HEALTH SCIENCES

JAYENDRA Z. PATEL

Development of Potent and Selective Inhibitors of Enzymes Involved in Endocannabinoid Inactivation

Publications of the University of Eastern Finland Dissertations in Health Sciences



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ABSTRACT

This doctoral dissertation describes the design, synthesis and structure-activity relationships (SARs) of novel 1,3,4-oxadiazol-2-one, 1,2,5-thiadiazole carbamate, and loratadine analogues as promising inhibitors of the endocannabinoid hydrolases, such as FAAH, ABHD6, and MAGL. Inhibition of endocannabinoid hydrolases by specific inhibitors enhance endocannabinoid signaling, which in turn has been linked to various therapeutic effects of cannabinoids. In recent years, inhibitors of FAAH, MAGL and ABHD6 have shown potential to treat several diseases, such as pain, inflammation, CNS disorders and metabolic disorders without association of cannabimimetic side effects.

By adopting a hybridization approach, the earlier discovered 1,3,4-oxadiazol-2-ones were combined with nonsteroidal anti-inflammatory drugs (NSAIDs) to produce chiral 1,3,4-oxadiazol-2-ones of which JZP-327A was found to be a slowly reversible, potent FAAH inhibitor with an IC₅₀ value of 11 nM. Systematic structural variations of 1,2,5-thiadiazole carbamates resulted in the identification of JZP-430 as an irreversible, potent ABHD6 inhibitor with an IC₅₀ value of 44 nM. By increasing the electrophilicity on the carbonyl functionality of loratadine, several novel MAGL inhibitors were identified of which JZP-361 acted as the most potent MAGL inhibitor having an IC₅₀ value of 46 nM. Interestingly, JZP-361 exhibited slowly reversible MAGL inhibition and it retained its histamine H₁ antagonistic affinity. The optimization of 1,3,4-oxadiazol-2-one by carrying out systematic structural variations led identification of JZP-169 as a potent ABHD6 inhibitor (IC₅₀ value of 216 nM) having an irreversible mode of inhibition. All these compounds showed selectivity over endocannabinoid hydrolases, cannabinoid receptors, and other studied off-targets.

In summary, the results presented here demonstrate that 1,3,4-oxadiazol-2-one is a promising scaffold for the development of potent and selective inhibitors of serine hydrolases: JZP-327A for FAAH and JZP-169 for ABHD6. Interestingly, 1,3,4-oxadiazol-2-ones showed reversible inhibition mechanism against FAAH whereas they inhibited ABHD6 in an irreversible manner. The ABHD6 inhibitor JZP-430 may serve as an important lead compound when developing novel drugs for metabolic disorders. Finally, as a dual blocker of MAGL activity and the histamine H₁ receptor, the loratadine analogue JZP-361 may have potential to treat inflammation in chronic diseases, such as asthma.

National Library of Medicine Classification: QU 143, QV 126, QV 744

Medical Subject Headings: Chemistry, Pharmaceutical; Chemistry Techniques, Synthetic; Endocannabinoids; Enzyme Inhibitors; Hydrolases; Amidohydrolases; Monoacylglycerol Lipases; Drug Discovery; Oxadiazoles; Thiadiazoles; Loratadine/analogs & derivatives; Carbamates; Structure-Activity Relationship



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TIIVISTELMÄ

Tässä väitöskirjassa kuvataan uusia 1,3,4-oksadiatsol-2-oni, 1,2,5-tiadiatsolikarbamaatti- ja loratadiinijohdoksia lupaavina FAAH-, ABHD6- ja MAGL–inhibiittoreina sekä esitetään niiden suunnittelu, synteesi ja rakenne-aktiivisuussuhteet (SAR). Endokannabinoidijärjestelmän hydrolaasien toiminnan estäminen erityisillä ensyymiinhibiittoreilla lisää endokannabinoidien signalointia, millä on yhteys kannabinoidien terapeuttisiin vaikutuksiin. Viime vuosina FAAH-, MAGL- ja ABHD6-inhibiittoreilla on osoitettu olevan terapeuttista tehoa useiden sairauksien, kuten kivun, tulehduksen ja aineenvaihdunnan häiriöiden hoidossa ilman kannabimimeettisiä sivuvaikutuksia.

Yhdistämällä aiemmin tunnettu 1,3,4-oksadiatsol-2-oni -rakenne tulehduskipulääkkeeseen (NSAID) saatiin aikaan kiraalisia 1,3,4-oksadiatsol-2-oni –rakenteita, joista yhdisteen JZP-327A havaittiin olevan hitaasti palautuva tehokas FAAH-estäjä IC50-arvolla 11 nM. Systemaattinen 1,2,5-tiadiatsolikarbamaatti-rakenteiden rakenne-aktiivisuus tarkastelu johti yhdisteeseen JZP-430, joka on tehokas palautumaton ABHD6-inhibiittori IC50-arvolla 44 nM. Lisäämällä loratadiinin karbonyyliryhmän elektrofiilisyyttä saatiin aikaiseksi useita uusia MAGL-estäjiä, joista JZP-361 oli voimakkain MAGL-estäjä IC50-arvolla 46 nM. Tutkimuksessa JZP-361:n todettiin olevan mekanismiltaan hitaasti palautuva MAGL-estäjä ja se säilytti affiniteetin H1-reseptoriin. 1,3,4-Oksadiatsol-2-oni -rakennetta optimoitiin valmistamalla systemaattisesti erirakenteisia johdoksia, joista yhdisteen JZP-169 todettiin olevan tehokas palautumaton ABHD6-estäjä (IC50-arvo 216 nM). Kaikkien näiden vhdisteiden todettiin olevan selektiivisiä endokannabinoidijärjestelmän hydrolaaseihin, kannabinoidireseptoreihin ja muihin tutkittuihin kohteisiin verrattuna.

Yhteenvetona voidaan todeta, että 1,3,4-oksadiatsol-2-oni –rakenne on käyttökelpoinen runkorakenne kehitettäessä tehokkaita ja selektiivisiä seriinihydrolaasiestäjiä, kuten työssä kehitetyt FAAH-estäjä JZP-327A ja ABHD6-estäjä JZP-169. Mielenkiintoinen tulos oli, että 1,3,4-oksadiatsol-2-oni –rakenteesta voitiin kehittää sekä palautuva FAAH-estäjä että palautumaton ABHD6-estäjä. Työssä löydetty ABHD6-estäjä JZP-430 voi toimia tärkeänä johtoyhdisteenä kehitettäessä uusia lääkehoitoja aineenvaihdunnan häiriöihin. Lopuksi, MAGL-entsyymin että histamiini-H1-reseptorin salpaaja, loratadiinianalogi JZP-361, voi olla käyttökelpoinen kroonisten tulehdussairauksien, kuten astman hoidossa.

Luokitus: QU 143, QV 126, QV 744

Yleinen suomalainen asiasanasto: orgaaninen kemia; lääkekemia; kemiallinen synteesi; endokannabinoidit; entsyymit; inhibiittorit; lääkesuunnittelu; rakenne; aktiivisuus



To My Family



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Kuopio, September 2015

Jayendra Z Patel

List of the original publications

This dissertation is based on the following original publications, denoted in the text as Roman numerals I-IV:

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- III Patel J Z, Ahenkorah S, Vaara M, Staszewski M, Adams Y, Laitinen T, Navia-Paldanius D, Parkkari T, Savinainen J R, Walczyński K, Laitinen J T and Nevalainen T J. Loratadine Analogues as MAGL Inhibitors. *Bioorg. Med. Chem. Lett. 25: 1436-1442, 2015.*
- IV Patel J Z, van Bruchem J, Laitinen T, Kaczor A A, Navia-Paldanius D, Parkkari T, Savinainen J R, Laitinen J T and Nevalainen T J. Revisiting 1,3,4-Oxadiazol-2-ones: Utilization in the Development of ABHD6 Inhibitors.
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APPENDICES: ORIGINAL PUBLICATIONS (I-IV)

Abbreviations

AA	arachidonic acid	BSA	bovine serum albumin
AA-5-HT	arachidonoyl serotonin	BuChE	butyrylcholinesterase
ABHD	α/β -hydrolase domain	СВ	cannabinoid
	containing	CB ₁	cannabinoid receptor
ABPP	activity-based protein		subtype-1
	profiling	CB ₂	cannabinoid receptor
ACN	acetonitrile		subtype-2
AChE	acetylcholinesterase	CCI	chronic constriction
ADMK	arachidonoyl		injury
	diazomethyl ketone	CDTA	calcium-dependent
AEA	N-arachidonoyl-		transacylase
	ethanolamine	CEs	carboxylesterases
	(anandamide)	CFA	complete Freund's
1(3)-AG	1(3)-arachidonoyl-		adjuvant
	glycerol	СНО	Chinese hamster ovary
2-AG	2-arachidonoyl-	<i>p</i> -CMB	<i>p</i> -chloromercuribenzoic
	glycerol		acid
2-AGE	2-arachidonyl	CNS	central nervous system
	glyceryl ether	COX	cyclooxygenase
	(noladin ether)	CRC	concentration response
AMC-AA	7-amino-4-methyl		curve
	coumarin arachidon-	СҮР	cytochrome P450
	amide	Cys	cysteine
AS	amidase signature	DAG	diacylglycerol
ATK	arachidonyl trifluoro-	DAGL	diacylglycerol lipase
	methyl ketone	DCM	dichloromethane
ATP	adenosine triphosphate	DEA	docosatetraenoyl-
			ethanolamide

XVIII

DHAP	dihydroxyacetone	FC	flash chromatography
	phosphate	FDA	Food and Drug
DMAP	4-(dimethylamino)-		Administration
	pyridine	FPs	fluorophosphonates
DMF	N,N-dimethylform-	GDE1	glycerophosphodiester
	amide		phosphodiesterase 1
DMSO	dimethylsulfoxide	GK	glycerol kinase
DPP-IV	dipeptidyl peptidase-4	G-1-P	glycerol-1-phosphate
ECS	endocannabinoid	GPCR	G-protein coupled
	system		receptor
EDTA	ethylenediamine-	GPO	glycerol phosphate
	tetraacetic acid		oxidase
EET-EA	epoxyeicosatrienoic	HEK293	human embryonic
	ethanolamide		kidney 293
EMT	endocannabinoid	hERG	human ether-a-go-go
	membrane transporter		related gene
ERG	electron releasing	HETE-EA	hydroxyeicosatetra-
	group		enoic acid ethanol-
EtOAc	ethyl acetate		amide
EtOH	ethanol	HPLC	high-performance
EWG	electron withdrawing		liquid chromatography
	group	HRP	horseradish peroxidase
FAAH	fatty acid amide	IC ₅₀	half maximal inhibitory
	hydrolase		concentration
FAAH-2	fatty acid amide	i.p	intraperitoneal
	hydrolase-2	IPA	isopropanol
FABP	fatty acid binding	Ki	inhibition constant
	protein	LOX	lipoxygenase
FAPA	fatty acid primary	LPS	lysophosphatidylserine
	amide	MAFP	methyl arachidonoyl
			fluorophosphonate

MAG/MAGL	monoacylglycerol	2-OG	2-oleoylglycerol
	lipase	PAP	phosphatidic acid
MeOH	methanol	PBS	phosphate-buffered
mSH	metabolic serine		saline
	hydrolase	PEA	N-palmitoylethanol-
MTI	mild thermal injury		amine (palmitoyl-
4-MUO	4-methylumbelli-		ethanolamide)
	ferone oleate	PE	petroleum ether
NAAA	N-acylethanolamine-	PG	prostaglandin
	hydrolyzing acid	PG-EA	prostaglandin-
	amidase		ethanolamide
NADA	N-arachidonyl	PGE2	prostaglandin E2
	dopamine	PG-GE	prostaglandin-
NAE	N-acylethanolamine		glyceryl ester
NAGIy	N-arachidonoyl-	PHARC	polyneuropathy,
	glycine		hearing loss, ataxia,
NAM	N-arachidonyl-		retinitis pigmentosa,
	maleimide		cataract
NAPE-PLD	N-acyl phosphatidyl-	PI-PLC	phosphatidyl inositides
	ethanolamine phos-		(PI)- selective phospho-
	pholipase D		lipase C
NAT	N-acyltaurine	PL	phospholipase
NAPE	N-arachidonoyl	PLC	phospholipase C
	phosphatidylethanol-	PLD	phospholipase D
	amine	PPAR	peroxisome proliferator
NEM	N-ethylmaleimide		activated receptor
NSAID	non-steroidal anti-	PTPN22	protein tyrosine phos-
	inflammatory drug		phatase, nonreceptor
OEA	N-oleoylethanolamine		type 22
		SAR	structure-activity
			relationship

XIX

S.E.M.	standard error of the	ТНС	(-)-∆ ⁹ -tetrahydro-
	mean		cannabinol
Ser	serine	THF	tetrahydrofuran
[³⁵ S]-GTPγS	guanosine 5'-O-(3-	THL	tetrahydrolipstatin
	[³⁵ S]-thiotriphosphate		(orlistat)
SH	serine hydrolase	TMS	tetramethylsilane
SHIP	SH2 (Src Homology 2)	∆TM-rFAAH	transmembrane
	domain-containing		domain deleted rat
	inositol 5'-phosphtase		FAAH
SNL	spinal nerve ligation	Tris-HCI	tris(hydroxymethyl)-
sPLA2	soluble form of		aminomethane
	phospholipase A2		hydrochloride
TAMRA-FP	carboxytetramethyl-	TRPV	transient receptor
	rhodamine fluoro-		potential vanilloid
	phosphonate	TSRI	The Scripps
TEA	triethylamine		Research Institute
TGH	triacylglycerol	UEF	The University of
	hydrolase		Eastern Finland

ΧХ

1 Introduction

Enzymes are biological catalysts responsible for mediating a wide variety of metabolic processes. They accelerate both the rate and specificity of metabolic chemical reactions and convert substrates into products. Both the specificity and their mechanistic diversity make enzymes appealing targets also for drug development (Patrick 2009) and today a large number of commonly used drugs are enzyme inhibitors. Currently, structure-based drug design is used, in which the three-dimensional (3D) structure of an enzyme's active site is used in the design of inhibitors of catalytic activity. A molecule acting as an inhibitor binds to the active site of enzyme in either a reversible or irreversible manner and when it is there, it is able to prevent a substrate from entering the enzyme's active site and/or hamper the enzymatic activity in some other way (Thomas 2007). Reversible inhibitors bind to enzymes noncovalently via ionic bonds, hydrogen bonds and hydrophobic interactions i.e. they do not undergo chemical reactions when bound to enzyme but have a tendency to form an equilibrium system with the enzyme and can thereby readily be removed by dilution or dialysis. There are some exceptions, a few reversible inhibitors bind by forming weak covalent bonds. In contrast, irreversible inhibitors predominantly form strong covalent bonds by modifying an essential residue present in the enzyme and thus their effect cannot be reversed. The reversible inhibitors can be classified as a competitive, uncompetitive, and noncompetitive while the irreversible inhibitor can be further classified as group-specific covalent modifying agents, affinity labels, transition state analogs, and suicide inhibitors (mechanism-based inhibitors). Earlier drug discovery attempts were focussed mainly on the development of compounds undergoing noncovalent interaction with targets. This is because of possible off-target interactions through the covalent bond which might result in long lasting detrimental effects. Nevertheless, it is possible to minimize undesirable interactions by modulating the electrophilic warhead reactivity of designed inhibitor. Indeed, several drugs forming covalent bonds with their targets with an acceptable side effect profiles have recently been approved by U.S. Food and Drug Administration (USFDA) (Mah et al., 2014). Additionally, a few more compounds are currently in the preclinical or earlystage clinical phase (Singh et al., 2011, Mah et al., 2014). In fact, many essential and life saving drugs including aspirin, penicillin, omeprazole and clopidogrel were discovered later to act via the covalent mechanism (Mah et al., 2014, Bauer 2015).

The serine hydrolase (SH) superfamily comprises ~240 enzymes in humans and generally they contain a conserved nucleophilic Ser residue which attacks ester, amide and thioester groups of substrate in order to facilitate hydrolysis. In general, SHs are categorized into two subfamilies - the serine proteases (~125 members, such as trypsin, thrombin, subtilisin enzymes) and the metabolic SHs (mSHs, ~115 members) (Simon & Cravatt 2010, Long &

Cravatt 2011). The mSHs include small-molecule hydrolases, such as amidases (e.g. FAAH), lipases (e.g. MAGL, ABHD6, ABHD12), esterases (e.g. acetylcholinesterase (AChE)), and peptidases (e.g. dipeptidyl peptidase-4 (DPP-IV)). Over 60% of mSHs adopt an α/β -fold and normally use a Ser-His-Asp triad for catalysis (Holmquist 2000) while some mSHs employ diverse folds and a catalytic dyad (e.g., Ser-Lys or Ser-Asp) or triad (e.g., Ser-Ser-Lys) (Dodson & Wlodawer 1998, Patricelli & Cravatt 1999). SHs play crucial roles in many biological processes, and inhibitors of several SHs are approved for clinical use, such as in type 2 diabetes (T2D), Alzheimer's disease (AD), and infectious diseases (Bachovchin & Cravatt 2012). However, there are still several poorly characterized SHs with respect to their substrate, inhibitors and (patho)physiological functions (Lord et al., 2013). Due to presence of nucleophilic serine residue, the functional state of most SHs can be assessed by using active-site directed affinity labels such as fluorophosphonates (FPs) (Liu et al., 1999, Simon & Cravatt 2010). At present, several selective inhibitors have been discovered by competitive ABPP (Bachovchin & Cravatt 2012). The competitive ABPP is also used extensively in the present study to confirm the selectivity of developed inhibitors toward specific endocannabinoid hydrolase (FAAH, MAGL, ABHD6/12) over other serine hydrolases (SHs) of mouse brain membrane proteomes (see subsection 5.3.12).

The biological significance of SHs has motivated numerous researchers to develop inhibitors for enzymes in this class, both for use as chemical tools to study enzyme function and as potential new therapeutic agents. This doctoral dissertation focuses on the serine hydrolases that belong to the so-called endocannabinoid system (ECS). Specific blockade of the endocannabinoid hydrolases, fatty acid amide hydrolase (FAAH), monoacyl glycerol lipase (MAGL), α/β -hydrolase domain containing 6 and 12 (ABHD6/12) prolongs the life-time of the endocannabinoid hydrolases are considered as compounds with great therapeutic potential to treat many common pathological conditions, such as pain, inflammation, central nervous system (CNS) disorders and metabolic disorders, none of the inhibitors have yet reached the clinics.

In this doctoral thesis, the design, synthesis and *in vitro* evaluation of compounds acting as potent and selective inhibitors of FAAH, MAGL and ABHD6 have been carried out. Moreover, a completely novel scaffold for the development of ABHD6 inhibitors has been identified and a dual-acting pharmacological tool possessing both MAGL inhibitory and antihistaminergic activities has been discovered. The inhibitors described in this thesis have great potential in clarifying the therapeutic potential of the ECS, for example, they can be considered as valuable leads for further drug development.

2 Review of the Literature

2.1 EMERGENCE OF THE ENDOCANNABINOID SYSTEM (ECS)

Twenty-five years of extensive research on the endocannabinoid system (ECS) has greatly improved our understanding of this highly complex signalling network. The ECS comprises two G-protein coupled cannabinoid receptors (GPCRs), their endogenous ligands (also known as endocannabinoids), and the proteins responsible for endocannabinoid biosynthesis, transportation and biodegradation.



Figure 1. Activation of cannabinoid receptors by Δ^9 -THC and endocannabinoids: AEA and 2-AG.

The hemp plant, *Cannabis sativa*, has been used for recreational and medicinal purposes for thousands of years, and it is known to affect mood, memory, cognition, pain etc. In 1964, researchers from Israel isolated the main psychoactive substance of *Cannabis sativa*, Δ^{9} -tetrahydrocannabinol (Δ^{9} -THC; Figure 1) (Gaoni & Mechoulam 1964). Later, in the early 1990s, the psychoactive effect of THC was found to be mediated through the activation of two GPCRs that were subsequently named cannabinoid receptor subtype-1, CB₁ (Devane *et al.*, 1988, Matsuda *et al.*, 1990) and cannabinoid receptor subtype-2, CB₂ (Munro *et al.*, 1993) (Figure 1). The CB₁ receptor is expressed mainly in the CNS, but is also present in peripheral tissues, including the lungs, liver, kidneys and adipose tissue (Pagotto *et al.*, 2006). The CB₂ receptor is expressed primarily by immune cells, including microglia in the brain (Munro *et al.*, 1993, Onaivi 2006). It has been found that the majority of the CNS effects of THC are

mediated by the CB₁ receptor (Zimmer *et al.*, 1999, Ledent *et al.*, 1999) while the immunosuppressive effects of cannabinoids are attributable to activation of the CB₂ receptor (Lunn *et al.*, 2006).

Soon after the discovery of the CB₁ receptor, the first endogenous cannabinoid ligand named as *N*-arachidonoylethanolamine (anandamide, AEA; Figure 1) was isolated in 1992 from porcine brain (Devane *et al.*, 1992). A second endocannabinoid, 2-arachidonoylglycerol (2-AG; Figure 1), was identified three years later from canine gut and rat brain (Mechoulam *et al.*, 1995, Sugiura *et al.*, 1995). AEA and 2-AG belong to a much larger class of lipids, so-called *N*-acylethanolamines (NAEs) and monoacylglycerols (MAGs), respectively. The lipids belonging to these classes differ in the length and degree of unsaturation of their acyl chains. Although, AEA and 2-AG have structural similarities and they bind to both cannabinoid receptors, they possess different efficacies *in vitro*: AEA acts as a partial agonist while 2-AG is considered to be a full agonist (Sugiura *et al.*, 1995, Stella *et al.*, 1997, Sugiura *et al.*, 1999, Sugiura *et al.*, 2000, Gonsiorek *et al.*, 2000, Savinainen *et al.*, 2001). Furthermore, the brain levels of 2-AG have been found to be 170-fold higher than those of AEA (Sugiura *et al.*, 1995, Stella *et al.*, 1997).

In addition to AEA and 2-AG, several other endocannabinoids have been described such as 2-arachidonyl glyceryl ether (2-AGE, noladin ether) (Sugiura *et al.*, 1999, Hanus *et al.*, 2001), *N*-arachidonoyl dopamine (NADA) (Bisogno *et al.*, 2000), *N*-stearoylethanolamine (Maccarrone *et al.*, 2002), *O*-arachidonoyl ethanolamine (virodhamine) (Porter *et al.*, 2002), homo-γ-linolenoyl ethanolamide and docosatetraenoylethanolamide (DEA) (Hanus *et al.*, 1993), although their biological activity and metabolism have not been fully clarified. Moreover, several endogenous fatty acid derivatives like *N*-palmitoylethanolamine (PEA) (Sheskin *et al.*, 1997, Lambert *et al.*, 1999), *N*-oleoylethanolamine (OEA) (Rodriguez *et al.*, 2001, Oveisi *et al.*, 2004, Nielsen *et al.*, 2004) and *N*-arachidonoylglycine (Sheskin *et al.*, 1997, Huang *et al.*, 2001) have been identified and termed as ''endocannabinoid-like'' compounds since they lack direct cannabinoid affinity. However, at present, out of all these endocannabinoid lipids, only AEA and 2-AG have been well-explored (Howlett *et al.*, 2002, Piomelli 2003, Pacher *et al.*, 2006).

Endocannabinoid biosynthesis

AEA and 2-AG are not prestored in vesicles but are produced "on demand" in postsynaptic nerve terminals and released only when and where necessary in response to physiological or pathological stimuli (Marsicano *et al.*, 2003, Alger & Kim 2011). Biochemical studies have revealed numerous biosynthetic pathways for AEA and 2-AG.

Biosynthetic pathways for AEA



Figure 2. AEA biosynthesis. Abbreviations: CDTA, calcium-dependent transacylase; NAPE-PLD, *N*-arachidonoyl-phosphatidylethanolamine-specific phospholipase D; ABHD4, α/β -hydrolase domain containing 4; sPLA₂, soluble form of phospholipase A2; GDE1, glycerophosphodiester phosphodiesterase 1; PLC, phospholipase C; PTPN22, protein tyrosine phosphatase, non-receptor type 22; SHIP, SH2 (Src Homology 2) domain-containing inositol 5'-phosphatase; R₁, R₂ and R₃, fatty acids.

A number of pathways have been proposed for AEA biosynthesis from its phospholipid precursor, *N*-arachidonoyl-phosphatidylethanolamine (NAPE) (Figure 2). NAPE, in turn, is believed to originate from the calcium-dependent transacylase (CDTA) enzymatic transfer of

arachidonic acid (AA) from the *sn*-1 position of phosphatidylcholine (PC) to the primary amine of phosphatidylethanolamine (PE; Step-1) (Cadas *et al.*, 1996, Cadas *et al.*, 1997). Several alternative AEA biosynthesis pathways have been proposed: (i) direct synthesis of AEA from NAPE by NAPE-specific phospholipase D (NAPE-PLD; Step-2) enzyme (Okamoto *et al.*, 2004); (ii) sequential *O*-deacylation of NAPE by α/β -hydrolase domain containing 4 (ABHD4; Steps 3–4) (Simon & Cravatt 2006), followed by the phosphodiesterase-mediated hydrolysis by the glycerophosphodiester phosphodiesterase 1 (GDE1; Step 5) (Simon & Cravatt 2008) (iii) *O*-deacylation of NAPE by a soluble form of phospholipase A2 (sPLA₂.Step 3), followed by hydrolysis of the phosphodiester bond by a lyso-PLD enzyme (step 6) (Sun *et al.*, 2004) ; and finally, (iv) conversion of NAPE to phospho-AEA by a phospholipase C (PLC)-like enzyme (step 7) followed by dephosphorylation by protein tyrosine phosphatase, non-receptor type 22 (PTPN22) (Liu *et al.*, 2006) or SH2 (Src Homology 2) domain-containing inositol 5'-phosphatase (SHIP) (step 8) (Liu *et al.*, 2008).

Biosynthetic pathways for 2-AG



Figure 3. 2-AG biosynthesis. Abbreviations: PI-PLC, phosphatidylinositides (PI)-selective phospholipase C; PAP, phosphatidic acid (PA) phosphohydrolase; DAGL, diacylglycerol lipase; R₁, fatty acid.

The main pathway for 2-AG formation is from the hydrolysis of diacylglycerols (DAGs) in the presence of DAG lipase isoenzymes (DAGL α and DAGL β ; Step-3; Figure 3) (Bisogno *et al.*, 2003). DAGs, in turn, can be produced via the hydrolysis of either phosphatidylinositides (PI), catalyzed by a PI-selective phospholipase C (PI-PLC; Step-1) (Stella *et al.*, 1997, Kondo *et al.*, 1998), or from phosphatidic acid (PA), catalyzed by a PA phosphohydrolase (PAP; Step2) (Bisogno *et al.*, 1999). Two other pathways have been proposed: (i) the sequential actions of a phosphatidylinositol-preferring phospholipase A₁ (PLA₁), producing the 2-arachidonoyl-lysophosphatidylinositol (lyso-PI) intermediate, followed by lysoPI-specific phospholipase C (lyso-PLC) generating 2-AG (Steps 4-5) (Ueda *et al.*, 1993); (ii) the conversion from 2-arachidonoyl lysophosphatidic acid (LPA) to 2-AG by phosphatase (Step 6) (Nakane *et al.*, 2002).

