

Inhibitors of the Endocannabinoid-Degrading Enzymes, or how to Increase Endocannabinoid's Activity by Preventing their Hydrolysis

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Abstract: Endocannabinoids are lipid transmitters binding and activating the cannabinoid receptors. Both cannabinoid receptors and endocannabinoids, such as 2-arachidonoylglycerol and anandamide, have been shown to control numerous physiological and pathological processes, including in the central nervous system. Thus regulating endocannabinoid levels *in-vivo* represents an interesting therapeutic perspective in several CNS-related diseases. To date four enzymes - Fatty Acid Amide Hydrolase (FAAH), *N*-Acylethanolamine-hydrolyzing Acid Amidase (NAAA), Monoacylglycerol Lipase (MAGL), α/β -Hydrolase Domain 6 (ABHD6) – were shown to control endocannabinoid levels in tissues or in intact cells. While the searches for NAAA and ABHD6 inhibitors are still in their beginning, a growing number of selective and potent inhibitors are now available to inhibit FAAH and MAGL activities. Here, based on the literature and patent literature, we review the compounds of the different chemical families that have been developed to inhibit these enzymes, with a special emphasis on FAAH and MAGL inhibitors.

Keywords: Fatty acid amide hydrolase, FAAH, inhibitors, inflammation, monoacylglycerol lipase, MAGL, *N*-acylethanolamine-hydrolyzing acid amidase, palmitoylethanolamide, ABHD12, ABHD6.

I. INTRODUCTION

The effects of cannabinoids (from natural, synthetic and endogenous origin) are mainly mediated by two G protein-coupled receptors, the cannabinoid receptors CB₁ and CB₂. These two receptors are activated by endogenous bioactive lipids named endocannabinoids, which are produced in an activity-dependent manner (i.e. following cell stimulation) from phospholipid precursors present in the cell membranes. So far, two types of endocannabinoids, based on an arachidonic acid moiety, have been fully characterized. Indeed, *N*-arachidonylethanolamine (anandamide, AEA) is a member of the large family of *N*-acylethanolamines, whereas 2-arachidonoylglycerol (2-AG) is an acylglycerol. The activity of these lipid mediators at the cannabinoid receptors is terminated essentially following their hydrolysis by several lipases [1]. Initially, Fatty Acid Amide Hydrolase (FAAH) [2] and Monoacylglycerol Lipase (MAGL) [3, 4], were described as the main enzymes regulating the activity of AEA and 2-AG, respectively. More recently, additional endocannabinoid-degrading enzymes were also shown to have key roles in the endocannabinoid system. Thus, a second FAAH-related mechanism was recently described for essentially regulating oleamide.[5] Although it shares only 20 % homology with FAAH, this novel enzyme, only found in humans, was called FAAH-2. Two serine hydrolases,

α/β -hydrolase 6 (ABHD6) and α/β -hydrolase 12 (ABHD12) [6], were also recently discovered and described as complementary 2-AG-degrading enzymes in the brain [7]. Interestingly, MAGL, ABHD6 and ABHD12 are present in different subcellular locations suggesting distinct roles in controlling 2-AG activities. In addition, another enzyme, called *N*-Acylethanolamine-hydrolyzing Acid Amidase (NAAA), was found to regulate the levels of *N*-acylethanolamines [8].

Because the activation of the cannabinoid receptors results in multiple beneficial effects, numerous CB₁ and CB₂ agonists are being described since the characterization of Δ^9 -THC structure and of its activity at cannabinoid receptors. However direct and constant activation of the receptors resulting from this strategy presents several drawbacks, including receptor desensitization, and numerous CNS-related side effects for the CB₁ receptor agonists. Conversely, increasing selectively the levels of an endocannabinoid is expected to result in a subset of the effects induced by the agonist but with more limited side effects [9]. Therefore there is a strong rationale for the preparation of potent and selective inhibitors of endocannabinoid degradation.

Thus, following a brief summary of the enzymes' characteristics, we will review the different classes of inhibitors described in the patent literature. The main focus will be on the compounds able to inhibit FAAH and MAGL since those are the primary enzymes controlling the endocannabinoid levels. However, we will also briefly describe the novel inhibitors developed to block the activity of NAAA.

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II. FATTY ACID AMIDE HYDROLASE

II.1. Pharmacology of FAAH Inhibition or why Inhibiting FAAH Activity?

FAAH is a membrane-bound serine hydrolase which belongs to a distinct class of enzymes characterized by the amidase signature (AS). It possesses an atypical catalytic triad consisting in Ser-Ser-Lys (instead of the classical Ser-His-Asp) which confers to FAAH the ability to hydrolyse amide bonds of various endogenous bioactive lipids [10, 11].

Essentially known for being the main anandamide-degrading enzyme [12], FAAH also hydrolyses and thus regulates the endogenous levels of other bioactive lipids (Fig. (1)). Indeed, several *N*-acylethanolamines (NAEs) - including *N*-palmitoylethanolamine (PEA), which does not activate CB receptors but induces anti-inflammatory responses *via* the PPAR receptors, and the satiating agent *N*-oleoylethanol-amine (OEA) - also undergo a FAAH-dependent catabolism [13, 14]. Furthermore, the levels of other classes of amide-derived lipids, like the *N*-acyl taurines (NATs), which activate transient receptor potential (TRP) ions channels [15], and fatty acid primary amides (FAPAs) such as the sleep-inducing lipid oleamide [12, 16, 17], are also regulated by FAAH. Interestingly, while the levels of these mediators are increased upon FAAH inhibition the levels of two other bioactive lipids, *N*-arachidonoyldopamine (a TRPV1 agonist) [18] and *N*-arachidonoylglycine (the putative GPR18 receptor endogenous agonist) [19], are decreased following FAAH inhibition.

Considering all its different substrates, FAAH inhibition will result in numerous effects, with several of them not me-

diated by anandamide or by cannabinoid receptors. These non-CB1 and non-CB2 effects can be mediated by G protein-coupled receptors (e.g. GPR18, GPR119), ion channels (e.g. TRPV1) or nuclear receptors (e.g. PPAR) [20, 21]. To date, only the consequences of increasing the levels of the *N*-acylethanolamines are relatively well characterized both in cells and *in-vivo*. Anandamide was shown to be involved in number of physiological processes [22-25], including appetite regulation [26], pain [27, 28] and inflammation, but also various CNS and psychiatric disorders [29], with anxiety and depressive disorders being the most studied. Concerning other substrates, oleamide was shown to be involved in sleep induction [30, 31], PEA is widely described as analgesic and anti-inflammatory molecule [27, 28, 32] and OEA as satiating factor and as analgesic [33]. OEA regulates feeding and body weight through activation of the nuclear receptor PPAR-alpha [34]. Today, FAAH inhibition is generally considered as a sound therapeutic strategy in the treatment of pain and inflammation [20, 35-39] as well as anxiety and depression [40-42].

II.2. FAAH, Structure and Function

Soon after the discovery of FAAH, X-ray crystallographic studies were performed to further understand its mode of action, but also to improve the research and the development of its inhibitors. Thus the rat isoform (*r*FAAH) was crystallised in presence of methylarachidonoylfluorophosphonate (MAFP), an irreversible and non selective inhibitor, and the structure was solved with a 2.8 Å resolution [43]. This three dimensional structure revealed the presence of several domains implicated in distinct functions. *i*) A large

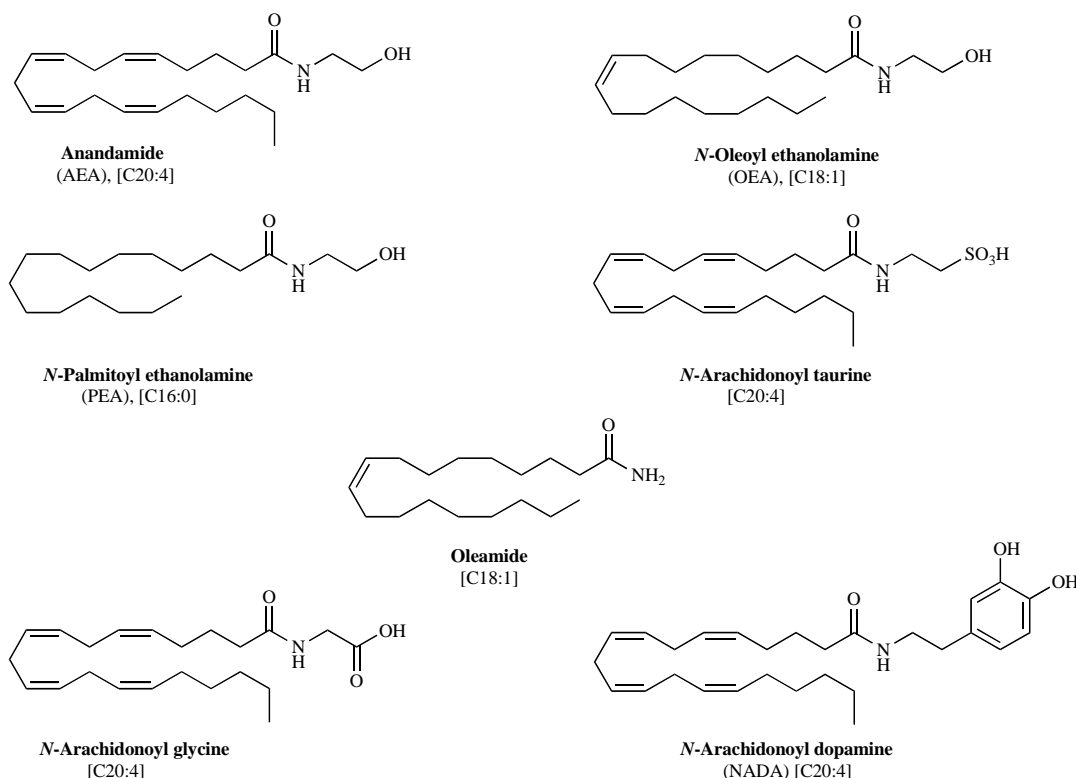


Fig. (1). Endogenous bioactive lipids regulated by FAAH.

domain composed by hydrophobic and basic residues covers the active site and allows the enzyme to anchor to the membrane. *ii*) Near this hydrophobic plateau, lies a channel responsible for the entry of the substrate. Commonly called the membrane access channel (MAC), this cavity allows a direct access for the lipid substrate from the membrane to the active site. *iii*) Close to the active site, a hydrophobic cavity is present. This acyl-chain binding pocket (ABP) interacts with the side-chain of the substrate. *iv*) Finally, the cytosolic port (CP) was found to interact with the polar head of the substrate and is connected with the cytosol. Moreover, the active site is able to accommodate a water molecule to hydrolyse the acyl-enzyme complex. As a result, the hydrophobic moiety and the hydrophilic one are released towards the MAC and CP, respectively. (For complete reviews, see [44, 45]) Further helping the drug development of FAAH inhibitors, an engineered "humanised" rat FAAH (*h/r*FAAH) [46] was produced and successively co-crystallised with three compounds representative of the major FAAH inhibitors classes [47-50]. This constitutes an interesting tool, as a large number of FAAH inhibitors present differences in activity depending on the origin of the enzyme, that is mouse FAAH (*m*FAAH) or human FAAH (*h*FAAH). An alternative strategy was to develop an homology model of *h*FAAH based on the reported X-ray structure of rat FAAH (*r*FAAH) [51]. This model was validated by docking the selective inhibitor PF-750 resulting in similar interactions to those found in the co-crystal structure of PF-750 into *h/r*FAAH.

A large number of FAAH inhibitors have been described over the years, starting from natural substrates analogues to well-adapted traditional types of serine hydrolase inhibitors [52-55]. Indeed, a wide variety of electrophilic functions has been used to target the enzyme's active site, generating large sets of structure-activity relationships aiming at improving, not only the potency, but also the selectivity of the inhibitors.

Among the numerous templates described in the patent literature (Fig. (2)), three main chemical families have been extensively studied. Below, we will review them in their order of development, starting from the α -keto heterocycles [56], then the carbamate-based inhibitors [57] and finally, the urea-derived inhibitors [58].

