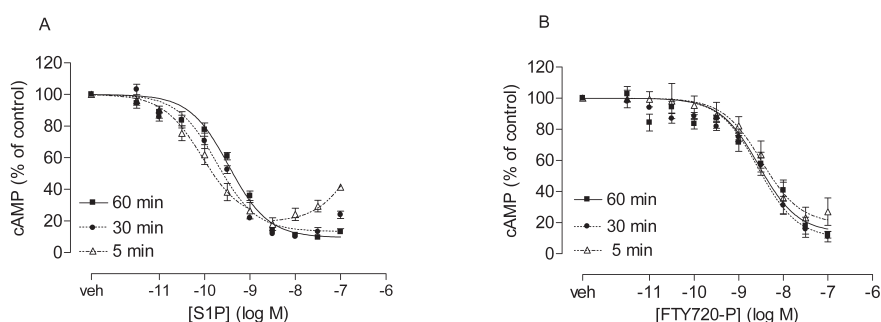


Figure 2. Inhibition of forskolin-induced cAMP accumulation at various stimulation times with S1P (A) or FTY720-P (B). Values are represented as % of forskolin-induced cAMP production of vehicle control ± s.e.m. (n=4).

used (Figure 2A, Table 2). Interestingly, at the shorter stimulation times after an initial decrease an increase in forskolin-induced cAMP accumulation was observed at high concentrations of SIP (> 10 nM).

In contrast to SIP the potency of FTY720-P was not changed at shorter stimulation times (Figure 2B, Table 2). However, all three VPC compounds also had an increased potency at shorter stimulation times (Table 2).

The efficacy of FTY720-P at shorter time points did not change, thus FTY720-P remains a full agonist at the SIP_3 receptor at shorter time points.

Table 2. Potencies for SIP, FTY720-P, VPC23019, VPC23153 and VPC24191 at the SIP_3 receptor for inhibition of forskolin-induced cAMP accumulation during various stimulation times. Efficacy values were not altered when using different stimulation times (n=4). *p<0.05 compared to 60 min.

	SIP	FTY720-P	VPC23153	VPC23019	VPC24191
Stimulation time	pEC50	pEC50	pEC50	pEC50	pEC50
5 min	10.1 ± 0.1*	8.4 ± 0.1	6.3 ± 0.1*	7.2 ± 0.0*	7.2 ± 0.0*
15 min	9.8 ± 0.1*	8.3 ± 0.2	-	-	-
30 min	9.7 ± 0.1*	8.4 ± 0.1	-	-	-
45 min	9.3 ± 0.1	8.3 ± 0.1	-	-	-
60 min	9.4 ± 0.1	8.3 ± 0.1	5.8 ± 0.2	6.6 ± 0.1	6.7 ± 0.1

Effect PTX on SIP and FTY720-P-induced inhibition of forskolin-induced cAMP accumulation

The observed increase in cAMP levels at higher SIP concentrations could be due to G_s -activation. To reveal the full effect of possible G_s -activation, instead of a combination between G_i and G_s -mediated effects, cells were incubated with PTX to block G_i -activation. Overnight incubation with PTX (100 ng/ml) abolished the inhibitory effect of SIP on forskolin-induced cAMP formation in these cells (data not shown). Additionally, in the PTX treated cells, SIP concentration-dependently stimulated cAMP accumulation, but not in mock transfected cells (Figure 3A). The SIP-induced cAMP accumulation after PTX treatment was further increased upon shorter incubation times with SIP. Stimulation with FTY720-P after PTX treatment did not stimulate cAMP accumulation in these cells (Figure 3B).

Correlation between efficacies of SIP_3 receptor ligands in various assays

The efficacies of the tested ligands in the cAMP assay and receptor internalization assay were very similar (Figure 4C, Table 1). However, the efficacies of the tested ligands for inducing elevations of $[Ca^{2+}]_i$ were generally lower than in the cAMP assay and receptor internalization (Figure 4A,B, Table 1).

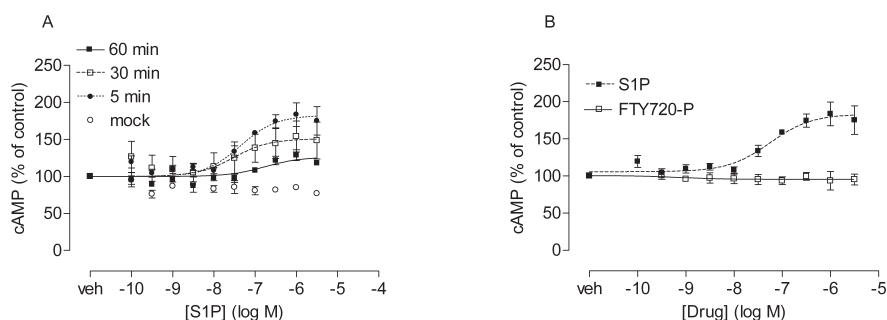


Figure 3. Effect of overnight PTX (100 ng/ml) incubation on S1P-induced changes in forskolin-induced cAMP accumulation at various stimulation times (A) and FTY720-P or S1P-induced changes in forskolin-induced cAMP accumulation at 5 min stimulation (B). The S1P curve in (A) is the same as in (B). Values are represented as % of forskolin-induced cAMP production of vehicle control \pm s.e.m. ($n=3-4$).

Discussion

Recently, several new S1P ligands with some specificity for the SIP_3 receptor have become commercially available but they have not been intensively characterized yet. In this study we, therefore, investigated the pharmacological properties of these ligands using different signal transduction assays. Specifically, we determined the potency and efficacy of all ligands with regard to their ability to inhibit forskolin-induced cAMP accumulation, to increase $[Ca^{2+}]_i$ and to internalize the SIP_3 receptor. In CHO-FlpIn cells expressing the human SIP_3 receptor.

In our experiments, both S1P and FTY720-P inhibited forskolin-induced cAMP accumulation, increased $[Ca^{2+}]_i$ and induced SIP_3 receptor internalization.

All these responses were mimicked by VPC23019, VPC23153 and VPC24191 (Table 1). In our assay system, VPC23019 thus behaved as a partial agonist rather than an antagonist. Although this may seem strange, differences in efficacy between varying cellular responses (Niedernberg et al., 2003) and/or varying cellular systems have been described before for other GPCRs and can e.g. be caused by differences in receptor expression levels (Sato et al., 2007).

Comparing the potency of the different VPC compounds tested in the cAMP accumulation versus the calcium assay revealed no obvious differences although the potency determined in the cAMP assay consistently tended to be higher than in the calcium assay. This tendency in difference in potency between the cAMP and calcium assay was also observed for SIP and was even more pronounced for FTY720-P. Consequently, the potency of SIP and FTY720-P to inhibit forskolin-induced cAMP accumulation differs only by 0.4 log units, while their potency to increase $[Ca^{2+}]_i$ differed by 1.6 log units (Table 1). For all compounds the potency to induce SIP_3 internalization most closely resembled the potency determined in the calcium assay.

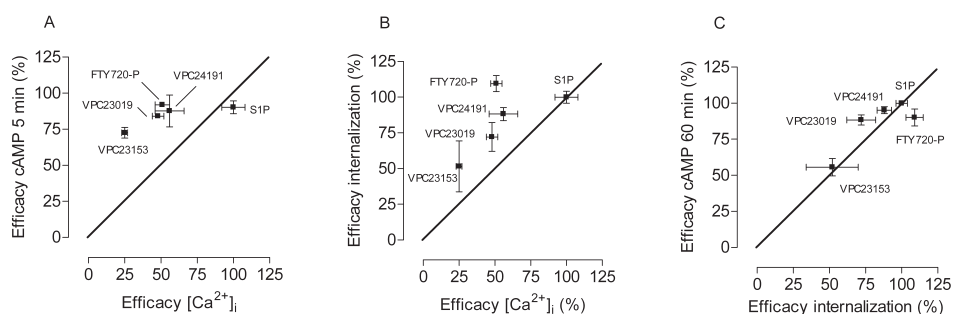


Figure 4. Comparison of efficacy between inhibition of forskolin-induced cAMP accumulation, elevation of $[Ca^{2+}]_i$ and induction of internalization for all used SIP_3 receptor ligands. Efficacies of the 5 min ligand stimulation times for the cAMP assay were compared to the efficacies in the calcium assay, while the 60 ligand stimulation times for the cAMP assay were compared to the internalization assay, since these combinations mostly represent similar assay conditions. The angled lines represent a line of identity rather than correlation lines.