Endocannabinoid biodegradation

Once synthesized, the endocannabinoids (AEA and 2-AG) are released from the cell to activate their target cell on presynaptic nerve terminals, and are then rapidly removed from the extracellular space by a selective cellular reuptake system followed by intracellular enzymatic degradation through two different pathways, hydrolysis and oxidation.

Cellular reuptake has been postulated to occur through endocannabinoid membrane transporter (EMT) (Mechoulam & Deutsch 2005, Moore *et al.*, 2005). The involvement of intracellular transporters of AEA, called fatty acid binding proteins (FABP5 and FABP7), have also been suggested (Kaczocha *et al.*, 2009, Maccarrone *et al.*, 2010).



Figure 4. Hydrolytic degradation of AEA and 2-AG by some specific enzymes.

The enzymatic degradation of AEA and 2-AG is caused mainly by FAAH (Cravatt *et al.*, 1996) and MAGL (Dinh *et al.*, 2002, Dinh *et al.*, 2004), respectively (Figure 4). Fatty acid amide hydrolase-2 (FAAH-2) (Wei *et al.*, 2006, Kaczocha *et al.*, 2010) and *N*-acylethanolamine-hydrolyzing acid amidase (NAAA) (Tsuboi *et al.*, 2005, Tsuboi *et al.*, 2007, Zhao *et al.*, 2007) have also been claimed to be involved in AEA degradation. However, the main substrate for FAAH-2 is oleamide (Wei *et al.*, 2006) while NAAA hydrolyzes mainly PEA (Tsuboi *et al.*, 2005). In addition to MAGL, 2-AG degradation can be performed by a number of enzymes, such as FAAH (Goparaju *et al.*, 1998), neuropathy target esterase (NTE) (van Tienhoven *et al.*, 2002), and hormone-sensitive lipase (HSL) (Belfrage *et al.*, 1977). Moreover, two additional serine hydrolases, α/β -hydrolase domain containing 6 (ABHD6) and 12 (ABHD12) have been identified as 2-AG degrading enzymes (Figure 4) (Blankman *et al.*, 2007, Savinainen *et al.*, 2012). Additionally, under some specific conditions, 2-AG can be recognized as a substrate

of human carboxylesterases 1 and 2 (CES1 and CES2) (Xie *et al.*, 2010) while both AEA and 2-AG are potential substrates of butyrylcholinesterase (BuChE) (Romani *et al.*, 2011, Barricklow & Blatnik 2013).



Figure 5. Oxidative degradation of AEA and 2-AG by some specific enzymes.

The endocannabinoids are susceptible to oxidation catalyzed by lipoxygenases (LOXs), cyclooxygenases (COXs), and cytochromes P450 (CYPs) (Kozak & Marnett 2002, Snider et al., 2010, Rouzer & Marnett 2011) (Figure 5). Lipoxygenases, such as 12-LOX and 15-LOX, oxidize both AEA and 2-AG to HETE-EA and HETE-G, respectively, and of these, 15-HETE-G acts as a peroxisome proliferator-activated receptor alpha (PPAR α) agonist (Rouzer & Marnett 2011) while 12-HETE-EA is considered to be an endocannabinoid since it can activate cannabinoid receptors (van et al., 2002). If one considers the isoenzymes of COX, then only COX-2 can catalyze the dioxygenation of both AEA and 2-AG to the corresponding endoperoxides as direct precursors of prostaglandin ethanolamides (prostamides, PG-EAs) and prostaglandin glyceryl esters (PG-GEs). Since PG-EAs and PG-GEs have no affinity for the cannabinoid or prostanoid receptors, the endocannabinoid oxygenation by COXs is considered as a termination step (Matias et al., 2004, Gatta et al., 2012). P450-mediated endocannabinoid metabolism has been demonstrated since AEA, but not 2-AG, can be converted into four different epoxy analogues termed as epoxyeicosatrienoic ethanolamides (EET-EAs) and/or hydroxyl analogue, 20-hydroxyeicosatetraenoic acid ethanolamide (20-HETE-EA), by the 4F2, 3A4 and 2D6 classes of CYP P450 (Rouzer & Marnett 2011).

2.2 ENDOCANNABINOID HYDROLASES

2.2.1 Fatty acid amide hydrolase (FAAH)

FAAH is the principal enzyme responsible for the hydrolysis of AEA to AA and ethanolamine (Figure 6). FAAH (EC 3.5.1.4), a 63 kDa membrane-bound protein of 579 amino acid residues, belongs to the amidase signature (AS) family of enzymes (Giang & Cravatt 1997). Mutagenesis and enzymological studies have confirmed the presence of the unusual catalytic triad of Ser²¹⁷-Ser²⁴¹-Lys¹⁴² in the FAAH active site. The degradation step involves nucleophilic attack of the Ser²⁴¹ residue on the carbonyl group of the substrate while Lys¹⁴² participates as acid/base catalyst to activate nucleophilic Ser²⁴¹ (Figure 7). The Ser²¹⁷ is also essential since it has been shown that its mutation to alanine will lower FAAH activity (Patricelli & Cravatt 1999, Patricelli *et al.*, 1999, Patricelli & Cravatt 2000, Bracey *et al.*, 2002, McKinney & Cravatt 2003).







Figure 7. Mechanism for AEA hydrolysis by FAAH.

In addition to AEA, FAAH also hydrolyses several other bioactive amide and ester lipids thereby controlling their endogenous levels. For example with respect to the amides, FAAH regulates (i) *N*-acylethanolamines (NAEs), such as PEA and OEA, both of which are agonists of PPAR α and involved in the regulation of inflammation and feeding, respectively (Fowler *et al.*, 2001, Bisogno *et al.*, 2002); (ii) *N*-acyltaurins (NATs) activating the transient receptor potential (TRP) family of calcium channels, including TRPV1 and TRPV4 (Saghatelian *et al.*, 2006); (iii) fatty acid primary amides (FAPAs), such as oleamide which induces sleep (Patricelli *et al.*, 1998). Esters, such as 2-AG are hydrolyzed by FAAH, although to a much lesser extent (Goparaju *et al.*, 1998). Consequently, FAAH inhibition increases the levels of

N-arachidonoyl dopamine, a TRPV1 agonist, and *N*-arachidonoylglycine, a GPR118 agonist (Hu *et al.*, 2009, Bradshaw *et al.*, 2009). Hence, inhibition of FAAH results in several effects which are not only mediated through the cannabinoid receptors but also via non-cannabinoid receptors, such as GPR118, ion channels (e.g. TRPV1) or nuclear receptors (like PPAR α).

At present, several X-ray crystal structures of FAAH are available. Bracey et al. reported the first X-ray crystal structure of the transmembrane domain truncated rat FAAH (∆TMrFAAH) in complex with an irreversible inhibitor, methyl arachidonoyl fluorophosphonate (MAFP (2), see Table 1, subsection 2.3.1) (Bracey et al., 2002). Later, researchers engineered a "humanized" rat FAAH (h/rFAAH) and obtained a second crystal structure in a complex with N-phenyl-4-(3-quinolinylmethyl)-1-piperidinecarboxamide (PF-750 (49), see Table 10, subsection 2.3.4). This was followed by several other h/rFAAH structures bound with different FAAH inhibitors such as N-3-pyridinyl-4-[[3-[[5-(trifluoromethyl)-2pyridinyl]oxy]phenyl]methyl]-1-piperidinecarboxamide (PF-3845 (51), see Table 10, subsection 2.3.4) (Ahn et al., 2009b), 1-oxo-1-[5-(2-pyridyl)oxazol-2-yl]-7-phenylheptane (OL-135 (12), see Table 3, subsection 2.3.2) and other related α -ketoheterocycles (Mileni *et al.*, 2009, Mileni et al., 2010b, Otrubova et al., 2013), and [3-(3-carbamoylphenyl)phenyl] Ncyclohexylcarbamate (URB597 (23), also known as KDS-4103, see Table 4, subsection 2.3.3) (Mileni et al., 2010a).

The FAAH knockout mouse has been developed. These animals display elevated AEA levels and exhibit several distinctive properties: (i) an analgesic phenotype in both the carrageenan model of inflammatory pain and the formalin model of spontaneous pain (Lichtman *et al.*, 2004b), (ii) reductions in inflammatory responses (Massa *et al.*, 2004, Cravatt *et al.*, 2004, Karsak *et al.*, 2007), (iii) improvements in sleep and memory (Huitron-Resendiz *et al.*, 2004, Varvel *et al.*, 2007). AEA has been shown to be involved in a number of physiological processes and several reports support the proposal that FAAH inhibition represents a promising novel approach in the treatment of pain, inflammation, anxiety, and depression (Fowler *et al.*, 2001, Labar & Michaux 2007, Ahn *et al.*, 2009a, Blankman & Cravatt 2013). This has stimulated the search for and the development of several classes of compounds acting as reversible or irreversible FAAH inhibitors; these compounds will be discussed in section 2.3.

2.2.2 Other AEA degrading enzymes (FAAH-2 and NAAA)

Fatty acid amide hydrolase-2 (FAAH2)

FAAH-2, a second amidase signature enzyme, is present in humans, but not in lower placental mammals, including mouse and rat (Wei *et al.*, 2006). FAAH-2 is a~60 kDa protein which shares the Ser-Ser-Lys catalytic triad with FAAH. However, it shares only 20%

homology with FAAH. FAAH-2 displays a substrate preference for primary fatty acid amides, such as oleamide, over NAEs, like AEA.

N-Acylethanolamine-hydrolyzing acid amidase (NAAA)

NAAA, a~40 kDa protein of 359 amino acid residues, belongs to choloylglycine hydrolase family having catalytic triad of Cys¹³¹-Asn²⁹²-Asp¹⁵⁰. NAAA is highly expressed by both rodents and humans in immune cells, specifically in macrophages. In contrast to FAAH, NAAA is most active at acidic pH (Ueda *et al.*, 2001). Cys¹³¹ residue acts as a catalytic nucleophile in NAAA rather than the Ser moiety in FAAHs. The preferred substrate of NAAA is PEA (the levels of which are increased during inflammation) over AEA. Since NAAA is highly expressed in macrophages, it is considered as an interesting target to treat inflammation (Tsuboi *et al.*, 2005).

2.2.3 Monoacylglycerol lipase (MAGL/MGL)

MAGL is the main enzyme involved in the 2-AG hydrolysis to AA and glycerol (Figure 8) in the brain. MAGL, a 33-kDa membrane-associated protein with 303 amino acid residues, is classified as a member of α/β -hydrolase family. MAGL active site contains the classical catalytic triad (Ser¹²²-His²⁶⁹-Asp²³⁹) and the lipase motif GXSXG typical of serine hydrolases. The nucleophilic Ser¹²² is activated by the Asp²³⁹ and the His²⁶⁹ residues (Karlsson *et al.*, 1997). By undertaking ABPP of mouse brain 2-AG hydrolases, it was found that nearly 85% of 2-AG hydrolase activity is attributable to MAGL while remaining 15% is accounted for by two additional serine hydrolases, ABHD6 and ABHD12 (Blankman *et al.*, 2007). It is important to note that in addition to metabolizing 2-AG, all of these three enzymes are involved in the control of other bioactive MAGs levels. For example, MAGL inhibition increases levels of 2oleoylglycerol (2-OG), an agonist of GPR119 (Hansen *et al.*, 2011).



Figure 8. Inactivation of 2-AG by MAGL, ABHD6 and ABHD12 to arachidonic acid (AA) and glycerol

In comparison to FAAH, the first crystal structure of human MAGL (hMAGL) in its apo form and in complex with the covalent inhibitor SAR629 (118) (see Table 24, subsection 2.4.6), a derivative of earlier patented triazolo-carboxamide series, was only reported in 2010 by Sanofi-Aventis (Bertrand *et al.*, 2010). Subsequently, a few more crystal structures of MAGL have been published (Labar *et al.*, 2010, Schubert *et al.*, 2009, Schalk-Hihi *et al.*, 2011). Like AEA, 2-AG has also been found to be involved in numerous physiological processes, and several reports suggest MAGL inhibition as a promising strategy to treat pain, inflammation, vomiting, nausea, cancer etc (Bisogno *et al.*, 2009, Kinsey *et al.*, 2011, Alhouayek *et al.*, 2011, Sticht *et al.*, 2012, Nomura *et al.*, 2010, Mulvihill & Nomura 2013). At present, several classes of MAGL inhibitors targeting either the cysteine residues (Cys²⁰¹, Cys²⁰⁸, and Cys²⁴²) located close to the catalytic site or the nucleophilic Ser¹²² residue of catalytic triad of MAGL have been described and will be reviewed in section 2.4.

2.2.4 Other 2-AG degrading enzymes (ABHD6 and ABHD12)

α/β -Hydrolase domain containing 6 (ABHD6)

ABHD6 accounts for minor degradation of 2-AG (~4%) to arachidonic acid (AA) and glycerol (Figure 8) (Blankman *et al.*, 2007). ABHD6, a 30-kDa integral membrane protein with 337 amino acid residues, belongs to the α/β -hydrolase family. The ABHD6 active site contains the classical catalytic triad (Ser¹⁴⁸-Asp²⁷⁸-His³⁰⁶) which is predicted to face into the cell interior (Blankman *et al.*, 2007, Savinainen *et al.*, 2012). This kind of alignment suggests that ABHD6 might be well-suited to guard the intracellular pool of 2-AG. Recently, a homology model of ABHD6 has been created and this may facilitate the development of ABHD6 inhibitors (Bowman & Makriyannis 2013). Recently, several reports have highlighted the potential importance of ABHD6 inhibitors in the treatment of inflammation, metabolic disorders, and epilepsy (Tchantchou & Zhang 2013, Alhouayek *et al.*, 2013, Thomas *et al.*, 2013, Naydenov *et al.*, 2014) but to date, only a few ABHD6 inhibitors have been described (see section 2.5).

α/β -Hydrolase domain containing 12 (ABHD12)

ABHD12 is responsible for only about 9% of total degradation of 2-AG to AA and glycerol (Figure 8) (Blankman *et al.*, 2007). ABHD12 is a ~ 45 kDa glycoprotein whose active site probably contains the classical catalytic triad (Ser²⁴⁶-Asp³³³-His³⁷²) which is predicted to face the lumen and/or extracellular space. It has been speculated that ABHD12 may be involved in metabolizing the extracellular pool of 2-AG (Blankman *et al.*, 2007, Savinainen *et al.*, 2012). Genetically modified ABHD12 deficient mice were found to develop age-dependent symptoms that resembled a human neurodegenerative disorder, PHARC (polyneuropathy, hearing loss, ataxia, retinitis pigmentosa and cataract) (Fiskerstrand *et al.*, 2010). Very recently, Blankman *et al.* revealed that ABHD12 controls brain lysophosphatidylserine (LPS) pathways and that these are deregulated in a murine model of the neurodegenerative disease PHARC (Blankman *et al.*, 2013). At present, no X-ray crystal structure of ABHD12 is available which makes it a challenging target for the development of novel inhibitors. However, recently triterpenoid-based compounds were identified as ABHD12 inhibitors (Parkkari *et al.*, 2014).

2.3 INHIBITORS OF FAAH

2.3.1. Substrate-based FAAH inhibitors

Substrate-based FAAH inhibitors are also described as the first generation FAAH inhibitors. The common design strategy utilized was the derivatization of a substrate (such as AEA) by making the carbonyl group more electrophilic/polar to facilitate the nucleophilic attack of reactive residues of an enzyme such as Ser²⁴¹. The concept was to mimic the tetrahedral intermediate state of an enzyme-substrate complex by forming stable enzyme adducts or hemiketal between the enzyme and the designed inhibitor (Figure 9).



Figure 9. Formation of tetrahedral intermediate between FAAH and AEA / FAAH inhibitor (1)

Arachidonyltrifluoromethyl ketone (ATFMK or ATK, 1) was one of the first reported trifluoromethyl ketone analogues, being the most active reversible FAAH inhibitor at that time (IC₅₀ = 1.9 μ M, Table 1) (Koutek *et al.*, 1994). The discovery of ATFMK (1) was followed by the identification of several other anandamide and oleamide analogues (Table 1) and of these, the highly potent irreversible FAAH inhibitor methyl arachidonyl fluorophosphate (MAFP, 2) is still used widely as a non-specific pharmacological tool. (IC₅₀ = 2.5 nM) (Deutsch et al., 1997). Subsequently, Boger et al. synthesized oleoyltrifluoromethyl ketone (3), originally reported as an inhibitor of oleamide hydrolase, as a potent and competitive reversible FAAH inhibitor (IC₅₀ = 0.082 µM) (Boger et al., 1999). This research group also highlighted the importance of the presence, position, and stereochemistry of the $\Delta^{9,10}$ *cis* double bond of oleamide. A close analogue of MAFP (2), arachidonylsulfonyl fluoride (4) was shown to be as potent as MAFP (2) (IC₅₀ = 0.1 nM) but it was a more selective FAAH inhibitor with respect to its binding to the other studied off-targets (Segall et al., 2003). Arachidonoyl diazomethyl ketone (ADMK, 5) has been found to inhibit reversibly detergent-solubilized FAAH (IC₅₀ = 2-6 μ M) whereas it behaves as an irreversible inhibitor of native membrane FAAH (IC₅₀ = 0.5 µM) (De et al., 1997, Edgemond et al., 1998). N-arachidonoyl amino acids, such as arachidonoyl serotonin (AA-5-HT, 6), N-arachidonoylglycine (NAGly), N-arachidonoyl

Compd	Structure	ΙC ₅₀ [μΜ]		References	
		FAAH	MAGL	-	
1 ATFMK (ATK)		1.9 ^a	66 ^b	Koutek <i>et al.,</i> 1994 Saario <i>et al.,</i> 2004	
2 MAFP	O P-F OMe	0.0025 ^a 0.0010 ^c	0.0022 ^b	Deutsch <i>et al.,</i> 1997 Segall <i>et al.,</i> 2003 Saario <i>et al.,</i> 2004	
3	9 10 CF ₃ 0	$K_i = 0.082^c$	NA ^d	Boger <i>et al.,</i> 1999	
4	O S S O O	0.0011 ^c	NA	Segall <i>et al.,</i> 2003	
5 ADMK	N ₂	2-6 ^e 0.5 ^f	NA	De <i>et al.,</i> 1997 Edgemond <i>et al.,</i> 1998	
6 AA-5-HT	O HO NH	12-26 ^g 5.6 ^h	NI ⁱ	Bisogno <i>et al.</i> , 1998	

Table 1. Inhibitory	values for the	substrate-based	compounds	1-6 against FAA	H and MAGL.
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Enzymatic assays were carried out using ^{a,f} rat brain homogenate protein and [³H]-AEA as the substrate; ^b solubilized COS-7 membrane extracts from cells transiently transfected with human FAAH cDNA and 2-AG as the substrate; ^c solubilized rat liver plasma membrane extracts and ¹⁴C-oleamide as the substrate; ^d Not analyzed. Enzymatic assays were carried out using ^e partially purified protein from mouse neuroblastoma N18TG2 or RBL-1 cells or porcine brain microsomal and [³H]-AEA as the substrate; ^g mouse neuroblastoma N18TG2 cells and [¹⁴C]-AEA as the substrate; ^h rat basophilic leukemia (RBL-2H3) cells and [³H]-AEA as the substrate. ⁱ No inhibition.

However, these substrate-based FAAH inhibitors tend to lack selectivity i.e. they also inhibit other enzymes such as MAGL (Table 1). They also bind to the cannabinoid or TRPV1 receptors or have other unknown targets (Koutek *et al.*, 1994, Patterson *et al.*, 1996, Deutsch *et al.*, 1997, Segall *et al.*, 2003, Saario *et al.*, 2004). Nevertheless, both ATK (1) and MAFP (2) are considered as useful pharmacological tools and valuable reference compounds.

2.3.2 a-Ketoheterocycle-based inhibitors of FAAH

The α -ketoheterocycle-based compounds are potent and competitive reversible FAAH inhibitors synthesized mainly by Dale Boger's group. The design of these inhibitors was based on the earlier reported protease inhibitors (Edwards *et al.*, 1995). Extensive work on modifications of oleoyltrifluoromethyl ketone (3) was carried out by (i) replacing the trifluoromethyl group with various carbonyl activating heterocycles and, (ii) optimising the fatty acid chain. This led the identification of hundreds of potent and reversible FAAH inhibitors; some example structures 7-10 are shown in Table 2 (Boger *et al.*, 2000). The reversible, competitive inhibition has been proposed to occur via hemiketal formation with the nucleophilic Ser²⁴¹ of FAAH (Figure 10). Several α -ketoheterocycle-based FAAH inhibitors (Boger *et al.*, 1999, Boger *et al.*, 2000) were later screened using the ABPP. Compound 11 was identified as a potent and highly selective FAAH inhibitor while others were found to react with 45- and 50-kDa brain enzyme KIAA1363 and 60-kDa heart enzyme triacylglycerol hydrolase (TGH) (Leung *et al.*, 2003).



Figure 10. Hemiketal adduct formed between FAAH and an α -keto heterocycle inhibitor

Table 2. Inhibition activity of the α -ketoheterocycle-based compounds 7-11 against FAAH and MAGL.

Compd	Structure	K _i (IC ₅₀) [μM]		References	
		FAAH ^a	MAGL		
7		0.017	NA ^b	Boger <i>et al.,</i> 2000	
8		0.0023 (0.013) ^c	NA	Boger <i>et al.,</i> 2000	


^a Enzymatic assays were carried out by using solubilized rat liver plasma membrane extracts and ¹⁴Coleamide as the substrate. ^b Not analyzed. ^c Enzymatic assays were carried out by using solubilized COS-7 membrane extracts from cells transiently transfected with human FAAH cDNA and ¹⁴C-oleamide as the substrate. ^d IC₅₀ values determined by using the ABPP assay. ^e No inhibition.

A systematic exploration around the oxazole-based compound 7 was carried out which led to the discovery of 1-oxo-1-[5-(2-pyridyl)oxazol-2-yl]-7-phenylheptane (OL-135, 12) as a potent FAAH inhibitor (Ki = 4.7 nM, Table 3) (Boger et al., 2005). In the ABPP assay, this compound showed selectivity for FAAH over other serine hydrolases of different mouse tissues, such as brain, heart, and kidney. After *i.p.* administration in mice, OL-135 (12) significantly elevated the endogenous levels of the AEA (and other lipids such as OEA and PEA) in the brain and spinal cord, and possessed an analgesic effect in multiple pain models (Lichtman et al., 2004a, Chang et al., 2006). However, subsequently it was shown that OL-135 (12) inhibited several carboxyesterases (CEs) (Zhang et al., 2007). Then after, Boger's group conducted further variations on OL-135 (12) by modifying its C2 acyl side chain and/or heterocyclic core to afford several potent FAAH inhibitors 13-20 (Table 3) (Romero et al., 2006, Romero et al., 2007, Hardouin et al., 2007, DeMartino et al., 2008, Kimball et al., 2008, Garfunkle et al., 2008, Ezzili et al., 2011, Otrubova & Boger 2012, Otrubova et al., 2013, Otrubova et al., 2014a, Duncan et al., 2014). Among these, oral administration of (S)-enantiomer 17 (CE-12) significantly elevated the brain AEA levels and attenuated mechanical (> 6 h) and cold (> 9 h) allodynia for sustained periods. Interestingly, compounds 18 and 19 displayed interactions with an additional active site residue Cys²⁶⁹. Compound 18 was reported to increase the levels of endogenous substrates to a greater extent and for a much longer duration (> 6 h) than OL-135 (12) and it reversed cold allodynia in the chronic constriction injury (CCI) model of neuropathic pain in mice for a sustained period (> 6 h). Janssen also described several modifications of OL-135 (12) in an attempt to improve its aqueous solubility and pharmacokinetic (PK) profile (Apodaca et al., 2007a, Apodaca et al., 2007b, Timmons et al., 2008). In contrast to the reports from Boger's group, 2-ketooxazoles containing polar head groups (such as -SO₂ group in compound 21) in their C5-side chain were reported as FAAH inhibitors (Table 3) (Rusch et al., 2012).

Compd	Structure	FAAH K _i (ΙC ₅₀) [μΜ]	References
12 OL-135	$ \begin{array}{c} \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\$	0.0047^{a} (0.0090) ^b ^c IC ₅₀ = 0.002 (FAAH) and 0.6 (TGH)	Boger et al., 2005
13		0.0004^{a} (0.001) ^b ^c IC ₅₀ = 0.0065 (FAAH) and 0.02 (TGH)	Romero <i>et al.,</i> 2006 Romero <i>et al.,</i> 2007
14		0.00075^{a} (0.0029) ^b cIC ₅₀ = 0.0007 (FAAH) and 1.2 (TGH)	Hardouin <i>et al.,</i> 2007
15	N N N N N N N N N N N N N N N N N N N	0.0002^{a} (0.00026) ^b ^c IC ₅₀ = 0.001 (FAAH) and 0.026 (TGH)	Kimball <i>et al.,</i> 2008
16		0.00029^{a} (0.0012) ^b ^c IC ₅₀ = 0.001 (FAAH) and 0.014 (TGH)	Garfunkle <i>et al.,</i> 2008
17 CE-12 (<i>S</i>)-enantiomer		0.0044 ^a (0.0058) ^b	Ezzili <i>et al.,</i> 2011
18	N N Br	0.0031^{a} $^{c}IC_{50} = 0.20$ (FAAH) and 0.9 (TGH)	Otrubova <i>et al.,</i> 2013 Otrubova <i>et al.,</i> 2014
19		0.0015^{a} ^c IC ₅₀ = 0.090 (FAAH) and 0.450 (TGH)	Otrubova <i>et al.,</i> 2013 Otrubova <i>et al.</i>
			2014
20 <i>R</i> and <i>S</i> enantiomers	C C C C N N N N N N N N N N N N N N N N	~0.018 ^a	Duncan <i>et al.,</i> 2014
21	n-C ₁₁ H ₂₃ N O O S O	0. 364 ^d	Rusch <i>et al.,</i> 2012

Table 3. Inhibitory activity of the α -ketoheterocycle-based compounds 12-21 against FAAH.

Enzymatic assays were carried out by using ^a purified recombinant rat FAAH expressed in *Escherichia coli* (*E. coli*) and ¹⁴C-oleamide as the substrate; ^b lysate of hFAAH-transfected COS-7 cells and ¹⁴C-oleamide as the substrate. ^c IC₅₀ values determined by the ABPP assay. ^d Enzymatic assays were carried out by using rat brain microsome FAAH and *N*-(2-hydroxyethyl)-4-pyren-1-ylbutanamid as the substrate.

To summarize, α -ketoheterocycle-based FAAH inhibitors have predominantly been explored by academic researchers. Due to their excellent potency and high FAAH selectivity with respect to possible off-targets, compounds 17 and 18 are considered to be valuable pharmacological tools. They could also serve as potential candidates for further drug development.

