

Histamine  $H_4$  receptor ligands reaching clinical development.

# Major advances in the development of histamine H<sub>4</sub> receptor ligands

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The search for new and potent histamine H<sub>4</sub> receptor ligands is leading to a steadily increasing number of scientific publications and patent applications. Several interesting and structurally diverse compounds have been found, but fierce IP competition for a preferred 2-aminopyrimidine scaffold is becoming apparent. Recent investigations into the role of the histamine H<sub>4</sub>R in (patho)physiology and the use of H<sub>4</sub>R ligands in *in vivo* disease models reveal enormous potential in the field of inflammation and allergy, among others. The development of ligands that display activity at two or more histamine receptor (HR) subtypes is another clinical opportunity that is currently being explored. Taken together, the histamine H<sub>4</sub>R field is gearing up for clinical studies and has the potential to deliver another generation of blockbuster drugs.

### Introduction

The histamine receptor (HR) research field has historically been fuelled by breakthrough discoveries at an interval of 17 years, that is, the discovery of two distinct HR subtypes ( $H_1R$  and  $H_2R$ ) in 1966 [1], the discovery of the  $H_3R$  in 1983 [2] and the identification of the sequence of  $H_4R$  in genome databases in 2000 [3–8]. This subfamily of G-protein-coupled receptors (GPCRs) has obtained a reputation of being very rewarding by the blockbuster status that was reached by  $H_1R$  and  $H_2R$  targeted drugs (for treating allergic conditions and gastric ulcers, respectively). Currently, the pharmaceutical industry is intensively exploring the clinical potential of  $H_3R$  and  $H_4R$  ligands [9,10]. Although the pharmacological effects of the  $H_3R$  were already known soon after its discovery in 1983, this receptor subtype was only embraced by the industry after the cloning of the human  $H_3R$  ( $H_3R$ ) cDNA by Lovenberg *et al.* [11,12]. Following this discovery, several groups identified the  $H_4R$  sequence in human genome databases on the basis of its homology with the  $H_3R$  (31% at the protein level, 54% within the transmembrane domains). To date, two additional non-7-TM  $H_4R$  splice variants have been identified [13]. Interestingly, these splice variants do not seem to bind histaminergic ligands, but affect the functionality of the full-length (functional) human  $H_4R$  ( $H_4R$ ) isoform, probably by GPCR hetero-dimerization [13]. Major species differ-

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#### **Prof Dr Rob Leurs**

obtained his PhD in pharmacochemistry from the VU University Amsterdam in 1991. As a postdoctoral fellow (1992–1993) at INSERM (Unite de Neurobiologie and Pharmacologie, Paris), he was involved in the cloning of genes encoding histaminergic



and serotonergic receptors. Thereafter, he was awarded with a five-year fellowship (1993–1998) of the Royal Netherlands Academy of Arts and Sciences. He was appointed as assistant and full professor in Medicinal Chemistry in 1998 and 2000.

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ences have been identified by the cloning of the genes for mouse, rat, guinea-pig, pig and, more recently, the dog and monkey  $H_4R$  [14,15].

The  $H_4R$  receptor is mainly expressed in bone marrow and peripheral leukocytes, and mRNAs of the  $hH_4$  receptor are detected in for example, mast cells, dendritic cells, spleen and eosinophils [3–7]. The  $H_4$  receptor has pronounced effects on the chemotaxis of several cell types that are associated with immune and inflammatory responses. Based on these early results from  $H_4R$  characterization, both industry and academia initiated an effective search for potent and selective  $hH_4R$  ligands and started to explore the therapeutic potential of such compounds. This has led to a steady output of both  $H_4R$ -related patents and publications (Fig. 1).

# H<sub>4</sub>R agonists

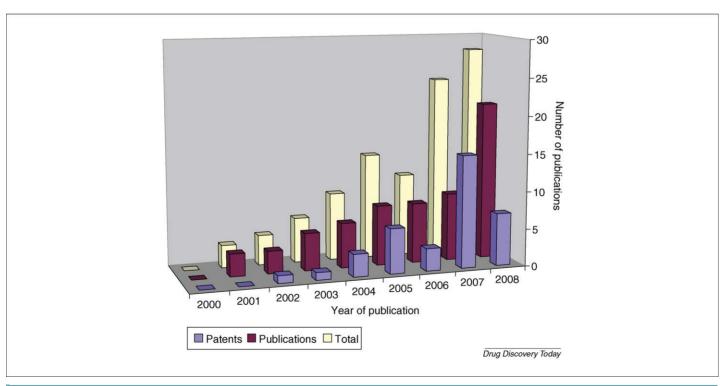
Most imidazole-containing  $H_3R$  ligands that were developed in the 1980s and 1990s turned out to have significant affinity for the  $H_4R$  as well [9]. For example, the  $H_3R$  reference agonist, (R)- $\alpha$ -methylhistamine (**2**), was found to be only 40-fold selective for the  $H_3R$  [9]. Another frequently used  $H_3R$  agonist, immepip (**3**), shows only 40-fold selectivity for the  $H_3R$  [16]. Structural modifications of immepip resulted in the potent and selective  $H_3R$  compounds, immethridine [17] and methimepip (**4**) [16], with the latter having a 2000-fold selectivity at  $H_3R$  over  $H_4R$ , illustrating that small structural changes of the ligands can differentiate binding to the different HR subtypes. Clobenpropit (**5**) was originally designed as a  $H_3R$  antagonist, but was later also identified as a high affinity  $H_4R$ 