To exclude that the observed differences in potency between the assays are the result of methodological differences (e.g. elevations in $[Ca^{2+}]_i$ are measured instantly upon ligand addition, while effects on cAMP accumulation are measured after 60 min stimulation) the effect of reducing the stimulation time in the cAMP accumulation assay was investigated. Interestingly, the potency of SIP to inhibit forskolin-induced cAMP accumulation was significantly increased upon shorter stimulation times. In contrast, the potency for FTY720-P was not altered with time. Thus, when comparing the potency of SIP and FTY720-P at shorter incubation times, the difference in potency between these compounds in inducing $[Ca^{2+}]_i$ elevations and inhibiting cAMP accumulation was comparable (approximately 1.5 log unit in each assay). In addition, all VPC compounds also showed an increase in potency upon shorter incubation times (Table 2).

Momentarily it is not clear what causes the shift in potency of S1P and/or the VPC compounds in the cAMP assay in time. Probably, it is due to a 'disappearance' of these ligands from the medium, making them less available to the receptor (see also General discussion). Whether this is caused by enzymatic break-down of these ligands, a possible uptake of these ligands or other processes is yet unknown. From these data it can thus be concluded that care should be taken when comparing potencies of ligands using different signal transduction assays especially when different incubation times are used. Because the potency determined in the cAMP assay at the shorter incubation times increased for the majority of compounds, all ligands under these conditions more potently induced changes in cAMP accumulation than in $[Ca^{2+}]_i$.

Another remarkable effect was observed when using shorter S1P stimulation times in the cAMP accumulation assay. We noticed that at these shorter incubation times the initial decrease in forskolin-induced cAMP accumulation was at high concentrations of S1P (>10 nM) followed by an increase in cAMP accumulation. In contrast to S1P, FTY720-P did not increase cAMP accumulation at higher concentrations. Because, CHO cells have been reported to endogenously express S1P receptors (Holdsworth et al., 2005), the observed S1P effect could be due to stimulation of endogenous S1P receptors. However, as the effect was not observed in mock transfected cells, an effect via endogenously expressed receptors is unlikely. The S1P-induced increase in cAMP accumulation was even more pronounced after PTX incubation indicating that in this system S1P activates not only G_i -coupled but also G_s -coupled pathways leading to an increase in cAMP accumulation via stimulation of adenylyl cyclase. Similar observations have been described for the M_2 acetylcholine receptors but only at high receptor density (Michal et al., 2001). In their paper the authors propose that in high receptor expression systems G_i -protein activation gets saturated at higher ligand concentrations resulting in the activation of G_s -proteins by the additionally activated receptors.

Another explanation for the effects we observed in this study could be, that the rise in cAMP accumulation is caused by a G_q -mediated response via calcium-dependent adenylyl cyclases. Based upon our data, this cannot be excluded. Although FTY720-P did not have stimulatory effects on cAMP accumulation the efficacy of this compound to induce elevations of $[Ca^{2+}]_i$ is also much lower than that of S1P. Compared to S1P, FTY720-P is a partial agonist in the calcium assay whereas it behaves as a full agonist at all stimulation times in the cAMP assay.

Because FTY720-P does not activate SIP_2 receptors, the observed difference in efficacy of SIP versus FTY720-P in the calcium assay could still be explained by a synergistic functioning of the SIP_3 receptor and an endogenous SIP_2 receptor. In mock transfected cells, SIP induced small increases in $[Ca^{2+}]_i$, whereas FTY720-P had no effect, which also argues in favour of SIP partly acting via an endogenously expressed receptor (Chapter 6). In addition to FTY720-P, there is also quite a difference in efficacy for VPC23153 and VPC23019 between the two assays (Table 1). It can thus be concluded that the majority of compounds is more efficacious in inhibiting forskolin-induced cAMP accumulation than in increasing $[Ca^{2+}]_i$. In addition, there is a striking similarity between the efficacies to induce internalization and the efficacies to inhibit cAMP accumulation. Therefore most likely, a higher efficacy is required for calcium signaling compared to inhibition of cAMP accumulation and/or internalization.

However, an alternative explanation based on a relatively new concept in GPCR pharmacology could be that some of the tested compounds differentially activate the signal transduction pathways investigated. Classical pharmacology assumed that all ligands activating a given GPCR yield similar responses on the same signal transduction pathways which only differ with regard to agonist potency and efficacy. However, recent evidence from several GPCRs demonstrates that some agonists preferentially elicit a certain response whereas others, acting on the same receptor, preferentially elicit another response, and that such differences cannot be explained in terms of agonist potency or efficacy (for an overview see Urban et al., 2007). These observations have led to the new concept that each agonist can induce a different conformational state of the receptor which results in preferential coupling to specific G-proteins and subsequent activation of specific signal transduction pathways. This phenomenon has been described in many different systems, and has been termed in as many different ways, including ligand-directed signaling, protean agonism, functional selectivity, conformational selection or ligand-directed trafficking (Simmons, 2005; Urban et al., 2007).

Interestingly, for the SIP_1 receptor, some observations suggesting that different ligands induce different receptor conformations have already been described. SIP_1 receptors internalized upon SIP or SEW2871 stimulation and were eventually recycled to the plasma membrane, while internalized upon stimulation with the immunosuppressive prodrug FTY720 these receptors were broken down in lysosomes (Oo et al., 2007). This difference in receptor fate has been related to

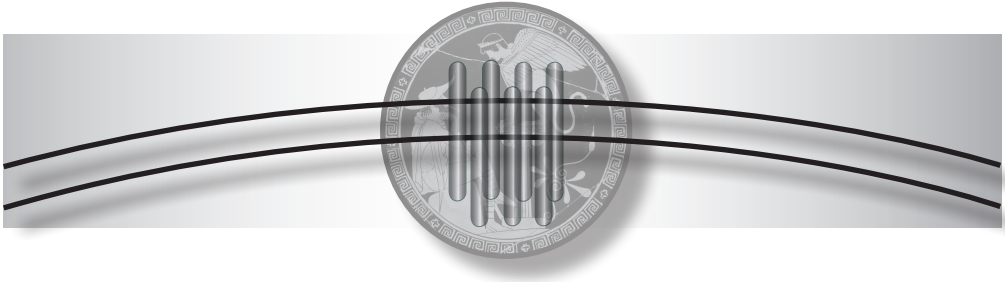
polyubiquitinylation of the SIP₁ receptor induced by FTY720-P, but not by SIP (Oo et al., 2007). In Chapter 6 we also show some data suggesting conformational differences between SIP and FTY720-P as a particular mutation in the second intracellular loop of the SIP₁ receptor did affect SIP but not FTY720-induced G_i-signaling. Although ligand-directed signaling may explain some of the effects observed in our study further investigation is necessary before drawing any definite conclusions. In addition, as neither the order of potency nor the order of efficacy of the tested ligands was markedly different between the assays, strong indications for ligand-directed signaling are missing.

Overall it is clear that several of the used SIP receptor ligands show some distinctions in SIP₃-receptor mediated responses. The majority of ligands had lower potencies and efficacies in increasing [Ca²⁺]_i compared to inhibiting cAMP accumulation. Furthermore SIP, in contrast to FTY720-P, seemed to additionally activate the G_s-pathway. Although these results may point in the direction of ligand-directed signaling definite proof to confirm this is still missing.