2.3.3 Carbamates as FAAH inhibitors

In 2003, Kathuria *et al.* utilized carbaryl, an anti-cholinesterase agent, and its positional isomer in the development of FAAH inhibitors. They produced highly potent compounds, such as [1,1'-biphenyl]-3-yl cyclohexylcarbamate (URB524, 22) and [1,1'-biphenyl]-3-yl cyclohexylcarbamate (URB597, also known as KDS-4103, 23) that had IC₅₀ values of 0.063 and 0.0046 µM, respectively (Table 4) (Kathuria *et al.*, 2003). *P.o.* or *i.p.* administration of URB597 (23) exerted a combination of anxiolytic-like, anti-depressant-like, and analgesic effects by inhibiting FAAH both centrally and peripherally (Gobbi *et al.*, 2005, Piomelli *et al.*, 2006, Russo *et al.*, 2007), and it also reduced carrageenan induced edema (Jayamanne *et al.*, 2006). However, the data related to its efficacy on putative anti-anxiety and anti-depressive properties are controversial (Kathuria *et al.*, 2003, Naidu *et al.*, 2007). Later, it was shown that URB597 (23) also inhibited several CEs (Zhang *et al.*, 2007).



Figure 11. The O-biaryl group of URB597 (23) serves as a leaving group, resulting in the carbamylation of the FAAH enzyme

The discovery of URB597 (23) prompted other researchers to explore carbamate-based FAAH inhibitors. Modulation at the *N*-portion of URB597 (23) resulted in FAAH inhibitors (24-26, Table 4) with improved potency (Alexander & Cravatt 2005, Mor *et al.*, 2008). JP83 (24) and JP104 (25) were FAAH selective in the nervous system, but they bound to and interfered with several enzymes in peripheral tissues. Alexander *et al.* also provided experimental evidence that carbamates, such as URB597 (23) could inhibit FAAH via covalent carbamylation of Ser²⁴¹ of FAAH (Figure 11). The details of the modifications made on the distal and proximal phenyl rings of URB524 (22) and URB597 (23) were also reported (Table 4) (Tarzia *et al.*, 2003, Mor *et al.*, 2004, Tarzia *et al.*, 2006, Clapper *et al.*, 2009, Clapper *et al.*, 2010, Moreno-Sanz *et al.*, 2013). This led to the identification of URB694 (27), a potent FAAH inhibitor having less reactivity

with off-target CEs, and URB937 (28), a peripherally restricted potent FAAH inhibitor which nonetheless exerted antinociceptive effects in mice and rats. Moreover, cyclohexylcarbamic acid 3'-carbamoyl-5-hydroxybiphenyl-3-yl ester (29) was identified as the most potent brainimpermeable FAAH inhibitor disclosed to date having an IC₅₀ value of 0.0005 µM.

Compd	Structure	ΙC ₅₀ [μΜ]		References
		FAAH	MAGL	-
22 URB524	C o J K	0.063ª	NI ^b	Kathuria <i>et al.,</i> 2003
23 URB597 (KDS- 4103)		0.0046 ^a	NI ^{b,*}	Kathuria <i>et al.,</i> 2003
24 JP83		0.014 ^c	NI	Alexander 2005
25 JP104	O NH2 O NH2 O N	0.0073°	NI	Alexander 2005
26 URB880		0.00063ª	100 ^d	Mor <i>et al.,</i> 2008
27 URB694	HOLOGIA	0.030 ^a	NA ^e	Clapper <i>et al.,</i> 2009
28 URB937		0.0028 ^a	NA	Clapper <i>et al.,</i> 2010
29		0.0005ª	NA	Moreno-Sanz <i>et al.,</i> 2013

Table 4. Inhibitory activity of the carbamate compounds 22-29 against FAAH and MAGL.

Enzymatic assays were carried out by using ^a rat brain homogenates and [³H]-AEA as the substrate. ^b No inhibition ^{*} (according to the ABPP assay (was found to inhibit CEs)). Enzymatic assays were carried out by using ^c purified recombinant enzyme and ¹⁴C-oleamide as a substrate; ^d purified rat recombinant MAGL and 2-OG as the substrate. ^e Not analyzed.

Researchers from the University of Kuopio (now the University of Eastern Finland, UEF) have identified several carbamate-based potent FAAH inhibitors, such as 30-34 as having good selectivity over MAGL (Table 5) (Myllymäki *et al.*, 2007, Minkkilä *et al.*, 2009a, Myllymäki *et al.*, 2009, Käsnänen *et al.*, 2010). This includes the first report of chiral FAAH inhibitors where the *R*-enantiomer of 3-(5-methyl-4,5-dihydrooxazol-2-yl)phenyl cyclohexylcarbamate (FAAH IC₅₀ = 0.0068 μ M) was found 10-times more potent than its corresponding *S*-enantiomer (see compounds 32 and 33).

Compd	Structure	IC ₅₀ [μM] or at 100 μM	% inhibition	References
		FAAH ^a	MAGL	-
30		0.028	46% ^b	Myllymäki <i>et al.,</i> 2007
31	S J J O J H	0.0063	2.1 ^c	Minkkilä <i>et al.,</i> 2009
32		0.0068	14% ^c	Myllymäki <i>et al.,</i> 2009
33		0.073	13% ^c	Myllymäki <i>et al.,</i> 2009
34		0.0007	NI ^{b,d}	Käsnänen <i>et al.,</i> 2010

Table 5. Inhibitory activity of the carbamate compounds 30-34 against FAAH and MAGL.

Enzymatic assays were carried out by using ^a brain homogenates and [³H]-AEA as the substrate; ^b rat brain membranes and 2-AG as the substrate; ^c human recombinant MAGL and tritium-labelled 2-oleoylglycerol [glycerol-1,2,3-³H] (i.e. [³H]-2-OG) as the substrate. ^d No Inhibition

Several carbamates have been evaluated by pharmaceutical companies and also in collaboration with academic groups (Table 6). For example, Gattinoni *et al.* determined that enol carbamates acted as reversibly acting potent FAAH inhibitors (ST-4070 (35) and 36) (Gattinoni *et al.*, 2010a). Oral administration of ST-4070 (35) at 30 mg/kg produced a similar anti-anxiety effect as diazepam in a mouse model. Bristol-Myers Squibb (BMS) described derivatives of bisarylimidazoles and oximes, such as compounds 37 (BMS-1) and 38, as potent inhibitors of FAAH (Sit *et al.*, 2007, Sit *et al.*, 2010). Interestingly, Gattinoni *et al.* reported oxime carbamates, such as compound 39 (ST4020), having an inversed oxime carbamate bond compared to the BMS compounds (Gattinoni *et al.*, 2010b). Oral

administration of ST4020 (39) (10 mg/10 mL/kg) reduced anxiety but did not affect the locomotor activity in the elevated plus maze. ST4020 (39) also significantly reduced mechanical hyperalgesia in a rat model of neuropathic pain induced by vincristine (Minetti *et al.*, 2009). Niphakis *et al.* published SARs of *O*-hydroxyacetamide carbamates among which 2-(methylamino)-2-oxoethyl 4-(4-chlorophenethyl)piperidine-1-carboxylate (SA-57, 40) showed excellent potency towards mouse and human FAAH with IC₅₀ values of 0.0032 and 0.0019 μ M, respectively. SA-57 (40) exhibited selectivity for FAAH at low concentrations but at higher concentrations (at 10 μ M) the drug inhibited MAGL and ABHD6. It was postulated that SA-57 could be a useful pharmacological tool to study the behavioral effects of complete FAAH blockade in combination with varying degrees of MAGL and ABHD6 inhibition (Niphakis *et al.*, 2012).

Compd	Structure	ΙC ₅₀ [μΜ]		References
		FAAH	MAGL	-
35 ST-4070	C C C N	0.009^{a} (K _i = 0.054) ^c	NI ^b	Gattinoni <i>et al.,</i> 2010
36	NC	0.013 ^a (K _i = 0.046) ^c	NI	Gattinoni <i>et al.,</i> 2010
37 BMS-1	N N N N N N N N N N N N N N N N N N N	0.002 ^c	NA ^d	Sit <i>et al.,</i> 2007
38		0.00015 ^c	NA	Sit <i>et al.,</i> 2010
39 ST4020	S S S S S S S S S S S S S S S S S S S	0.0044ª	NI	Gattinoni <i>et al.</i> , 2010
40 SA-57	CI C	0.0032 (mFAAH) ^e 0.0019 (hFAAH) ^e	0.41 (mMAGL) ^e 1.4 (hMAGL) ^e	Niphakis <i>et al.,</i> 2012

Table 6. Inhibitory values for the carbamate compounds 35-40 against FAAH and MAGL.

^a Enzymatic assays were carried out by using mouse brain homogenates and [³H]-AEA as the substrate. ^b No inhibition observed in mouse brain supernatants using [³H]-2-OG as a substrate. ^c Enzymatic assays were carried out by using homogenates of crude membranes prepared from H4 cells that express transfected human FAAH (H4-FAAH) cells and [³H]-AEA as the substrate. ^d Not analyzed. ^e human and mouse FAAH, MAGL, and ABHD6 orthologues assayed as recombinantly expressed proteins in transfected human embryonic kidney 293 (HEK293) cells and 2-AG as the substrate.

Butini *et al.* reported series of phenylpyrrole-based carbamates as potent FAAH inhibitors. Within the series compound 41 (ST3913, Table 7) showed excellent inhibition of FAAH (IC₅₀ 0.6 nM) along with selectivity over other studied potential off-targets (Butini *et al.*, 2012). Moreover, compound ST3913 (41) was orally efficacious (30 mg/kg dose) producing an analgesic effect, and it did not exert any adverse effects. Later, they undertook structural modifications of ST3913 (41) and identified analogue 42 as a potent, reversible and noncompetitive mouse FAAH (mFAAH) inhibitor (Table 7) (Butini *et al.*, 2013).

Compd	Structure	Ki (IC ₅₀)	K _i (IC ₅₀) [nM]		References
		mFAAH ^a	hFAAH ^b	mMAGL℃	
41 ST3913	H ₂ N{O	0.16 (IC ₅₀ = 0.60)	NA ^d	NI ^e	Butini <i>et al.,</i> 2012
42		13	26 (IC ₅₀ = 44)	NA ^d	Butini <i>et al.,</i> 2013

Table 7. Inhibitory activity of the carbamate compounds 41-42 against FAAH and MAGL.

Enzymatic assays were carried out by using ^a mouse brain homogenates and [³H]-AEA as the substrate; ^b hFAAH and [³H]-AEA as the substrate; ^c mouse brain supernatants and [³H]-2-OG as the substrate. ^d Not analyzed. ^e No inhibition.

A carbamate analogue 43, structurally related to macamides, was shown to be an irreversible inhibitor of FAAH having an IC₅₀ value of 0.153 μ M (Table 8) (Wu *et al.*, 2013). Terwege *et al.* published detailed SARs of ω -heteroarylalkylcarbamate as FAAH inhibitors among which ω -indolylalkylcarbamate-based compound 44 was identified as a potent FAAH inhibitor (Terwege *et al.*, 2014). Very recently, *O*-(triazolyl)methyl carbamates were reported as a novel and potent class of FAAH inhibitors. In this class, compound 45 had a single-digit nanomolar IC₅₀ value for both rat and human FAAH (Colombano *et al.*, 2015). Moreover, compound 45 showed a remarkable improvement over the *O*-arylcarbamate URB524 (22) in terms of its rat plasma stability.

Compd	Structure	IC50 []	uM]	References
		FAAH	MAGL	
43	N N N N N N N	0.153ª	NA ^b	Wu <i>et al.,</i> 2013
44	K N N N O N N	0.0036 ^c	0.22 ^d	Terwege <i>et al.,</i> 2014
45	Meo N N N N N N N N N N N N N N N N N N N	0.004 ^e	NA	Colombano <i>et al.,</i> 2015

Table 8. Inhibitory activity of the carbamate compounds 43-45 against FAAH and MAGL.

^a Enzymatic assays were carried out by using human recombinant hFAAH and 7-amino-4-methyl coumarinarachidonamide (AMC-AA) as the substrate. ^b Not analyzed. Enzymatic assays were carried out by using ^c rat brain microsome and *N*-(2-hydroxyethyl)-4-pyren-1-ylbutanamide as the substrate; ^d human recombinant MAGL and 1,3-dihydroxypropan-1-yl 4-pyren-1-ylbutanoate as the substrate; ^e brain homogenates and [³H]-AEA as the substrate.

In summary, systematic structural modifications of *O*-arylcarbamate-based irreversible FAAH inhibitors have led to the development of highly potent inhibitors with improved selectivity (e.g. URB597 (23) v_s URB694 (27)) and biological stability (e.g. URB597 (23) v_s compound 45). The carbamate scaffold has been proven to be a valuable tool that makes possible different mechanistic approaches when studying the *in vivo* effects of FAAH blockade in the CNS or even solely in the periphery.

2.3.4 Urea derivatives as FAAH inhibitors

The putative endocannabinoid transport inhibitor LY2183240 (46) was identified as a potent albeit non-selective FAAH inhibitor ($IC_{50} = 13 \text{ nM}$, Table 9) (Alexander & Cravatt 2006). From the results of a mass spectrometry study it, was concluded that LY2183240 (46) bound covalently to FAAH by carbamylation of Ser²⁴¹ of FAAH with tetrazole moiety acting as a leaving group. Maione *et al.* synthesized tetrazole urea-based, potent FAAH inhibitors 47 and 48 (Table 9) which showed selectivity over other tested endocannabinoid targets, such as MAGL, DAGL, CB₁ and CB₂ (Maione *et al.*, 2008). There are also several other reports describing tetrazole urea analogues as potent FAAH and/or MAGL inhibitors (Ortar *et al.*, 2008, Ortar *et al.*, 2013, Holtfrerich *et al.*, 2013).

Compd	Structure	ΙC ₅₀ [μΜ]		References
		FAAH	MAGL	
46 LY2183240	O N N-N N-N	0.013 ^a	0.0053ª	Alexander and Cravatt 2006
47 OMDM119		0.0027 ^b	10 ^c	Maione <i>et al.</i> , 2008
48 OMDM122	$\mathbb{I}_{\mathcal{N}}^{\mathcal{O}} = \mathbb{I}_{\mathcal{N}}^{\mathcal{O}} = \mathbb{I}_{\mathcal$	0.0021 ^b	>10 ^c	Maione <i>et al.,</i> 2008

Table 9. Inhibitory activity of the tetrazole urea-based compounds 46-48 against FAAH and MAGL.

 IC_{50} values were measured by ^a competitive ABPP assays using mice brain membrane proteomes; ^b using rat brain membranes and [¹⁴C]-AEA as the substrate; ^C using COS cell cytosolic fractions and [³H]-2-AG as the substrate.

Pfizer and/or the Scripps Research Institute (TSRI) have produced numerous potent and highly selective FAAH inhibitors (49-57, Table 10). Both PF-750 (49) and PF-622 (50) inhibited FAAH in a time-dependent manner by covalently modifying the nucleophile Ser²⁴¹ residue (Ahn et al., 2007). Compound 51 (PF-3845) elevated brain levels of AEA for up to 24 h (Ahn et al., 2009b). It also produced significant CB₁/CB₂-dependent antihyperalgesic effects in the complete Freund's adjuvant (CFA) model of inflammatory pain. Another compound they discovered, PF-465 (52) showed anti-inflammatory activity in a rat CFA model (Johnson et al., 2009). In 2011, they provided details about the highly potent and selective FAAH inhibitor PF-04457845 (54) (Johnson et al., 2011). Oral administration of PF-04457845 (54) at 0.1 mg/kg resulted in efficacy comparable to that obtained with naproxen at 10 mg/kg in a rat model of inflammatory pain. Although being well tolerated in the phase I trials, during phase II trials PF-04457845 (54) was found to be ineffective since it did not reduce osteoarthritis (OA) pain in the patients' knees (Huggins et al., 2012). Pfizer undertook several modifications of PF-04457845 (54) by replacing its methylenepiperidine core with different spirocyclic cores as well as simultaneous optimization at its other modular units i.e. a tail group, an optional hydrocarbon linker and a urea head group; this led the discovery of compounds 55 and 56 (Meyers et al., 2011a). Although selective towards FAAH in the ABPP assay, these compounds were not efficacious in vivo. Hence further variations were carried out aimed at improving the drug's PK properties which resulted in the identification of PF-04862853 (57) as a potent and selective FAAH inhibitor (Meyers et al., 2011b). PF-04862853 (57) was finally taken to the phase I clinical trials as it exhibited good PK profile and it was effective in a pain model.

Compd	Structure	$k_{\text{inact}}/K_{\text{i}}(\text{M}^{-1}\text{s}^{-1})^{\text{a}}$		References
		rFAAH	hFAAH	-
49 PF-750	CL) CN TH	104	$791 \\ IC_{50} = 0.0162 \\ \mu M^b$	Ahn <i>et al.</i> ,2007
50 PF-622		154	NA ^c IC _{50 =} 0.0330 μM ^b	Ahn <i>et al.,</i> 2007
51 PF-3845	F ₃ C	3900 <i>K</i> i = 0.23 μΜ	12,600	Ahn <i>et al.</i> ,2009
52 PF-465	S N N N N N N N N N N N N N N N N N N N	1060	2931	Johnson <i>et al.</i> ,2009
53 PF-946		729	4474	Johnson <i>et al.</i> ,2009
54 PF-04457845	F ₃ C Core N Head Linker	32400	40300	Johnson <i>et al.</i> ,2011
55	F ₃ C	NA	1570	Meyers <i>et al.</i> ,2011a
56	$F_3C - C_N - C_N$	NA	3040	Meyers <i>et al.</i> ,2011a
57 PF-04862853		5820	4190	Meyers <i>et al.</i> ,2011b

Table 10. Inhibitory activity of the piperazine/piperidine urea-based compounds 49-57 against FAAH.**

^{**} In the ABPP most of the compounds exhibited high selectivity for FAAH over other serine hydrolases. ^a Determination of inhibitory potencies $k_{inact}/K_i(M^{-1}s^{-1})$ was carried out by GDH-coupled FAAH assay (De Bank *et al.*, 2005). Briefly, the production of ammonia generated from the hydrolysis of oleamide by FAAH was monitored. GDH catalyzes the condensation of ammonia and α -ketoglutarate to glutamate with a

concomitant conversion of NADH to nicotinamide adenine dinucleotide, oxidized form (NAD⁺), which is spectrophotometrically measured at 340 nm. ^b IC_{50} values were measured for hFAAH and rFAAH (1h pre-incubation with test compound) by GDH-coupled FAAH assay. ^c Not analyzed.

Johnson & Johnson provided details of a series of thiadiazolopiperazinyl aryl and heterocyclic ureas with good PK profiles as a novel class of potent FAAH inhibitors such as compounds 58 (JNJ-1661010) and 59 (JNJ-40355003), respectively (Table 11) (Keith et al., 2008, Keith et al., 2012). JNJ-1661010 (58) exhibited analgesic activity in two models of neuropathic pain i.e. mild thermal injury (MTI) and spinal nerve ligation (SNL) (Karbarz et al., 2008). Oral administration of JNJ-40355003 (59) in rats, dogs and monkeys elevated the plasma and brain concentrations of three FAAs, namely AEA, PEA, and OEA. The elevation of FAAs in a monkey which expressed FAAH-2, strongly suggested that FAAH-2 was not contributing greatly to the breakdown of fatty acid ethanolamides in the plasma. Later, they identified two potent FAAH inhibitors compound 60 and JNJ-42119779 (61) (Tichenor et al., 2012, Keith et al., 2014b). Amgen described a series of novel, noncovalent FAAH inhibitors, such as compound 62 (Table 11) having in vitro potency comparable to known covalent FAAH inhibitors and good PK profiles along with reasonable CNS penetration (Gustin et al., 2011). Takeda reported a series of piperazine ureas and identified compounds 63 and 64 as potent FAAH inhibitors having good brain permeability (Table 11) (Kono et al., 2013, Kono et al., 2014). Oral administration of compound 63 was efficacious in the acetic acid-induced writhing test in mice and compound 64 showed robust and dose-dependent analgesic efficacy in animal models of both neuropathic and inflammatory pain.

Compd	Structure	IC ₅₀ (nM) ^a		References
		rFAAH	hFAAH	
58 JNJ-1661010		34	33	Keith <i>et al.</i> ,2008
59 JNJ- 40355003		33	1.4	Keith <i>et al.</i> ,2012
60		0.7	0.5	Tichenor <i>et al.</i> ,2012
61 JNJ- 42119779		9	8	Keith <i>et al.</i> ,2014

Table 11. Inhibitory activity of the piperazine/piperidine urea-based compounds 58-64 against FAAH.

62	N H2 N N N N N N N N N N N N N N N N N N N	43 ^b	14 ^b	Gustin <i>et al.</i> ,2011
63	F F HN O'N	0.46 ^c	0.43 ^c	Kono <i>et al.</i> ,2013
64		0.28 ^d	0.72 ^d	Kono <i>et al.</i> ,2014

Values were measured using ^a a 1h preincubation and [³H]-AEA as a substrate; ^b purified rat or human FAAH (with a 2h preincubation) and AMC-AA as the substrate; ^c human and rat FAAH enzyme fractions and [³H]-AEA as the substrate (with 30 min preincubation); ^d purified rat or human FAAH enzyme fractions (with 30 min preincubation) and AMC-AA as the substrate

Recently, Otrubova *et al.* identified several pyrazole-based ureas such as compound 65 (Table 12) as potent irreversible inhibitors of FAAH (Otrubova *et al.*, 2014b). BIAL-Portela & Ca. disclosed in a patent that a urea-based compound 66 was a potent, peripherally selective and metabolically stable FAAH inhibitor (Rosa *et al.*, 2014).

Compd	Structure	K _i (nM) rFAAH	References
65		0.1 ^a	Otrubova <i>et al.</i> 2014b
66	H_2N H H N H	0.7 % ^{b,c}	Rosa <i>et al</i> ., 2014

Table 12. Inhibitory activity of the urea-based compounds 65 and 66 against FAAH.

^a Values were measured with a 3h preincubation using the ABPP assay. ^b Values were measured with brain, liver or lung protein and [³H]-AEA as the substrate. ^c Remaining activity after 8 h in mouse liver (oral administration of compound 66 (3 mg/kg)).

Pfizer designed novel FAAH inhibitors by replacing piperidine of PF-3845 (51) with azetidine (exemplified by compound 67, Table 13) (Wang *et al.*, 2009a). The *in vitro* ABPP of both *cis* and *trans* stereoisomers of 67 showed similar selectivity profiles as observed for PF-3845 (51). However, they failed to achieve comparable *in vivo* efficacy to PF-3845 (51). Vernalis produced a chiral azetidine urea-based compound VER-156084 (68) having an *S*-configuration (Hart *et al.*, 2009). Its *R*-isomer was inactive at inhibiting both enzymes. However, VER-156084 (68) suffered from a poor PK profile (Roughley *et al.*, 2012).

Compd	Structure	$k_{\text{inact}}/K_{i}(M^{-1}s^{-1})^{a}$		References
		rFAAH	hFAAH	-
67 (cis and trans isomers)	F ₃ C N N N N N N N N N N N N N N N N N N N	<i>cis</i> 5390 <i>trans</i> 2380	<i>cis</i> 9000 <i>trans</i> 9340	Wang <i>et al.</i> , 2009
68 VR-156084		0.078 ^b	0.531 ^b	Hart <i>et al.,</i> 2009 Roughley <i>et al.,</i> 2012

Table 13. Inhibitory activity of the azetidine urea-based compounds 67 and 68 against FAAH.

Inhibitory activies have been determined by using ^a rat or human FAAH and AMC-AA as the substrate; ^b rat or human FAAH (pre-incubation 3h) and AMC-AA probe.

In general, piperidine/piperazine urea-based irreversible FAAH inhibitors have been mainly developed by pharmaceutical companies. It appears that these types of compounds offer an excellent balance of potency, selectivity, and PK profiles, and hence, have been considered as promising drug candidates.

2.3.5 1,3,4-Oxadiazol-2-ones as FAAH inhibitors

Commercially available hormone sensitive lipase (HSL) inhibitor, CAY10499 (69, Table 14) was identified as a potent inhibitor of both FAAH and MAGL (Minkkilä *et al.*, 2009b). Since inhibition of both FAAH and MAGL was attributed to the 1,3,4-oxadiazol-2-one pharmacophore, numerous 1,3,4-oxadiazol-2-ones were synthesized and identified as potent and reversible inhibitors of both FAAH and MAGL (Minkkilä *et al.*, 2009b, Käsnänen *et al.*, 2013). BIAL - Portela & Ca. described several 1,3,4-oxadiazol-2-ones as potent FAAH inhibitors. For example, catechols 70 and 71 at doses of 10 mg/kg were found to be peripherally restricted FAAH inhibitors and it was proposed that these compounds could be useful in the treatment of certain cardiovascular disorders such as hypertension and heart failure (Kiss *et al.*, 2011). In short, 1,3,4-oxadiazol-2-one represents a promising scaffold for further exploration.

Compd	Structure	ΙC ₅₀ [μΝ	/I]	References
		FAAH	MAGL	-
69 CAY10499		0.086ª	0.092 ^b	Minkkilä <i>et al.</i> ,2009
70		0.004°	NA ^d	Kiss <i>et al.</i> ,2011
71		0.006 ^c	NA	Kiss <i>et al.</i> ,2011

Table 14. Inhibitory activity of the 1,3,4-oxadiazol-2-ones 69-71 against FAAH and MAGL.

Values were measured with ^a rat brain homogenate (10 min preincubation) and [³H]-AEA as the substrate; ^b human recombinant MAGL (hMAGL) activity with 2-AG as the substrate. ^c Values were measured with rat brain (without cerebellum) (15 min preincubation) and [³H]-AEA as the substrate. ^d Not analyzed.

2.3.6 β-Lactam-based FAAH inhibitors

Table 15. Inhibitory activities of β -lactam compounds 72 and 73 against FAAH and MAGL.



^a Values were measured with hFAAH (10 min preincubation) and [³H]-AEA as the substrate. ^b hMAGL activity (10 min preincubation) was measured using [³H]-OG as the substrate.

Urbach *et al.* have described the properties of β -lactams/2-azetidinones as inhibitors of hFAAH (Urbach *et al.*, 2008). The best compound in this series, 3-(pent-4-en-1-yl)-1-(pent-4-enoyl)azetidin-2-one (72), had an IC₅₀ value of 4.5 µM and ~146 fold selectivity for hFAAH versus hMAGL (Table 15). Later, they identified several potent inhibitors of hFAAH, such as compound 73 which exhibited an IC₅₀ value in the nanomolar range and higher selectivity (~800-fold) over MAGL (Feledziak *et al.*, 2009). Interestingly, these kinds of analogues were found to have a reversible mechanism of hFAAH inhibition as determined by time-dependent preincubation and rapid dilution studies and confirmed in docking analyses in a homology model of FAAH. Due to their reversible mode of FAAH inhibition, the β -lactam class of compounds should receive further attention.