II.3. FAAH Inhibitors

II.3.1. α -keto Heterocycle-based FAAH Inhibitors

Activated ketones were disclosed very early in the development of FAAH inhibitors. Disclosed for inhibiting serine proteases [59, 60], a first series was described by Dale Boger's group, who generated a series of arachidonoyl- and oleoyl-based α -keto heterocycles [56]. The inhibition is based on the attack by FAAH's active serine on the electrophilic carbonyl of the inhibitor. The resulting reversible tetrahedral intermediate was recently observed in OL-135 – *h/r*FAAH co-crystals [48]. After studying a large range of heterocycles, α -keto oxazoles and α -keto oxazolopyridines were identified as the most efficacious moieties. Therefore, Boger's group refined its inhibitors and reported more potent heterocyclic inhibitors exemplified with OL-135 (Fig. (3), **1**, $IC_{50} = 2.1$ nM and > 100 μ M on *m*FAAH and on *m*MAGL, respectively), a potent and highly selective pharmacological tool commonly used and based on the pyridyl oxazole template.

From this study, an optimal C6 linker length with a phenyl end-group was chosen to replace the fatty acid chain. Beside the acyl chain, the authors also explored the impact of the heterocycle nature. Thereby, analogues were synthesised, based on 2-keto-1,3,4-oxadiazole scaffold with a subnanomolar activity (**2**, Fig. (3), $K_i = 290$ pM on *r*FAAH) [61] and more recently, 2-keto-1,2,4-oxadiazole scaffold (**7**, $K_i = 920$ pM and 340 pM on *h*FAAH and *r*FAAH, respectively)

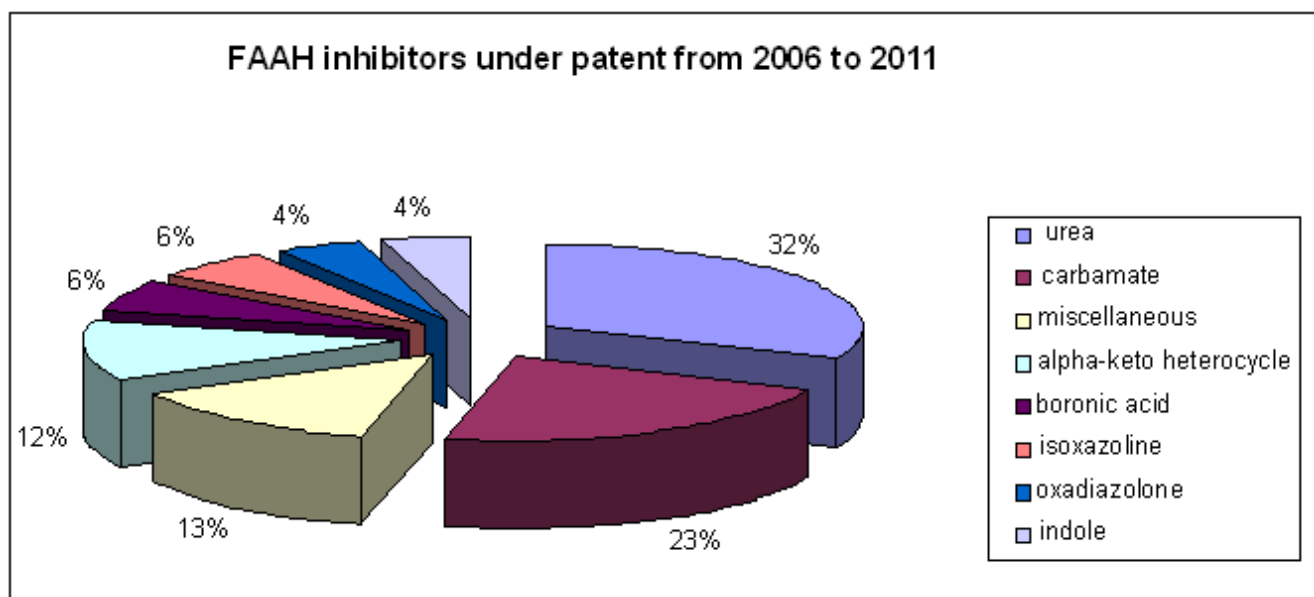


Fig. (2). Pie-chart, based on the chemical family, of the FAAH inhibitors families found in the patent literature (2006-2011).

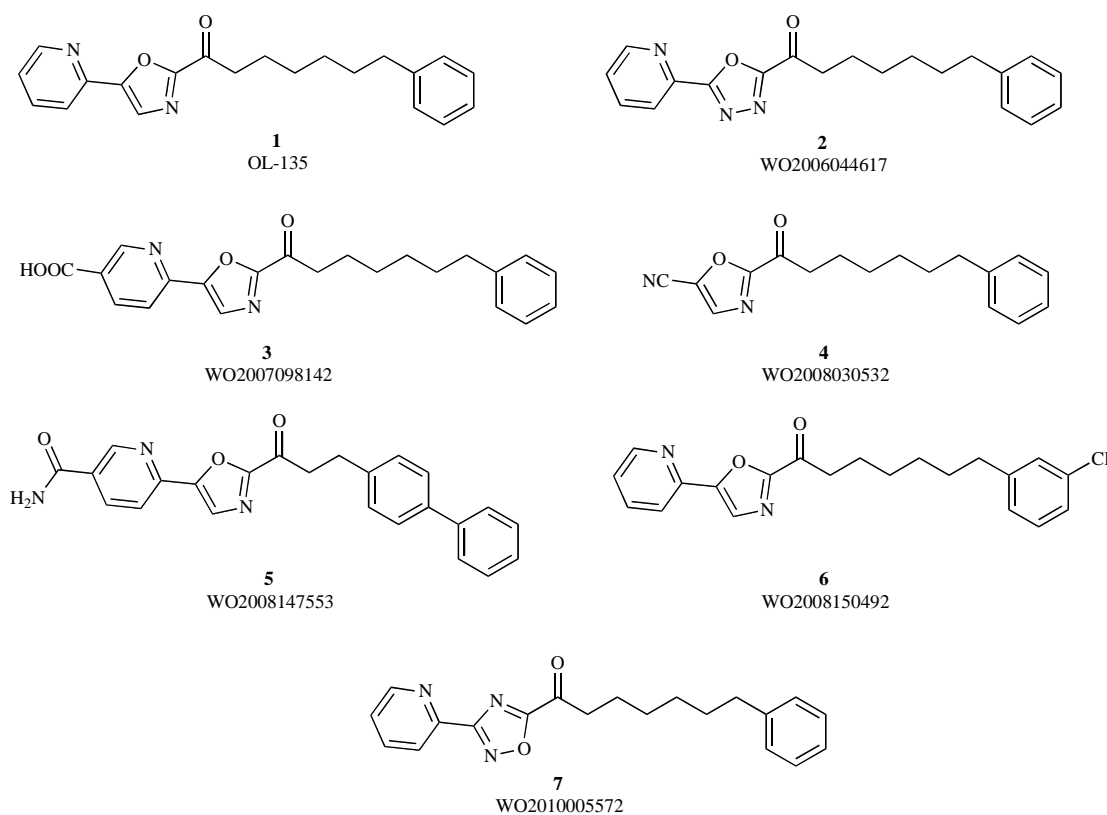


Fig. (3). FAAH inhibitors based on α -keto heterocycle templates described by Boger *et al.*

[62]. Similarly to the C6 alkylphenyl chain, the H-bond acceptor pyridine group appeared to be responsible for the higher potency. Boger's group also investigated the substitution of either the pyridine or oxazole ring [63, 64]. Again highly potent inhibitors were obtained featuring subnanomolar activities (Fig. (3), **3**, $K_i = 200$ pM and 2 nM on *h*FAAH and *r*FAAH, respectively; **4**, $K_i = 900$ pM on *h*FAAH). Focusing on the lipophilic portion that binds the ACB pocket, a series of bioisosteres was synthesized. For instance, equal activity to the C6 alkylphenyl chain was found for the C2 alkylbiphenyl chain, resulting in the inhibitor **5** (Fig. (3), $K_i = 400$ pM and 500 pM, on *h*FAAH and *r*FAAH, respectively) [65]. Finally, the same authors also studied the substitution of the phenyl ring at the end of the acyl side-chain, resulting in inhibitors such as **6** (Fig. (3), $K_i = 400$ pM on *r*FAAH) [66].

Meanwhile, another study on α -keto heterocycles was undertaken by the researchers of Janssen Pharmaceuticals. Based on the structure of OL-135 (**1**, Fig. (3)), including the oxazole ring, they oriented their efforts toward the insertion of a piperidinyll scaffold which allows for a wide diversity of substitutions. They published two patents containing SAR studies, and exemplified here with compounds **8** (Fig. (4), 400 pM and 4.7 nM on *h*FAAH and *r*FAAH, respectively) [67] and **9** (Fig. (4), $K_i = 2$ nM and 2 nM on *h*FAAH and *r*FAAH, respectively) [68].

II.3.2. Carbamate-based FAAH Inhibitors

Carbamate-based FAAH inhibitors were inspired by the structures of previously reported inhibitors of serine hydro-

lases. This function is usually used to inhibit serine proteases in an irreversible manner. Indeed, the tetrahedral intermediate evolves towards a stable acyl-enzyme complex. This mechanism of action was put forth, first by MS analyses [69] and then by X-ray structures [50], in studies involving URB597 (or KDS-4103, **10**, Fig. (5)) the lead compound of this class of inhibitors [70, 71]. URB597 is largely used as pharmacological tool, both *in-vitro* and *in-vivo* ($IC_{50} = 4.6$ nM on *r*FAAH). In addition, analogues of **10** were synthesized in order to increase their stability towards oxidative metabolism. Indeed, hydroxylation of the C4 position was observed following *in-vivo* administration of **10**. Thereby, this position was blocked by adding various substituents, like the gem-dimethyl found in **11** (Fig. (5)) [72], allowing a significant reduction of anandamide hydrolysis (50 % of control, at 30 nM).

Additional series of analogues were also synthesized, including compound **12** (Fig. (5), $IC_{50} < 0.1$ μ M) [73]. This compound is interesting as it exhibits a good oral bioavailability (C_{max} in plasma 2-fold higher compared to **10**) and a low CNS uptake (C_{max} in brain 10-fold lower compared to **10**) which may thus allow to target selectively FAAH in the periphery while not affecting FAAH activity in the CNS. This feature may be useful to treat pain and inflammation disorders by acting at peripheral sites, without inducing potential CNS (side)effects. Another carbamate derivative, compound **13** (Fig. (5)), $IC_{50} < 0.1$ μ M [74], also possesses a good oral bioavailability and its administration results in enhanced OEA and PEA levels in blood (7.55 and 8.85 ng/mL compared to 4.18 and 4.17 ng/mL for OEA and PEA,

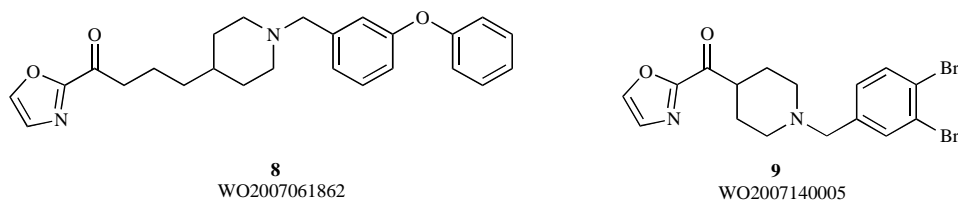


Fig. (4). FAAH inhibitors based on α -keto heterocycle templates described by Janssen Pharmaceutica.