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General discussion



Chapter 8

General discussion

Sphingosine-1-phosphate (S1P) regulates various cellular processes via a family of five specific G-protein coupled receptors (GPCRs), named S1P₁₋₅ receptors (Chun et al., 2002). Being discovered less than two decades ago, many questions and “riddles” about the role and biology of these receptors in health and disease are still unanswered. Because S1P receptors are a relatively newly discovered family of receptors, the tools available to investigate these receptors such as subtype-selective agonists and/or antagonists are very limited. In addition, the available S1P ligands are generally not that well characterized. The first part of this thesis describes the development and/or pharmacological characterization of some tools which should facilitate the investigation of the molecular pharmacology of S1P receptors. In the second part of the thesis the signaling and regulation of S1P receptors is described in cell lines stably expressing the receptor, primary vascular cells or whole blood vessels. As a main focus in our department is the cardiovascular system, these studies have been restricted to the S1P₁, S1P₂ and S1P₃ receptors, as these subtypes are suggested to be the most important S1P receptors in the cardiovascular system (Michel et al., 2007).

Part I: Tools

As already stated above, selective S1P ligands are relatively few, and most of the newly developed ligands are often not that well characterized pharmacologically. For example suramin has been reported to act as an antagonist specific for the S1P₃ subtype (Salomone et al., 2003). However, this compound has many other effects on cellular signaling, e.g. suramin has been described as a kind of general inhibitor of GPCR signaling because it inhibits receptor-G-protein coupling (Beindl et al., 1996). In this thesis (Chapter 2), we pharmacologically characterized BML-241, a compound that was suggested to be an interesting lead compound for the development of S1P₃ antagonists. In the original publication this compound was found to inhibit

SIP-induced elevations in $[Ca^{2+}]_i$ in cells expressing the SIP_3 receptor but not in cells expressing the SIP_1 receptor (Koide et al., 2002). However, BML-241 was only tested at one relatively high concentration of 10 μ M in that study. One should thus be careful to draw any conclusions based upon these results. For example, elevations of $[Ca^{2+}]_i$ via SIP_1 receptor activation have been thought to be a result of the $\beta\gamma$ -subunits of G_i -proteins (Okamoto et al., 1998), while in case of the SIP_3 receptor, G_q -proteins are involved (Kon et al., 1999). An alternative explanation for the effects observed for BML-241 could thus be that BML-241 behaves as a G_q -inhibitor instead of a SIP_3 receptor inhibitor. In our study we confirmed the inhibitory effect of high BML-241 concentrations on the SIP_3 receptor-mediated increases in $[Ca^{2+}]_i$. However, BML-241 also inhibited SIP_2 -, α_{1A} -adrenergic- and purinergic receptor-mediated effects on $[Ca^{2+}]_i$ and reduced the α_1 -adrenergic-mediated contraction of rat mesenteric artery (Chapter 2). The effects observed on α_{1A} -adrenergic receptor signaling were not the result of a direct interaction of BML-241 with this receptor as, based upon competition binding studies, BML-241 lacked high affinity for this receptor. Perhaps even more importantly, BML-241 did not antagonize the SIP_3 receptor-mediated inhibition of forskolin-induced cAMP accumulation, demonstrating that it does not inhibit the receptor itself but rather certain signaling responses which can be elicited by the SIP_3 receptor. From our study we thus conclude that BML-241 indeed inhibits SIP_3 mediated effects on $[Ca^{2+}]_i$ but that it is most likely a low potency, non-selective inhibitor of increases in $[Ca^{2+}]_i$, rather than a specific antagonist at the SIP_3 receptor.

In our own search for specific SIP receptor ligands, we have performed some medium throughput screening. These studies were done in close collaboration with Dr. I. de Esch from the department of Medicinal Chemistry from the Free University (Amsterdam) and Dr. J. Brussee and Dr. M.W. Beukers from the division of Medicinal Chemistry from Leiden University (Leiden). In this screening approach two different strategies to select a series of compounds were used. One strategy was based on a pharmacophore model for the SIP_1 receptor whereas the other was a selection of compounds with high structural diversity. The first strategy identified a SIP_1 selective lead compound whereas in the second strategy a SIP_3 -selective lead compound was identified. Based upon the lead compounds two new series of ligands with close homology to the lead compounds were selected to determine some structure activity relationships for these compounds. For confidentiality reasons the screening efforts and results are no part of this thesis but are promising

for further development of selective $S1P_1$ and/or $S1P_3$ ligands.

Another important tool in research, besides the ligands, is the existence or development of innovative methods to study GPCR signaling. In this respect, Hoffmann et al. (2005) have published an elegant method using a fluorescein arsenical hairpin binder (FLAsH)-based fluorescence resonance energy transfer (FRET) approach to study the activation-induced conformational changes in the adenosine A_{2A} receptor. In their studies a FLAsH-motif of six amino acids was inserted into the third intracellular loop of the receptor, and a cyan fluorescent protein (CFP) was attached to the C-terminal tail of the receptor. When the CFP and the FLAsH are in close proximity, CFP can act as a fluorescent donor and thereby transfer energy to the acceptor (FLAsH). Upon activation, a conformational change is induced in the receptor which may subsequently change the orientation and distance between the donor and acceptor thus altering the energy transfer between these two. In collaboration with Dr. C. Hoffmann from the Institute of Pharmacology and Toxicology of the University of Würzburg (Würzburg, Germany) we applied the FLAsH-based FRET approach to the $S1P_1$ -receptor. Usually, to allow the best signal to occur, the receptor constructs need to be optimized for the relative distance and orientation of the two fluorophores. This procedure aims to place the two fluorophores within the Förster-distance and thus allowing FRET to occur. Since structural data are only available for rhodopsin, this process is still laborous and needs to be individually adjusted for each receptor.

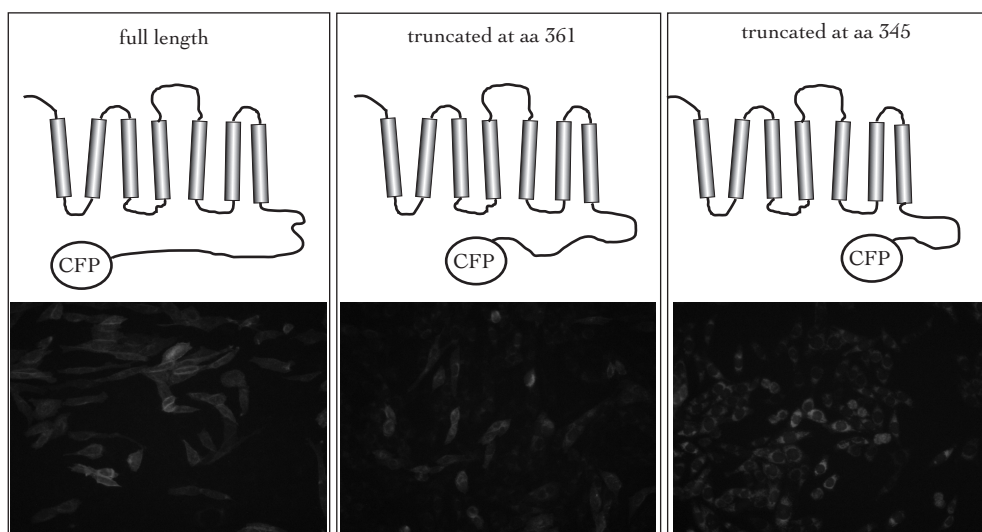


Figure 1. Constructs of CFP-tags added to truncated $S1P_1$ receptor with corresponding fluorescent images CHO-FlpIn cells expressing these constructs.