2.3.7 FAAH inhibitors lacking electrophile

Renovis investigated a series of 2,5-disubstituted benzoxazole and tetrahydropyridopyridine analogues and claimed compounds 74 and RN-450 (75) were potent and reversibly acting FAAH inhibitors (Table 16) (Gowlugari et al., 2012, Estiarte et al., 2012). Very recently, researchers from Janssen Pharmaceutical produced a series of 1-aryl-2-[(6-aryl)pyrimidin-4yl]aminoethanols as competitive inhibitors of FAAH (Keith et al., 2014a). The best compound of the series (R)-2-({6-[3-chloro-4-(trifluoromethyl)phenyl]pyrimidin-4-yl}amino)-1phenylethan-1-ol (JNJ-40413269, 76) showed excellent inhibitory potency towards FAAH with optimal PK properties. It was also efficacious in the SNL model of neuropathic pain in the rat. Merck has also developed the oxazole-based compound MK-4409 (77) as a potent, selective, and reversible noncovalent modifying FAAH inhibitor (Chobanian et al., 2014). MK-4409 (77) showed a promising preclinical profile and excellent efficacy in pain models such as CFA and SNL. Infinity Pharmaceuticals patented several isoxazoline heterocyclic FAAH inhibitors lacking electrophile (Behnke et al., 2010). In summary, this unique class of compounds lacking electrophile has shown encouraging outcomes in preclinical studies and holds potential for further development as a drug candidates.

Compd	Structure	ΙC ₅₀ [μΜ]		References
		rFAAH	hFAAH	
74		NA ^a	0.0012 ^b	Estiarte <i>et al.</i> ,2012

Table 16. Inhibitory activities of the compounds 74-77 lacking electrophile against FAAH.



^a Not analyzed. Values were measured ^b with human FAAH expressed in HEK293 cells with a 3h preincubation and AMC-AA as the substrate; ^c by using HTS assay: rat brain homogenates or human FAAH expressed in Chinese hamster ovary (CHO) cells (with a 2 min preincubation) and AMC-AA as the substrate; ^d with human FAAH or rat FAAH (1h preincubation) and [³H]-AEA as the substrate; ^e cell lysate or microsome fractions with the fluorescent substrate AMC-AA or [³H]-AEA.

2.3.8 Marketed drugs and its derivatives as FAAH inhibitors

Several non-steroidal anti-inflammatory drugs (NSAIDs), such as ibuprofen, flurbiprofen, naproxen and carprofen have been utilized in the development of FAAH inhibitors (like compound 78, Table 17) (De Wael *et al.*, 2010), dual FAAH-COX inhibitors (compounds 132-137, Table 29, section 2.6) (Holt *et al.*, 2007, Favia *et al.*, 2012, Fowler *et al.*, 2013, Cipriano *et al.*, 2013) or multi-target compounds (like 145 and 146, Table 29, section 2.6) (Rose *et al.*, 2014). Attempts to reveal new properties of already marketed compounds have resulted in the identification of phenmedipham (79) and amperozide (80) as inhibitors of human FAAH, (Vincent *et al.*, 2009). In addition, some ester analogues of paracetamol have been identified as FAAH inhibitors (see compound 81) (Onnis *et al.*, 2010). In summary, screening of marketed drugs has led to the successful identification of several potent FAAH inhibitors. Many of them are being studied for their dual blockade of FAAH-COX or as multiple target acting compounds.

Compd	Structure	IC50	[µM]	References
		FAAH	MAGL	
78	S S S	1.38 ^a	NA ^b	De Wael <i>et al.</i> ,2010
79		0.377 ^c	NA	Vincent <i>et al.</i> ,2009
80 Phenmedipham	I ho ho	1.34 ^c	NA	Vincent <i>et al.</i> ,2009
81 Amperozide		0.10 ^d	1.9 ^e	Onnis <i>et al.,</i> 2010

Table 17. Inhibitory values for the marketed drugs and its derivatives 78-81 against FAAH and MAGL.

^a Inhibitory activities were measured by using recombinant human FAAH (10 min preincubation) and [³H]-AEA as a substrate. ^b Not analyzed. Values were measured by using ^c rat brain homogenates (HTS assay) or human FAAH expressed in CHO cells (with a 2 min preincubation) and AMC-AA as the substrate; ^d rat brain homogenate (10 min preincubation) and [³H]-AEA as the substrate; ^e human recombinant MAGL (hMAGL) activity with 4-nitrophenylacetate as the substrate.

2.3.9 Miscellaneous FAAH inhibitors

Several substituted 2-thioxoimidazolidin-4-ones (thiohydantoin) and imidazolidine-2,4diones (hydantoin) have been reported to be reversible and competitive FAAH inhibitors; examples are compounds 82 and 83, which showed pl₅₀ values of 5.12 and 5.94, respectively (Table 18) (Muccioli et al., 2006). Minkkilä et al. described boronic acid, 2H-isoxazol-5-one and carbamoyltriazole containing analogues 84-86 (Table 18), respectively, as potent FAAH inhibitors having selectivity over MAGL (Minkkilä et al., 2008, Minkkilä et al., 2009b). Infinity Pharmaceuticals has been granted several patents describing boronic acid analogues as potent FAAH inhibitors (Adams et al., 2008, Behnke et al., 2009, Castro et al., 2010b, Castro et al., 2010a). Abbott has described the identification through HTS of the benzothiazole analogue 87 (Table 18) which they claimed to be a potent, selective and reversible FAAH inhibitor (Wang et al., 2009b). In addition, several diverse amide analogues 88-91 (Table 18) have been described as potent FAAH inhibitors (Scott et al., 2011, Andrzejak et al., 2011, Kharul et al., 2013, Tourteau et al., 2014) and of these, compound 89 (AZ513) acted as a noncovalent, reversible, and noncompetitive inhibitor of FAAH. The structural features of compound 90 resemble a well-known FAAH inhibitor, PF-04457845 (54). The sulfonyl fluoride-based analogue 5-(4-hydroxyphenyl)pentanesulfonyl fluoride (AM3506, 92, Table

18) exhibited good selectivity for FAAH over other serine hydrolases in the ABPP assay and it normalized cardiovascular function in hypertension without causing adverse metabolic effects (Godlewski *et al.*, 2010, Alapafuja *et al.*, 2012). The rapid dilution assay and mass spectrometry analysis suggested that AM3506 inhibited FAAH in an irreversible manner by forming a covalent bond with Ser²⁴¹ of FAAH.

Compd	Structure	ΙC ₅₀ [μΜ]		References
		FAAH	MAGL	_
82		7.58 ^a (p <i>I</i> ₅₀ = 5.12)	NA ^b	Muccioli <i>et al.</i> ,2006
83		1.15 ^a (p <i>I</i> ₅₀ = 5.94)	NA	Muccioli <i>et al.</i> ,2006
84	OH B OH	0.0091 ^a	7.9 ^c	Minkkilä <i>et al.</i> ,2008
85	Y N N	0.00045 ^a	6 ^c	Minkkilä <i>et al.</i> ,2009
86		0.030 ^a	4.6 ^c	Minkkilä <i>et al.</i> ,2009
87		0.018 ^d	NA	Wang <i>et al.</i> ,2009
88	NH NH NH	0.088ª	NI ^e at 30 µM	Andrzejak <i>et al.</i> ,2011
89 AZ513		0.027 ^a (0.551) ^f	NA	Scott <i>et al.</i> ,2011

Table 18. Inhibitory values for the miscellaneous compounds 82-92 against FAAH and MAGL.



^a Values were measured by using rat brain homogenate protein (10 min preincubation) and [³H]-AEA as the substrate. ^b Not analyzed. Inhibitory activity was measured by using ^c human recombinant MAGL and 2-AG as the substrate; ^d rat FAAH enzyme expressed in HEK293 cells and [³H]-AEA as the substrate; ^e human recombinant MAGL and [³H]-2-OG as the substrate; human FAAH expressed in CHO cells (with a 30 min preincubation) and AMC-AA as the substrate; ^g human FAAH as enzyme source and AMC-AA as the substrate; ^h rat FAAH as the enzyme source and AMC-AA as the substrate; ⁱ recombinant human FAAH preparation (expressed in *E. coli*) and [³H]-AEA as the substrate. For the inhibition assay ^j rat Δ TM FAAH was expressed in *E. coli* cells (with a 15 min preincubation) and AMC-AA was used as the substrate; ^k recombinant hexahistidine-tagged human MAGL (hMAGL) was expressed in *E. coli* cells (with a 15 min preincubation) and AMC-AA as the substrate.

2.4 INHIBITORS OF MAGL

To date, numerous classes of MAGL inhibitors targeting either the nucleophilic cysteine residues (Cys²⁰¹, Cys²⁰⁸, and Cys²⁴²) located close to the catalytic site or the nucleophilic Ser¹²² residue of MAGL have been described.

2.4.1 Maleimide-based MAGL inhibitors

Inhibition of MAGL by *p*-chloromercuribenzoic acid (*p*-CMB), mercury chloride (HgCl₂), and *N*-ethylmaleimide (NEM, 93) pointed to the existence of essential thiol containing residues in the active site of MAGL (Table 19) (Tornqvist & Belfrage 1976, Sakurada & Noma 1981). Saario *et al.* developed several NEM analogues, among which *N*-arachidonylmaleimide (NAM, 94) was found to be the most potent MAGL inhibitor (IC₅₀ = 0.14 μ M) (Saario *et al.*, 2005). They also investigated binding of NAM with a comparison model of MAGL and provided an initial insight into the location of the cysteine residues (Cys²⁰⁸ and Cys²⁴²) near the binding site. Site mutagenisis and mass spectrometry studies revealed that NAM formed a Michael conjugate addition product with Cys²⁴² of human MAGL thereby causing its inhibition (Zvonok *et al.*, 2008). Matuszak *et al.* also synthesized *N*-substituted maleimide derivatives among which 1-biphenyl-4-ylmethylmaleimide (95) inhibited MAGL with an IC₅₀ value of 790 nM in an irreversible manner (Matuszak *et al.*, 2009).

Compd	Structure	ΙC ₅₀ [μΙ	M]	References
		MAGL	FAAH	
93 NEM		53ª	NA ^b	Saario <i>et al.</i> ,2005
94 NAM		0.140 ^a (1.10) ^c	3.3 ^d	Saario <i>et al.</i> ,2005
95		0.79 ^c	16.6 ^d	Matuszak <i>et al.</i> ,2009

Table 19. Inhibitory activities of the maleimide compounds 93-95 against MAGL and FAAH.

^a Inhibitory activity has been determined by using rat cerebellar membranes and 2-AG as the substrate. ^b Not analyzed. Inhibitory activity has been determined by using ^c human recombinant MAGL and [³H]-2-OG as the substrate; ^d hFAAH and [³H]-AEA as the substrate.

2.4.2 Natural compounds as MAGL inhibitors

King *et al.* screened a natural compound library and identified two bioisosteric terpenes, pristimerin (96) and euphol (97) as being potent and reversible MAGL inhibitors ($IC_{50} = 93$ nM and 315 nM, respectively) (Table 20) (King *et al.*, 2009b). Moreover, with the help of site mutagenesis studies, they demonstrated that pristimerin and euphol could interact with Cys²⁰⁸ and Cys²⁰¹ respectively. Interestingly, both pristimerin and euphol were also found to inhibit ABHD6 with IC₅₀ values of 98 nM and 9 μ M, respectively. However, they did not affect the activity of ABHD12.

Compd	Structure	I C ₅₀ [µ	IM]	References
		rMAGL ^a	rFAAH⁵	
96 Pristimerin	HO TO TO T	0.093	NIc	King <i>et al.,</i> 2009

Table 20. Inhibitory activity of the triterpenoid compounds 96 and 97 against MAGL and FAAH.



Inhibitory activity has been determined by using ^a purified *E. coli* recombinant rat MAGL and 2-OG as the substrate; ^b rat brain FAAH and [³H]-AEA as the substrate. ^c No inhibition. ^d Not analyzed.

2.4.3 Disulfide-based MAGL inhibitors

Labar *et al.* found disulfiram (98), a marketed drug used for decades to treat alcoholism, and related analogues (such as 99) could act as inhibitors of MAGL (Table 21). They inhibited MAGL in an irreversible manner through an interaction with Cys²⁰⁸ and/or Cys²⁴² of hMAGL (Labar *et al.*, 2007). Later, Kapanda *et al.* published a detailed SAR study of bis(dialkylaminethiocarbonyl)disulfide derivatives and identified them as potent hMAGL inhibitors having high selectivity over FAAH such as compound 100 (Kapanda *et al.*, 2009). Their MAGL activity was attributed to the presence of both thiocarbonyl groups and disulfide moiety which interacted with Cys²⁰⁸ and Cys²⁴² of hMAGL.

Compd	Structure	ΙC ₅₀ [μ	IM]	References
		hMAGLa	hFAAH ^b	
98 Disulfiram	N S S N	0. 36	NAc	Labar <i>et al.</i> ,2007
99	S S S S N S S S N S	0.13	NA	Labar <i>et al.</i> ,2007
100		0.109	> 1000	Kapanda <i>et al.</i> ,2009

Table 21. Inhibitory activity of the disulfide compounds 98-100 against MAGL and FAAH.

Inhibitory activity was determined using ^a human recombinant MAGL and [³H]-2-OG as the substrate; ^b hFAAH and [³H]-AEA as the substrate. ^c Not analyzed.

2.4.4 Isothiazolinone-based MAGL inhibitors

King *et al.* described a new family of isothiazolinone-based MAGL inhibitors (101-103, Table 22) (King *et al.*, 2009a). Kinetic and site-directed mutagenesis studies showed that octhilinone (101) inhibited MAGL through a partially reversible mechanism that involved a specific interaction with Cys²⁰⁸.

Table 22. Inhibitory activity of the isothiazolinone and benzisothiazolinone compounds 101-103 against MAGL and FAAH.

Compd	Structure	IC ₅₀	[µM]	References
		MAGL	FAAH	
101 Octhilinone		0. 088	NA ^b	King <i>et al.</i> ,2009
102	S S S S S S S S S S S S S S S S S S S	0.043	NA	King <i>et al.</i> ,2009
103	N	0.059	NA	King <i>et al.</i> ,2009

^a Inhibitory activity was determined using purified rat MAGL and 2-OG as the substrate. ^b Not analyzed.

2.4.5 Carbamate-based MAGL inhibitors

Hohmann *et al.* stated that the carbamate-based compound URB602 (104) was a selective MAGL inhibitor ($IC_{50} = 28 \mu$ M) (Table 23) (Hohmann *et al.*, 2005). However, subsequently several research groups showed that it also inhibited FAAH with almost similar potency ($IC_{50} = 17 \mu$ M) (Vandevoorde *et al.*, 2007). Later, Szabo *et al.* investigated a series of URB602 analogues and of these, compound 105 displayed 74% inhibition of MAGL at 100 μ M compared to 27% for the parent compound URB602 (Szabo *et al.*, 2011).

Long *et al.* used competitive ABPP to screen a structurally diverse library of carbamates and discovered 4-bisarylcarbinol analogue JZL184 (106) as a potent MAGL inhibitor having good selectivity (>100-fold) over other serine hydrolases (Table 23) (Long *et al.*, 2010). Moreover, they identified 4-aryloxybenzyl-based analogue JZL195 (107) as a dual FAAH-MAGL

inhibitor. JZL184 (106) treated mice had increased brain 2-AG levels and exhibited a broad array of CB₁-dependent behavioral effects, including analgesia, hypothermia and hypomotility (Long *et al.*, 2009a). JZL184 (106) irreversibly inhibited MAGL via carbamoylation of the enzyme's serine nucleophile (Long *et al.*, 2009b). However, due to the residual low-level cross-reactivity with FAAH and peripheral carboxylesterases (CESs), JZL184 (106) may not be a suitable compound for investigating in biological studies with high doses or chronic treatment. JZL195 (107) was found to produce a CB₁ agonist like effect in the mouse tetrad test. JZL195 (107) also evoked THC like responses in a drug discrimination behavior test and these could be reversed by treatment with a CB₁ antagonist (Long *et al.*, 2009a, Long *et al.*, 2009b, Schlosburg *et al.*, 2010). Chang *et al.* produced an *O*-hexafluoroisopropyl (HFIP) carbamate analog of JZL184 (106), KML29 (108, Table 23) which potently and selectively inhibited MAGL *in vitro* and *in vivo* with minimal cross-reactivity with central and peripheral serine hydrolases, including no detectable activity against FAAH (Chang *et al.*, 2012). Furthermore, a dual FAAH-MAGL inhibitor JZL195 (107) was converted to an HFIP carbamate analog, leading to identification of potent MAGL inhibitor JW642 (109).

Kapanda *et al.* reported 2,4-dinitroaryldithiocarbamate-based compound 110 (CK16) to be a potent MAGL inhibitor, and compound 111 (CK37) as a dual FAAH-MAGL inhibitor (Table 23) (Kapanda *et al.*, 2012). They found that these compounds behaved as irreversible inhibitors of MAGL, interacting with Cys²⁰⁸ or Cys²⁴² and Ser¹²² residues of the enzyme. When tested in a murine melanoma cell line that endogenously expresses MAGL, CK16 (110) only slightly increased 2-AG levels in contrast to the robust increase observed with CK37 (111). Niphakis *et al.* reported that MJN110 (112, Table 23), an *N*-hydroxysuccinimidyl (NHS) carbamate analog of JZL184 (106) /KML29 (108), possessed superior potency and comparable selectivity to the previously developed MAGL inhibitors, such as KML29 (108), with both acute and chronic dosing regimens (Niphakis *et al.*, 2013). Moreover, MJN110 (112) could reverse established mechanical allodynia in a rat model of diabetic neuropathy, pointing to a potential clinical application for MAGL inhibitors in the treatment of chronic pain caused by diabetes.

Table 23. Inhibitory activities of the carbamate compounds 104-112 against MAGL, ABHD6 and FAAH.

Compd	Structure	IC ₅₀	[µM]	References	
		MAGL	ABHD6	FAAH	_
104 URB602	₩	28ª	NA ^b	17 ^c	Hohmann <i>et al.,</i> 2005 Vandevoorde <i>et</i> <i>al.,</i> 2007

105	HO H O O	26% ^d	NA	NA	Szabo <i>et al.</i> ,2011
106 JZL184		0.010 ^e	3.27 ^e	4.7 ^e	Long <i>et al.</i> ,2009 Long <i>et al.</i> ,2010
107 JZL195	N_{0}	0.013 ^e	0.050 ^e	0.019 ^e	Long <i>et al.</i> ,2009 Long <i>et al.</i> ,2010
108 KML29	$ \begin{array}{c} O \\ O \\ O \\ HO \\ HO \\ O \\ O \\ O \\ O \\ O$	0.015 ^e	4.87 ^e	>50 ^e	Chang <i>et al.</i> ,2012
109 JW642	Contraction N CF3	0.0076 ^e	0.107 ^e	31 ^e	Chang <i>et al.</i> ,2012
110 CK16	NO2 NNNS NO2	0.354 ^f	NA	>1000 ^g	Kapanda <i>et al.</i> ,2012
111 CK37	$\sum_{N \in \mathbb{N}} N^{N} \sum_{N \in \mathbb{N}} N^{NO_2}$	0.154 ^f	NA	1.73 ^g	Kapanda <i>et al.</i> ,2012
112 MJN110		0.0095 ^e	0.260 ^e	>100 ^e	Niphakis <i>et al.</i> ,2013

^a Inhibitory activity was determined by using cytosolic fractions from rat brain and 2-OG as the substrate. ^b Not analyzed. ^c Inhibitory activity was determined by using rat brain homogenate and [³H]-AEA as the substrate. ^d Remaining activity at 100 µM compared to 73% for the URB602 (human recombinant MAGL used as the enzyme source and 4-nitrophenylacetate (4-NPA) as the substrate). ^e Inhibitory activity was determined by using competitive ABPP: reductions in FP-Rh labeling of MAGL, ABHD6, and FAAH were quantified following preincubation of proteomic samples with the test compound. Brain membrane proteomes were used to measure the inhibition of mouse and rat enzymes, whereas human enzymes were evaluated from proteomes of transiently transfected HEK293T cells. Inhibitory activity was determined by using ^f human recombinant MAGL and [³H]-2-OG as the substrate; ^g human recombinant FAAH and [³H]-AEA as the substrate.

2.4.6 Urea-based MAGL inhibitors

Sanofi-Aventis possess several patents describing triazolopyridine carboxamide derivatives as dual FAAH-MAGL inhibitors (such as 113 and 114, Table 24) (Even & Hoornaert 2008a, Even & Hoornaert 2008b) or MAGL inhibitors (such as 115 and 116, Table 24) (Hoornaert 2008c). Zvonok *et al.* identified the 2,5-LY2183240 isomer 5-[(biphenyl-4-yl)methyl]-*N*,*N*-dimethyl-*2H*-tetrazole-2-carboxamide (AM6701, 117) as a potent inhibitor of MAGL (Table 24) (Zvonok *et al.*, 2008). Later Bertrand *et al.* from Sanofi-Aventis published the first crystal structure of human MAGL in its apo form and in a complex with the covalent inhibitor SAR629 (118, Table 24), a derivative of an earlier patented triazolo-carboxamide series (Bertrand *et al.*, 2010).

Morera *et al.* reported that benzotriazol-1-yl carboxamide derivatives functioned as dual FAAH-MAGL inhibitors (such as 119) or MAGL inhibitors (like compound 120, ML30) (Table 24) (Morera *et al.*, 2012). Our group revealed the importance of the 1,2,4-triazole leaving group in piperidine/piperazine ureas if one wanted to achieve excellent potency and selectivity (Aaltonen *et al.*, 2013, Korhonen *et al.*, 2014). Compound 121 (JJKK-048, Table 24) was found to be potent inhibitor of both human and rodent MAGL (IC₅₀ < 0.4 nM). The 1,2,4-triazole analogue of JZL184 (106) (compound 122, Table 24) was also identified as a potent MAGL inhibitor with good selectivity over FAAH.

Compd	Structure	IC ₅₀	References	
		MAGL	FAAH	
113		0.004ª	0.010 ^b	Even 2008a
114	CF3 ON	0.002ª	0.002 ^b	Even 2008b
	$ \begin{array}{c} $			

Table 24. Inhibitory activity of the urea compounds 113-122 against MAGL and FAAH.

0.004^a NAc Hoornaert 2008c 0.025^a Hoornaert 2008c NA 0.0017^d NA Zvonok et al.,2008 (0.0009)^e 0.0011^f NA Bertrand et (219 pM)⁹ al.,2010 0.003^h 0.002ⁱ Morera et al., 2012

120 0.0054^h 0.562ⁱ Morera et al., 2012 $(0.0044)^{f}$ ML30 (0.0019)^g (0.0015)^j 121 214 pM^f 4.8^k Aaltonen et al.,2013 **JJKK-048** (275 pM)^g (363 pM)^j 122 0.0007^j 1.6^k Korhonen et al.,2014

Inhibitory activity was determined by using ^a mice brain homogenate (cerebellum and medulla eliminated) and [³H]-2-OG as the substrate; ^b mouse brain homogenate (cerebellum and medulla eliminated) and [³H]-AEA as the substrate. ^c Not analyzed. ^d Inhibitory activity was determined by using fluorescence-based assay by using either rMAGL or ^e hMAGL and AMC-AA as the substrate. Inhibitory activity was determined by using ^f rat or ^g mouse brain membranes and 2-AG as the substrate; ^h human MAGL and [³H]-2-OG as the substrate; ⁱ human FAAH and [³H]-AEA as a substrate; ^j HEK293 cell lysates overexpressing hMAGL and 2-AG as the substrate; ^kCOS-7 cells transiently overexpressing human recombinant FAAH and [³H]-AEA as the substrate.

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115

116

117

118

119

SAR629

AM6701

2.4.7 Miscellaneous MAGL inhibitors

Bisogno *et al.* reported on the modification of tetrahydrolipstatin (THL, orlistat) and identified OMDM169 (123) as a reversible MAGL inhibitor (Table 25) (Bisogno *et al.*, 2009). However, this compound was also found to inhibit other enzymes such as pancreatic lipase, FAAH and DAGL. Janssen Pharmaceuticals have been awarded three patents for amide analogues as MAGL inhibitors (Bian *et al.*, 2010, Chevalier *et al.*, 2010, Flores *et al.*, 2010). Very recently, López-Rodríguez *et al.* described compound 124 (Table 25) as a potent (IC₅₀ = 0.18 μ M) and reversible MAGL inhibitor having good selectivity towards the other endocannabinoid targets and several other tested receptors and enzymes (Hernández-Torres *et al.*, 2014). Compound 124 also improved the clinical progression of the disease in a mouse model of multiple sclerosis (MS) without inducing undesirable CB₁ -mediated side effects.

Compd	Structure	ΙC ₅₀ [μΜ]		References
		hMAGL ^a	hFAAH ^b	-
123 OMDM169		0.89 ^a	3 ^b	Bisogno <i>et al.</i> , 2009
124	Ph O O O O O O O O O O O O O O O O O O O	0.24 ^c	18 ^d	Hernández-Torres <i>et al.,</i> 2014

Table 25. Inhibitory activity of the compounds 123 and 124 against MAGL and FAAH.

Inhibitory activity was determined by using ^a human recombinant MAGL and [³H]-2-AG as the substrate; ^b rat brain as enzyme source and [¹⁴C]-AEA as the substrate; ^c cytosolic and membrane homogenates of rat brain as MAGL source and [³H]-2-OG as the substrate; ^d rat brain homogenate as the enzyme source and [³H]-AEA as the substrate.

2.5 INHIBITORS OF ABHD6

To date, very little success has been achieved in the identification of potent and selective ABHD6 inhibitors (see subsections 2.5.1-2.5.3). There are several examples of non-selective ABHD6 inhibitors e.g. MAFP (2, Table 1), THL, RHC-80267, and the triterpene pristimerin (Navia-Paldanius *et al.*, 2012).

2.5.1 Carbamate-based ABHD6 inhibitors

In 2007, Li *et al.* screened a library of carbamate derivatives using a functional proteomic strategy and identified compound WWL70 (125) as a potent ($IC_{50} = 70$ nM) and selective inhibitor of ABHD6 (Table 26) (Li *et al.*, 2007). WWL70 (125) exerted anti-inflammatory and beneficial effects in metabolic disorders (Thomas *et al.*, 2013, Tchantchou & Zhang 2013). A brain penetrating and selective ABHD6 inhibitor, WWL123 (126, Table 26) (Bachovchin *et al.*, 2010), an isoster analogue of WWL70 (125), showed an antiepileptic effect in pentylenetetrazole (PTZ)-induced epileptiform seizures and spontaneous seizures in R6/2 mice (Naydenov *et al.*, 2014).

Table 26. Inhibitory values for the carbamate compounds 125 and 126 against ABHD6, MAGL and FAAH.

Compd	Structure	ΙC ₅₀ [μΜ]			References
		ABHD6	MAGL	FAAH	
125 WWL70		0.070 ^a (0.085) ^b	NIc	NI	Li <i>et al.</i> , 2007 Navia-Paldanius <i>et</i> <i>al.</i> , 2012
126 WWL123		0.43 ^a	NI	NI	Bachovchin <i>et al.,</i> 2010

^a Inhibitory activity was determined by using transiently transfected COS-7 cells with the human ABHD6 cDNA and using competitive ABPP reductions in FP-Rh labeling of ABHD6 was quantified (in ABPP assay, these compounds are ABHD6 selective). ^b Glycerol liberated from 1-AG hydrolysis was determined with a sensitive fluorescent glycerol assay using lysates of HEK293 cells expressing hABHD6 (Navia-Paldanius *et al.*, 2012). ^c No inhibition via ABPP.