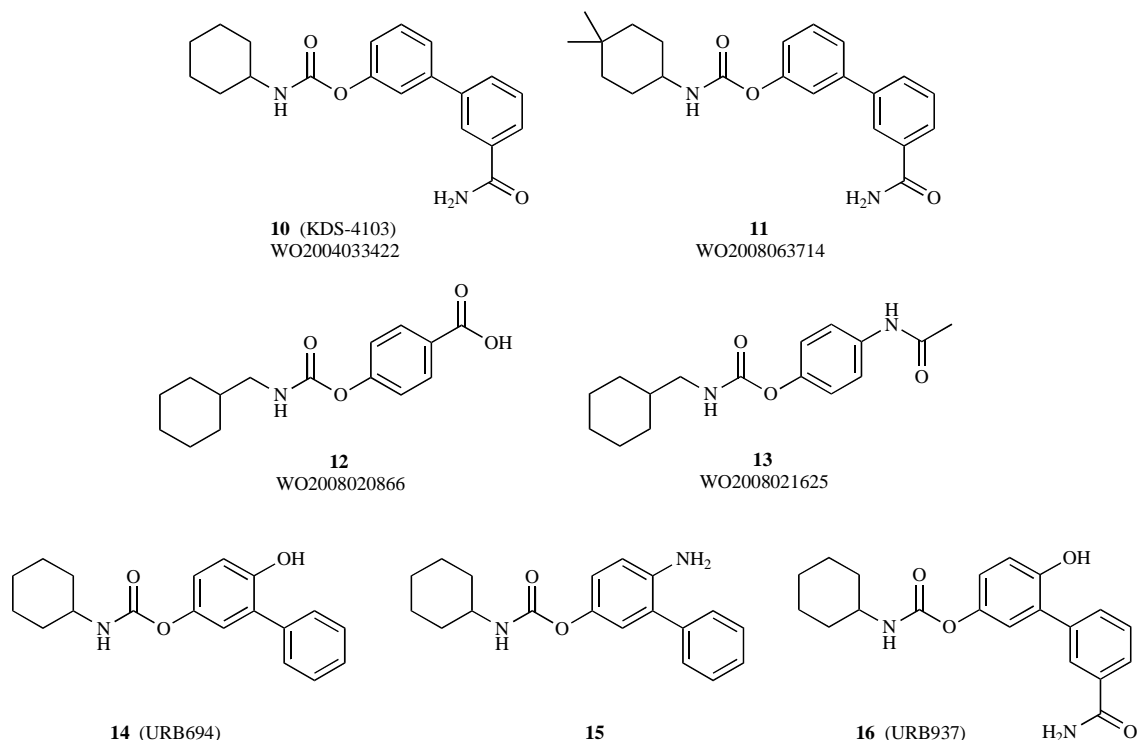


Fig. (5). Carbamate-type FAAH inhibitors developed by Piomelli *et al.* and Kadmus Pharmaceuticals.

respectively). The same authors described some other inhibitors but neither their potency nor *in-vivo* activity were disclosed [75, 76]. Piomelli and co-workers also developed new KDS-4103-based inhibitors with the aim to reduce its activity towards liver carboxylesterases. It is indeed known that **10** (URB597, Fig. (5)), while being quite selective, has several off-targets [77, 78], including carboxylesterases, which could prevent its further development. Thus, URB694 (**14**, Fig. (5), $IC_{50} = 30.0$ nM) and the aniline analogue **15** (Fig. (5), $IC_{50} = 27.2$ nM) [79], both possessing electron-donating substituents on the phenyl ring that reduce the electrophilicity of the carbonyl, were described. This decreased electrophilicity resulted in more selective compounds that retained their good activity against FAAH *in-vitro* and *in-vivo* [80]. Therefore, novel URB694-based inhibitors could soon be developed with improved selectivity for FAAH. Note that recently, a first URB694 derivative, URB937, (**16**, Fig. (5), $IC_{50} = 26.8$ nM) was disclosed to selectively inhibit FAAH in the periphery [81].

In addition, *O*-phenylcarbamates were also described by Astellas (**17**, Fig. (6), $IC_{50} = 12$ nM) and Myllymaeki and co-workers (**18**, Fig. (6), $IC_{50} = 240$ pM, *r*FAAH) [82, 83].

Due to their properties towards FAAH, carbamate-based inhibitors were also largely investigated by Sanofi-Aventis. Numerous series were described, based on various *O*- and *N*-substituents including alkyl, piperazinyl, azetidinyll or thiazolyl, as illustrated in Fig. (7) [84-91]. These inhibitors were all described for having an analgesic activity and their inhibition potencies against *m*FAAH are summarized in Fig. (8).

Two other carbamate-based families were developed at Sigma-Tau Pharmaceuticals: one is based on an enol carbamate template (**27** and **28**, Fig. (9), IC_{50} and K_i both < 10 nM, *m*FAAH) [92], and the other is based on an oxime carbamate (**29**, Fig. (9), IC_{50} and K_i both < 10 nM, *m*FAAH) [93]. These compounds are described as selective for FAAH over various cannabinoid-related targets (< 60 % versus CB_1 , CB_2 , TRPV1, NAPE-PLD, AMT, DAGL, MAGL, at a concentration equal to 1000-fold their IC_{50} against FAAH).

Interestingly, **27-29** inhibit FAAH in a reversible manner which is quite unexpected for carbamate-type inhibitors. *In-vivo*, compound **28** exhibits analgesic activity and reduces anxiety without affecting locomotor activity, whereas **29** was able to reduce anxiety as well as the hyperalgesia in a model of neuropathic pain.

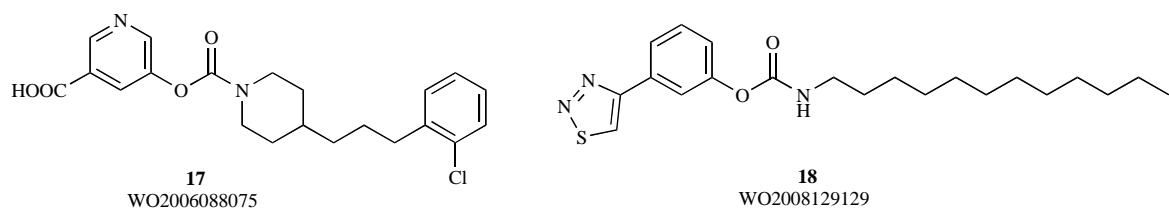


Fig. (6). Carbamate-type FAAH inhibitors described by Astellas Pharma and Saario *et al.*

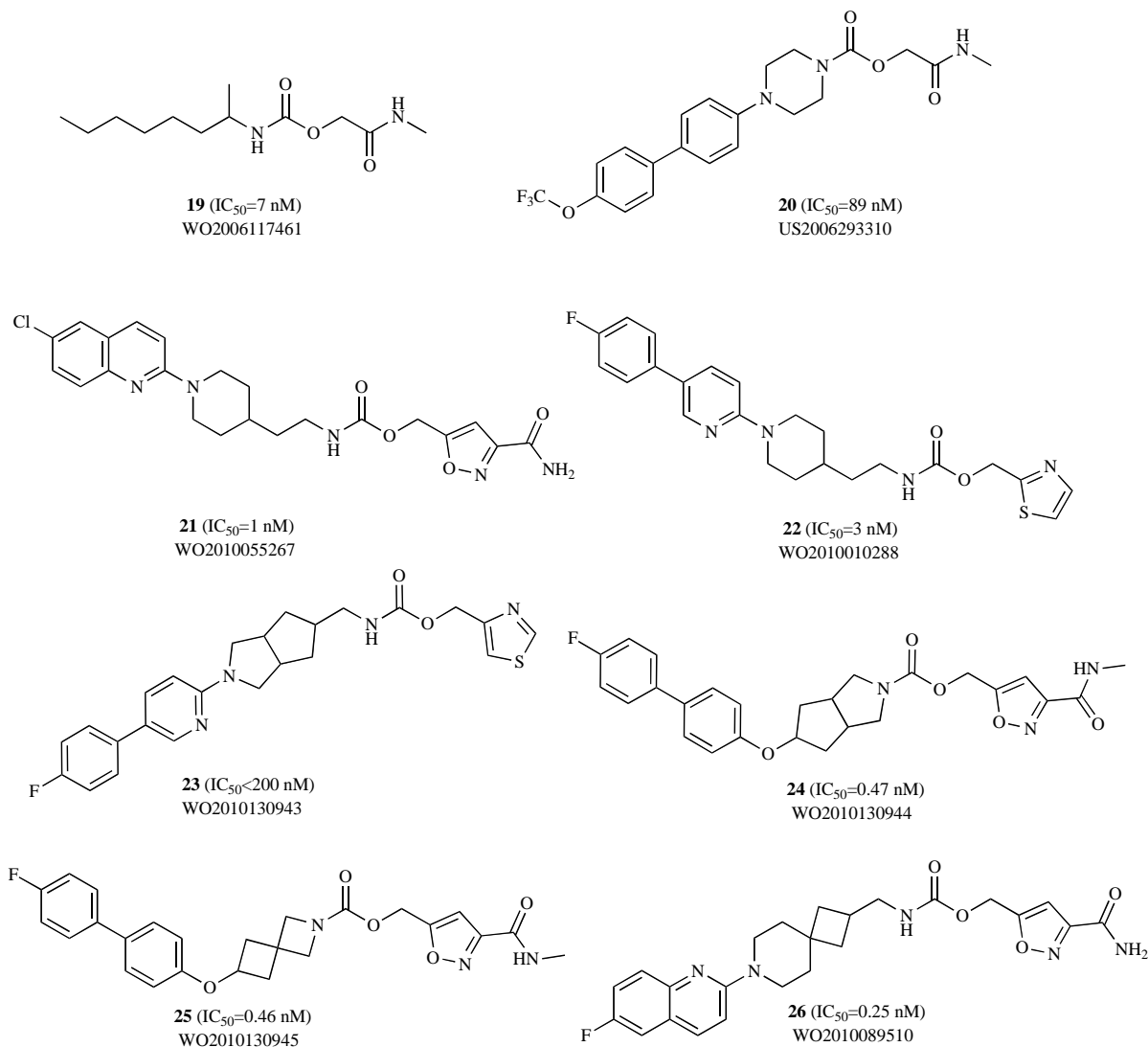


Fig. (7). Carbamate-type FAAH inhibitors described by Sanofi-Aventis.

II.3.3. Urea-based FAAH Inhibitors

Due to its high resistance towards chemical and biological hydrolysis, the urea function is usually not considered as a pharmacophore to inhibit serine hydrolases. However, it was shown that adding a good leaving group, such as an aniline function, transforms urea into a more reactive moiety, which could then function as enzyme inhibitor. Urea-based FAAH inhibitors originated from high-throughput screening studies of industrial chemical libraries. Both Janssen Pharmaceuticals and Takeda companies described compounds

based on a piperazinyl urea moiety (**30** and **31**, Fig. (12)). Compound **30** exhibits an IC₅₀ value of 16 nM or 50 nM depending on the source of enzyme (human and rat, respectively) [94], while for **31** 100 % of enzyme (*r*FAAH) inhibition was obtained at 1 μM [95].

However, the development of urea-based FAAH inhibitors really started with the discovery at Pfizer of PF-622 and PF-750 (**32** and **33**, Fig. (10)), $k_{inact}/K_i = 621 \text{ M}^{-1} \cdot \text{s}^{-1}$ and $791 \text{ M}^{-1} \cdot \text{s}^{-1}$ on *h*FAAH, respectively [46, 58].

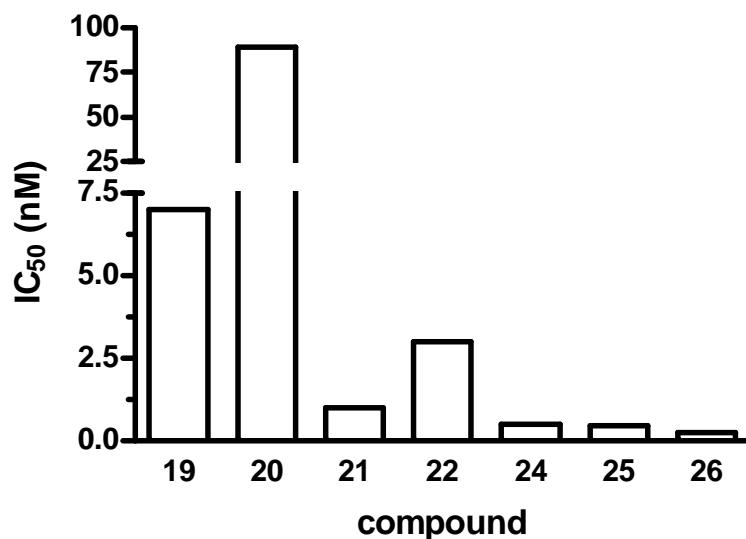


Fig. (8). IC₅₀ values of the carbamate-based inhibitors from Sanofi-Aventis depicted in Fig. (7).