To do so we used two rounds of optimization. In the first set of experiments a C-terminal CFP tag was added to the S1P₁ receptor in three different positions and the membrane expression of the tagged construct was studied. As the orientation of the CFP tag to the FIAsh motif is of major importance, besides the full length construct also two C-terminal truncated S1P₁ receptor constructs, either truncated

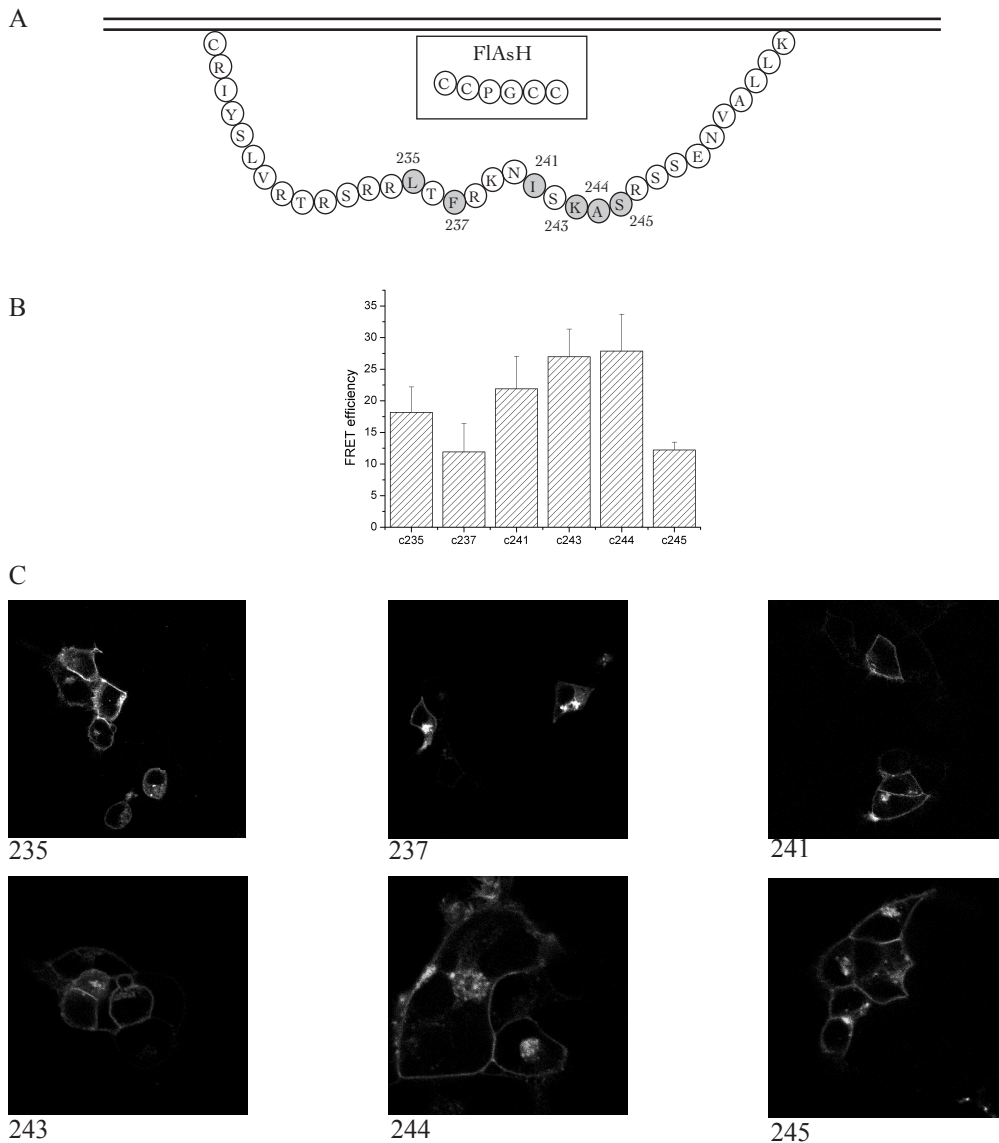


Figure 2. Amino acid sequence of the third intracellular loop of the S1P₁ receptor (A). Grey circles represent the amino acid after which the FIAsh construct has been inserted. FRET efficiencies of S1P₁ receptor constructs with a truncated C-terminus (at amino acid 361) and inserted FIAsh (numbers correspond to A) (B). FRET efficiency is measured by determining the change in FRET in time after removing the FIAsh ligand. Fluorescent images showing membrane expression of the corresponding S1P₁ receptor constructs in CHO-FlpIn cells (C). Receptors are visualized via the C-terminal CFP-tag.

at the amino acid 361 (SIP₁C³⁶¹trunc) or amino acid 345 (SIP₁C³⁴⁵trunc) position, have been created and the CFP tag has been added to them (Figure 1). Two of the constructs, the full length and the SIP₁C³⁶¹ truncated receptor showed plasma membrane expression whereas the SIP₁C³⁴⁵ truncated receptor was predominantly located in the cytosol (Figure 1). In a second set of experiments the FLAsH motif (CCPGCC) was inserted at amino acid position 239 in the third intracellular loop of the full length and SIP₁C³⁶¹ truncated receptor. The selection of these sites was reasoned to be suitable, since they closely correspond to sites which had been used for the adenosine A_{2A} receptor. The SIP₁C³⁶¹ truncated receptor showed some FRET efficiency, which is measured by removing the FLAsH ligand and subsequently determined the change in FRET in time, but this efficiency was not changed by the addition of SIP (data not shown). Therefore, we made six additional constructs in which the FLAsH motif was inserted at different positions in the third intracellular loop of the SIP₁C³⁶¹ truncated CFP-tagged receptor (Figure 2A). Although all constructs were expressed at the cell membrane (Figure 2C) and they all showed some FRET efficiency (Figure 2B), the FRET ratio was not changed in the presence of SIP (Figure 3), SEW2871, FTY720 or FTY720-P (data not shown). Probably due to the fact that the third intracellular loop of the SIP₁ receptor compared to the adenosine A_{2A} receptor is much shorter, it is possibly less flexible. Consequently, the change in position of intracellular loop 3 (with the FLAsH element) compared to the C-terminus (with the CFP) upon activation could be too moderate to result in any changes of the FRET signal. Since these results for the SIP₁ receptor were not that promising we did not pursue this project any further. An interesting finding was also that the C-terminus of the SIP₁ receptor between amino acid 361 and amino

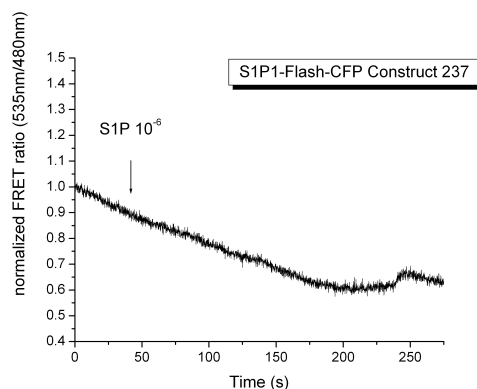


Figure 3. FRET ratio of the SIP₁-CFP-237 receptor construct. SIP (1 μ M) is added at the indicated time and the FRET ratio (535nm/480nm) is followed in time.

acid 345 seem to have an important role in receptor surface expression.

One major advantage of FLAsH, compared to other fluorescent tags as e.g. green fluorescent protein (GFP), is its small size. Fluorescent tags are often used to investigate receptor regulatory processes such as internalization, recycling and down regulation. However, because of the relative large size of these fluorescent tags, they can affect expression, signaling and other receptor-mediated processes (McLean & Milligan, 2000). Since we were interested in investigating regulatory processes of S1P receptors such as internalization, we set up a method that allowed the detection of receptors at the cell surface on a medium throughput scale (96 wells plate) (Chapter 3). This immunocytochemical method was based on the addition of a small N-terminal HisG-tag to the receptor and the subsequent detection of this tag with an anti-HisG antibody and a e.g. fluorescent second antibody. The small N-terminal tag did not affect membrane expression and signaling properties of the receptor. The method was optimized using the S1P₁ receptor as a model receptor comparing several first and second antibodies, ranging from absorbance to fluorescent or luminescent read outs. The combination of a specific first antibody for the HisG-tag and a fluorescent second antibody (AlexaFluor® 488) proved most reliable and sensitive. The developed method was suitable to investigate the agonist-induced internalization of the S1P₁ receptor but also of other S1P receptors (Chapter 5) and could be performed in a 96 well plate and quantified using a microplate reader.