2.5.2 Triazole urea-based ABHD6 inhibitors

Hsu *et al.* recently claimed that (2-substituted)-piperidyl-1,2,3-triazole ureas (127-129) were ABHD6 inhibitors (Table 27) (Hsu *et al.*, 2013). KT182 (127) and KT203 (128) showed exceptional potency and selectivity in Neuro2A cells (<5 nM), and at equivalent doses in mice (1 mg/kg), these compounds acted as systemic and peripherally restricted ABHD6 inhibitors, respectively. Moreover, KT185 (129) was found to be an orally bioavailable ABHD6 inhibitor having excellent selectivity against other brain and liver serine hydrolases *in vivo*.

Compd	Structure	ΙC ₅₀ [μΜ]			References
		ABHD6	MAGL	FAAH	
127 KT182		0.0017 ^a (0.015) ^b	NIc	NI	Hsu <i>et al.</i> ,2013
128 KT203		0.00082 ^a (0.0039) ^b	NI	NI	Hsu <i>et al.</i> ,2013
129 KT185		0.0013ª (0.0136) ^b	NI	NI	Hsu <i>et al.</i> ,2013

Table 27. Inhibitory activity of the carbamate compounds 127-129 against ABHD6, MAGL and FAAH.

Inhibitory activity was determined by using ^a competitive ABPP in mouse brain membrane proteomes. The reductions in HT-01 (Hsu *et al.*, 2012) probe labeling of ABHD6 were quantified after preincubation of a proteomic sample with the inhibitor; ^b recombinant mouse ABHD6 protein overexpressed in HEK293T cells and 2-AG as a substrate. ^c No inhibition in competitive ABPP of mouse brain membrane proteomes.

2.5.3 Miscellaneous ABHD6 inhibitors

Marrs and colleagues described UCM710 (130) as a dual inhibitor of ABHD6 (IC₅₀ = 2.4 μ M) and FAAH (IC₅₀ = 4 μ M) (Table 28) (Marrs *et al.*, 2011). Very recently, Janssen *et al.* described a glycine sulfonamide analogue LEI-106 (131, Table 28) as a potent and selective dual inhibitor of sn-1-diacylglycerol lipase α (DAGL- α) and ABHD6 (Janssen *et al.*, 2014).

Table 28. Inhibitory activity of the carbamate compounds 130 and 131 against ABHD6, MAGL and FAAH.

Compd	Structure	IC ₅₀ [μM]	References	
		ABHD6	MAGL	FAAH	
130 UCM710	C_6H_{13} H_{7} O O	2.4 ^a	NI ^b	4 ^c	Marrs <i>et al.,</i> 2011

131
LEI-106

$$K_i = 0.8^d$$
 NI^e NI^e Janssen *et al.*, 2014
 $K_i = 0.8^d$ NI^e VI^e Janssen *et al.*, 2014

^a Inhibitory activities were determined by using homogenates prepared from COS-7 cells heterologously expressing ABHD6 and [³H]-2-AG as the substrate. ^b No inhibition IC₅₀ > 1mM (homogenates prepared from COS-7 cells heterologously expressing MAGL and [³H]-2-AG as the substrate). Inhibitory activities were determined by using ^c homogenates prepared from COS-7 cells heterologously expressing FAAH and [³H]-AEA as the substrate; ^d HEK293T cells transiently transfected with hABHD6 and 2-AG as the substrate. (Note: K_i (DAGL_a) = 0.7 µM. This was determined using HEK293T cells transiently transfected with hDAGL-a and *para*-nitrophenylbutyrate (PNP-butyrate) as the substrate).

2.6 COMPOUNDS WITH DUAL OR MULTIPLE TARGETS

Several research groups have reported dual FAAH-COX inhibitors, such as compounds 132-137 (Table 29) (Holt et al., 2007, Favia et al., 2012, Fowler et al., 2013, Cipriano et al., 2013, De Vivo et al., 2014a, De Vivo et al., 2014b). Instead of targeting one enzyme, dual inhibition has been considered as a potential approach to achieve a sufficent degree of analgesia at a lower dose (Naidu et al., 2009, Sasso et al., 2012). Indazole-5-carboxylic acid derivatives 138 and 139 were identified as dual cytosolic phospholipase $A_2\alpha$ (cPLA₂ α)-FAAH inhibitors (Forster *et al.*, 2010, Zahov et al., 2011). The authors speculated that dual inhibition could lead to improved analgesic and anti-inflammatory properties in comparison with selective cPLA₂ α and FAAH inhibition. As described above in sections 2.3 and 2.4, several compounds such as CAY10499 (69, Table 14), JZL195 (107, Table 23), 113, 114 and 119 (Table 24) act as dual FAAH-MAGL inhibitors. Furthermore, a series of dual acting FAAH-MAGL inhibitors has been investigated and of these, (2R)-(–)-oxiran-2-ylmethyl(4-benzylphenyl)acetate (140, Table 29) potently inhibited human recombinant MAGL ($IC_{50} = 2.4 \mu M$), rat brain MAGL hydrolysis $(IC_{50} = 0.68 \mu M)$, and rat brain FAAH $(IC_{50} = 0.29 \mu M)$ (Cisneros *et al.*, 2012). This compound acted as a noncompetitive, reversible MAGL inhibitor. Dual acting FAAH-cholinesterase (ChE) inhibitors 141 and 142 have also been identified (Table 29). By targeting both the cholinenergic and endocannabinoid systems, it has been proposed that they may be able to prevent the progression of Alzheimer's disease (AD) (Rampa et al., 2012). AA-5-HT (6), a dual FAAH-TRPV1 blocker, was found to be more efficacious at inhibiting pain and anxiety in rodents than either a high-potency, FAAH-selective inhibitor or a TRPV1-selective inhibitor (Costa et al., 2010, Micale et al., 2009). Maione et al. reported a similar study using the FAAH-TRPV1 blocker OMDM198 (143, Table 29) and the FAAH inhibitor-TRPA1 agonist, OMDM202 (144, Table 29), where they found these compounds had improved analgesic effects similar to those observed with AA-5-HT (Maione et al., 2013). Very recently, NSAIDserotonin conjugates ibuprofen-5-HT (145) and flurbiprofen-5-HT (146) were developed and they were able to inhibit three targets, FAAH, TRPV1, and COX2 (Table 29). This kind of triple inhibition may offer effective pain relief with a high therapeutic index (Rose et al., 2014).

Compd	Structure	IC ₅₀ [µM] or	References	
·		FAAH	target(s)	
132 ibu-am5	L C N N	4.7 ^{a,b} 2.5 ^{a,c}	NI (at 300 µM) for MAGL ^d 60 for COX-1 ^e NI for COX-2 ^f	Holt <i>et al.</i> ,2007 Fowler <i>et al.</i> ,2013
133	CI C	22 ^g	NA ^h for MAGL 74.3 for COX-1 ^e 72.3 for COX-2 ^f	Favia <i>et al.</i> ,2012
134 Flu-AM1		0.44 ^a	NA for MAGL 6.6 for COX-1 ^e 42 for COX-2 ^f	Cipriano <i>et al.</i> ,2013
135 Nap-AM1		0.74 ^a	NA for MAGL 56 for COX-1 ^e >100 for COX-2 ^f	Cipriano <i>et al.</i> ,2013
136	H O O O O	0.015 ⁱ	0.3 for COX-1 ^e 0.32 for COX-2 ^f	De Vivo <i>et al.,</i> 2014a
137 (+)-isomer	~~~~~ И состать он	0.094 ⁱ 0.097 ^j	0.00001 for COX-1 ^e 0.010 for COX-2 ^f	De Vivo <i>et al.,</i> 2014b
138	Сения Ссоон	0.16 ^k	0.005^{I} for cPLA ₂ a	Forster et al., 2010
139	F COOH	0.090 ^k	0.015^{I} for cPLA ₂ a	Zahov <i>et al.,</i> 2011
140	O'CLI.~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	0.29 ^a	0.64 ^m for MAGL 2.4 ⁿ for hrMAGL	Cisneros <i>et al.,</i> 2012
141	of the property of the states	0.050°	0.075 ^p for AChE 0.0016 ^q for BuChE	Rampa <i>et al.,</i> 2012
142	CN ON ON ON ON THE	0.040°	0.090 ^p for AChE 0.0017 ^q for BuChE	Rampa <i>et al.,</i> 2012
143 OMDM198	CF ₃ N O CF ₃ CF ₃	3.36°	1 ^r for hTRPV1	Maione <i>et al.</i> , 2013

Table 29. Inhibitory activity of the compounds 132-146 acting at dual or multiple targets.

144 OMDM202		0.38°	1 ^r for rTRPA1	Maione <i>et al.</i> , 2013
145 Ibuprofen-5- HT	L C L NH	5 ^s	10 ^t for COX2 6 ^u for TRPV1	Rose <i>et al.,</i> 2014
146 Flurbiprofen- 5-HT	F S S S S S S S S S S S S S S S S S S S	15 ^s	8 ^t for COX2 9 ^u for TRPV1	Rose <i>et al.</i> , 2014

Values were measured by using a rat brain homogenate protein and [3H]-AEA as the substrate at b pH = 8 or ^c pH = 6; ^d rat brain cytosolic fractions or human recombinant MAGL and [³H]-2-OG as the substrate; ^e ovine COX-1 or ^f human recombinant COX-2 enzyme and AA as the substrate; ^g 30 min preincubation with protein and [³H]-AEA as the substrate. ^h Not analyzed. Values were measured by using ⁱ rat brain homogenate and [3H]-AEA as the substrate; ^j human FAAH expressed in HEK293 cells with a 50 min preincubation and [³H]-AEA as the substrate; ^k rat brain microsomes and N-(2-hydroxyethyl)-4-pyren-1ylbutanamide as the substrate; ¹ cPLA₂a isolated from human platelets and 1-stearoyl-2-arachidonoyl-snglycero-3-phosphocholine sonicated with 1,2-dioleoyl-sn-glycerol as the substrate; ^m cytosolic and membrane homogenates of rat brain and [³H]-2-OG as the substrate; ⁿ lysates of human recombinant MAGL and 4-nitrophenyl acetate as the substrate; ° rat brain FAAH and [14C]-AEA as the substrate; P human recombinant AChE using acetylthiocholine iodide as the substrate; ^q human serum BuChE using butyrylthiocholine iodide as the substrate; ^r intracellular Ca²⁺ elevation mediated by TRPV1 and TRPA1 channels overexpressed in HEK293 cells; ^s human recombinant FAAH and AMC-AA as the substrate; ^t human recombinant COX2 and 10-acetyl-3,7-dihydroxyphenoxazine (ADHP) as the substrate; "BEAS-2B human TRPV1 over-expressing cells and Fluo-4 AM, a fluorogenic calcium indicator using the Fluo-4 Direct Kit (Invitrogen).

3 Therapeutic Utility of Endocannabinoid Hydrolase Inhibitors

During the past few years, major progress has been made in the development of endocannabinoid hydrolase inhibitors. The field has evolved from substrate-based potent albeit non-selective FAAH inhibitors to diverse classes of FAAH inhibitors having excellent potency and significantly improved selectivity profiles (Seierstad & Breitenbucher 2008, Vandevoorde 2008, Petrosino & Di Marzo 2010, Deng 2010, Minkkilä et al., 2010, Otrubova et al., 2011, Feledziak et al., 2012, Blankman & Cravatt 2013, Bisogno & Maccarrone 2013). Several α -ketoheterocyclic, carbamate and urea-based FAAH inhibitors have been used widely in diverse preclinical models to validate the therapeutic potential of FAAH inhibition in pain, inflammation, CNS disorders etc. Some of these compounds have even progressed to clinical trials. Compounds, such as OL-135 (12, Table 3) and URB597 (23, Table 4) are known to cause significant elevations of the endogenous levels of the AEA (as well as OEA and PEA) and they have been found to be highly effective in different pain models, including inflammatory and neuropathic models (Lichtman et al., 2004b, Chang et al., 2006, Piomelli et al., 2006, Jayamanne et al., 2006, Russo et al., 2007). Analogues of OL-135 (12) such as CE-12 (17, Table 3) and 18 (Table 3) have shown extended duration of FAAH inhibition thereby achieving long lasting analgesic effects (Ezzili et al., 2011, Otrubova et al., 2013). A peripherally restricted inhibitor URB937 (28, Table 4) along with its improved selectivity and biological stability was found to be effective in pain models such as SNL (Clapper et al., 2010). Several other in vivo efficacious compounds ST4070 (35, Table 6) (Gattinoni et al., 2010a) and ST4020 (39, Table 6) (Minetti et al., 2009) have emerged as potential anxiolytic drugs while ST3913 (41, Table 7) has elicited an analgesic effect (Butini et al., 2012). Some of the most remarkable in vivo efficacious FAAH inhibitors are PF-3845 (51, Table 10) (Ahn et al., 2009b), PF-465 (52, Table 10) (Johnson et al., 2009), PF-04457845 (54, Table 10) (Johnson et al., 2011), and PF-04862853 (57, Table 10) (Meyers et al., 2011b). Both PF-3845 (51) and PF-465 (52) have been shown to increase AEA levels and demonstrated to possess anti-inflammatory activity in a rat CFA model. PF-04457845 (54) was well tolerated in the phase I clinical study, however, it failed to show efficacy in patients in phase II clinical trials (Huggins et al., 2012). Other ongoing or completed studies with PF-04457845 (54) include trials evaluating the potential of these types of drugs for treating Tourette syndrome, cannabinoid dependence, fear conditioning, sleep and acute and chronic pain (https://clinicaltrials.gov/ct2/results?term=PF-04457845&Search=Search). PF-04862853 (57) is currently under consideration for human trials. All of the three compounds JNJ-1661010 (58, Table 11) (Keith et al., 2008), JNJ-40355003 (59, Table 11) (Keith et al., 2012) and JNJ-42119779 (61, Table 11) (Keith et al., 2014b) have been reported to elevate the levels of AEA, PEA, and OEA. JNJ-1661010 (58) exerted analgesic activity in both MTI and SNL models of neuropathic

pain (Karbarz et al., 2008), and JNJ-40355003 (59) treated rats exposed to chronic stress did not develop any significant increases in anxiety-like behavior (Hill et al., 2013) while JNJ-42119779 (61) showed modest efficacy in a neuropathic pain model. Takeda's compounds 63 and 64 (Table 11) possessed analgesic efficacy in animal models of pain (Kono et al., 2013, Kono et al., 2014). AM3506 (92, Table 18) has been reported to reduce blood pressure, heart rate and to improve cardiac contractility in hypertensive animals through activation of CB1 receptors, which correlates with the blockade of FAAH activity and the subsequent increase in tissue levels of AEA (Godlewski et al., 2010). In contrast to the above FAAH inhibitors, compounds lacking a reactive carbonyl group such as JNJ-40413269 (76, Table 16) and MK-4409 (77, Table 16) (Chobanian et al., 2014, Keith et al., 2014a) has shown promising preclinical profiles and excellent efficacy in pain models such as SNL. In addition to the above classes, 1,3,4-oxadiazol-2-one and β -lactam have emerged as promising scaffolds for the development of potent and selective FAAH inhibitors (see subsections 2.3.5 and 2.3.6). Several other candidates are under clinical investigations but their chemical structures have not been disclosed. For example, Janssen's JNJ-42165279 recently completed phase 1 clinical studies (https://clinicaltrials.gov/ct2/show/NCT01650597?term=FAAH&rank=6) and Vernalis' V158866 has passed through First-in-Human (FIH) study а (https://clinicaltrials.gov/ct2/show/NCT01634529?term=FAAH&rank=8).

In contrast to FAAH blockers, very few potent MAGL inhibitors have been tested (Minkkilä et al., 2010, Feledziak et al., 2012, Blankman & Cravatt 2013, Kapanda et al., 2013). However, some compounds have shown promising results in preclinical models of pain, inflammation, vomiting, cancer, etc. (Mulvihill & Nomura 2013). The earlier developed maleimide, dithio-, triterpenoid- and octhilinone analogues targeted the cysteine residues near to the catalytic site of MAGL. However, these analogues are likely to react with other cysteine containing proteins. The appearance of X-ray structures of MAGL has facilitated the development of piperidine/piperazine containing carbamates/ureas targeting mainly the nucleophilic Ser¹²² residue of MAGL. Brain 2-AG levels were elevated in JZL184 (106, Table 23) treated mice and the drug exhibits a broad array of CB₁-dependent behavioral effects, including analgesia, hypothermia and hypomotility (Long et al., 2009a). It is known that genetic inactivation of MAGL or chronic treatment with the irreversible MAGL inhibitor JZL184 (106) causes a prompt increase in 2-AG levels thereby leading functional antagonism, pharmacological tolerance, and receptor desensitization that eventually weakens the antinociceptive properties of the inhibitor (Lichtman et al., 2010, Chanda et al., 2010, Schlosburg et al., 2010). Functional antagonism associated with chronic MAGL blockade may be avoided by partially inhibiting MAGL. Thus, partial blockade of MAGL by treatment with a low dose of JZL184 (106) has maintained the antinociceptive and anxiolytic-like responses after chronic treatment (Busquets-Garcia et al., 2011). In KML29 (108, Table 23) treated mouse, 10-fold elevations in 2-AG (and reductions in AA) have been detected without any accompanying alterations in the FAAH substrates, AEA, OEA, or PEA (Chang et al., 2012). MJN110 (112, Table 23) has been reported to be able to reverse established mechanical allodynia in a rat

model of diabetic neuropathy (Niphakis *et al.*, 2013). In comparison to the irreversible MAGL inhibitors described above, the first potent and selective reversible MAGL inhibitor 124 (Table 25) has been found to be effective in the treatment of multiple sclerosis (MS) without any sign of CNS side effects (Hernández-Torres *et al.*, 2014).

At present, very limited pharmacological data of ABHD6 inhibitors are available. ABHD6 is considered as an emerging target to treat inflammation, metabolic disorders, and epilepsy (Tchantchou & Zhang 2013, Alhouayek *et al.*, 2013, Thomas *et al.*, 2013, Naydenov *et al.*, 2014). Chronic treatment with WWL70 (125, Table 26) has exerted anti-inflammatory and neuroprotective effects in a mouse model of traumatic brain injury (TBI) (Tchantchou & Zhang 2013). Inhibition of ABHD6 with WWL70 (125) has also been reported to protect mice against high-fat-diet-induced obesity and glucose intolerance (Thomas *et al.*, 2013). WWL123 (126, Table 26), a brain penetrating selective ABHD6 inhibitor, exerted an antiepileptic effect in pentylenetetrazole (PTZ)-induced epileptiform seizures and also against spontaneous seizures in R6/2 mice (Naydenov *et al.*, 2014). Furthermore, ABHD6 blockade retained its antiepileptic activity even with chronic dosing and no tolerance was observed in contrast to the situation reported after chronic treatment with CB1 receptor agonists (Blair *et al.*, 2009) and MAGL inhibitors (Schlosburg *et al.*, 2010).

The development of selective ABHD12 inhibitors is eagerly awaited. However, a genetic mutation of ABHD12 has been associated with neurodegenerative disease, and hence, it is possible that inhibitors of ABHD12 may cause long term adverse effects.

As mentioned in subsection 2.2.1, inhibition of FAAH not only causes an increase in the levels of AEA but the concentrations of other lipids are also affected; for example levels of OEA and PEA increase which results in several effects mediated by non-cannabinoid receptors such as GPR118, TRPV1 or PPAR α . These kinds of effects might enhance or reduce the favourable properties of FAAH, for example, activation of PPARs could also modulate inflammatory effects whereas activation of TRPV1 could evoke effects that are opposite to those mediated by cannabinoid receptor activation. As discussed in section 2.6, compounds with either dual targets, such as FAAH-COX inhibitors, cPLA₂ α -FAAH inhibitors, FAAH-ChE inhibitors, FAAH-TRPV1 blocker, FAAH inhibitor/TRPA1 agonist or compounds acting on multiple targets, such as FAAH, TRPV1, and COX2 might be beneficial since this may be one way to reduce any potential side effects.

4 Aims of the Study

This thesis belongs to the medicinal chemistry research field and it involves the development of potent and selective inhibitors of endocannabinoid hydrolases (FAAH, MAGL and ABHD6). The main objectives of this research work have been as follows:

- 1. To design, synthesize and identify novel 1,3,4-oxadiazol-2-ones as potent and selective inhibitors of FAAH and ABHD6.
- 2. To exploit 1,2,5-thiadizole carbamate scaffold in the identification of novel potent and selective ABHD6 inhibitors.
- 3. To identify potential lead structures and utilize them in the development of potent and selective MAGL inhibitors.
5 Experimental Section

5.1 MATERIALS AND METHODS

Reagents and solvents were purchased from commercial suppliers and were used without further purification. Most of the reactions were monitored by thin-layer chromatography (TLC) with suitable UV visualization. Purification was carried out by flash chromatography (FC) or by CombiFlash Companion (Teledyne Isco, USA). ¹H NMR and ¹³C NMR were recorded on a Bruker Avance AV 500 (Bruker Biospin, Switzerland) spectrometer and processed from the recorded FID files with TOPSPIN 2.1 software or ACD/NMR Processor Academic Edition. ESI-MS spectra were acquired using an LCQ quadrupole ion trap mass spectrometer equipped with an electrospray ionization source (Thermo LTQ, San Jose, CA, USA). Elemental analyses were performed on a ThermoQuest CE instrument (EA 1110 CHNS-O) or a Perkin-Elmer PE 2400 Series II CHNS-O Analyzer. For a racemic compound and its pure enantiomers, optical rotation angles ($[\alpha]_D$) were measured using a Jasco model P-2000 polarimeter. The enantiomers of selected racemic compounds described in subsection 5.2.1 were separated by preparative high-performance liquid chromatography (HPLC; Shimadzu LC- 10Avp, Fennolab, Fenno Medical Oy) on a chiral column (Lux 5µ Cellulose-2, 250 x 21.2 mm, Phenomenex).

5.2 DESIGN AND GENERAL SYNTHESIS PROCEDURES

The general synthetic pathways of 1,3,4-oxadiazol-2-ones as inhibitors of FAAH and ABHD6, 1,2,5-thiadiazole carbamates as inhibitors of ABHD6, and loratadine analogues as inhibitors of MAGL, are briefly described below. The detailed synthetic procedures have been reported in the original publications (I-IV).

5.2.1 Chiral 1,3,4-oxadiazol-2-ones (I)

Earlier our group has defined 1,3,4-oxadiazol-2-ones (i, Figure 12) as potent and reversible inhibitors of both FAAH and MAGL (Minkkilä *et al.*, 2009b, Käsnänen *et al.*, 2013). However, in the ABPP assay, they showed poor selectivity for the serine hydrolases in the mouse brain membrane proteomes (unpublished work). To improve the selectivity, it was decided to adopt a new approach where the 1,3,4-oxadiazol-2-one scaffold (i) was combined with non-steroidal anti-inflammatory drugs (NSAIDs, ii, Figure 1), ibuprofen (Fowler *et al.*, 2013, Holt *et al.*, 2007, De Wael *et al.*, 2010) and carprofen (Favia *et al.*, 2012) which both known to have

activity towards FAAH and/or COX isoenzymes. Ibuprofen and carprofen both have a chiral center and the *R*-enantiomer of ibuprofen is known to be slightly more potent than the *S*-enantiomer with respect to its ability to inhibit FAAH (Fowler *et al.*, 1999). Hence, it was thought that by introducing a chiral center into the 1,3,4-oxadiazol-2-ones, it would be possible to study its effect on the selectivity towards either FAAH or MAGL. Figure 12 illustrates the hybridization approach where our earlier disclosed 1,3,4-oxadiazol-2-ones (i) are combined with derivatives of phenylalkanoic acid present in marketed NSAIDs (ii) to afford novel chiral 1,3,4-oxadiazol-2-ones (iii).



Figure 12. Chiral 1,3,4-oxadiazol-2-ones via the hybridization approach.

The general synthetic routes of 1,3,4-oxadiazol-2-ones 147-170 are listed below (Schemes 1 & 2, Table 30). The detailed synthetic procedures for the compounds 147-170 are described in the original publication (I). Briefly, an appropriate ketone or aldehyde was condensed with methyl carbazate to afford Schiff's base which was reduced *in situ* by either sodium borohydride or 10% Pd/C to the corresponding hydrazine carboxylate derivative. Finally, a cyclization in the presence of phosgene gave the desired 1,3,4-oxadiazol-2-ones 147-170.



Scheme 1. Synthesis of 1,3,4-oxadiazol-2-ones 147-151 and 154-170. Reagents and conditions: (a) $NH_2NHCOOR^2$ ($R^2 = Me$), MeOH, 4 Å molecular sieves, 3-4 drops of AcOH, reflux, 2-16 h; (b) $NaBH_3CN$, MeOH, methanolic HCl, 0-25 °C, 24-72 h or 10% Pd/C, H₂ (atm), 20-25 °C, 6-12 h, MeOH; (c) $COCl_2$, CH_2Cl_2 , pyridine, 0-25 °C, 6-16 h.



Scheme 2. Synthesis of 1,3,4-oxadiazol-2-ones 152 and 153. Reagents and conditions: (a) $NH_2NHCOOR^2$ ($R^2 = Me$), acetone (for 152) and MeOH (for 153), 4 Å molecular sieves, 3-4 drops of AcOH, reflux, 2-16 h; (b) & (c) same as Scheme 1.

Table 30. List of the synthesized 1,3,4-oxadiazol-2-ones 147-170 as per Schemes 1 and 2.



 R^2 = Me for compounds **147-170**

Compd	R ¹	R ³	Compd	R^1	R ³	_
147	<i>p</i> -isobutyl	Me	159	(-)- <i>p</i> -Ph	Ме	-
148	<i>p</i> -Ph	Me	160	(+)- <i>p</i> -Ph	Me	
149	<i>m</i> -Ph	Me	161	(-)- <i>p</i> -Ph ₂ N	Me	
150	<i>p</i> -OPh	Me	162	(+)- <i>p</i> -Ph ₂ N	Me	
151	<i>m</i> -OPh	Me	163	Н	Н	
152			164	p-isobutyl	Н	
153			165	<i>p</i> -Ph	Н	
154	Н	Me	166	<i>m</i> -Ph	Н	
155	Н	Ph	167	<i>o</i> -OPh	Н	
156	<i>p</i> -Ph₂N	Me	168	<i>m</i> -OPh	Н	
157	(-)-p-isobutyl	Me	169	<i>p</i> -OPh	Н	
158	(+)-p-isobutyl	Me	170	p-Ph ₂ N	Н	

5.2.2 1,2,5-Thiadiazole carbamates (II)

In 2010, Helquist and colleagues reported 1,2,5-thiadiazole carbamates (i, Figure 13) as potent inhibitors of lysosomal acid lipase (LAL, also known as LIPA) (Rosenbaum *et al.*, 2010). They noted that 1,2,5-thiadiazole carbamate bound covalently to LAL by carbamylation of Ser of LAL with 1,2,5-thiadiazole alcohol group serving as the leaving group. Till date, numerous carbamate compounds have been reported as inhibitors of endocannabinoid hydrolases (Mor *et al.*, 2004, Li *et al.*, 2007, Long *et al.*, 2010, Bachovchin *et al.*, 2010, Chang *et al.*, 2012, Niphakis *et al.*, 2013) (for recent reviews, see (Minkkilä *et al.*, 2010, Feledziak *et al.*, 2012, Kapanda *et al.*, 2013, Blankman & Cravatt 2013). Hence, it was intended to utilize the 1,2,5-thiadiazole carbamate scaffold for the development of inhibitors of the endocannabinoid hydrolases. Since only limited structure-activity relationship (SAR) studies based on this scaffold have been reported (Rosenbaum *et al.*, 2010), it was decided to carry optimization by introducing different cyclic amines were introduced as potential leaving groups (ii, Figure 13).