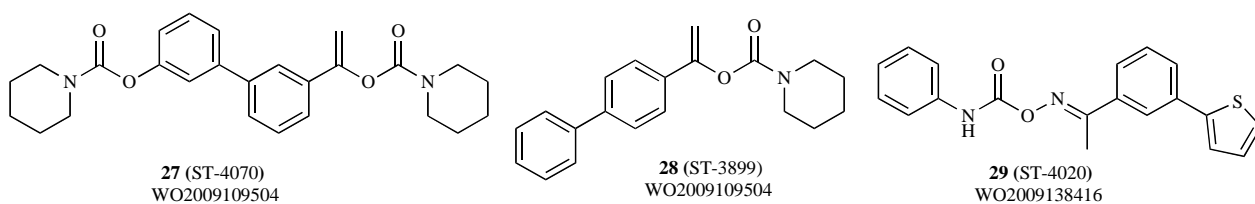


Fig. (9). Carbamate-type FAAH inhibitors described by Sigma-Tau Pharmaceuticals.

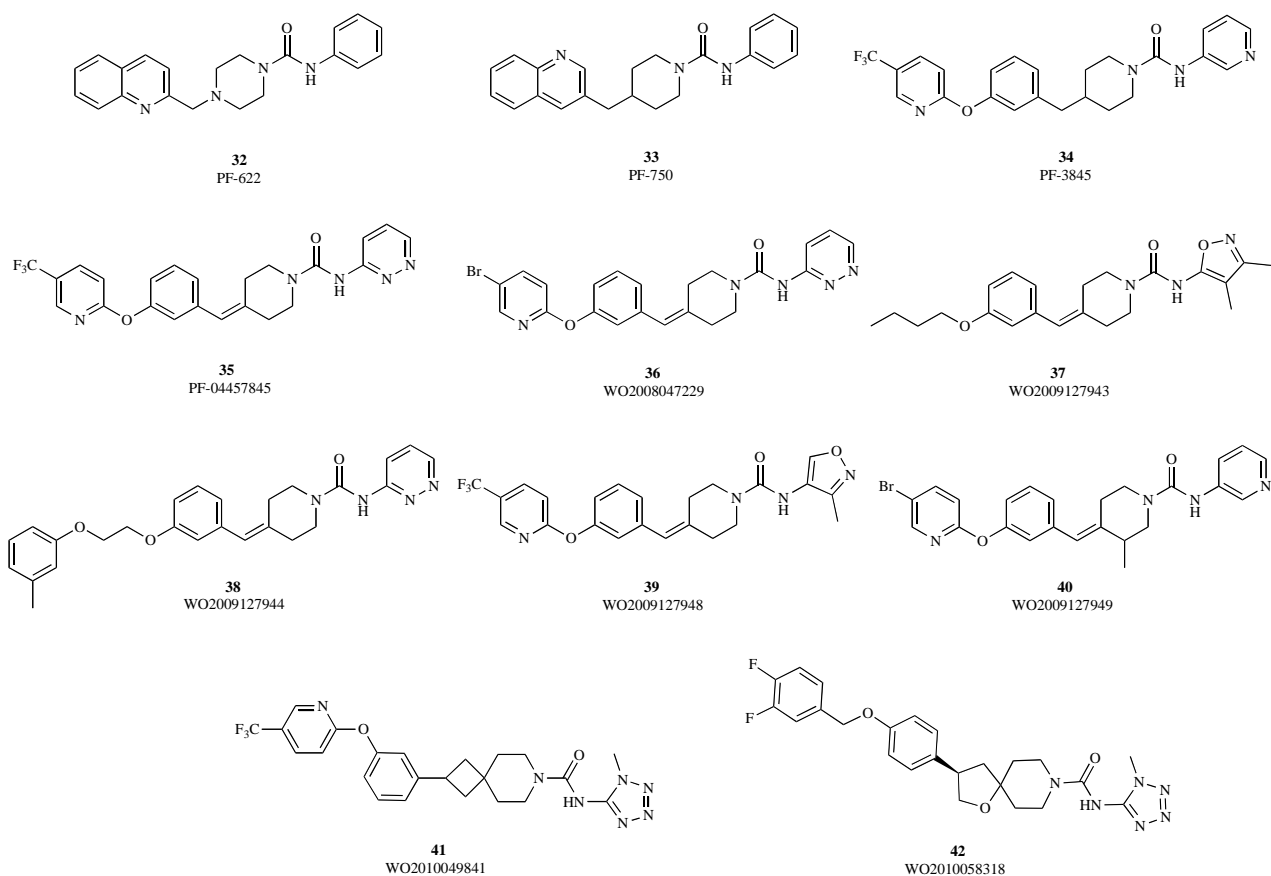


Fig. (10). Evolution of Pfizer's piperidinyl urea-type FAAH inhibitors.

Indeed, after a high-throughput screening study to improve either drug-like pharmacokinetic properties and/or selectivity, Pfizer published a new kind of mechanistic class of inhibitors, based on the piperidinyl urea scaffold, showing a combination of potency and excellent selectivity [58]. Of note, the development of this class of inhibitors at Pfizer benefited from a *h/r*FAAH three dimensional structure [46]. Thus, the stable acyl-enzyme complex was identified by MALDI-MS analyses and then by X-ray structures with *h/r*FAAH, which confirmed the addition of the inhibitor to the active serine and the irreversible mechanism of inhibition. These studies led to a new chemical family obtained by replacing, not only the quinoline group of **32** and **33** with a biaryl ether group, but also the aniline leaving group with a 3-aminopyridine one. These changes resulted in the inhibitor PF-3845 (**34**, Fig. (10)) which possesses much higher activity ($k_{inact}/K_i = 14,310 \text{ M}^{-1}\cdot\text{s}^{-1}$) [47]. Pfizer further described in several patents the synthesis and pharmacological evaluations of numerous urea-based inhibitors (e.g. compounds **35-40**, Fig. (10)) [96-100]. One line of research was to design more rigid compounds using a methylenepiperidine scaffold while incorporating a polar moiety to improve the pharmacokinetic parameters. This led to the clinical candidate PF-04457845 (**35**, Fig. (10), $k_{inact}/K_i = 40,300 \text{ M}^{-1}\cdot\text{s}^{-1}$) which contains a pyridazinyl moiety instead of the 3-aminopyridine one. This inhibitor exhibits a high selectivity for FAAH, excellent potency and good pharmacokinetic profile. The same company also prepared a new series of rigid piperidines where the methylene group was replaced with a C4-spirocycle (**41** and **42**, Fig. (10)) [101, 102]. The later inhibitors possess, similarly to the inhibitors containing a methylene unit, a high potency against FAAH activity.

It is difficult to directly compare the inhibitory activity of Pfizer's ureas with the other types of inhibitors since their potency is expressed in the literature as k_{inact}/K_i values instead of the IC_{50} values traditionally reported. However, the use of k_{inact}/K_i values appears more suitable than IC_{50} values when studying irreversible inhibitors. Note that in the same assay, the well-known URB597 (**10**, KDS-4103, Fig. (5)) shows a k_{inact}/K_i value of $1,590 \text{ M}^{-1}\cdot\text{s}^{-1}$ Fig. (11). Based on all the reported assays, piperidinyl ureas appeared to be more selective and efficacious than KDS-4103 [47].

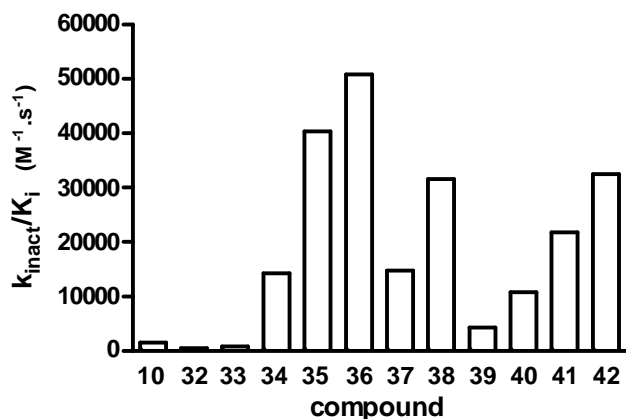


Fig. (11). Potency of Pfizer's ureas compared to the carbamate-based inhibitor KDS-4103 (**10**, Fig. (5)). Note that a highly potent inhibitor is characterized by a high k_{inact}/K_i value (and a low IC_{50} value).

While other urea-derivatives have been investigated by several pharmaceutical companies, piperazinyl urea remains the most common template. *i)* Sanofi-Aventis described two compounds exhibiting dual inhibition against both *m*FAAH and *m*MAGL (**43** and **44**, Fig. (12)) [103, 104]. *ii)* Janssen Pharmaceuticals reported two additional series of compounds (**45-49**, Fig. (12) and **53**, Fig. (13)). Compound **46** (Fig. (12), $IC_{50} = 19 \text{ nM}$ and 6 nM on *h*FAAH and *r*FAAH) [105] is derived directly from JNJ-1661010 (**45**, Fig. (12)) structure, whereas compounds **47-49** [106] contained a biaryl ether motif. When administered *in-vivo*, **47** (Fig. (12), $IC_{50} = 8 \text{ nM}$ and 10 nM on *h*FAAH and *r*FAAH, 20 mg/kg , po) showed analgesic effects in a model of mechanical allodynia. Its activity was improved by replacing the benzisoxazole moiety with an isoxazolopyridine resulting in compound **48** described as a subnanomolar inhibitor [107]. Inspired from the structure of **48**, **49** (Fig. (12), $IC_{50} = 1 \text{ nM}$) conserves a biaryl ether moiety, but a pyridine replaces the isoxazolopyridine motif [108]. *iii)* Other piperazinyl urea-based FAAH inhibitors were shown to possess *in-vivo* efficacy. For instance, Astella published a series of inhibitors, illustrated with **50** (Fig. (12), $IC_{50} = 160 \text{ pM}$, *r*FAAH), which seem to be useful in the context of overactive bladder [109], and Takeda published two families of compounds based on an isoxazole or pyridine moiety, exemplified here with **51**, which exhibited analgesic effect at 10 mg/kg [110], and **52**, which was proposed to treat sleep disorders [111]. *iv)* More recently, azetidiny ureas were described as FAAH inhibitors. Vernalis described compound **53** Fig. (13) which exhibits an IC_{50} value of 3 nM on *h*FAAH [112]. Janssen Pharmaceuticals investigated also azetidiny ureas with compound **54** Fig. (13), which presents an IC_{50} value of 1 nM both on *h*FAAH and *r*FAAH [113], and the recently published rigid spirocyclic compound **55** (Fig. (13), 10 nM and 20 nM on *h*FAAH and *r*FAAH, respectively) [114].

II.3.4. Boronic acid-based FAAH Inhibitors

Recently, a new kind of FAAH inhibitors using a boronic acid as the electrophilic function was reported. This function has been already described for inhibiting serine proteases in a reversible manner [115]. Indeed, boron's ability to go up from a trigonal planar geometry to a tetrahedral geometry allows boronic acids to form a transient and reversible tetrahedral intermediate with the nucleophilic serine. Both Infinity Pharmaceuticals (**56**, Fig. (14)) [116] and Minkkila and co-workers (**57**, Fig. (14)) [117] published in 2008, the first arylboronic acids described as FAAH inhibitors. These two inhibitors exhibited nanomolar activities ($K_i \leq 10 \text{ nM}$ for **56** and $IC_{50} = 9.1 \text{ nM}$ for **57**) and a reversible inhibition of the enzyme. In the case of compound **56**, kinetic data consolidated the hypothesis of a reversible inhibition. Furthermore, supplementary investigations were undertaken to unravel the interactions between the inhibitor and the enzyme. Both the molecular modelling studies and mutagenesis studies of *r*FAAH demonstrated that the phenyl moiety of these inhibitors interacts with the enzyme's hydrophobic channel.