Part II: S1P receptor signaling and regulation

In the cardiovascular system, several S1P receptors are expressed, most importantly the S1P₁, S1P₂ and S1P₃ receptor. Many things are still unclear about their regulation and signaling. Measuring the effect of exogenously added S1P on vascular preparations in an organ bath set up is difficult due to technical difficulties. However, recently some S1P receptor ligands, selective and non-selective, have become commercially available and enabled a detailed study on the role of S1P receptors in vascular responses. The study described in this thesis (Chapter 4) investigated the effects of some S1P ligands on contraction/relaxation of aortic rings and on S1P receptor-mediated signaling in vascular cells (smooth muscle and endothelial cells) isolated from the aorta.

We were the first to show that all the S1P receptor ligands induce an endothelium-dependent S1P₁ and/or S1P₃ receptor-mediated relaxation of aortic rings most likely

via an activation of endothelial nitric oxide synthase (eNOS) and the subsequent production of the vasodilator NO. When studying the two main cell types of the aorta separately, all ligands affected the forskolin-induced cAMP accumulation in rat aortic endothelial cells (RAECs). The majority of ligands inhibited cAMP accumulation in these cells, except for VPC23153 which increased the cAMP accumulation. In addition, the $S1P_1/S1P_3$ antagonist VPC23019 competitively antagonized the effect of FTY720-P on cAMP accumulation in RAECs. Since both, ligands increasing and decreasing endothelial cAMP, were associated with vasorelaxation, the vasodilation appears to be independent of endothelial cAMP content. Besides the effects on cAMP accumulation, S1P and FTY720-P at high concentrations also moderately elevated $[Ca^{2+}]_i$ in RAECs but it is not known whether these effects are indeed S1P receptor mediated. Because cultured RAECs predominantly show $S1P_1$ -receptor mediated signaling these cells most express only the $S1P_1$ receptor, a finding that was supported by qPCR data.

In vascular smooth muscle cells (VSMCs), FTY720, FTY720-P and SEW2871 slightly inhibited forskolin-induced cAMP accumulation, but the inhibitory effect was too small to allow reliable quantification. Of all tested ligands, only the one with $S1P_2$ agonistic activity (S1P) elevated $[Ca^{2+}]_i$ in these cells. In addition, the $S1P_2$ receptor antagonist JTE013 partly inhibited this effect. qPCR showed that the $S1P_1$, $S1P_2$ and $S1P_3$ receptors were expressed in these cells, but apparently signaling in these cells occurs predominantly via the $S1P_2$ receptor. This led to the overall conclusion that in the two main cell types of a vessel, being smooth muscle cells and endothelial cells, different S1P receptors are likely to cause opposite effects on vascular tone. Mainly the $S1P_1$ receptor, but possibly also the $S1P_3$ receptor, in endothelial cells (ECs) induces a relaxant effect upon S1P stimulation, while mostly stimulation of the $S1P_2$ receptor in VSMCs causes a contractile effect. The net effect of S1P ligands in a vessel is, therefore, most likely a result of both effects, depending on receptor expression and or regulation.

As found for other GPCRs, S1P receptors are also subject to regulatory processes. Using the immunocytochemical method we have developed (Chapter 3), the internalization of S1P receptors which are expressed in the cardiovascular system was compared (Chapter 5). Some interesting differences were found between the $S1P_1$ and $S1P_2$ receptor on one hand and the $S1P_3$ receptor on the other hand. The $S1P_3$ receptor internalized faster, required lower concentrations of S1P and was internalized more efficiently. The differences in internalization between the

S1P receptor subtypes could only partly be explained by differences in signaling between these receptors. Pertussis toxin treatment showed the involvement of G_i-protein activation for the S1P₃ receptor but not for the S1P₁ and S1P₂ receptor. However, cAMP signaling itself or calcium signaling was not involved in internalization of each of the three S1P receptors. Since [Ca²⁺]_i is increased via S1P₃ receptor stimulation to a much larger extent than via S1P₁ receptor stimulation, we hypothesized that this may play a role in internalization. However, increasing [Ca²⁺]_i in S1P₁ expressing cells or blocking [Ca²⁺]_i elevations in S1P₃ expressing cells did not change the internalization characteristics of either receptor. The observed differences in internalization between the S1P receptors can thus not be completely explained based upon our study. In this respect it is interesting to mention that it has been demonstrated by others that the S1P₃ receptor in contrast to the S1P₁ receptor, does not require phosphorylation to be internalized (Rutherford et al., 2005). Probably this can explain the fast internalization of the S1P₃ compared to the S1P₁ and S1P₂ receptor.

As mentioned before, the signaling properties of the S1P₁, S1P₂ and S1P₃ receptor are different. The S1P₁ receptor exclusively activates G_i-proteins whereas the S1P₂ and S1P₃ receptor can activate various different G-proteins. One conserved domain that has been shown to play an important role in G-protein activation of GPCRs is the DRY motif, which is located at the boundary of the third transmembrane domain and the second intracellular loop. Interestingly, the S1P receptors show a remarkable variance in this highly conserved motif. The S1P₁ receptor is the only receptor that contains an ERY motif, while the other S1P receptors contain either an ERH, ERF or ERS motif. We therefore investigated whether the tyrosine residue of the D/ERY motif is involved in restricting S1P₁ receptor coupling (Chapter 6). Mutating the tyrosine residue to a histidine or alanine residue markedly decreased the potency of S1P to inhibit cAMP accumulation whereas the potency of FTY720-P was unchanged. The effect on the potency of the S1P₁ selective agonist SEW2871 mimicked that of S1P, since mutations in the D/ERY motif also decreased the potency of SEW2871. Mutations in the ERH motif of the S1P₂ and S1P₃ receptor did not induce many changes in signaling, apart from the S1P₂ H¹³¹Y mutant receptor which showed an enhanced increase in cAMP accumulation upon S1P stimulation. Overall, we conclude that tyrosine residue of the D/ERY motif of the S1P₁ receptor does not restrict the receptor from promiscuous coupling or influence constitutive receptor activity as seen for some other GPCRs. However, the mutations somehow

influence the conformational state of the receptor thereby affecting G_i -signaling in response to S1P and SEW2871 but not to FTY720-P. Besides investigating the signaling properties of the mutant receptors we also studied internalization of these receptors in additional experiments. Pilot experiments for the $S1P_1$ receptor mutants indicated that the effects of the mutations observed on G_i -signaling were comparable to the effects found on internalization (data not shown). For the $S1P_2$ and $S1P_3$ mutant receptors no differences in S1P-induced internalization between wild type and mutant receptors were observed (Figure 4). Because the results on

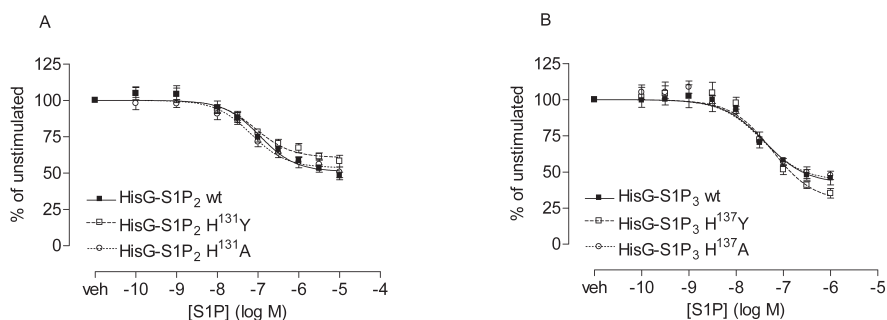


Figure 4. Concentration-dependent S1P-induced internalization of the $S1P_2$ (A) and $S1P_3$ (B) receptor wt and D/ERY-mutants. Stimulations were carried out at 37 °C for 30 min. Values are expressed as percentage of fluorescent signal measured for unstimulated cells which was set at 100% and are presented as means \pm s.e.m. (n=4-13).

internalization were similar to the results obtained for cAMP accumulation for the $S1P_1$ and $S1P_3$ receptor; this could suggest a link between the two pathways.