Figure 13. Optimization of 1,2,5-thiadiazole carbamates.

The general synthetic procedure of 1,2,5-thiadiazole carbamate 171-204 is presented below (Scheme 3, Tables 31 & 32). The detailed synthetic procedures are described in the original publication (II).

In accordance with the procedure described in the literature, commercially available 3,4dichloro-1,2,5-thiadiazole was coupled with the appropriate secondary amine to obtain the corresponding monochloro 1,2,5-thiadiazole derivative, which was then converted to 1,2,5thiadiazole alcohol via treatment with aqueous alkali. Finally, coupling with the appropriate carbamoyl chloride gave the desired 1,2,5-thiadiazole carbamates 171-204.



Scheme 3. Synthesis of 1,2,5-thiadiazole carbamate derivatives 171-204. Reagents and conditions: (a) 110-120 °C, 2-6 h or K_2CO_3 , DMF, 100-110 °C, 6-10 h; (b) aq. NaOH or KOH, DMSO, reflux, 1-6 h; (c) dry THF, KOtBu, 0-25 °C, 16-24 h.

Table 31. List of the synthesized 1,2,5-thiadiazole carbamates 171-179 as per Scheme 3 (cyclic amine at the main core).





Table 32. List of the synthesized 1,2,5-thiadiazole carbamates 180-204 as per Scheme 3 (non-cyclic amine at the main core).



Compd	R ² R ¹	N. N.	Compd	R ¹ N,55	N. N.
180	_N,~~	SN SN	193	N _p rt	N
181	N _r s ^c	N	194	N _r r ^r	N.N.
182	N _p or	N N	195		N N N
183	N _r r ^r	N	196	N _p r ^c	N
184	Npri	N N	197		N N
185		N	198	O N _c ^s ^s	N N
186	O ₂ N	N	199		Solo N
187		N.	200		NNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN
188		N.N.	201	N _p r ^e	N Yry



5.2.3 Loratadine analogues (III)

At present, several different classes of MAGL inhibitors have been reported and of these, some piperidine/piperazine-based carbamates/ureas have been described as the most potent and selective MAGL inhibitors (see subsections 2.4.5 and 2.4.6). Compounds JZL184 (106) (Long et al., 2010), KML29 (108) (Chang et al., 2012), CK16 (110) (Kapanda et al., 2012), MJN110 (112) (Niphakis et al., 2013), 115 and 116 (Hoornaert 2008c), SAR629 (118) (Bertrand et al., 2010, Aaltonen et al., 2013), ML30 (120) (Morera et al., 2012), JJKK-048 (121) (Korhonen et al., 2014) and 122 (Korhonen et al., 2014), share common structural features such as two aromatic rings, a piperidine/piperazine linker, and a polar tail. These features are crucial in achieving good potency and selectivity towards MAGL over other possible off-targets, particularly FAAH and/or ABHD6 (Kapanda et al., 2012, Long et al., 2010, Chang et al., 2012, Minkkilä et al., 2010, Feledziak et al., 2012, Blankman & Cravatt 2013, Kapanda et al., 2013). Based on the structural similarity with the above mentioned class of MAGL inhibitors, it was decided to select the well-known histamine H₁ antagonist loratadine (205), for further optimization (Figure 14). As the carbamate functionality present in loratadine was not reactive, a small series of loratadine analogues 206-217 was synthesized with a more electrophilic carbamate/urea moiety.



Figure 14. Structural similarities between known MAGL inhibitors, such as JZP184 (106), KML29 (108) and 116, with the well-known drug loratadine (205).

The general synthetic procedure for obtaining the loratadine analogues 206-217 is presented below (Scheme 4, Table 33). Additionally, the 1,2,4-triazole urea analogue 219 of the pharmaceutical drug, cyproheptadiene (218), was also synthesized (Scheme 5). The detailed synthetic procedures are described in the original publication (III).

In agreement with the literature procedure, commercially available loratadine was converted into desloratadine by the addition of aqueous alkali. Finally, desloratadine was coupled with the appropriate hetetocyclic urea or carbamoyl chloride or isocyanate to obtain the desired loratadine urea or carbamate analogues 206, 207, 209-213, 215-217. For the synthesis of the urea analogue 208, desloratadine was first treated with triphosgene followed by in situ coupling with 1*H*-1,2,3-triazolo[4,5-*b*]pyridine. The dithiocarbamate derivative 214 was obtained by the reaction with a dithiocarbamic acid salt; prepared *in situ* via the reaction of desloratadine with CS₂ in the presence of triethylamine, with 2,4-dinitrofluorobenzene using the well-known Sanger reaction.



Scheme 4. Synthesis of loratadine analogues 206-217. Reagents and conditions: (a) Aq. KOH, EtOH, reflux, 16-18 h; (b) DIPEA, RCOR (R = 206a, 207a, 209a, 211a, respectively), DCM, 22-25 °C, 24-26 h; (c) (i) pyridine, DCM, triphosgene, 0-25 °C, 2-3 h; (ii) carbamoyl chloride from (i) followed by RH (R = 208a), DIPEA, DMAP, DCM, 0-25 °C, 24-26 h; (d) DIPEA or pyridine, RCOCI (R = 210a, 212a, 213a, 216a, 217a, respectively), DCM, 0-25 °C, 16-24 h; (e) (i) CS₂,

 Et_3N , 22-25 °C, 16 h; (ii) R-F (R = 2,4-dinitrophenyl), DMF, 22-25 °C, 24 h; (f) RNCO (R = cyclohexyl), THF, reflux, 4 h.

Table 33. List of the intermediates 206a-213a, 216a and 217a, and synthesized compounds 206-217 as per the Scheme 4.





The 1,2,4-triazole urea bearing cyproheptadine 219 was synthesized through a three-step procedure (Scheme 5). First, the commercially available cyproheptadine (218) was converted into an ethyl ester analogue by reaction with ethylchloroformate which was subsequently treated with alkali to obtain the amine. Finally, the amine was coupled with 1,1'-carbonyl-di-(1,2,4-triazole) (CDT) to produce the desired triazole urea analogue 218.



Scheme 5. Synthesis of triazole analogue of cyproheptadine 219. Reagents and conditions: Reagents and conditions: (a) Toluene, CICOOEt, reflux, 3 h; (b) Aq. KOH, EtOH, reflux, 16-18 h; (c) DIPEA, RCOR (R = 206a), DMSO, 22-25 °C, 24 h.

5.2.4 3-Benzyl-1,3,4-oxadiazol-2-ones (IV)

Several research groups including our own have described the use of the 1,3,4-oxadiazol-2one scaffold (Figure 15, general structures i and ii) in the development of serine hydrolase inhibitors (Ben Ali *et al.*, 2006, Minkkilä *et al.*, 2009b, Point *et al.*, 2012, Delorme *et al.*, 2012, Kiss *et al.*, 2011, Käsnänen *et al.*, 2013, Patel *et al.*, 2013, Savinainen *et al.*, 2014). Recently, we have shown that the removal of the methyl group at the 3-position of 1,3,4-oxadiazol-2-ones led to the identification of 3-benzyl-1,3,4-oxadiazol-2-ones (Figure 15, general structure iii) as dual FAAH-MAGL inhibitors (Patel *et al.*, 2013). While screening 1,3,4-oxadiazol-2-ones against other possible targets, it was observed that compound 163 (see Table 30, subsection 5.2.1) achieved nearly 40% inhibition of hABHD6 at 1 μ M and importantly no inhibition was observed towards either hFAAH or hMAGL at 10 μ M (Patel *et al.*, 2013). This finding prompted the design and synthesis of another series of 3-benzyl-1,3,4-oxadiazol-2-ones.



Figure 15. Advancement in 1,3,4-oxadiazol-2-ones (i-iii).

The synthetic procedures of 1,3,4-oxadiazol-2-ones (220-223, 225 and 235-247, Table 34) are similar as described in the original publication (I). Compound 224 was disclosed in our

earlier report (Savinainen *et al.*, 2014) while the detailed synthetic procedures of analogues (226-234, Table 34) is described in the original publication (IV).

The targeted naphthalene analogues (220 and 221, see Scheme 6) were synthesized using the appropriate naphthaldehyde in a similar manner as described for Scheme 1 (Note: $R^2 = Me$). The targeted 3-benzyl-1,3,4-oxadiazol-2-ones 222, 223, 225 and 235-247 (Note: $R^2 = Me$, except compounds 244 and 246 where $R^2 = Et$ and Ph, repectively, and for all compounds $R^3 = H$ in Scheme 1) were obtained through a similar procedure as described in Scheme 1 (see section 5.2.1). Nitro-containing 1,3,4-oxadiazol-2-ones (223-225, 244 and 246) obtained through Scheme 1 were reduced by 10% Pd/C to the corresponding amino derivatives (226-228, 245 and 247, respectively, see Scheme 7). Selected amino derivatives (226-228) were coupled with either acetyl chloride or benzoyl chloride to produce the desired acetamides 229-231 and benzamides 232-234, respectively.



Scheme 6. Synthesis of 1,3,4-oxadiazol-2-ones 220 and 221. Reagents and conditions: see Scheme 1.



For compounds: **223-225**, **226-228**, **229-231**, **232-234**: R² = Me; **244** & **245**: R² = Et; **246** & **247**: R² = Ph

Scheme 7. Synthesis of 1,3,4-oxadiazol-2-ones 226-234, 245 & 247. Reagents and conditions: (a) 10% Pd/C, H₂ (atm), 20-25 °C, 4-8 h, MeOH; (b) MeCOCI or PhCOCI, DCM, Et₃N, 0-25 °C, 16-24 h

Compd	R ¹	R ²	Compd	R ¹	R ²	Compd	R ¹	R ²
220			230	<i>m</i> -NHCOMe	Ме	240	<i>m</i> -Cl	Ме
221			231	<i>p</i> -NHCOMe	Ме	241	<i>m</i> -OMe	Ме
222	<i>o</i> -Ph	Me	232	o-NHCOPh	Me	242	<i>m</i> -OBn	Ме
223	o-NO ₂	Me	233	<i>m</i> -NHCOPh	Me	243	<i>p</i> -OBn	Ме
224	<i>m</i> -NO ₂	Me	234	<i>p</i> -NHCOPh	Me	244	<i>m</i> -NO ₂	Et
225	p-NO ₂	Me	235	<i>m</i> -CN	Me	245	<i>m</i> -NH ₂	Et
226	o-NH ₂	Ме	236	p-CN	Ме	246	$m-NO_2$	Ph
227	<i>m</i> -NH ₂	Ме	237	<i>m</i> -COOMe	Ме	247	<i>m</i> -NH ₂	Ph
228	p-NH ₂	Ме	238	p-COOMe	Ме			
229	o-NHCOMe	Ме	239	<i>m</i> -CF ₃	Ме			

5.3 IN VITRO BIOLOGICAL SCREENING

Substrates such as *N*-arachidonoylethanolamine (AEA), 2-arachidonoylglycerol (2-AG), and 1(3)-arachidonoylglycerol (1(3)-AG) were purchased from Cayman Chemical (Ann Arbor, MI, USA). Tritium-labelled *N*-arachidonoyl ethanolamine [ethanolamine 1-³H] (i.e. ³H-AEA) was procured from American Radiolabeled Chemicals Inc. (St Louis, MO, USA). Hydrolase inhibitors were from the following sources: 1-oxazolo[4,5-*b*]pyridin-2-yl-1-dodecanone (CAY10435), *N*-phenyl-4-(quinolin-2-ylmethyl)piperidine-1-carboxamide (PF-750, 49) were obtained from Cayman Chemicals; MAFP (2), URB597 (23), WWL70 (125) and THL (orlistat) from Sigma-Aldrich; 5-ethoxy-3-(3-phenoxyphenyl)-1,3,4-oxadiazol-2(3*H*)-one (ATM-114), JZL195 (107) and JJKK-048 (121) were synthesized in house.

5.3.1 *In vitro* assay for FAAH activity

FAAH inhibitory values of the synthesized compounds 147-217 and 219-247 described in sections 5.2.1-5.2.4 were determined using membranes of COS-7 cells expressing human recombinant FAAH (hFAAH-COS-7) which has the ability to hydrolyze tritium-labelled

Table 34. List of the disclosed 1,3,4-oxadiazol-2-ones 220-247.

anandamide [ethanolamine 1-³H] (i.e. [³H]-AEA) (Figure 16), essentially as in the previously described radioactivity-based enzymatic assay (Saario *et al.*, 2006).



Figure 16. Schematic representation of radioactivity-based FAAH assay. FAAH causes hydrolysis of [³H]-AEA to AA and radioactive ethanolamine; in the presence of an inhibitor of FAAH this process is either blocked or slowed down.

The assay buffer was 50 mM tris(hydroxymethyl)aminomethane hydrochloride (Tris-HCl) (pH 7.4); 1 mM ethylenediaminetetraacetic acid (EDTA) and the test compounds as well as reference standard/s were dissolved in DMSO (the final DMSO concentration was max 5% v/v). The incubations were performed in the presence of 0.5% (w/v) bovine serum albumin (BSA), essentially fatty acid free. Test compound or vehicle (as control, 5 µL) was preincubated with protein (1 µg, 55 µL) for 10 min at 37 °C (60 µL). The protein concentration was determined by the method of Bradford with BSA as a standard (Bradford 1976). At the 10 min time point, 20 µM AEA was added so that its final concentration was 2 µM in 100 µL (containing 10 nM of [³H]-AEA having specific activity of 60 Ci/mmol and concentration of 1 mCi/mL). The incubations proceeded for 10 min at 37 °C. Ethyl acetate (EtOAc, 400 µL) was added at the 20 min time point to stop the enzymatic reaction. Additionally, 100 µL of 50 mM Tris-HCl, pH 7.4; 1 mM EDTA was added. Samples were centrifuged at 13000 rpm for 4 min at RT, and aliquots (100 µL) from the aqueous phase containing [ethanolamine 1-³H] were measured for radioactivity by liquid scintillation counting (Wallac 1450 MicroBeta; Wallac Oy, Finland). Non-specific binding was subtracted from all the values. Those inhibitors causing more than 50% FAAH inhibition at 10 µM were tested with at least five to six different concentrations (10-9 M to 10-4 M) to obtain a dose response curve and the IC₅₀ values calculated from non-linear regressions using GraphPad Prism 5.0 for Windows.

5.3.2 In vitro assay for MAGL activity

The ability of synthesized compounds 147-217 and 219-247 described in sections 5.2.1-5.2.4 to inhibit MAGL was determined using lysates of human MAGL (hMAGL-HEK) overexpressing HEK cells which has the capability to hydrolyze 2-arachidonoylglycerol (2-AG) (Figure 17), this being assayed by the previously described HPLC-based enzymatic method (Saario *et al.*, 2004, Minkkilä *et al.*, 2008).



Figure 17. Schematic representation of HPLC-based MAGL assay. MAGL causes hydrolysis of 2-AG to AA and glycerol; in the presence of an inhibitor of MAGL this process is either blocked or slowed down.

Incubations were conducted in 50 mM Tris–HCI (pH 7.4), 1 mM EDTA with 0.5% (w/v) BSA. Test compounds were dissolved in DMSO (the final DMSO concentration was not more than 5% v/v). Test compound or vehicle (as control, 5 μ L) was preincubated with protein (2.5 μ g, 55 μ L) for 10 min at 37 °C (60 μ L). The hrMAGL concentration chosen was within the initial rate velocity for the enzyme reaction. At the 10 min time point, 2-AG (125 μ M) was added to achieve a final concentration of 50 μ M with the final incubation volume of 100 μ L. The incubations proceeded for 10 min at 37 °C. To stop the enzymatic reaction against 2-AG, 400 μ L of cold 11 mM phosphate buffer (H₃PO₄) in acetonitrile (ACN) (the pH of the samples was simultaneously decreased to 3.0) added to stabilize 2-AG against acyl migration to 1(3)-AG. All samples were centrifuged at 13000 rpm for 4 min at RT. The formation of arachidonic acid (AA) and depletion of 2-AG (and 1(3)-AG) were measured by HPLC according to the previously reported method with minor modifications (Saario *et al.*, 2004).

The analytical HPLC system consisted of an Agilent (Agilent Technologies, Santa Clara, California, USA) G1322A vacuum degasser, G1312A binary pump, G1313A autosampler, G1316A Colcom thermostatted column compartment, G1315B diode array UV/Vis detector (set at 211 nm) and 35900E interface module. The separations were accomplished on a Zorbax XDB-C18 Rapid Resolution HT column (4.6 x 50 mm, 1.8 µm) (Agilent, U.S.A). The injection volume was 20 µL. A mobile phase mixture of 34% H₃PO₄ (30 mM, pH 3.0) in ACN was used at a flow rate of 2 mL/min at 50 °C. Retention times were 2.57 min for 2-AG, 2.81 min for 1(3)-AG and 4.41 min for AA. The relative concentrations of 2-AG, 1(3)-AG and AA were determined by the corresponding peak areas. This was justified by the equivalence of response factors for the studied compounds, and was supported by the observation that the sum of the peak areas was constant throughout the experiments. Any inhibitor causing more than 50% MAGL inhibition at 10 µM was tested with at least five to six different

concentrations (10⁻⁹ M to 10⁻⁴ M) to obtain a dose response curve and the IC₅₀ values calculated from non-linear regressions using GraphPad Prism 5.0 for Windows.

5.3.3 Determination of ABHD6/12 activity

ABHD6/12 inhibitory values of the selected synthesized compounds described in sections 5.2.1-5.2.4 were determined using lysates of HEK293 cells expressing human ABHD6 (hABHD6) or human ABHD12 (hABHD12) and its ability to hydrolyze 1(3)-AG (or 2-AG) to AA and glycerol (Figure 18) where the production of the latter was quantified by a sensitive fluorometric enzymatic method according to the validated method described previously (Navia-Paldanius *et al.*, 2012, Aaltonen *et al.*, 2013).



Figure 18. Schematic representation of a sensitive fluorometric glycerol assay for ABHD6 and ABHD12. ABHD6/12 causes hydrolysis of 1(3)-AG (or 2-AG) to AA and glycerol. The glycerol production was coupled via a three-step enzymatic cascade to hydrogen peroxide (H₂O₂) dependent generation of resorufin whose fluorescence was monitored. Abbreviations: GK, glycerol kinase; G-1-P, glycerol-1-phosphate; ATP, adenosine triphosphate; GPO, glycerol phosphate oxidase; DHAP, dihydroxyacetone phosphate; HRP, horseradish peroxidase.

Incubations were conducted in the assay buffer consisting of 50 mM Tris–HCI (pH 7.4), 1 mM EDTA with 0.5% BSA (w/v, essentially fatty acid free), 5 mM MgCI₂, 100 mM NaCI. Cellular

Iysates (99 µL, 0.3 µg/well, minimum amount required to obtain a optimal signal) were preincubated for 30 min at RT with 1 µL DMSO (control) or test compound. Thereafter, the glycerol assay mix containing 1(3)-AG or 2-AG (100 µL, 12.5 µM final concentration) was added and glycerol production monitored kinetically for 90 min at RT. Glycerol is converted to glycerol-1-phosphate (G-1-P) by glycerol kinase (GK) in the presence of adenosine triphosphate (ATP). Glycerol phosphate oxidase (GPO)-catalyzed oxidation of G-1-P generates H₂O₂ which in the presence of horseradish peroxidase (HRP) converts AmplifuTM Red into the fluorescent product resorufin. Resorufin fluorescence (λ_{ex} 530; λ_{em} 590 nm) was kinetically monitored using a Tecan Infinite M200 plate reader (Tecan Group Ltd., Männedorf, Switzerland). The assays routinely contained 0.5% (w/v) BSA (essentially fatty acid free) as a carrier for lipophilic compounds. 1-AG was used instead of 2-AG, as this is the preferred endocannabinoid isomer for hABHD6 and hABHD12 (Navia-Paldanius et al., 2012). The IC₅₀ values at the time-point 90 min were calculated after nonlinear fitting of the inhibitor dose-response curves. To monitor assay performance, assay blanks without enzyme, cellular background (HEK293 cell lysates) as well as a glycerol standard were included for each tested substrate. Fluorescence of the assay blank was subtracted before calculation of the final results. Inhibitors causing more than 50% ABHD6/12 inhibition at 1 or 10 µM were tested with at least four to five different concentrations (10⁻⁹ M to 10⁻⁴ M) to obtain dose response curve and the IC₅₀ values calculated from non-linear regressions using GraphPad Prism 5.0 for Windows.

5.3.4 CB1 and CB2 receptor assays

Cannabinoid receptor-mediated G-protein activity of the selected compounds described in sections 5.2.1-5.2.4 was measured according to earlier reported methods (Savinainen *et al.*, 2003, Savinainen *et al.*, 2005). CB₁ receptor-dependent activity was determined in rat cerebellar membranes and CB2 receptor-dependent activity in membranes prepared from CHO cells stably transfected with the human CB2 receptor using guanosine 5'-O-(3-[³⁵S]-thio)-triphosphate ([³⁵S]-GTP_YS) membrane-binding assay.

5.3.5 FAAH reversibility studies and kinetics

To determine the reversibility of FAAH inhibition by enantiomers 157, 159 and 161 (see section 5.2.1), the effect of preincubation time and the recovery of enzymatic activity after a rapid and large dilution (500-fold) of enzyme-inhibitor complex was studied, as described previously (Minkkilä *et al.*, 2009a). In the pre-incubation mixture, the amount of hFAAH-COS-7 cell membranes was 20-fold higher than the amount required for the activity assay, and the inhibitor concentration was 20-fold higher than the IC₅₀ value (i.e. 11 nM (157), 23 nM (159), 98 nM (161)), and the concentrations of selected reference inhibitors were as follows, 17 nM for CAY10435, an established reversible inhibitor (Leung *et al.*, 2003), and 321

nM for PF-750 (49), a known irreversible inhibitor (Ahn *et al.*, 2007). hFAAH-COS-7 cell membranes (40 µg) were preincubated with the inhibitor (or DMSO) in 50 mM Tris-HCI (pH 7.4) and 1 mM EDTA at 37°C for 60 min. Then 1.8 µL of enzyme-inhibitor mixture was taken and immediately diluted 500-fold into the assay buffer containing 30 µM AEA (total volume 900 µL). Incubations were continued at 37°C and 100 µL samples were taken at 0,10, 20, 30, 40, 50, 60 and 70 min time points. The enzymatic reaction was stopped by adding 400 µL of cold 11 mM H₃PO₄ in ACN. Tubes were centrifuged for 4 min at RT 14000 rpm. Then 150 µL aliquots were taken and 50 µL of 30 mM H₃PO₄ (pH 3.0) was added. The relative concentration of arachidonic acid was analyzed by the HPLC method (see *in vitro* assay for MAGL activity).

Kinetic studies of enantiomers 157, 159 and 161

In the time-dependency studies, hrFAAH-COS-7 cell membranes (1 μ g, 55 μ L) were preincubated at different time points of 0, 10 or 60 minutes with an inhibitor (or DMSO, 5 μ L) in 50 mM Tris-HCI (pH 7.4) and 1 mM EDTA at 37°C for 10 minutes and then 40 μ L at the appropriate AEA dilution was added giving a final volume of 100 μ L. The incubation was continued for 10 min at 37°C. The reaction was stopped and the samples were analyzed as described in the FAAH reversibility assay.

5.3.6 MAGL reversibility studies

The reversibility of MAGL inhibition by compound 206 (see section 5.2.3) was determined in the presence of reference compounds; MAFP (2), an established irreversible inhibitor (Savinainen *et al.*, 2010) and ATM-114, a known reversible inhibitor (Savinainen *et al.*, 2010), through a 96-well plate format based fluorescent assay by quantifying the glycerol liberated from the 1(3)-AG by a sensitive fluorometric enzymatic method according to the validated method described previously (Aaltonen *et al.*, 2013). The reversibility of compounds (in volume of 0.5 µL) to inhibit hMAGL in HEK cells (in volume of 4.5 µL) was tested after a rapid 40-fold dilution of the preincubated enzyme-inhibitor complex with the substrate solution (195 µL). The inhibition of enzymatic hydrolysis and the reversibility of responses were followed at 0, 10, 30 and 60 min time-points at 25 °C. The statistically significant difference in the IC₅₀ concentrations between 10 and 60 min time-points was taken as an indication of compound reversibility whereas a stable IC₅₀ value between 0 and 60 min was considered as irreversible behaviour.

5.3.7 ABHD6 reversibility studies

The reversibility of ABHD6 inhibition by compounds 204 and 227 (see sections 5.2.2 and 5.2.4) was determined in the presence of reference compound WWL70 (125), an established irreversible inhibitor, in a 96-well plate format based fluorescent assay by quantifying

glycerol with a sensitive fluorometric enzymatic method according to the validated method described for the determination of reversibility for hMAGL (see subsection 5.3.6) (Aaltonen *et al.*, 2013).

5.3.8 Cyclooxygenase (COX) isoenzymes assay

The ability of compounds 157 and 158 (see section 5.2.1) to inhibit COX isoenzymes were determined by the earlier reported method with minor modifications (Meade *et al.*, 1993, Onnis *et al.*, 2010). Briefly, buffer (1 μ M hematin, 2 mM phenol, 5 mM EDTA, substrate [AA or 2-AG, 10 μ M final concentrations], 0.1 M Tris-HCI, pH 7.4) and test compound (in DMSO) was added to an oxygen electrode chamber (Oxygraph System, Hansatech Instruments, King's Lynn, U.K.). A baseline was established over a period of 5 min before reactions were started by the addition of COX (ovine COX-1 or human recombinant COX-2, as appropriate), and the oxygen consumption was measured in 10 second blocks. Data are presented as the change in oxygen consumption from the point of addition of the enzyme.

5.3.9 Lysosomal lipase (LAL) assay

The ability of selected synthesized compounds described in section 5.2.2 to inhibit LAL was determined using a previously described fluorescence-based enzymatic method (Figure 19) (Rosenbaum *et al.*, 2010).



4-Methylumbelliferone oleate (4MUO)

Figure 19. Schematic representation of the fluorescence-based assay for LAL. LAL causes hydrolysis of 4-methylumbelliferone oleate (4-MUO) to 4-methylumbelliferone. The generation of 4-methylumbelliferone fluorescence was monitored.