Infinity Pharmaceuticals further covered this area with three patents in 2009 and 2010. Thus, they developed several series of FAAH inhibitors based on various substituted arylboronic acids, exemplified with compounds **58** [118] and **59** [119] (Fig. (14), $K_i \leq 10 \text{ nM}$), and also based on a tetrahydro

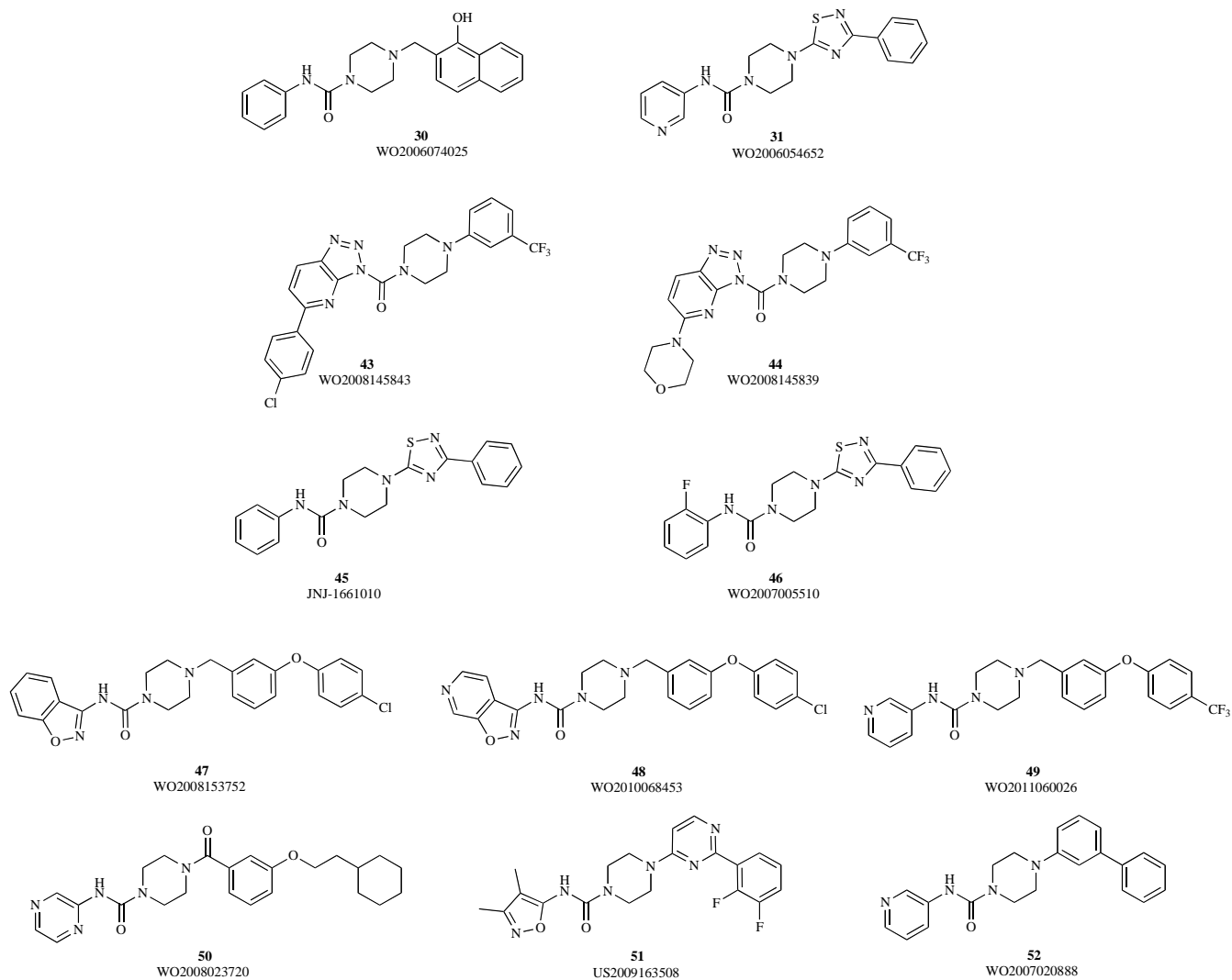


Fig. (12). Urea-type FAAH inhibitors.

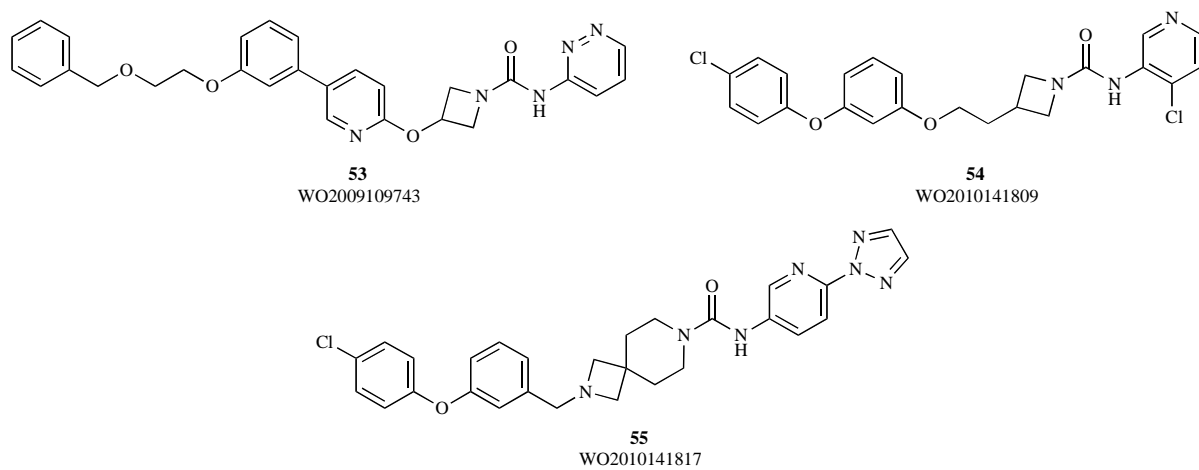


Fig. (13). Urea-type FAAH inhibitors from Vernalis and Janssen Pharmaceuticals.

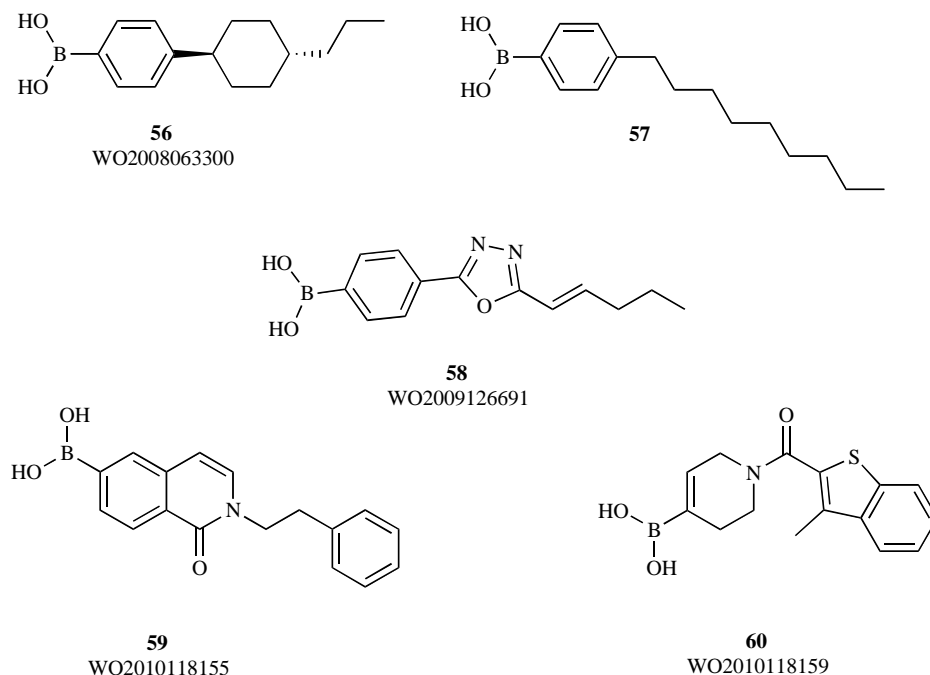


Fig. (14). Boronic acids developed as FAAH inhibitors.

pyridine boronic acid (*i.e.* compound **60**, Fig. (14), $K_i \leq 10$ nM) [120].

II.3.5. FAAH Inhibitors of Miscellaneous Structures

Whereas initially, carbamate and activated ketone-based inhibitors were mainly described, a wider diversity of templates has been explored since 2006. Some indole and pyrrole derivatives were investigated by Ironwood Pharmaceuticals (Microbia) which published several patents from 2006 to 2011 [121-125]. Therein, different series of compounds were disclosed that were, not only FAAH inhibitors, but that also interact with one or several targets involved in inflammation and pain (e.g. COX-1 or COX-2). For example, compounds **61** [122], **62** [123] and **63** [125] (Fig. (15)) were found to inhibit *h*FAAH at submicromolar concentrations.

Bial (Portela & C^o) reported several series of oxadiazolones (**64**, **65** and **66**, Fig. (16)) as FAAH inhibitors being selective for peripheral over CNS located FAAH [126-128].

Indeed, following administration of the inhibitors to mice (30 mg/kg, p.o.) the residual activity in liver was found to be very low (5 %, 6 % and 29 % compared to the control, for **64**, **65** and **66** respectively) whereas it was almost completely conserved in brain (83 %, 86 % and 84 % compared to the control, for **64**, **65** and **66** respectively).

Merck designed various heterocycle-based FAAH inhibitors like imidazole, pyrazole or oxazole cycles. Thus, two series of imidazole derivatives were described which inhibit *h*FAAH with nanomolar and subnanomolar activities (Fig. (17), compounds **67** and **68**, IC_{50} values of 6.3 nM and 0.2 nM, respectively) [129, 130]. In addition, a pyrazole series was published, illustrated with **69** (Fig. (17), IC_{50} value of 0.47 nM) [131] and also an oxazole one, represented by **70** (Fig. (17) [132]. The inventors described the later inhibitor as

having good cell permeability ($IC_{50} = 5$ and 20 nM, in cell lysate and in whole cell, respectively).

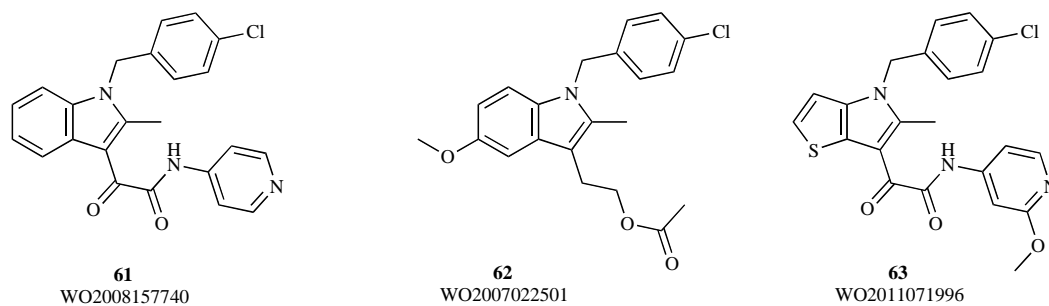
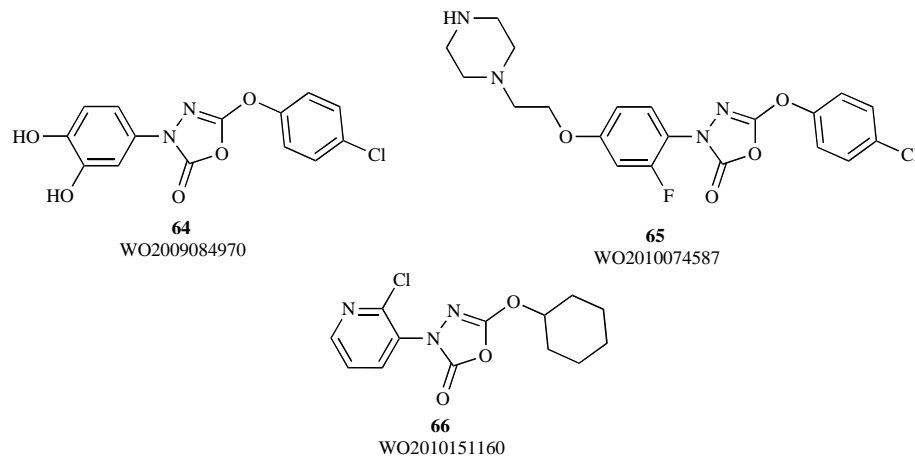
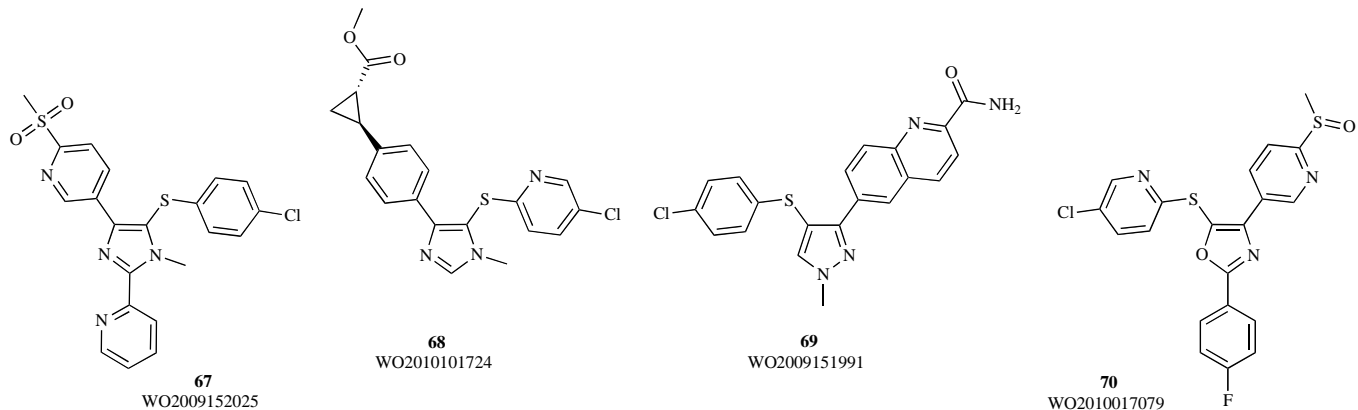
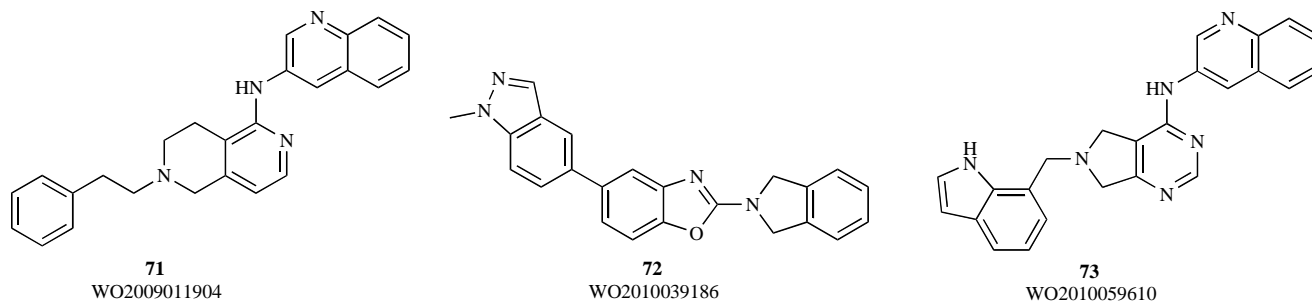
Renovis published several lipophilic and polycyclic compounds as FAAH modulators. For instance, compounds **71** [133] and **72** [134] exhibited nanomolar IC_{50} values (Fig. (18), $IC_{50} < 100$ nM and $IC_{50} = 1.2$ nM for **71** and **72**, respectively) while compound **73** (Fig. (18)) [135] showed more than 75 % of FAAH inhibition at 1 μ M compared to the control.