Recently, several new S1P receptor ligands have become commercially available. In the final part of this thesis, we investigated the signaling of these ligands via the $S1P_3$ receptor (Chapter 7). As this receptor can activate both, G_i and G_q -coupled pathways, the potency and efficacy of a series of compounds on inhibition of the forskolin-induced cAMP production (G_i -mediated) and elevation of $[Ca^{2+}]_i$ (G_q -mediated) was determined. In addition, the potency and efficacy of these compounds to induce $S1P_3$ receptor internalization was investigated. Our study found some interesting differences in the ability of some ligands to activate either G_i or G_q -coupled pathways. For example, the phosphorylated and active form of FTY720, FTY720-P was a full agonist when inhibiting the forskolin-induced cAMP accumulation whereas it had only partial agonistic activity in inducing elevations in $[Ca^{2+}]_i$. Similarly, the $S1P_1$ and $S1P_3$ agonist VPC23153, proved more efficient in inducing G_i -mediated inhibition of cAMP accumulation than in G_q -mediated elevation of $[Ca^{2+}]_i$ via the $S1P_3$ receptor.

In general, the efficacy of the tested ligands to induce S1P₃ receptor internalization matched more closely with their efficacy to inhibit cAMP accumulation than with their efficacy to elevate [Ca²⁺]_i. Another interesting finding that could point in the direction of ligand-directed signaling of the S1P₃ receptor is the finding that at high concentrations S1P but not FTY720-P induces an increase in cAMP accumulation most likely via the activation of G_s-proteins. Besides our findings suggesting ligand-directed signaling at the S1P₃ receptor, it also becomes clear from our study that care should be taken before drawing any definite conclusion about ligand-directed signaling. Most importantly, when comparing potencies determined in signal transduction assays which differ in incubation time it is important to be sure that the ligand concentration is stable in time. Pilot experiments in our laboratory indicated that the concentration of S1P itself declines in time (Figure 5), probably due to degradation and/or uptake, resulting in an underestimation of the potency determined in the cAMP assay when using longer incubation times. In addition, in case the cells used to investigate ligand-directed signaling contain endogenous receptors that also can be activated by some of the ligands under investigations one should also be careful before drawing any conclusions. In conclusion, our study indicates that S1P compared to FTY720-P differentially activates S1P₃-mediated cAMP and calcium signaling pathways probably because both ligands induce different conformations of the S1P₃ receptor.

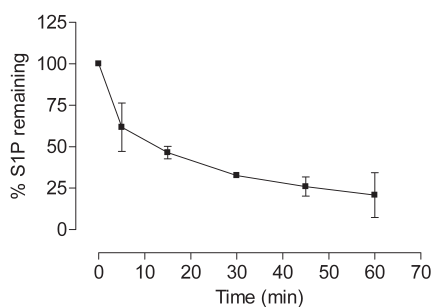


Figure 5. A pilot experiment where disappearance of S1P from stimulation buffer was measured using an HPLC based method (Butter et al., 2005). 2 ml stimulation buffer was incubated with CHO-FlpIn cells expressing the S1P₃ receptor for the indicated times at 37 °C in a 6 wells plate. S1P concentrations were measured using an HPLC method. Data was normalized to the starting S1P concentration as measured by HPLC.

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Summary

Sphingosine-1-phosphate (S1P) is a bioactive lipid that regulates many cellular functions. Currently, five G-protein coupled receptors (GPCRs) are known which are specifically activated by S1P, resulting in a wide range of cellular effects.

Chapter 1 of this thesis gives a general introduction into GPCRs with a special focus on S1P receptors. Their signaling and effects on a molecular level are described, together with the regulation of GPCRs upon stimulation. Furthermore, the role of S1P receptors in the cardiovascular system is described. Also, various tools to study receptor signaling are discussed.

Chapter 2 studies a recently described possible lead compound for a new S1P₃ antagonist. However, our study on this compound, BML-241, shows that besides the established inhibition of S1P₃-mediated changes in $[Ca^{2+}]_i$, this compound does not affect S1P₃ receptor-mediated decreases in cAMP accumulation. Furthermore, BML-241 inhibited changes in $[Ca^{2+}]_i$ mediated via other GPCRs, including the α_{1A} -adrenoceptor. The inhibitory effect on $[Ca^{2+}]_i$ is not mediated via a direct interaction with the receptor as BML-241 did not displace parazosin from the α_{1A} -adrenoceptor. From these results we concluded that BML-241 is a low potency inhibitor of $[Ca^{2+}]_i$ responses instead of a specific S1P₃ receptor antagonist.

In *chapter 3* we describe a new method to study S1P receptor internalization. The method is based on stable cell lines expressing a N-terminally HisG-tagged receptor, which can be visualized using various combinations of antibodies leading to diverse read outs. The best combination turned out to be a specific 1st antibody against the HisG-tag combined with a fluorescent 2nd antibody. The intensity of fluorescence is a direct measure for the level of receptor expression on the membrane. The developed method is quantitative and sensitive enough to measure differences in internalization induced by several ligands. It is also suitable to measure a large number of data points in a short time using a microplate reader.

Chapter 4 studies the vasoactive effects of S1P in a blood vessel. The effects of several S1P receptor ligands on the relaxation of a isolated rat aorta rings were combined with those on the signaling pathways of the two main cell types (smooth muscle

cells (VSMCs) and endothelial cells (ECs)). These effects were correlated to the specific receptor subtypes these ligands bind to and to their relative expression as established by qPCR. All tested compounds induced relaxation of rat aorta rings. In the VSMCs, mostly S1P induced elevation of $[Ca^{2+}]_i$, which in a vessel would result in contraction. In the ECs, most compounds inhibited cAMP accumulation, while hardly any effect was seen on $[Ca^{2+}]_i$ levels. From these data we concluded that in VSMCs, effects occur mostly through the S1P₂ receptor, while in the ECs, effects are a result of activation of the S1P₁ receptor. Therefore, since most compounds, with the exception of S1P itself, lack S1P₂ agonism, their net effect in the rat aorta will be relaxation.

The method described in chapter 3 has been utilized to study internalization of the S1P₁₋₃ receptors, which is described in *chapter 5*. There were striking differences between internalization of the S1P₃ receptor compared to the S1P₁ and S1P₂ receptor. Internalization of the S1P₃ receptor occurs within minutes, while the S1P₁ and S1P₂ receptor internalization requires approximately half an hour. Furthermore, the S1P₃ receptor internalizes at lower S1P-concentrations and shows a higher reduction of membrane receptors. Also, internalization for the S1P₃ receptor is influenced by G_i-inactivation, which does not influence internalization of the other two receptors.

In *chapter 6* the importance of the D/ERY motif in G-protein selectivity of S1P receptors is investigated. The influence of the 3rd amino acid in the D/ERY domain of the S1P₁₋₃ receptor was studied using site-directed mutagenesis with subsequent assessment of the effects of these mutations on the signaling via typical G_i and G_q-coupled pathways. It was found that for the S1P₁ receptor, mutations of this amino acid induced a ligand-dependent effect, in that for S1P and SEW2871, but not for FTY720-P, the G_i-mediated signaling was reduced. However, these mutations did not affect S1P-mediated changes in $[Ca^{2+}]_i$ in these cells. Mutating the ERH motif of the other two receptors did not induce major changes in signaling except for the S1P₂ receptor with an ERY motif instead of its native ERH motif. This mutated receptor showed an elevated S1P-induced increase of cAMP accumulation. From these data we concluded that the tyrosine residue of the D/ERY motif is not involved in restricting the S1P₁ receptor from activating more than one G-protein. However, since the mutation in this receptor differentially affected S1P and SEW2871 on one hand and FTY720-P mediated effects on the other hand we speculate that these

ligands probably induce different conformational states of the receptor which are differentially affected by the introduced mutations.