In short, purified human LAL overexpressed in Pichia pastoris (phLAL, 0.01 U/mL, 105 U/mg) was mixed with test compounds at 10 μ M and preincubated for 20 minutes at 37°C. The reaction was started by addition of 4-methylumbelliferone oleate (4-MUO), which was cleaved by enzymatic activity to 4-methylumbelliferone. The reaction was allowed to proceed for 1 h at 37°C, and enzymatic activity was quantified by 4-methylumbelliferone fluorescence (λ_{ex} 355 nm/ λ_{em} 450 nm) monitoring using a SpectraMaxM2 fluorometer (MDS

Inc., Toronto, Canada). Background fluorescence was subtracted from all the values, and results were normalized to the DMSO control value.

5.3.10 Determination of histamine H1 antagonistic activity

Compounds 206 and 219 described in section 5.2.3 were tested for histamine H₁ antagonistic effects *in vitro*, following standard methods, using the guinea pig ileum (Arunlakshana & Schild 1959). For tested compounds, concentrations up to 2 x 10^{-4} mol/dm³ were prepared using DMSO as a solvent. Further dilutions were prepared in water. Aqueous solutions were prepared for pyrilamine. The compounds were tested a minimum of two times on the two different animal preparations and at a minimum of three different concentrations.

Male guinea pigs weighing 300-400 g were sacrificed by a blow to the head. The ileum was excised and placed in phosphate buffer at room temperature (pH 7.4) containing (mM) NaCl (136.9); KCI (2.68); NaHPO₄ (7.19). After flushing the intraluminal contents, segments of about 2 cm long were cut and mounted for isotonic contractions in water jacketed 20 mL organ baths filled with oxygenated ($O_2:CO_2 = 95:5$, v/v) Krebs buffer containing (mM) NaCl (117.5); KCI (5.6); MgSO₄ (1.18); CaCl₂ (2.5); NaH₂PO₄ (1.28); NaHCO₃ (25); glucose (5.5) and indomethacin (1 x 10⁻⁶ mol/L) at 37 °C under a constant load of 0.5 g. After a 30 min equilibration period with washings every 10 mins, a sub-maximal priming dose of histamine (1 mM) was given and washed out (standard washing procedure: 3 changes of buffer during 30 mins). After washing, the antagonistic activity of given compounds was measured by recording a concentration response curve (CRC) for histamine in the presence of the tested compounds which was added 5 mins before histamine. This procedure was repeated with higher concentrations of the compounds. The antagonism was of a competitive nature causing a parallel shift of the CRC. The pA₂-values were calculated according to Arunlakshana and Schild (Arunlakshana et al., 1959). The pA₂ values were compared with the affinity of pyrilamine and loratadine.

5.3.11 Data analysis

The results from the enzyme inhibition experiments are presented as mean ± 95% confidence intervals of at least two independent experiments performed in duplicate. The results from enzyme reversibility studies are presented as mean ± S.E.M. of at least three independent experiments performed in duplicate. The inhibitor dose–response curves (DRCs) and IC₅₀ values were calculated from nonlinear regressions using GraphPad Prism 5.0 for Windows (GraphPad Software, San Diego, CA (USA): www.graphpad.com) and Matlab.

5.3.12 Activity-based protein profiling (ABPP) assay

The selected compounds described in sections 5.2.1-5.2.4 were tested using *in vitro* ABPP assay for its selectivity towards specific endocannabinoid hydrolyzing enzyme/s (i.e. FAAH, MAGL, ABHD6 and 12) over other serine hydrolases of mouse whole brain proteomes prepared from brain tissue of 4-week-old male mice (Figure 20), in accordance with the previously reported assay (Navia-Paldanius *et al.*, 2012, Aaltonen *et al.*, 2013). The ethical approval was obtained from the local welfare officer of UEF.



Figure 20. Schematic representation of a competitive ABPP of mouse brain membrane proteomes in the absence (Panel A) or presence of the test inhibitor (Panel B). The structure of the TAMRA probe also shown.

ABPP was conducted using carboxytetramethylrhodamine fluorophosphonate (TAMRA-FP; ActivX Fluorophosphonate Probes (FPs), Thermo Fisher Scientific Inc., Rockford, IL) probes. Briefly, brain membranes (100 μ g) diluted in phosphate-buffered saline (PBS) were treated for 1 h with vehicle (DMSO) or the selected inhibitors (with appropriate concentrations ranging from 0.010-10 μ M) at 25 °C. Then, serine hydrolases were labeled with 0.5 μ L of 100 μ M TAMRA-FP for 1 h at 25 °C (final probe concentration 2 μ M). The reaction was quenched by addition of 2 x gel loading buffer, after which 10-20 μ g protein was loaded per lane and the proteins were resolved in 10% SDS-PAGE together with molecular weight standards.

TAMRA-FP labeling was visualized (λ_{ex} 552; λ_{em} 575 nm) using a fluorescent scanner (FLA-3000 laser fluorescence scanner, Fujifilm, Tokyo, Japan).

5.4 SINGLE CRYSTAL X-RAY DIFFREACTION ANALYSIS

Enantiomerically pure solid compound 159 (see section 5.2.1) was crystallized by slow evaporation of its dichloromethane (DCM)/ethanol (EtOH) solution at 278 K. Subsequently, a suitable crystal was immersed in cryo-oil, mounted in a Nylon loop, and measured at a temperature of 120 °K. Data concerning the X-ray diffraction was collected on a Bruker Kappa Apex II Duo diffractometer using CuK α radiation (λ = 1.54178 Å). The APEX2 program package was used for cell refinements and data reductions. The structures were solved by direct methods using the SHELXS-97 programs with the WinGX graphical user interface. A semi-empirical absorption correction (SADABS) was applied to all data. Structural refinements were carried out using SHELXL-2013. All hydrogen atoms in enantiomer 159 were positioned geometrically and constrained to ride on their parent atoms, with C-H = 0.95-1.00 Å, and Uiso = 1.2-1.5 Ueq (parent atom).

Analysis of the absolute structure of enantiomer 159 using likelihood methods (Hooft, Straver & Spek, 2008) was performed using PLATON. The absolute structure parameter y was calculated to yield 0.06(8) indicating that the absolute structure (*S*) has probably been determined correctly. The method calculated that the probability that the structure is inverted was smaller than 4 x 10⁻²⁸. The Flack *x* parameter was refined to give a value of 0.08(9) that also corresponds to a correct absolute structure determination.

6 Results and Discussion

The *in vitro* biological evaluation of 1,3,4-oxadiazol-2-ones (147-170, 220-247), 1,2,5thiadiazole carbamates (171-247), loratadine (205) and its analogues (206-217), and cyproheptadine analogue 219 was initiated by screening the inhibitory activities against hFAAH and hMAGL at 10 μ M concentrations (see subsections 5.3.1-5.3.2 for assay protocols). Selected compounds were further screened at 1 or 10 μ M concentrations against hABHD6 and hABHD12 using the assay protocols discussed above (see subsection 5.3.3). The IC₅₀ values were calculated for those compounds having \geq 50% inhibition at the mentioned concentration against specific enzymes. The detailed results are discussed in the original publication (I-IV) and only a brief overview is presented here.

Some of the potent inhibitors found through the above screening procedures were further tested towards cannabinoid receptors (see subsection 5.3.4 for assay protocol) and other possible off-targets, such as COX isoenzymes for 1,3,4-oxadiazol-2-ones (see subsection 5.3.8 for assay protocol), LAL for 1,2,5-thiadiazole carbamates (see subsection 5.3.9 for assay protocol), histamine H₁ antagonistic affinity for loratadine analogues and cyproheptadiene analogue (see subsection 5.3.10 for assay protocols). The detailed results are discussed in the original publication (I-IV) and only outcome is revealed here. Finally, the best inhibitors in terms of potency and selectivity were tested further to determine their binding properties and/or inhibition mechanism towards specific enzyme (see subsections 5.3.5-5.3.7 for reversibility assay protocol), and selectivity over serine hydrolases of mouse brain membrane proteomes (see subsection 5.3.12). The detailed results are discussed in the original publication (I-IV) and only outcome is revealed here.

6.1 Chiral 1,3,4-oxadiazol-2-ones (I)

The inhibitory activity data of 1,3,4-oxadiazol-2-ones 147-170 tested at 10 μ M concentration towards hFAAH and hMAGL are presented in Table 35. All of these compounds achieved \leq 50% inhibition for hABHD6 and hABHD12 at 1 μ M. Hence, inhibition data towards these enzymes are not shown here, however, in the original publication (I) they were reported as supplementary information.

Table 35. Inhibitory activities of the 1,3,4-oxadiazol-2-ones 147-170 against FAAH and MAGL.



 R^2 = Me for compounds **147-170**

Compd	R ¹	R ³	p/ ₅₀ (range) $[IC_{50,} \mu M]^a$ or % inhibition at 10 μM	
			hFAAH	hMAGL
147	<i>p</i> -isobutyl	Ме	7.64 (7.63-7.65) [0.023]	48%
148	<i>p</i> -Ph	Ме	7.33 (7.32-7.33) [0.048]	34%
149	<i>m</i> -Ph	Ме	5.93 (5.92-5.94) [1.2]	42%
150	<i>p</i> -OPh	Ме	7.47 (7.45-7.49) [0.034]	5.68 (5.62-5.75) [2.0]
151	<i>m</i> -OPh	Ме	5.28 (5.25-5.32) [5.2]	14%
152			12%	6%
153			40%	NI ^c
154	Н	Ме	5.22 (5.20-5.24) [6.0]	NI
155	Н	Ph	NI	NI
156	p-Ph ₂ N	Ме	6.94 (6.92-6.96) [0.12]	6.20 (6.17-6.23) [0.63]
157 (JZP- 327A)	(-)- <i>p</i> -isobutyl	Ме	7.94 (7.91-7.97) [0.011]	16%
158	(+)-p-isobutyl	Ме	6.61 (6.57-6.66) [0.24]	4.87 (4.51-5.24) [4.0]
159	(-)- <i>p</i> -Ph	Ме	7.63 (7.60-7.66) [0.023]	18%
160	(+)- <i>p</i> -Ph	Ме	5.54 (5.50-5.58) [3.0]	49%
161	(-)- <i>p</i> -Ph ₂ N	Ме	7.0 (6.99-7.01) [0.098]	41%
162	(+)- <i>p</i> -Ph ₂ N	Ме	5.21 (5.03-5.40) [6.3]	6.47 (6.43-6.50) [0.35]
163	Н	Н	NI	NI
164	<i>p</i> -isobutyl	Н	7.05 (7.02-7.08) [0.091]	5.87 (5.86-5.89) [1.3]
165	<i>p</i> -Ph	Н	6.79 (6.78-7.79) [0.16]	5.79 (5.69-5.88) [1.6]
166	<i>m</i> -Ph	Н	46%	32%
167	<i>o</i> -OPh	Н	5.20 (5.15-5.26) [6.3]	NI
168	<i>m</i> -OPh	Н	6.01 (5.97-6.03) [0.98]	15%

169	<i>p</i> -OPh	Н	7.0 (6.89-7.14) [0.10]	6.40 (6.39-6.41) [0.40]
170	p-Ph ₂ N	Н	6.34 (6.30-7.37) [0.47]	6.60 (6.55-6.65) [0.26]
CAY10435 ^d			7.78 ± 0.09 [0.017] ^e	NA ^f
PF-750 (49) ^g			6.49 ± 0.09 [0.32] ^e	NA

^a p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean (range) from two independent experiments performed in duplicate. IC₅₀ values are calculated for those compounds having \geq 50% inhibition at the tested concentrations and are derived from the mean p*I*₅₀ values as shown in brackets. ^b The percentage (%) of inhibition is represented as the mean from two independent experiments performed in duplicates. ^c NI indicates no inhibition. ^d 1-oxazolo[4,5-*b*]pyridin-2-yl-1-dodecanone has a commercial supplier code CAY10435 and was used as a reference FAAH inhibitor as has been reported by Boger *et al.*, 2000. ^e p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean ± S.E.M. from three independent experiments performed in duplicates. ^f NA indicates not analyzed. ^g *N*-phenyl-4-(quinolin-2-yl-methyl)piperidine-1-carboxamide (PF-750, 49) was used as reference FAAH inhibitor and reported by Ahn *et al.*, 2007.

SARs of 1,3,4-oxadiazol-2-ones 147-170

As shown in Table 35, the hybridized 1,3,4-oxadiazol-2-ones of ibuprofen (147) and simplified flurbiprofen (148) showed excellent inhibitory activities against FAAH in low nanomolar range (IC₅₀ = 23 nM and 48 nM, respectively) while fenoprofen analogue (151) showed moderate inhibitory activity ($IC_{50} = 5.2 \mu M$). These compounds showed good selectivity towards FAAH since they only weakly inhibited MAGL at 10 µM concentration (48%, 34%, 14%, respectively). A similar trend in the inhibitory activities against FAAH was observed for the meta-substituted analogue of 148 (compound 149) and para-substituted analogue of 151 (compound 150) as meta-substitution (149) decreased and para-substitution (150) improved FAAH inhibition (IC₅₀ = 1.2 and 34 nM, respectively). Compound 149 showed very weak inhibition (42%) of MAGL at 10 µM concentration while compound 150 exhibited a weak inhibitory activity against MAGL (IC₅₀ = 2 μ M). The difference in the binding of compounds 149 and 150 toward FAAH and MAGL was attributable to the different substituents. The 1,3,4-oxadiazol-2-ones of acetone (152), cyclohexanone (153), acetophenone (154), and benzophenone (155) showed a dramatic loss of inhibitory activities against both FAAH and MAGL. The results of analogues 152 and 153 highlight the importance of the aromatic ring while compounds 154 and 155 emphasize the importance of proper fitting of a smaller methyl group compared to bulky phenyl ring at the 3-position of 1,3,4-oxadiazol-2ones to FAAH binding site. Compound 156 which had a bulkier hydrophobic substituent at the para-position of the phenyl ring exhibited slightly less potency (IC₅₀ = 0.12 μ M) toward FAAH. However, improvement in MAGL activity ($IC_{50} = 0.63 \mu M$) was observed.

Selected chiral 1,3,4-oxadiazol-2-ones 147, 148 and 156 were separated using a chiral column (see subsection 5.1). An improved FAAH inhibition was observed for (-)-enantiomers 157, 159 and 161, with IC₅₀ values of 11, 23 and 98 nM, respectively, compared to their racemic

analogues. At the same time, improved selectivity of these analogues over MAGL was observed. In contrast, (+)-enantiomers 158, 160 and 162 show decreased FAAH inhibition ($IC_{50} = 0.24$, 3.0 and 6.3 μ M, respectively) as compared to both racemic mixture and (-)-enantiomers though their potency to inhibit MAGL was slightly improved. For example, the IC_{50} value of the MAGL inhibition for the compound 162 was 0.35 μ M while its (-)-enantiomer 161 showed only 41% inhibition at 10 μ M concentration. Finally, the absolute configuration of the solid compound 159 was determined through the single-crystal X-ray diffraction and it was found to be an *S*-enantiomer (subsection 5.4). Since the configuration will remain the same for other (-)-enantiomers, we concluded that 157 and 161 have the same configuration i.e. *S*-configuration.

Compounds 163-170 lacking the methyl group at 3-position of 1,3,4-oxadiazol-2-one ring showed either a drastic drop or a complete loss of the inhibitory activities against FAAH and MAGL. Moreover, several of these analogues showed loss of selectivity over MAGL.

Overview towards other potential off-targets

The hybridized analogues of ibuprofen, i.e. (-) and (+)-enantiomers 157 (JZP-327A) and 158, were found to be inactive towards COX isoenzymes at 10 µM concentration. The best FAAH inhibitors, (-)-enatiomers 157 (JZP-327A) and 159 did not show any blockade or activation of cannabinoid receptors 10 µM concentration. Moreover, detailed inhibition mechanism studies of (-)-enantiomers 157 (JZP-327A), 159 and 161 suggested that they were tight binding, slowly reversible inhibitors of the hFAAH. Nonetheless, the mechanistic studies provided evidence of the noncompetitive nature of these compounds, in the molecular modeling studies they showed favorable interactions within the active site of hFAAH comprising the catalytic triad Ser²¹⁷-Ser²⁴¹-Lys¹⁴² including important hydrogen-bonding of the carbonyl oxygen to the oxyanion hole Ile²³⁸-Gly²³⁹-Gly²⁴⁰-Ser²⁴¹. Finally, when compound 157 (JZP-327A) was tested against other serine hydrolases labelled by TAMRA-FP in mouse brain membrane proteome using competitive ABPP, it exhibited a dose-dependent inhibition of FAAH, achieving complete inhibition at 1 µM. Moreover, the high selectivity of compound 157 (JZP-327A) towards FAAH was also evident as it did not show any inhibition against the other brain serine hydrolases including MAGL, ABHD6 and ABHD12 at 10 μ M concentration.

Conclusion

In short, with respect to the disclosed analogues, S-enantiomer of 3-(1-(4-isobutylphenyl)ethyl)-5-methoxy-1,3,4-oxadiazol-2(3*H*)-one (JZP-327A, 157), was found to be a potent FAAH inhibitor (IC₅₀ 11 nM) with a slowly reversible mode of inhibition. Moreover, it was found to be > 900 fold FAAH selective over the main putative off-targets, MAGL, cannabinoid receptors, and COX isoenzymes. Finally, in proteomic analysis,

compound 157 displayed high selectivity among the serine hydrolases of mouse brain membrane proteome.

6.2 1,2,5-thiadiazole carbamates (II)

The inhibitory activity data of 1,2,5-thiadiazole carbamates 171-204 screened at 1 and 10 μ M towards hABHD6 and hFAAH, respectively, are presented in Tables 36 and 37. All compounds, except compound 185 showed \leq 50% inhibition at 10 μ M concentration for hMAGL. Moreover, all of these compounds were found to be inactive toward hABHD12 at 1 μ M. Hence, inhibition data towards hMAGL and hABHD12 enzymes are not shown here, however, in original publication (II) they were reported as supplementary information.

Table 36. ABHD6 and FAAH inhibitory activities of the 1,2,5-thiadiazole carbamates 171-179 having cyclic amine at the main core.



Compd	R ¹ N.e		pI_{50} (range) $[IC_{50}, \mu M]^a$ or % inhibition at 1 μM^b	pI ₅₀ (range) [IC ₅₀ , µM] ^a or % inhibition at 10 µM ^b	
	R ^{2 , ²}	22	hABHD6	hFAAH	
171		STR.	7.28 (7.23-7.32) [0.052]	6.39 (6.29-6.49) [0.40]	
172		N N	7.07 (7.03-7.10) [0.085]	6.48 (6.41-6.55) [0.30]	
173	N rri	N.	6.58 (6.43-6.73) [0.26]	6.09 (6.01-6.18) [0.81]	
174	N.s.	N.	6.88 (6.80-6.95) [0.13]	6.25 (6.23-6.27) [0.56]	
175	N	Ph Yul Ph	41 %	5.83 (5.34-6.31) [1.47]	
176	N	^۲ رو کرو کرو	40 %	6.68 (6.51-6.84) [0.21]	

177	N	Solution Shares and Sh	15 %	6.49 (6.30-6.67)	[0.32]
178	N	n.N.	6.34 (6.22-6.45) [0.46]	7.77 (7.71-7.83)	[0.017]
179	N	N.	6.25 (6.19-6.31) [0.56]	7.51 (7.48-7.53)	[0.031]
WWL70 (125) ^c			$7.07 \pm 0.05 \ [0.085]^d$	30 %	
THL ^e			7.32 ± 0.11 [0.048]	NA ^f	
JZP-327A (157) ⁹			NI ^h	7.94 (7.91–7.97)	[0.011]

^{a,b} See footnotes of Table 35. ^c *N*-Methyl-*N*-[[3-(4-pyridinyl)phenyl]methyl]-4'-(aminocarbonyl)[1,1'biphenyl]-4-yl carbamic acid ester (WWL70, 125) used as reference ABHD6 inhibitor and reported by Li *et al.*, 2007. ^d p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean \pm S.E.M. from three independent experiments performed in duplicates and reported by Navia-Paldanius *et al.*, 2012. ^e THL, tetrahydrolipstatin (orlistat) and its ABHD6 inhibitory values are reported by Navia-Paldanius *et al.*, 2012. ^f NA indicates not analyzed. ^g *S*-(-)-3-(1-(4-isobutylphenyl)ethyl)-5-methoxy-1,3,4-oxadiazol-2(3*H*)-one (JZP-327A, 157) used as reference FAAH inhibitor reported by Patel *et al.* 2013. ^h NI indicates no inhibition.

Table 37. ABHD6 and FAAH inhibitory activities of the 1,2,5-thiadiazole carbamates 180-204 having non-cyclic amine at the main core.



180-204

Compd	R ¹ - N.خ		p1 ₅₀ ± SEM [IC ₅₀ , µM] ^a or % inhibition at 1 µM ^b	p1 ₅₀ (range) [IC ₅₀ , µM] ^c or % inhibition at 10 µM ^b
	R ² ²		hABHD6	hFAAH
180	 N ₅ 55	N N N	10 %	5.19 (5.17-5.20) [6.45]
181		222 N	7.66 ± 0.07 [0.022]	5.06 (5.05-5.07) [8.91]

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182	^		NIId	NI
102	N _r r ^e	N.		
183		N	NI	NI
184	N _p r ⁱ	N. N	6.33 ± 0.13 [0.47]	24 %
185	I I I I I I I I I I I I I I I I I I I	N	8.01 ± 0.03 [0.010]	7.20 (7.17-7.23) [0.063]
186		N N	5.90 ± 0.08 [1.25]	19 %
187		N N	5.92 ± 0.05 [1.20]	11 %
188		N.	NI	46 %
189		N.	15 %	19 %
190		N.N.	6.39 ± 0.03 [0.41]	16 %
191	F	N N	7.11 ± 0.07 [0.078]	22 %
192	F N _r r ^r	N.	7.22 ± 0.05 [0.060]	48 %
193	N _p ²	N	6.83 ± 0.04 [0.15]	21 %
194	N _r r ⁱ	N.N.	7.27 ± 0.07 [0.054]	9 %

	Ĩ. Î		17.0/	40.9/
195	N _r r ^r	NNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	17 %	40 %
196	N _e r ²	\square	6.58 ± 0.04 [0.26]	17 %
197		N N	6.71 ± 0.07 [0.19]	17 %
198	`o ₁	NNN N	11 %	18 %
	N _r rx	N		
199		N	6.04 ± 0.10 [0.91]	13 %
200		NNN NNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	NI	7 %
201	N _c ,	N.	7.36 ± 0.05 [0.044]	16 %
202		N N	7.37 ± 0.05 [0.043]	21 %
203		N	7.14 ± 0.06 [0.072]	13 %
204 (JZP- 430)	N _p , r ⁱ	N N	7.36 ± 0.05 [0.044]	18 %
	-			

^a p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean ± S.E.M. from three independent experiments performed in duplicates. IC₅₀ values are calculated for those compounds having \geq 50% inhibition at 1 µM for hABHD6 and are derived from the mean p*I*₅₀ values as shown in brackets. ^b The percentage (%) inhibition is represented as the mean from two independent experiments performed in duplicate. ^c pI₅₀ values (-log₁₀ [IC₅₀]) represent the mean (range) from two independent experiments performed in duplicate. IC₅₀ values are calculated for those compounds having \geq 50% inhibition at 10 µM for hFAAH and are derived from the mean p*I*₅₀ values as shown in brackets. ^d NI indicates no inhibition.

As shown in Table 36, 1,2,5-thiadiazole carbamates 171-174 having diverse cyclic amines at the main core were found to be slightly better inhibitors of ABHD6 compared to FAAH at the tested concentration. For example, analogue 171 showed excellent ABHD6 inhibition in the low nanomolar range (IC_{50} 52 nM) while moderate inhibitory activity was observed against FAAH (IC_{50} 0.40 μ M). With respect to the 1,2,5-thiadiazole carbamates 175-179 which had an intact piperidine at the main core and a different bulky substituent as a possible leaving group, analogues 175-177 showed moderate FAAH inhibition, while only weak inhibition of ABHD6 was observed at selected concentration. However, fused bicyclic analogues 178 and 179 showed improved FAAH inhibition (IC_{50} 0.46 and 0.56 μ M, respectively) while moderate inhibition ABHD6 was observed (IC_{50} 0.46 and 0.56 μ M, respectively). In general, several of analogues 171-179 were found to be potent ABHD6 inhibitors but no satisfactory selectivity over FAAH was observed (selectivity-ratio < 30-fold).

Subsequently, 1,2,5-thiadiazole carbamates 180-204 having non-cyclic amines at the main core were tested and the results are shown in Table 37. The *N*,*N*-dimethyl analogue 180 showed weak FAAH inhibition (IC₅₀ 6.45 μ M) while no inhibition was observed toward ABHD6. However, replacement of one methyl group of 180 with a phenyl group (compound 181) resulted in the creation of a highly potent ABHD6 inhibitor (IC₅₀ 22 nM) which had 404-fold selectivity over FAAH (IC₅₀ 8.9 μ M). However, the addition of another phenyl group into compound 181 (compound 182) resulted in a complete loss of activity. A similar result was observed with the *N*,*N*-diisopropyl analogue (compound 183) as it also showed a loss of activity. The alteration of the methyl group of compound 181 for an ethyl (compound 184) resulted in a ~20-fold drop in potency while selectivity was still evident as it possessed weak FAAH inhibition. The replacement of the phenyl ring of compound 181 into benzyl (compound 185) resulted in a 2-fold increase in ABHD6 inhibitory activity (IC₅₀ 10 nM). However, the loss of selectivity was observed as it also showed excellent FAAH inhibition (IC₅₀ 67 nM) as well as moderate MAGL inhibition (IC₅₀ 5.6 μ M).

Substitution at the *para* or *meta* position on the phenyl ring of compound 181 by different electron withdrawing groups (EWGs; i.e. compounds 186, 187, 189-192) or electron releasing groups (ERGs; i.e. compounds 193, 194, 196, 197) generally displayed a 4- to 55-fold drop of ABHD6 inhibitory activity while *ortho* substitution by either EWG (compound 188) or ERGs (compounds 195 and 198) resulted in a complete loss of activity. Although, the majority of these analogues still maintain significant selectivity over FAAH. Compound 199 which had a *meta*-phenyl substituent led to a 40-fold decline in the inhibitory activity against ABHD6. However, it still maintains marginal selectivity over FAAH. The bulky trimethyl substituted compound 200 showed complete loss of activity or selectivity were obtained with the *N*-methyl-*N*-aryl analogues 186-200, the phenyl ring of compound 181 was replaced by different cycloalkyl rings (compounds 201-203). These *N*-methyl-*N*-benzyl analogues (201-203)

resulted in approximately a 2-4-fold loss of ABHD6 inhibition, while interestingly no inhibition of FAAH was observed at 10 μ M. Finally, the lipophilicity of compound 203 (cLogP = 5.5) was reduced by replacing its piperidine ring with a morpholine ring (compound 204). Compound 204 exerted comparable ABHD6 inhibitory activity as compounds 201 and 202 but in addition it was less lipophilic (cLogP = 4.1).