Janssen Pharmaceuticals developed also a family of inhibitors based on a pyrimidine moiety with a C6 aryl group and a C4 amine function. Compounds **74** [136] and **75** (Fig. (19)) [137] featured nanomolar activities (Fig. (19), $IC_{50} = 1$ and 3 nM for **74** and $IC_{50} = 7$ and 240 nM for **75**, against *h*FAAH and *r*FAAH, respectively).

Very recently, Infinity Pharmaceuticals identified the isoxazoline heterocycle as a new template for FAAH inhibition. Four series of compounds were presented based on this new scaffold. The representative compounds **76**, **77**, **78** and **79** (Fig. (20)) were reported to have $K_i \leq 100$ nM and an irreversible mode of *h*FAAH inhibition [138-140]. Indeed, the authors reported evidence for a covalent FAAH inhibition *via* kinetic data, on one hand, and rapid dilution experiments, on the other hand, confirming the irreversible or slowly reversible inhibition. This mechanism of action was explained by the nucleophilic addition of the active Ser-241 on the isoxazoline C=N function followed by the elimination of the leaving group, *i.e.* Br or ArO substituent at C3.

II.3.6 Other Structures Not Covered by Patents

Since 2006, a number of inhibitors have been published in the literature but are not covered by patents. We have summarized here the main families. A family of 2-thioxo-

**Fig. (15).** 2-Methylindole-based inhibitors of FAAH.**Fig. (16).** Oxadiazolone-type FAAH inhibitors.**Fig. (17).** Central heterocycle-based FAAH inhibitors from Merck.**Fig. (18).** FAAH inhibitors from Renovis.

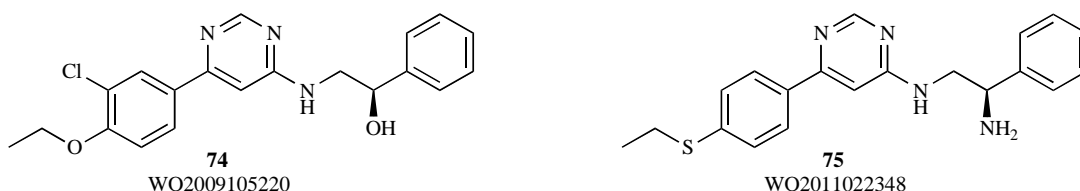


Fig. (19). Pyrimidine-based FAAH inhibitors from Janssen Pharmaceuticals.

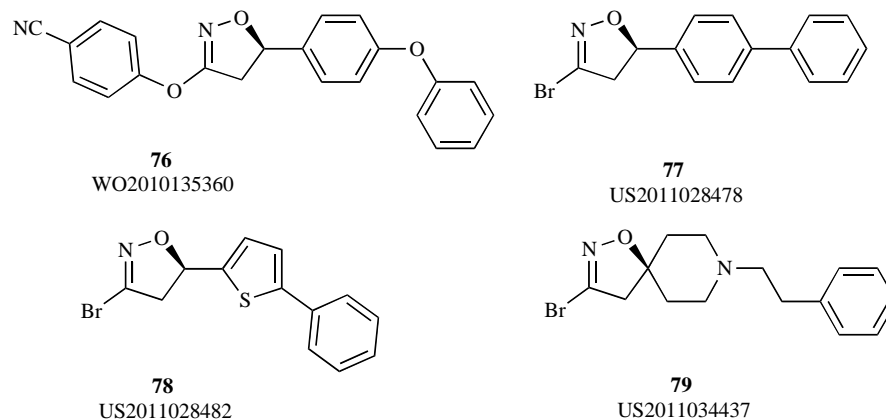


Fig. (20). Isoxazoline-based FAAH inhibitors described by Infinity Pharmaceuticals.

imidazolidin-4-ones was described to inhibit FAAH in a reversible and competitive manner (**80**, Fig. (21), $pI_{50} = 5.86$) [141]. Then, benzothiazole-based inhibitors were reported to reversibly inhibit FAAH with nanomolar activity (**81**, Fig. (21), $IC_{50} = 1.7$ nM) [142]. During the same year, two distinct series of paracetamol [143] and ibuprofen [144] analogues were disclosed to block FAAH activity with good to moderate potency (**82** and **83**, Fig. (21), $IC_{50} = 100$ nM and $pI_{50} = 5.86$, respectively). A unique series of 1-indol-1-ylpropan-2-ones was also described for a dual inhibition towards FAAH and cytosolic phospholipase $A_2\alpha$ (**84**, Fig. (21), $IC_{50} = 47$ nM and 2.2 μ M against FAAH and $cPLA_2\alpha$, respectively) [145]. In 2009, β -lactam-based inhibitors were disclosed to inhibit FAAH in a reversible manner without being processed by the nucleophilic serine (**85**, Fig. (21), $IC_{50} = 8$ nM) [51, 146]. Additionally, the first potent non-covalent and competitive inhibitors of FAAH were disclosed (**86**, Fig. (21), $IC_{50} = 36$ nM) [147].

II.4. Current Clinical Trials Involving FAAH Inhibitors

Based on the preclinical studies reported so far, the most promising therapeutic applications for FAAH inhibitors are to be found in the treatment of pain and mood, and sleep disorders. Recently, Pfizer undertook a phase II clinical trial with PF-04457845 (**35**, Fig. (10)) to evaluate its efficacy, safety and tolerability in knee osteoarthritis (NCT00981357). Another small scale clinical trial (NCT01092845) aimed at studying the effect of **35** on sleep. Indeed, a positive effect on sleep would represent a proof-of-concept for the CNS efficacy of the compound, and more largely of increasing AEA levels, in humans.

In addition, phase II clinical trials were also undertaken to evaluate SSR-411298, a FAAH inhibitor developed by Sanofi-Aventis, for treatment of major depressive disorders in the elderly patients (NCT00822744). To date, neither the inhibitor structure nor results were reported concerning these investigations. Note however that although the development of SSR-411298 in this indication has been abandoned, other indications (e.g. pain, NCT01439919) are being investigated.

Infinity Pharmaceuticals is developing IPI-940 (no structure available) in order to treat various types of pain. Phase I resulted in positive data, and IPI-940 is presented as a well-tolerated compound with good pharmacokinetic, pharmacodynamic and safety properties. Purdue Pharmaceutical Products is expected to initiate Phase II studies with this compound. Finally, Vernalis has its own FAAH inhibitor, V158866 ($IC_{50} = 24$ nM) entering Phase I clinical trials.

We are at the early stages of the clinical development of FAAH inhibitors; the results of the first Phase II trials are eagerly awaited to determine whether FAAH inhibition will prove to be a viable drug target.

III. MONOACYLGLYCEROL LIPASE

III. 1. MAGL, Structure and Mechanism of Action

Although the existence of a monoacylglyceride hydrolase in the adipose tissues was reported decades ago [148, 149], MAGL became more actively investigated after its role in controlling 2-AG (Fig. (22)) levels was demonstrated [3-4].

Molecular cloning allowed determining the catalytic triad of the enzyme and its classification as a member of the α/β

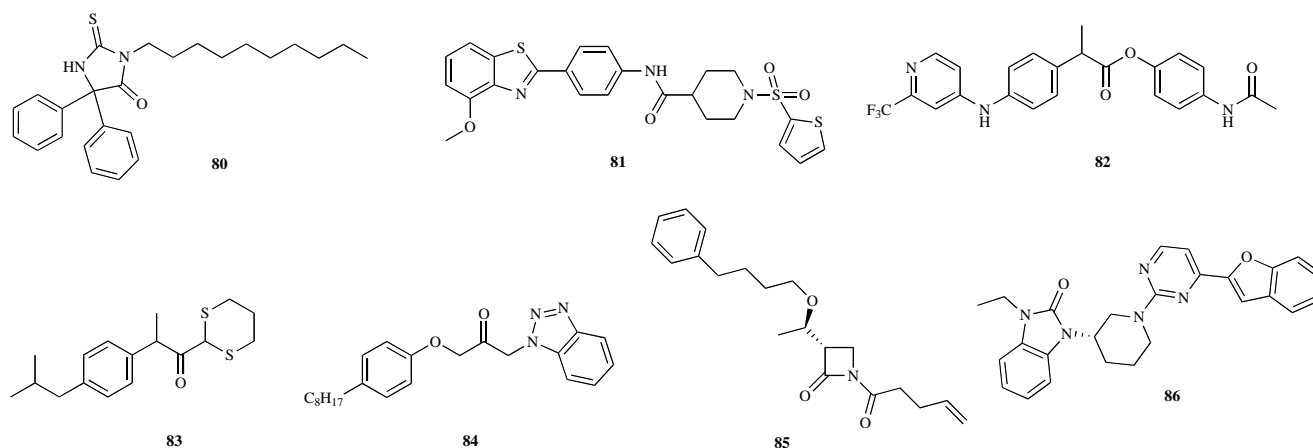


Fig. (21). FAAH inhibitors not covered by the patents.

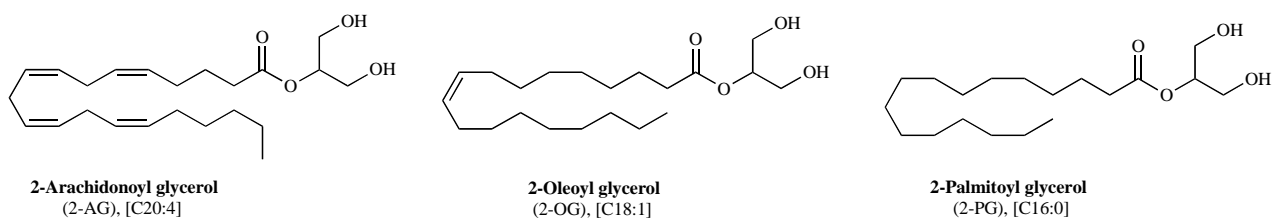


Fig. (22). Known endogenous substrates of MAGL.

hydrolase family [150], MAGL activity is governed by the classical Ser-His-Asp catalytic triad of the serine hydrolases. Additionally, four cysteine residues were shown to interact with some enzyme inhibitors [151-153]. Thus for instance, *N*-arachidonoylmaleimide, disulfiram and octhilonone were developed as MAGL inhibitors targeting those cysteine residues [151, 153, 154].