Several new SIP receptor ligands have recently become commercially available. In *chapter 7* we have studied their potency and efficacy in different signaling pathways via the SIP₃ receptor and in SIP₃ receptor internalization. In this regard, FTY720-P reacted as a full agonist when inhibiting cAMP accumulation, but was a partial agonist when inducing elevations in $[Ca^{2+}]_i$. Also, VPC23153 was more efficient in inhibiting cAMP accumulation than in elevating $[Ca^{2+}]_i$. Furthermore, a supposed antagonist for the SIP_{1,3} receptor, VPC23019, behaved as a partial agonist in all three tested assays, being $[Ca^{2+}]_i$ -signaling, cAMP-signaling and internalization. Another finding was that in high concentrations, SIP, in contrast to FTY720-P, also seemed to activate G_s-coupled pathways. We conclude from these data that several ligands show different response patterns via the SIP₃ receptor. This may be a clue towards ligand-directed signaling for the SIP₃ receptor, but that needs to be studied in more detail.

In *chapter 8*, besides the research in this thesis, other projects which did not result in a separate chapter are discussed. This includes our search for SIP receptor ligands in collaboration with the Free University and the Leiden University. The results were promising, but due to confidentiality reasons, this could not be further discussed. Another project involved a collaboration with the University of Würzburg, where they successfully developed a FRET method to directly measure ligand/induced conformational changes in the adenosine A_{2A} receptor. We have made efforts to apply this method to the SIP₁ receptor. We obtained a FRET signal between the C-terminal CFG-tag and a built-in FRET-motif in the 3rd intracellular loop (IL3), but the FRET ratio did not change upon ligand stimulation. This was possibly due to a shorter, and thus less flexible, IL3 of the SIP₁ receptor compared to the adenosine A_{2A} receptor. However, when truncating the C-terminus of the SIP₁ receptor, we found that the part between aa 361 and 345 plays an important role in receptor surface expression. A final finding was made when the potency of ligands was compared in signaling transduction assays which have different incubation times. Pilot experiments have shown that in time, the SIP concentration declined, possibly due to degradation and/or uptake. Thus one should be careful not to underestimate the potency of a ligand when longer incubation times are used.



Samenvatting

Sphingosine-1-fosfaat (S1P) is een lipide dat in het lichaam diverse functies reguleert op cellulair niveau. Dit gebeurt via vijf G-eiwit gekoppelde receptoren (GPCRs), een soort zintuigen van de cellen, genaamd S1P₁₋₅. In dit proefschrift wordt eerst de ontwikkeling van enkele methodes en technieken besproken, welke nodig zijn om farmacologisch onderzoek te verrichten naar deze receptoren. Het tweede deel beschrijft het onderzoek naar de werking en regulatie van de receptoren zelf.

Hoofdstuk 1 is een introductie over GPCRs in het algemeen en S1P receptoren in detail. Het bespreekt de signalering, regulatie en effecten op een moleculair en cardiovasculair niveau. Tevens worden algemene methoden en technieken besproken die gebruikt worden bij het bestuderen van de signalering via S1P receptoren.

In *hoofdstuk 2* wordt een recent beschreven, nieuw ontwikkelde stof, genaamd BML-241, nader onderzocht. Deze stof werd geselecteerd op zijn eigenschappen om calcium signalering via de S1P₃ receptor, maar niet de S1P₁ receptor te remmen. Dit zou dus een mogelijk begin kunnen zijn van een nieuw te ontwikkelen S1P₃ antagonist (remmer). Echter, wij laten zien dat deze stof ook calcium signalering remt via andere GPCRs. Dit effect wordt niet veroorzaakt door een directe interactie met deze receptoren aangezien BML-241 geen competitie aanging met prazosine voor binding aan de α_{1A} -adrenoceptor. Daarnaast heeft BML-241 geen invloed op remming van de cAMP productie via de S1P₃ receptor. We concluderen dan ook dat het geen specifieke S1P₃ receptor remmer kan zijn, maar dat het meer lijkt op een algemene remmer van calcium signalering.

Hoofdstuk 5 beschrijft de ontwikkeling van een nieuwe methode om internalisatie van GPCRs te bestuderen en kwantificeren. Bij deze methode hebben een sequentie van 7 aminozuren (6xHisG-tag) die met een specifiek antilichaam herkend kunnen worden aan het N-terminale uiteinde van de receptor bevestigd. Er zijn diverse combinaties van antilichamen getest die met behulp van verschillende uitleesmethoden gekwantificeerd kunnen worden gebruikt. De combinatie van het eerste antilichaam met een tweede antilichaam met een fluorescerend label leverde de meest bruikbare methode op. Deze methode bleek zeer nuttig om een groot aantal monsters per keer te bestuderen en kwantificeren.

Hoofdstuk 4 beschrijft het onderzoek naar de effecten van SIP op het cardiovasculaire systeem. Er is gekeken naar de contractie (samentrekking) en relaxatie (ontspanning) van de aorta (de grote lichaamsslagader) van een rat onder invloed van diverse liganden van de SIP_{1,3} receptoren, aangezien hiervan bekend is dat ze voorkomen in de aorta. Dit werd vervolgens gecombineerd met de effecten van deze stoffen op calcium en cAMP signalering in de twee meest voorkomende celtypen in de aorta, namelijk de endotheel cellen (ECs) (die de binnenkant van het bloedvat bekleden) en de gladde spiercellen (VSMCs). Dit werd tot slot gerelateerd aan de expressie van elk van de drie receptoren, gemeten via quantitative PCR. Er bleek dat in de VSMCs de effecten voornamelijk via de SIP₂ receptor worden veroorzaakt, wat resulteert in contractie. In de ECs worden de effecten veroorzaakt via de SIP₁ receptor, wat resulteert in relaxatie. Aangezien alle liganden met uitzondering van SIP zelf geen effecten via de SIP₂ receptor bewerkstelligen, zal het totale effect op de aorta dus een relaxatie zijn.

De methode die we hebben ontwikkeld in hoofdstuk 3 is toegepast om de internalisatie van de SIP_{1,3} receptoren te bestuderen, wat is beschreven in *hoofdstuk 5*. Er bleek een verschil te zijn in internalisatie van de SIP₃ receptor vergeleken met de SIP₁ en SIP₂ receptor. De SIP₃ receptor internaliseert al binnen enkele minuten na stimulatie, terwijl dit bij de andere twee receptoren ongeveer een half uur duurt. Bovendien bleek de internalisatie van de SIP₃ receptor ook nog eens te gebeuren bij een veel lagere SIP concentratie en er bleven veel minder SIP₃ receptoren in het celmembraan over dan SIP₁ of SIP₂ receptoren. Tevens bleek de internalisatie van de SIP₃ receptor te worden beïnvloed door het G_i-eiwit, wat bij de andere receptoren niet het geval was.