Overview towards other potential off-targets

Selected potent ABHD6 inhibitors were further screened at 10 µM towards the main offtarget LAL. Briefly, cyclic analogues (171, 172, 178 and 179) were found to inhibit LAL activity almost completely while among the non-cyclic analogues (181, 191, 194 and 201-204), Nmethyl-N-aryl analogues 181, 191 and 194, and N-methyl-N-cycloalkyl analogues 201-204 were found to be weak LAL inhibitors. Interestingly, the N-methyl-N-benzyl analogue 185 achieved > 99% inhibition. The IC₅₀ values were determined for those compounds having > 50% inhibition in the original publication (II). In the ABPP assay, using mouse brain membrane proteomes, cyclic analogues (171, 172, 178 and 179) as well as non-cyclic analogues (181, 185, 191, 194 and 201) tested at 1 µM were found to inhibit ABHD6 along with inhibition of an unidentified ~30 kDa serine hydrolase as an off-target. However, noncyclic analogues 202-204 were found to inhibit ABHD6 selectively at 1 µM. Finally, due to less lipophilicity of analogue 204 (JZP-430) compared to analogues 202 and 203, it was decided to test it in the ABPP assay with different concentrations varying from 0.25-5 μ M. It was found that 204 (JZP-430) inhibited ABHD6 dose-dependently, being effective even at 0.25 µM concentration. Selective inhibition of ABHD6 was detected even at the 1 µM concentration while negligible inhibition of FAAH was obtained with the 2.5 µM concentration. At a 20-fold (5 µM) concentration, partial inhibition of FAAH was detected. In short, when tested at a concentration below 2.5 µM, JZP-430 (204) appeared to be selective for ABHD6 over the other assayed brain serine hydrolases, including FAAH, MAGL and ABHD12. Compound JZP-430 (204) was inactive towards cannabinoid receptors at 10 µM. Moreover, in the reversibility study, it was found to inhibit ABHD6 in an irreversible manner. Furthermore, in the molecular modeling studies, compound JZP-430 (204) revealed favorable interactions within the active site of hABHD6 comprising the catalytic triad Ser¹⁴⁸- His³⁰⁶-Asp²⁷⁸ including important hydrogen-bonding of the carbonyl oxygen to the oxyanion hole Met¹⁴⁹-Phe⁸⁰.

Conclusion

In short, the best compound of the series 4-morpholino-1,2,5-thiadiazol-3-yl cyclooctyl(methyl)carbamate (JZP-430, 204), potently and irreversibly inhibited hABHD6 (IC₅₀ 44 nM) and was found to be > 200-fold selective over other potential off-targets such as FAAH, LAL and cannabinoid receptors. Additionally, proteomic analysis indicated that compound 204 has good selectivity i.e. it does not inhibit the other serine hydrolases of mouse brain membrane proteome.

6.3 Loratadine analogues (III)

The inhibitory activity data of loratadine (205) and its analogues 206-217, and cyproheptadine analogue 219 screened at 10 μ M towards hMAGL and hFAAH are presented in Table 38. Selected analogues 206 and 219 were also tested at 10 μ M against hABHD6 and hABHD12, and only the outcome towards these enzymes has been described. However, in the original publication (III) they were reported in detail.

Table 38. Inhibitory activities of the loratadine (205) and its analogues 206-217, and cycproheptadine analogue 219 against MAGL and FAAH.



Compd	R	pI_{50} (range) [IC ₅₀ , μM] ^a or % inhibition at 10 μM ^b			
		hMAGL	hFAAH		
205 Loratadine		NIc	11%		
206 (JZP-361)	N-N N	7.34 (7.21-7.47) [0.046]	5.14 (5.12-5.16) [7.24]		
207	N N N	6.68 (6.66-6.69) [0.208]	22%		
208	N N	7.01 (7.01-7.02) [0.098]	30%		
209	° ^S N ∧N	NI	7%		
210	$\sim 0 $ CF ₃ CF ₃	6.19 (6.11-6.26) [0.645]	15%		



^a p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean (range) from two independent experiments performed in duplicates. IC₅₀ values were calculated for those compounds having \geq 50% inhibition at 10 µM and were derived from the mean p*I*₅₀ values as shown in brackets. ^b The percentage (%) of inhibition is represented as the mean from two independent experiments performed in duplicates. ^c NI indicates no inhibition. ^d ({4-[bis-(benzo[*a*][1,3]dioxol-5-yl)methyl]-piperidin-1-yl}(1*H*-1,2,4-triazol-1-yl)methanone) (JJKK-048, 121) is used as MAGL reference standard and reported by Aaltonen *et al.*, 2013. ^e p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean ± S.E.M. from three independent experiments performed in duplicate. ^f *S*-(-)-3-(1-(4-isobutylphenyl)ethyl)-5-methoxy-1,3,4-oxadiazol-2(3*H*)-one (JZP-327A, 157) used as reference FAAH inhibitor reported by Patel *et al.* 2013.

As shown in Table 38, loratadine (205) was found to be inactive towards both MAGL and FAAH. However, replacement of the ethoxy group present in loratadine with the 1,2,4triazole group (compound 206) resulted in potent MAGL inhibitor (IC₅₀ = 46 nM) having ~150fold selectivity over hFAAH (IC₅₀ = 7.24 μ M). Two urea analogues, triazolopyridine and benzotriazole (compounds 207 and 208) were also found to be good MAGL inhibitors although 2-4 fold less potent than compound 206. However, the imidazole urea analogue 209 was found to be inactive. The carbamate analogues, hexafluoroisopropyl (HFIP) and succinimide (210 and 211) showed weak MAGL inhibition with IC₅₀ values of 0.64 and 2.57 μ M, respectively. Compounds 212-214 which possessed a nitro substituted phenyl ring showed a loss of MAGL inhibitory activity. This result indicates that these compounds had inappropriate dimensions due to their bulky phenyl group and were incapable of fitting into the MAGL binding pocket. As expected, analogues with the less electrophilic carbonyl groups (compounds 215-217) were found to be inactive towards MAGL. Moreover, most of the analogues 207-217 did not show any appreciable FAAH inhibition. The 1,2,4-triazole urea analogue of cyproheptadine (compound 219) was found to be equally potent with compound 206 as an MAGL inhibitor and had comparable selectivity over FAAH.

Overview towards other potential targets

1,2,4-Triazole urea analogues 206 and 219 caused moderate inhibition of ABHD6 with IC₅₀ values of 1.78 and 0.75 μ M, respectively. However, they were inactive towards ABHD12 at 10 μ M. Interestingly, they showed histamine H₁ antagonistic properties (pA₂ = 6.81 and 6.50, respectively). Compound JZP-361 (206) was found to be inactive towards cannabinoid receptors at 10 μ M. Moreover, in the reversibility study, it was found to inhibit hMAGL in a slowly reversible manner. Interestingly, in the molecular modeling studies, compound 206 was found to be capable of having favorable interactions within the active site of hMAGL comprising the catalytic triad Ser¹²²- His¹⁶⁹-Asp²³⁹ including important hydrogen-bonding of the carbonyl oxygen to the oxyanion hole Met¹²³-Ala⁵¹.

Conclusion

In conclusion, a small set of analogues was synthesized and from these, (4-(8-chloro-5*H*-benzo[5,6]cyclohepta[1,2-*b*]pyridin-11(6*H*)-ylidene)piperidin-1-yl)(1*H*-1,2,4-triazol-1-yl)methanone (JZP-361, 206) potently and reversibly inhibited hMAGL (IC₅₀ 46 nM) and was found to be > 150-fold selective for inhibiting hMAGL in preference to other potential off-targets, such as FAAH and cannabinoid receptors. Moreover, compound 206 retained histamine H₁ antagonistic affinity, hence, it emerged as a novel dual-acting pharmacological tool possessing both MAGL inhibitory and antihistaminergic activities.

The inhibitory activity data of 1,3,4-oxadiazol-2-ones 220-247 screened at 1 μ M towards ABHD6 and, 10 μ M towards hFAAH and hMAGL are presented in Table 39. All compounds were found to be inactive toward hABHD12 at 1 μ M. Hence, inhibition data towards this enzyme are not shown, however, in the original publication (IV) they were reported as supplementary information.

Table 39. Inhibitory activities of the 1,3,4-oxadiazol-2-ones 163-169* and 220-247* against FAAH, MAGL and ABHD6.



Compd	mpd R ¹ R ²		pI_{50} (range) [IC ₅₀ inhibition at 10 μ	ο, μ Μ] ^a or % M ^b	p1 ₅₀ (range) [IC ₅₀ , μM] ^a or % Remaining ABHD6 activity at 1 μM ^c
			hFAAH	hMAGL	hABHD6
163	Н	Me	NI ^d	NI	61%
164	p-isobutyl	Me	7.05 (7.02-7.08) [0.091]	5.87 (5.86-5.89) [1.3]	NI
165	<i>p</i> -Ph	Me	6.79 (6.78-7.79) [0.16]	5.79 (5.69-5.88) [1.6]	NI
166	<i>m</i> -Ph	Me	46%	32%	NI
167	<i>o</i> -OPh	Ме	5.20 (5.15-5.26) [6.3]	NI	60%
168	<i>m</i> -OPh	Me	6.01 (5.97-6.03) [0.98]	15%	70%
169	<i>p</i> -OPh	Ме	7.0 (6.89-7.14) [0.10]	6.40 (6.39-6.41) [0.40]	77%
220			35%	11%	64%
221			5.40 (5.26-5.54) [3.98]	NI	NI
222	<i>o</i> -Ph	Me	23%	8%	NI
223	o-NO ₂	Ме	36%	43%	57%

224	<i>m</i> -NO ₂	Me	7%	32%	65%
225	p-NO ₂	Ме	18%	36%	NI
226	o-NH ₂	Ме	8%	4%	NI
227 (JZP-169)	<i>m</i> -NH ₂	Ме	18%	21%	$6.66 \pm 0.04 \ [0.216]^{e}$
228	p-NH ₂	Ме	34%	23%	NI
229	o-NHCOMe	Ме	11%	3%	NI
230	<i>m</i> -NHCOMe	Ме	NI	12%	NI
231	p-NHCOMe	Ме	37%	16%	NI
232	o-NHCOPh	Ме	NI	NI	NI
233	<i>m</i> -NHCOPh	Ме	NI	48%	79%
234	<i>p</i> -NHCOPh	Me	6.21 (6.10-6.32) [0.62]	6.25 (6.21-6.29) [0.56]	88%
235	<i>m</i> -CN	Me	NI	23%	63%
236	p-CN	Me	NI	25%	NI
237	<i>m</i> -COOMe	Ме	20%	7%	6.35 (6.24-6.47) [0.45]
238	p-COOMe	Ме	5.17 (5.14-5.20) [6.76]	15%	NI
239	<i>m</i> -CF ₃	Ме	12%	48%	6.06 (5.94-6.19) [0.87]
240	<i>m</i> -Cl	Ме	37%	43%	52%
241	<i>m</i> -OMe	Ме	NI	6%	70%
242	<i>m</i> -OBn	Ме	5.82 (5.80-5.84) [1.51]	5.49 (5.48-5.50) [3.23]	53%
243	<i>p</i> -OBn	Me	6.39 (6.37-6.41) [0.41]	6.28 (6.25-6.31) [0.52]	NI
244	m-NO ₂	Et	NI	30%	63%
245	<i>m</i> -NH ₂	Et	10%	15%	NI
246	<i>m</i> -NO ₂	Ph	38%	5.48 (5.47-5.49) [3.31]	6.88 (6.7-6.75) [0.13]
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247	<i>m</i> -NH ₂	Ph	25%	5.1 (5.0-5.1) [7.94]	7.13 ± 0.05 [0.073] ^e
JZP-327A (157) ^f			7.94 (7.91-7.97) [0.011]	16%	NI
JZP-430 (204) ^g			18%	NI	$7.36 \pm 0.05 \ [0.044]^{e}$
JZP-361 (206) ^h			5.14 (5.12-5.16) [7.24]	7.34 (7.21-7.47) [0.046]	5.75 (5.66-5.84) [1.78]

^{*} Compounds 163-169 are from the section 6.1 and are reported in original publication (I) and included here to provide a better clarification of SARs while compound 224 was reported in our published work (Savinainen *et al.*, 2014). ^a p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean (range) from two independent experiments performed in duplicates. IC₅₀ values were calculated for those compounds causing \geq 50% inhibition at the stated concentration and were derived from the mean p*I*₅₀ values as shown in brackets. ^b The percentage (%) of inhibition is calculated from the mean of two independent experiments performed in duplicates. ^c Remaining activity (%) at 1 µM is given when it is <90%.^d NI indicates no inhibition. ^e p*I*₅₀ values (-log₁₀ [IC₅₀]) represent the mean ± S.E.M. from three independent experiments performed in duplicates. ^f *S*-(-)-3-(1-(4-isobutylphenyl)ethyl)-5-methoxy-1,3,4-oxadiazol-2(3*H*)-one (JZP-327A, 157) was used as reference FAAH inhibitor as reported by Patel *et al.* 2013. ^g 4-morpholino-1,2,5-thiadiazol-3-yl cyclooctyl(methyl)carbamate (JZP-430, 204) was used as reference ABHD6 inhibitor as previously reported (Patel *et al.*, 2015b). ^h (4-(8-chloro-5*H*-benzo[5,6]cyclohepta[1,2-*b*]pyridin-11(*6H*)-ylidene)piperidin-1yl)(1*H*-1,2,4-triazol-1-yl)methanone (JZP-361, 206) was used as a reference MAGL inhibitor as reported previously (Patel *et al.*, 2015a).

SARs of 1,3,4-oxadiazol-2-ones 220-247

As shown in Table 39, among the analogues 163-169 and 222, the unsubstituted analogue 163 was inactive towards both FAAH and MAGL. However, at 1 μ M concentration analogue 163 exhibited nearly 40% ABHD6 inhibition. Analogues with bulky substituents on the *para* position of the benzyl moiety, such as isobutyl (compound 164), Ph (compound 165), and OPh (compound 169) exhibited dual FAAH-MAGL inhibition. Shifting the substituent from *para* (compounds 165 and 169) to *meta* (compounds 166 and 168) positions resulted in a drop of FAAH inhibitory activity. However, this change resulted in a complete loss of MAGL inhibition. Similarly shifting of *para* substituents (compounds 165 and 169) to the *ortho* (compounds 167 and 222) position led to very weak or even complete loss of FAAH and/or MAGL inhibition. Of the bicyclic analogues, the 1-naphthyl analogue 220 was determined to be inactive towards FAAH while the 2-naphthyl analogue 221 (IC₅₀ 3.98 μ M) was a weak FAAH inhibitor. However, both of these analogues were found to be inactive towards MAGL while only 1-naphthyl analogue weakly inhibited ABHD6 at 1 μ M concentration.

Analogues with a smaller EWG nitro and ERG amino substituents (compounds 223-225 and 226-228, respectively) at different positions on the benzyl moiety were found inactive towards FAAH and MAGL. However, in these analogues, only the *meta*-amino analogue 227

showed significant ABHD6 inhibition with an IC₅₀ value of 216 nM. Protection of the amino group in compounds 226-228 with an acetyl (compounds 229-231) achieved no improvement in FAAH and MAGL inhibition, and loss of ABHD6 inhibition by meta substituted compound 230 revealed the importance of the free amino of compound 227. On the contrary, protection with the bulky benzoyl group of compounds 226-228 (compounds 232-234, repectively) resulted in enhanced FAAH and MAGL inhibition for para substituted analogue 234 over its ortho (compound 232) and meta (compound 233) counterparts, a consistent outcome previously observed with para bulky substituted compounds 165 and 169. The majority of the analogues having either smaller EWGs at either *meta* (compounds 235, 237, 239 and 240) or para positions (compound 236) or smaller ERG at the meta position (compound 241) were found to be inactive towards FAAH and MAGL although analogue 238 which had a para substituted EWG exhibited weak FAAH inhibition. Moreover, among these analogues, only meta-substituted analogues 237 and 239 showed weak ABHD6 inhibition with IC50 values of 0.45 and 0.87 µM, respectively. As expected, analogues 242 and 243 which had bulky benzyloxy substituents at their *meta* and *para* position respectively, resulted in an improved FAAH and MAGL inhibition.

Replacement of the methoxy group present at the 5-position of 1,3,4-oxadiazol-2-ones in compounds 224 and 227 with an ethoxy group (compounds 244 and 245) resulted in the loss of inhibitory activities towards all studied enzymes, the sole exception being compound 244 which at 1 μ M concentration weakly inhibited ABHD6. However, when a phenoxy group was incorporated (compounds 246 and 247), it was found that the compounds exhibited weak inhibition toward FAAH and moderate inhibition towards MAGL while excellent inhibition towards ABHD6 was detected i.e. IC₅₀ values of 0.13 and 0.071 μ M, respectively. Overall, majority of analogues with a bulky or smaller *meta* and *ortho* substituent displayed weak to noticeable ABHD6 inhibition.

Overview towards other potential off-targets

In the ABPP assay, using mouse brain membrane proteomes, compound 227 was found to inhibit ABHD6 selectively at 10 μ M without affecting the other brain serine hydrolases, including FAAH, MAGL and ABHD12. However, in the same assay, compound 247 targeted ABHD6 in a non-selective manner as two other serine hydrolases were found as an off-targets at 10 μ M. Compound 227 caused no blockade or activation of cannabinoid receptors at 10 μ M. The detail inhibition mechanism study of compound 227 (JZP-169) suggested that it bound to hABHD6 in an irreversible manner and the molecular modeling studies suggested favorable interactions with the active site of hABHD6 which contains the catalytic triad Ser¹⁴⁸-His³⁰⁶-Asp²⁷⁸ and where there is important hydrogen-bonding of the carbonyl oxygen to the oxyanion hole Met¹⁴⁹-Phe⁸⁰.

In summary, many analogues were prepared and of these, 3-(3-aminobenzyl)-5-methoxy-1,3,4-oxadiazol-2(3*H*)-one (JZP-169, 227) was found to be a potent (IC₅₀ 216 nM) and irreversible inhibitor of ABHD6. Moreover, it displayed no inhibition towards MAGL, FAAH, ABHD12, nor did it bind to cannabinoid receptors. Furthermore, in proteomic analysis, compound 227 inhibited only ABHD6 band without affecting the other serine hydrolases of mouse brain membrane proteomes.

7 General Discussion

7.1 SUMMARY OF THE DEVELOPMENT OF POTENT AND SELECTIVE INHIBITORS OF ENZYMES INVOLVED IN ENDOCANNABINOID INACTIVATION

Inhibition of FAAH has been shown to have therapeutic potential in several disorders such as pain, inflammation, anxiety and depression. The majority of FAAH inhibitors have been designed through structural modifications of other serine hydrolase inhibitors. In general, these compounds share a reactive carbonyl which undergoes a nucleophilic attack by the nucleophilic Ser²⁴¹ residue of FAAH. However, one recent report revealed that it is also possible to target an additional active site residue Cys²⁶⁹ of FAAH through which one can prolong FAAH inhibition and in turn extend the duration of the pharmacological effect. Moreover, some promising FAAH inhibitors even lack electrophile. At present, several chemical tools are available with which to study beneficial and detrimental effects of pharmacological modulation of FAAH. It is clear that much more work will be needed before it will be possible to transfer any potential FAAH inhibitor from the preclinical to the clinical stages. With respect to clinical utility, one major challenge for FAAH inhibitor design is to optimize the following requirements i.e. excellent potency, selectivity over other potential off-targets, reversibility or irreversibility, and efficacy toward human rather than non-human FAAH. However these hurdles can be overcome and a few FAAH inhibitors are currently undergoing clinical trials. Moreover, recent published reports have highlighted the importance of targeting FAAH simultaneously with other targets such as COX, TRPV, etc. to achieve desired effects with reduced side effects.

In contrast to FAAH inhibitors, the utility of MAGL inhibitors has only been validated in preclinical studies. Inhibition of MAGL is considered as a promising approach to treat pain, inflammation, vomiting, nausea and cancer. Some of the early developed inhibitors have several sulfhydryl-reacting compounds mainly targeting the cysteine residues (Cys²⁰¹, Cys²⁰⁸, and Cys²⁴²) located close to the catalytic site of MAGL. In recent years, a few potent and selective inhibitors targeting nucleophilic Ser¹²² of MAGL have been developed. It may be possible to avoid the undesired side effects associated with chronic treatment with irreversible MAGL inhibitors, only a limited number of potent and selective ABHD6 inhibitors have been identified while the search for the potent and selective inhibitors have been identified while the search for the potent and selective inhibitors have therapeutic potential in the treatment of neuroinflammation, epilepsy and metabolic disorders.

The present study was aimed at the development of potent and selective inhibitors of the endocannabinoid hydrolases. Previously known scaffolds were optimized to obtain inhibitors which not only blocked the target enzymes with improved potency but also possessed good selectivity with respect to the other off-targets. The research results of this study are discussed below based on the main objectives listed in section 4.

In the first subproject, the earlier disclosed potent but non-selective 1,3,4-oxadiazol-2-ones were hybridized with derivatives of phenylalkanoic acid present in marketed NSAIDs to develop novel chiral 1,3,4-oxadiazol-2-ones. The best compound identified in this series, JZP-327A, was able to inhibit FAAH in a reversible manner with IC₅₀ values in the nanomolar range as well as demonstrating high selectivity over other studied potential off-targets (other ECS members, COX-isoenzymes, and serine hydrolases of mouse brain membrane proteomes). Hence, JZP-327A is an important lead which should be tested in animal models to confirm its *in vivo* efficacy and to elucidate its PK profile.

The emerging therapeutic potential of ABHD6 in the treatment of inflammation, epilepsy and metabolic disorders, was the reason to develop inhibitors of ABHD6 as described in the second subproject. Only limited SARs had been carried out on the recently reported 1,2,5-thiadiazole carbamate-based LAL inhibitors. Hence, it was decided to develop novel ABHD6 inhibitors by undertaking systematic structural variations around the 1,2,5-thiadiazole carbamate scaffold. Many compounds were produced and of these, JZP-430 was identified as a potent ABHD6 inhibitor with an IC₅₀ value of 44 nM while displaying a good selectivity profile over the other studied potential targets such as other members of ECS, LAL and mouse brain membrane proteomes. Therefore, JZP-430 is an important lead with which to clarify the effects of ABHD6 inhibition in animal models.

The third subproject was based on structural similarity of piperidine/piperazine- based carbamate/urea inhibitors of MAGL with the well-known histamine H₁ antagonist, loratadine. A small set of carbamate/urea analogues was produced having a polar/electrophilic carbonyl group, and from these, the 1,2,4-triazole-based loratadine analogue, JZP-361, was discovered to be a potent MAGL inhibitor with an IC₅₀ value of 46 nM and it was observed to be selective over FAAH and cannabinoid receptors. In contrast to the earlier reported 1,2,4-triazole containing irreversible inhibitor JJKK-048, compound JZP-361 was found to inhibit MAGL in a slow reversible manner. Moreover, JZP-361 retained its histamine H₁ antagonistic affinity, making it an interesting dual-acting pharmacological tool to study the consequences of concomitant blockade of MAGL and histamine H₁ receptor activities.

The fourth subproject involved the further optimization of 1,3,4-oxadiazol-2-ones as ABHD6 inhibitors. The introduction of different substitutions on the benzyl moiety of 1,3,4-oxadiazol-2-ones led to the identification of JZP-169 as a potent ABHD6 inhibitor (IC₅₀ = 216 nM) while also displaying a good selectivity profile over the studied off-targets such as other members of ECS and serine hydrolases of mouse brain membrane proteomes. In contrast to reversible inhibition of FAAH achieved by the 1,3,4-oxadiazol-2-one-based compound JZP-327A, this new compound, i.e. JZP-169, inhibited ABHD6 in an irreversible manner.

All of the lead compounds described in this study achieved the goals set at the start of this research project. In addition to striking a good balance between potency and selectivity toward other potential off-targets, these compounds also exhibited optimal molecular weight and lipophilicity; the next step will be to investigate their PK profile and determine their *in vivo* efficacy. Clearly, further study will be needed to explore their therapeutic potential.

7.2 FUTURE PERSPECTIVES

Several developed potent and selective FAAH inhibitors have shown promising outcomes in preclinical studies and a few have even progressed to clinical trials. However, the disappointing results of PF-04457845 in phase II trials in patients suffering from osteoarthritis pain raised several questions about the clinical effectiveness of selective FAAH inhibitors. One major question still needing to be answered is whether administration of a FAAH inhibitor alone will be sufficient to achieve efficacious therapies or whether simultaneous modulation of several targets will be needed. Indeed, several promising results have been reported, at least in animal models, with dual FAAH-COX inhibitors and with a dual FAAH-TRPV1 blocker as well as with a compound having dual action as FAAH inhibitor/TRPA1 agonist. It is possible that irreversible MAGL inhibitors may cause excessive blockade of the MAGL leading to clinically unacceptable side effects. However, one recent study indicated that it may be advantageous to devise reversibly acting MAGL inhibitors for situations when rapid clearance of drug is needed and thus it may be beneficial to focus on the development of these kinds of compounds in the future. Although, ABHD6 inhibitors have shown promising results in combatting the metabolic syndrome, their utility as efficacious therapies still needs to be clarified in further studies.

8 Conclusions

The general aim of the studies described in this thesis was to develop potent and selective inhibitors of FAAH, MAGL and ABHD6. This goal was achieved by conducting the design, synthesis, structural characterization, and the *in vitro* pharmacological evaluation of these novel compounds. The following conclusions can be drawn from the results of this thesis:

- 1. Chiral 1,3,4-oxadiazol-2-ones have been identified as inhibitors of FAAH. Compound JZP-327A exhibited excellent FAAH potency and selectivity over other off-targets, and can be considered as a promising compound to be studied further in various *in vivo* tests, such as in models of pain and inflammation.
- 2. The 1,2,5-thiadiazole carbamate scaffold has been successfully utilized in the development of ABHD6 inhibitors. The excellent potency towards ABHD6 and the good selectivity over the potential off-targets found for the best compound in this series, JZP-430, make it a promising candidate for further investigation.
- 3. A close stuctural analogue of loratadine, JZP-361 was identified as a potent MAGL inhibitor. Since JZP-361 also retained its histamine H₁ antagonistic affinity, it can be considered as an intriguing pharmacological tool with which to study inflammation in chronic diseases, such as asthma. Finally, the structural similarity of loratadine and JZP-361 raises the possibility that some of the other known MAGL inhibitors may also possess H₁ antagonistic properties.
- 4. The 1,3,4-oxadiazol-2-one-based compound, JZP-169, was identified as a potent and selective ABHD6 inhibitor displaying good selectivity toward other off-targets. Furthermore, its irreversible inhibition of ABHD6 highlights that 1,3,4-oxadiazol-2-one is a versatile scaffold for the development of serine hydrolase inhibitors. Additionally, 1,3,4-oxadiazol-2-one scaffold constitutes a novel class of ABHD6 inhibitors.

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JAYENDRA Z. PATEL Development of Potent and Selective Inhibitors of Enzymes Involved in Endocannabinoid Inactivation



In recent years, inhibitors of FAAH, MAGL and ABHD6 have shown potential to treat several diseases, such as pain, inflammation, CNS disorders and metabolic disorders without association of cannabimimetic side effects. This doctoral thesis reports identification of novel 1,3,4-oxadiazol-2-one, 1,2,5-thiadiazole carbamate, and loratadine analogues as potent and selective inhibitors of FAAH, ABHD6, and MAGL, respectively. Moreover, 1,3,4-oxadiazol-2-one scaffold was successfully utilized in the development of ABHD6 inhibitors.



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