Very recently, the *h*MAGL's three dimensional structure was independently elucidated by two research teams, with a resolution of 2.2 Å [155] and 2.7 Å [156], respectively. The publication by Sanofi-Aventis described also a co-crystal between MAGL and SAR-629, one of their own MAGL inhibitors (87, Fig. (23)). *In silico* modelling of the tetrahedral intermediate between 2-AG and the active serine [155, 156], as well as the X-ray structure of MAGL-SAR-629 co-crystal [156], allowed important structural features to be established. i) At the surface of the enzyme, a large highly hydrophobic cavity which leads to the active site is present. This channel, made of several hydrophobic residues, appears to be suitable for interacting with the lipophilic chain of the substrate and seems to govern substrate specificity. ii) a lid (or cap) is present at the entrance of the channel. This lid is suggested to allow MAGL to interact with the cell membrane, thus helping in recruiting its lipophilic substrates from the membrane. Note that MAGL is found in both soluble and particulate fractions suggesting that the interaction between the lid and the membrane is reversible. iii) Closer to the active site a hydrophilic pocket is present and appears to be able to accommodate the substrate's glycerol moiety. This pocket, named "alcohol-binding pocket" [155] or "exit-hole" [156] by the two groups, contains three residues Ala51 (Ala61 in Sanofi's paper), His121 (His131) and Tyr194

(Tyr204), important for substrate recognition, and thus potential residues to be targeted by novel inhibitors. iv) Two non-catalytic cysteines, Cys201 (Cys211) and Cys242 (Cys252), which are supposed to be targeted by Michael-acceptor inhibitors, are in the vicinity of the catalytic site. Cys242 (Cys252) lies very close to the active Ser, deeply buried in the catalytic pocket, and Cys201 (Cys211) is farther from the active serine but remains accessible to inhibitors from the active site. On the contrary, Cys208 (Cys218) is described as pointing toward the outside of the enzyme. It is expected that these crystal structures will aid the development of novel MAGL inhibitors.

III.2. Pharmacology of 2-AG or why Inhibiting MAGL Hydrolase Activity?

2-AG is present at high levels in the brain, where it exerts an important role in controlling neurotransmitters release, and is also present in the periphery throughout the organism. (For a review see [157]) Indeed, beside its role as transmitter, 2-AG is an intermediate in lipid metabolism and, very likely, only a limited fraction of the 2-AG available acts as lipid mediator. Among the proposed roles for 2-AG, it was demonstrated to be involved in various processes like neuroprotection [158-160], appetite [161], cognitive and affective behaviours, or nociception [162] and inflammation, resulting from CNS and peripheral system locations, respectively. (for a complete review, see [157]) Furthermore, several studies demonstrated the involvement of 2-AG in controlling cell proliferation and invasion, suggesting that MAGL inhibitors could be relevant in cancer treatment [163, 164]. Preclinical studies suggested that MAGL inhibition could represent an interesting strategy for treating pain [20, 36, 165, 166, 167],

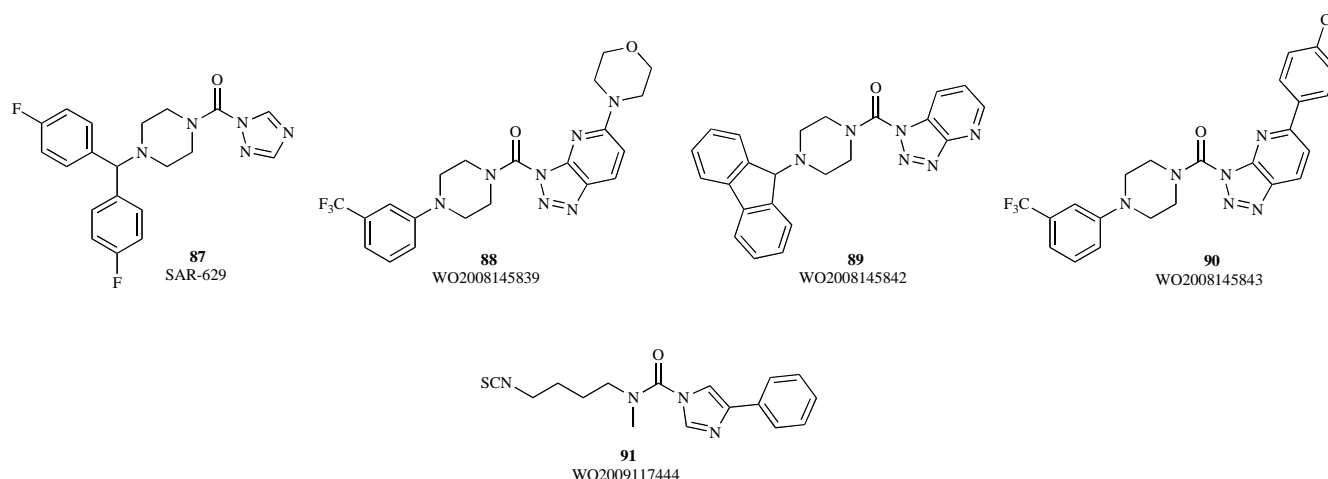


Fig. (23). Urea-type MAGL inhibitors.

inflammation [37, 166, 168], vomiting, nausea [169] and anxiety [40]. Investigations for MAGL inhibitors are more recent than those on FAAH, resulting in a limited number of inhibitors (see [54, 170]). Below we will review the available patents describing MAGL inhibitors.

III.3. MAGL Inhibitors

III.3.1. Urea-based MAGL Inhibitors

Sanofi-Aventis was the first pharmaceutical company which published MAGL inhibitors. Urea-based compounds bearing piperazinyl and triazole or triazolopyridine moieties as substituents were designed and developed.

Compounds **88** [104], **89** [171] and **90** [103] (Fig. (23)) were found to inhibit *m*MAGL at nanomolar concentrations (IC_{50} values of 4 nM, 4 nM and 2 nM for **88**, **89** and **90**, respectively) and to exhibit either selectivity for MAGL over FAAH, or dual nanomolar inhibition of both enzymes. Makriyannis' group at Northeastern University also disclosed several urea-based MAGL inhibitors such as **91** (IC_{50} = 42 nM against *h*MAGL) [172]. Note that, due to its isothiocyanate function, this compound could be useful as a covalent probe to explore MAGL properties, and notably the cysteine residues.

III.3.2. Carbamate-based MAGL Inhibitors

Piomelli's group reported the ability of a carbamate derivative, URB602 (**92**, Fig. (24)), to inhibit MAGL (IC_{50} value of 28 μ M) [173]. However, this compound lacks selectivity since it inhibits FAAH with a similar potency [174, 175].

Much more recently, by screening its own library of carbamates, Cravatt's group found piperidinyl and piperazinyl carbamates which were able to inhibit MAGL without affecting FAAH activity [176]. The authors demonstrated that an increased steric hindrance improved the selectivity toward MAGL. This work resulted in the design of a selective and potent MAGL inhibitor with the synthesis of compound JZL184 (**93**, Fig. (24)) [177, 178]. Indeed, with the incorporation of two oxygen atoms in the 3 and 4 positions of the

phenyl rings, they obtained an excellent selectivity in the range of 400-fold (IC_{50} values of 10 nM and 4690 nM for MAGL and FAAH, respectively). In addition, inspired by the selective piperazinyl urea-based FAAH inhibitor **32** (PF-622, Fig. (10)) and by **93**, Cravatt *et al.* developed a series of compounds, such as **94** (JZL195, Fig. (24)), which inhibited FAAH and MAGL with a similar potency without affecting other serine hydrolases (IC_{50} values of 13 and 19 nM for FAAH and MAGL, respectively) [9].

Today, compound **93** is extensively used as a reference pharmacological tool to study the effects of MAGL inhibition. Several publications report the use of **93** for increasing 2-AG levels and the resulting effects, for instance, in cancer pathogenesis [164], neuropathic pain [36], anxiety [40] and colon inflammation [168].

III.3.3. MAGL Inhibitors of Miscellaneous Structure

MAGL inhibitors based on an activated ketone were described by Makriyannis' group at Northeastern University. The α -keto oxadiazole derivatives, such as **97**, are quite active against MAGL, although they remain more active against FAAH (Fig. (25), IC_{50} = 71 nM and K_i = 17 nM against *h*FAAH) [179].

Janssen Pharmaceuticals published a series of three patents describing MAGL inhibitors based on an amide function. Each patent is illustrated with a lead compound (**98-100**, Fig. (25)) that inhibit MAGL with an IC_{50} value of 10.4 nM, 10 nM and 50 nM, respectively) which was tested in various *in-vitro* and/or *in-vivo* pharmacological evaluations [180-182]. Thus for instance, **98** was able to increase 2-AG levels in an *ex-vivo* preparation of rat brain. *In-vivo*, compound **98** (30 mg/kg, po) completely prevented the CFA-induced heat hypersensitivity and partially the CFA-induced pressure hypersensitivity. Compound **99** was similarly tested in various experimental models to assess its antinociceptive properties. The group also published crystal structures of different MAGL mutants and co-crystallised forms with compounds **99** and **100** with resolutions of 1.35 and 2.3 Å, respectively [183, 184].

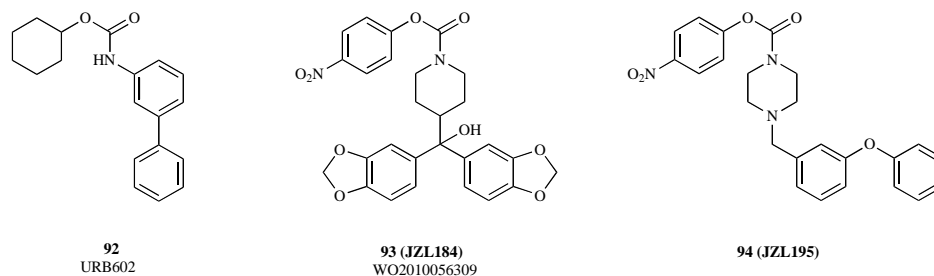


Fig. (24). Carbamate-type MAGL inhibitors.

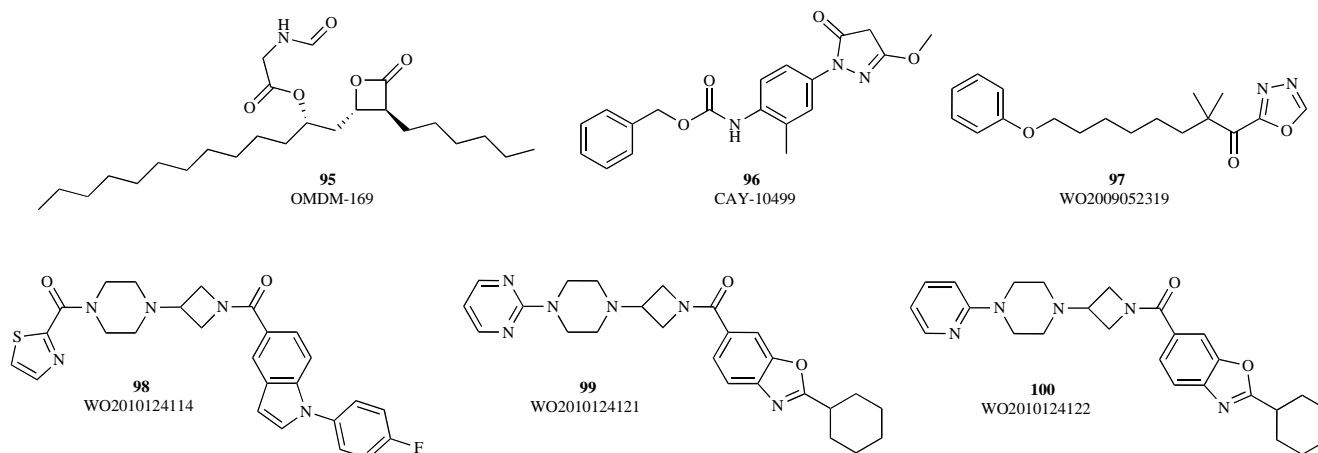


Fig. (25). MAGL inhibitors of various structures.