In *hoofdstuk 6* wordt het belang van een bepaald gebied (domein) van de SIP receptoren bestudeerd. Dit domein bestaat uit drie aminozuren, gecodeerd als D/ERY, en is bij diverse andere GPCRs al van belang gebleken voor signalering en constitutieve activiteit van de receptoren. Er is een opvallend verschil tussen de SIP receptoren op dit gebied. De SIP₁ receptor activeert namelijk alleen het G_i-eiwit en bevat een ERY domein, terwijl de andere receptoren meerdere G-eiwitten activeren en een ERH(SIP_{2,3}), ERF (SIP₄) of ERS (SIP₅) domein bevatten. Zou dus deze tyrosine (Y) verantwoordelijk zijn voor het activeren van slechts het G_i-eiwit? Dit is onderzocht door het ERY motief van de SIP₁ receptor te vervangen

door het ERH motief van de SIP_2 en SIP_3 receptor en vervolgens te kijken naar het effect op de signalering, als gevolg van de activatie van de betreffende G-eiwitten. De signalering voor de SIP_1 receptor via het G_i -eiwit bleek te zijn verminderd, maar slechts voor 2 agonisten (activators) namelijk SIP en SEW2871 (die specifiek de SIP_1 receptor activeert) en niet voor een andere agonist FTY720-P. Echter, er werd geen verschil gevonden voor een signalerings route via het G_q -eiwit. Voor de andere receptoren bleken er geen grote verschillen te zijn tussen de oorspronkelijke en de gemuteerde receptoren. De SIP_2 receptor met een ERY domein in plaats van een ERH domein bleek cAMP concentraties in de cel flink te verhogen. Uit deze data concluderen we dat er geen verband is gevonden tussen het D/ERY domein in de SIP receptoren en hun specifieke G-eiwit signalering. Echter, er is een opvallend verschil in respons tussen de liganden SIP en SEW2871 aan de ene kant en FTY720-P aan de andere. Het zou kunnen dat FTY720-P een andere receptor conformatie induceert dan SIP en SEW2871 waardoor de mutaties in het D/ERY domein ook een andere effect op de signalering hebben.

Recentelijk zijn nieuwe liganden beschikbaar gekomen. In *hoofdstuk 7* zijn eigenschappen van deze stoffen onderzocht in diverse signaleringsroutes via de SIP_3 receptor. Dit resulteerde in de ontdekking dat een bepaald ligand, VPC23019, dat werd beschreven als een antagonist zich gedroeg als een partiële agonist in de geteste signalerings routes. Als extra werd nog gevonden dat SIP, maar niet FTY720-P, G_s -gekoppelde signalering bleek te activeren via de SIP_3 receptor. We concluderen dan ook dat de geteste liganden via de SIP_3 receptor verschillende signaleringspatronen induceren, wat mogelijk duidt op een nieuw fenomeen ook wel ligand-gestuurde signalering noemen.

Hoofdstuk 8 wordt naast het onderzoek dat beschreven staat in dit proefschrift, ook het onderzoek bediscussieerd dat nite heeft geleid tot een apart hoofdstuk. Dit omvat onze speurtocht naar nieuwe SIP receptor liganden in samenwerking met de Vrije Universiteit en de Universiteit Leiden. Wegens geheimhouding mogen deze veelbelovende resultaten echter niet in detail worden besproken.

Een ander project was een samenwerking met de Universiteit in Würzburg, waar een elegante FLAsH-based FRET methode was ontwikkeld om conformatie veranderingen geïnduceerd door ligand binding in de adenosine A_{2A} receptor te meten. Wij hebben geprobeerd om deze methode ook toe te passen voor de

SIP₁ receptor. Ondanks dat we wel een FRET signaal konden meten tussen een C-terminale CFP en een FLAsH-motif in de 3^e intracellulaire loop (IL3), bleek deze niet te veranderen onder invloed van ligand binding. Een mogelijke oorzaak hiervoor is de veel kortere en dus minder flexibele IL3 van de SIP₁ receptor in vergelijking met de adenosine A_{2A} receptor. Wel werd tijdens trunceren van de C-terminus van de SIP₁ receptor duidelijk, dat deze tussen aminozuur 361 en 345 een rol speelt bij expressie van de receptor aan de membraan.

Tot slot bemerkten we nog tijdens het vergelijken van liganden met behulp van verschillende signaaltransductie assays, dat de incubatie tijd erg van invloed kan zijn op de potentie van het ligand. Proef-experimenten hebben aangetoond dat de SIP concentratie daalt in de tijd, mogelijk door degradatie of opname door de cel. Men moet dan ook oppassen met het vergelijken van ligand eigenschappen met behulp van verschillende assays, aangezien dit kan leiden tot een onderschatting van de werkelijke potency van het ligand.





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Jongsma M, Hendriks-Balk MC, Michel MC, Peters SL, Alewijnse AE. BML-241 fails to display selective antagonism at the sphingosine-1-phosphate receptor, SIP₃. *Br J Pharmacol*. 2006; **149**:277-282

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Jongsma M, Mathy MJ, Van Loenen PB, Hajji N, Michel MS, Peters SLM, Alewijnse AE. Pharmacological and biochemical characterization of the vasoactive properties of sphingosine-1-phosphate receptors in rat aorta. *Manuscript submitted*

Jongsma M, Michel MC, Peters SLM, Alewijnse AE. Differential agonist-induced internalization of sphingosine-1-phosphate receptor subtypes 1, 2 and 3. *Manuscript in preparation*

Jongsma M, Den Boon FS, Van Loenen PB, Michel MC, Peters SLM, Alewijnse AE. Involvement of the third amino acid of the D/ERY motif in G-protein selectivity of SIP receptors. *Manuscript in preparation*

Jongsma M, Van Unen J, Hajji N, Michel MC, Peters SLM, Alewijnse AE. Differential response patterns of several ligands at the sphingosine-1-phosphate receptor subtype 3. *Manuscript in preparation*



Dankwoord

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Een volledig nieuwe afdeling, waarin alles nog moest worden geregeld, besteld, getest en opgezet, was de omgeving waarin ruim vier jaar geleden zes nieuwe AIO's zijn gestart, waarvan ik er een was. Uiteraard wordt hierbij veel gevallen en weer opgestaan. In mijn opinie is het onjuist om pas achteraf te laten blijken dat mensen belangrijk voor je zijn op bepaalde momenten in je leven. Het is juist zeer wenselijk om ze ook op de betreffende momenten te laten weten dat je de ontvangen hulp en steun waardeert. Ik hoop dat ik dat dan ook gedaan heb en dat de mensen die hieronder hun naam zullen lezen dat kunnen beamen. Ik houd niet van volgorde, rangen en standen, maar voor mij is iedereen gelijk. Echter, in een dankwoord ontkom je niet aan een volgorde, maar voor mij is die dus willekeurig.

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laten terechtkomen. In mijn geval is je dat goed gelukt. Mijn postdoc plek op het NKI/AVL is in mijn ogen een directe verdienste van jou. Ik hoop dat je na jaren van investeren nu kunt gaan oogsten met mooie publicaties met de nieuwe lichte AIO's. Heel veel succes hierbij en ik hoop dat we elkaar in de toekomst nog kunnen helpen.

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Maikel





Curriculum Vitae

The author of this thesis was born on June 6th, 1977 in Diepenheim. At the age of 10, he moved to Hengelo where he graduated from his pre-university education at the R.K. Lyceum de Grundel at the age of 17. He started his study of Chemical Engineering at the University of Twente in 1994, from which he did not graduate. The following year, he started his Medical Laboratory Education at the University of Professional Education in Enschede. This was successfully concluded in 1999 including a 10 month internship, working on *Haemophilus influenzae* at the National Institute for Public Health and the Environment in Bilthoven, under supervision of Dr. P. van Ulsen. In that same year he started his Biology education at the University of Utrecht, which included a 9 month practical period at the University of Stellenbosch (South Africa) under supervision of Prof.dr. J.L. Snoep. There he worked on glucose-dependent growth of *Saccharomyces cerevisiae*. After graduation in December 2002, he started in the summer of 2003 to work in the lab of Prof.dr. M.C. Michel, under supervision of Dr. A.E. Alewijnse at the department of Pharmacology and Pharmacotherapy at the Academic Medical Center in Amsterdam. It was here that he performed the work described in this thesis. At the Keystone meeting in Taos (NM, USA) on Bioactive Lipids in the Lipidomics Era in 2007 he was awarded a Keystone Symposia Scholarship for his work on internalization of sphingosine-1-phosphate receptors. In August 2007 he started as a postdoc in the lab of Prof.dr. W.H. Moolenaar at the Netherlands Cancer Institute to work on lysophosphatidic acid receptors.