III.3.4. Other Recent Structures Not Covered by Patents

A β -lactone inspired from the serine hydrolase inhibitor, tetrahydropipstatin, was designed by Di Marzo's group. OMDM169 (**95**, Fig. (25)) inhibits *h*MAGL with an IC_{50} value of 0.89 μ M in a competitive manner. However, the authors also disclosed that compound **95** also inhibits *r*FAAH with an IC_{50} value of 3.0 μ M [165]. Also of interest, is the finding that the 5-methoxy-1,3,4-oxadiazol-2(3H)-one moiety (**96**, Fig. (25)) is also able to inhibit MAGL activity, thus offering an additional template for the development of inhibitors of the enzyme [185, 186].

IV. N-ACYLETHANOLAMINE ACID AMIDASE

IV.1. NAAA, Structure and Mechanism

Like FAAH, the *N*-Acylethanolamine-hydrolyzing Acid Amidase (NAAA) is also able to cleave amide bonds of saturated and unsaturated NAEs [8]. NAAA is thought to exert almost all of its hydrolytic activity towards PEA (Fig. (2)) since other NAEs are hydrolysed at much lower rates [187]. Moreover, it is notable that NAAA does not hydrolyse 2-AG. Although NAAA, like FAAH, exerts its activity towards NAEs, there is no sequence homology between these two enzymes, and whereas the optimum pH for FAAH activity is around 9, NAAA's activity is the highest at pH 5 [188, 189]. This is actually consistent with the subcellular localisation of NAAA in the lysosomes [189, 190]. Moreover, NAAA shares high sequence homology with the human acid ceramidase family, and its mode of action and structural

features are closer to those of this hydrolase family than to FAAH [191]. For instance, similarly to what is found for the cholesteryl glycerol hydrolase superfamily [192], and more precisely for the acid ceramidase family, the precursor form of NAAA is auto-catalytically cleaved into two subunits, α and β , at acidic pH. Then, this cleavage leads to the appearance of the unmasked *N*-terminal nucleophilic residue responsible for the catalytic activity of NAAA [193]. Wang *et al.* also identified Cys126 as the *N*-terminal residue and Cys126/Arg142/Asp154 as the residues constituting the catalytic triad of the human NAAA. As a consequence, the strategy for NAAA targeting is mainly based on the cysteine hydrolase activity of the enzyme, contrasting to the strategies used to target the serine hydrolases of the endocannabinoid system (i.e. FAAH, MAGL, and ABHD6). Because the discovery and initial characterisation of NAAA are quite recent, only a very limited number of studies have been published to date.

IV.2. Pharmacology of PEA or why Inhibiting NAAA Hydrolase Activity?

Several studies suggested that the role of NAAA is to regulate NAEs levels in macrophages and peripheral tissues [194]. As PEA is NAAA's primary substrate, its inhibition appears to be a relevant alternative to FAAH inhibition in the induction of anti-inflammatory [195], analgesic [27, 28] and neuroprotective effects [196]. Indeed, these effects can be mediated by PEA through receptors that are distinct from cannabinoid receptors (e.g. PPAR α) [197].

IV.3. NAAA Inhibitors

IV.3.1. Substrate-like NAAA Inhibitors

The initial studies on NAAA inhibitors consisted in the synthesis of substrate analogues. Thus esters (**101** and **106**, IC_{50} values of 19 and 10 μ M respectively, Fig. (26)), retroesters (**102**, IC_{50} value of 53.8 μ M, Fig. (26)), amides and retroamides (**103**, **104** and **105**, IC_{50} values of 31.8, 4.5 and 8.3 μ M respectively, Fig. (26)) of palmitic acid were developed [198-201].

Compounds **101**, **102** and **103** were tested against *r*FAAH and *r*NAAA (solubilised from the 12000*g pellet of rat lung homogenates) and were found to be selective at 100 μ M for NAAA versus FAAH (84, 71 and 77 % of NAAA inhibition versus 36, 0 and 8 % of FAAH inhibition, for compounds **101**, **102** and **103** respectively) [198, 199]. Similarly, compounds **104** and **105** do not inhibit FAAH at 100 μ M. The inhibition mechanism of **104** was further investigated and **104** was found to act by a reversible and non-competitive mechanism. This compound was also shown to inhibit NAAA in intact macrophages and in macrophage homogenates [200].

More recently, another study based on PEA analogues has been published. The authors used recombinant *r*NAAA expressed in HEK cells to test novel series of NAAA inhibitors. Among the assayed compounds, **106** (Fig. (26)) was found to be a selective and competitive NAAA inhibitor [201].

IV.3.2. β -lactone-based Inhibitors

To date, and to our knowledge, the only inhibitors known for inhibiting NAAA with a sub-micromolar activity are based on the β -lactone template.

Compounds **107** and **108** (Fig. (27)) [202] were found to inhibit *r*NAAA (recombinant HEK cells expressing *r*NAAA) in a non-competitive and reversible manner with IC_{50} values of 115 and 420 nM, respectively. As no crystal structure of NAAA is available, the authors built a model of NAAA catalytic site based on its high homology with conjugated bile acid hydrolase (CBAH). This model, which was validated by the docking of the tetrahedral intermediate between PEA and

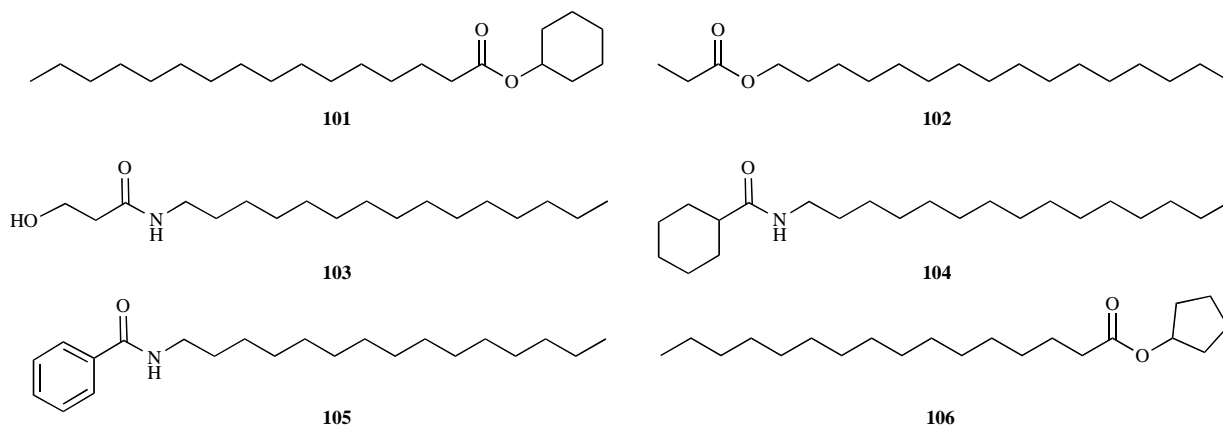


Fig. (26). Substrate-based NAAA inhibitors.

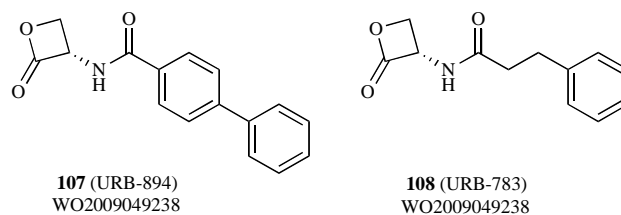


Fig. (27). β -lactone-type NAAA inhibitors.

Cys131 (*r*NAAA), is to date, the only tool available for designing new inhibitors of NAAA [203]. Compound **107** [204] as well as compound **108** [202] exhibited anti-inflammatory effects in various inflammation models where URB597 (KDS-4103 or **10**, Fig. (5)) had no effect, suggesting that NAAA is solely implicated.

V. CURRENT AND FUTURE DEVELOPMENTS

When looking at the variety of compounds described here, it is safe to say that we now have the tools to fully explore the consequences of FAAH and MAGL selective inhibition. The early thinking was that using an inhibitor would allow to increase local levels of endocannabinoids due to their on-demand production. However, it appears that the administration of a FAAH or MAGL inhibitor results in increased endocannabinoid levels throughout the body (see for instance [205, 206]). Although this results in a situation not that different from agonist administration, advantages of inhibiting the endocannabinoid hydrolysing enzymes still exist. First, by selectively inhibiting FAAH or MAGL only a subset of the effects obtained following agonists administration are observed. Thus when looking at the cannabinoid tetrad of effects [23] - i.e. antinociception, catalepsy, hypolocomotion, hypothermia - all the effects are present following CB_1 agonist administration, but only antinociception is induced upon FAAH inhibition. Another interesting point is that neither selective FAAH inhibition nor selective MAGL inhibition induce a cataleptic behaviour in mice. However, upon blockade of both enzymes catalepsy is present, as it is following CB_1 agonist administration (Table 1, and see [9]).

Table 1.

	FAAH Inhibition	MAGL Inhibition	FAAH/MAGL Dual Inhibition
Antinociception	+	+	+++
Catalepsy	-	-	++
Hypolocomotion	-	+	+
Hypothermia	-	-	-

Based on the published studies, it appears that FAAH inhibition generates less CNS-related side effects compared to MAGL inhibition. Thus for instance MAGL, but not FAAH, inhibition reduces locomotion. Another difference between MAGL and FAAH inactivation is the adaptations in CB₁ signalling observed following MAGL, but not FAAH, complete and chronic inhibition [207, 208]. These adaptations, resulting in functional antagonism of the endocannabinoid system, provoke a lower analgesic effect upon MAGL chronic inhibition compared to FAAH inhibition, even though acute MAGL inhibition induces similar effects than acute FAAH inhibition [208]. Based on these studies, it has been suggested, but has not been demonstrated yet, that partial blockade of MAGL could preserve its analgesic potential also during chronic administration.

It is also worth noting, that FAAH inhibition seems to be safe although a large number of bioactive lipids, besides NAEs, are hydrolysed by the enzyme. For instance, chronic inhibition of FAAH increases NAE levels, but also *N*-acyltaurines which are transient receptor potential channels agonists [206]. Conversely, brain levels of the GPR18 receptor endogenous agonist *N*-arachidonoylglycine are decreased following FAAH inhibition.[19] The question whether MAGL inhibition results in the exclusive modulation of monoacylglycerols (and corresponding fatty acids) [178, 205] remains open. Of great interest is the recent demonstration that MAGL-produced arachidonic acid is further metabolized in prostaglandins. Thus inhibition of MAGL results in increased levels of 2-AG, but also in decreased prostaglandins levels, further supporting MAGL as an interesting anti-inflammatory target [209].

Because NAEs (e.g. anandamide and *N*-palmitoylethanolamine) and 2-AG levels are profoundly affected throughout the body by FAAH and MAGL, respectively, inhibition, one could question the interest in pursuing inhibitors of the additional endocannabinoid hydrolysing enzymes (NAAA, ABHD6, ABHD12). One argument in favour of these enzymes can be found in the localisation of specific enzymes at the tissue and cell level. This results in enzymes controlling pools of signalling mediator. Thus, although MAGL controls 85 % of 2-AG hydrolysis in whole brain homogenates, ABHD6 selective inhibition [210-212] in intact neurons and in brain slices results in increased 2-AG levels and 2-AG induced-synaptic plasticity, respectively [7]. Note that it was also recently demonstrated that carboxylesterase-1 (CES-1) participates in the control of 2-AG metabolisms in macrophages [213]. When looking at NAAA, its inhibition in intact macrophages reduces AEA degradation to a similar, if not higher, extent than FAAH

inhibition [202, 214]. These two examples underscore the potential of targeting NAAA, ABHD6, and perhaps ABHD12, to more precisely fine-tune endocannabinoid levels in a subset of cells inside a tissue. Thus, in addition to advancing the development of FAAH and MAGL inhibitors from bench to bedside, efforts aiming at inhibiting the other endocannabinoid-hydrolyzing enzymes should bring exciting new developments in the endocannabinoid field.

CONFLICT OF INTEREST

The authors do not have any conflict of interest.

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