partial agonist [18]. The series of clobenpropit compounds have also been used in efforts to quantify the differences in  $H_3R$  and  $H_4R$  binding requirements [18].

The first reported  $H_4R$  agonist with moderate affinity and some selectivity over  $H_3R$  is OUP-16 (6) [19]. Later, at VU University Amsterdam, an in-house screening of histaminergic compounds from  $H_1R-H_3R$  medicinal chemistry programs revealed considerable affinity of 4-methylhistamine (4-MeHA, **7**) for the  $H_4R$  [20]. 4-MeHA acts as a full agonist and is at least 100-fold selective over all other HRs, including the  $H_2R$  for which this compound had originally been developed.

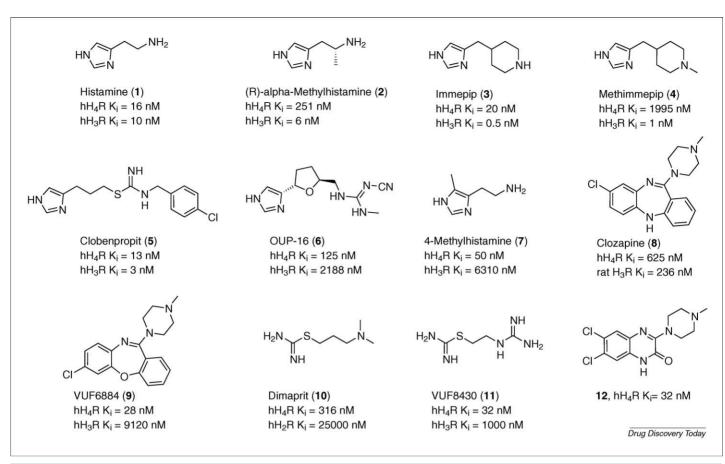
For the closely related H<sub>3</sub>R receptor, all reported agonists have an imidazole ring, but this heterocycle is not necessary for H<sub>4</sub>R agonism, as is illustrated by clozapine (**8**, Fig. 2), an antipsychotic drug used for the treatment of schizophrenia. Being extremely promiscuous in nature, this tricyclic compound binds many GPCRs including dopaminergic, serotonergic, muscarinic and adrenergic receptors [21]. Clozapine (**8**) has moderate affinity for the H<sub>4</sub>R, but some minor structural modifications led to a close analog (VUF6884, **9**) with good H<sub>4</sub>R affinity and 300-fold selectivity over the H<sub>3</sub>R. The rigid structure of this ligand has been used to map the H<sub>4</sub>R active site and to construct a pharmacophore model that was useful in the design of new H<sub>4</sub>R ligands [22,23].

Dimaprit (10) is another non-imidazole  $H_4R$  agonist, albeit with low affinity [20]. This alkylisothiourea was originally developed as a  $H_2R$  agonist but it possesses considerable  $H_4R$  affinity with an 80-fold selectivity for the  $H_4R$  over the  $H_2R$  [20]. Structural modification of 10 to increase its potency at the  $H_4R$  and its selectivity over



FIGURE

Number of  $H_4R$ -related publications (2000–2008). After the discovery of the receptor in 2000,  $H_4R$ -related research has rapidly gained momentum. Both the number of chemistry-related patent applications and publications show a steady annual increase over the past eight years indicating a gradual maturation of  $H_4R$ -related research. A relatively large number of patents covering the 2-aminopyrimidine scaffold was published in 2007, two years after the appearance of the first 2-aminopyrimidine patent application from Bayer Healthcare AG. (Searches were conducted by entering 'histamine H4' in pubmed (http://www.pubmed.com) and esp@cenet (http://www.espacenet.com) after which non- $H_4$ -related hits were removed (status: April 2009).)



#### FIGURE 2

Histamine H<sub>4</sub> receptor agonists and their affinities for the H<sub>3</sub> and H<sub>4</sub> receptors.

the other HR subtypes led to VUF8430 (**11**), a very useful full agonist with about 30-fold selectivity for the  $H_4R$  over the  $H_3R$  [24]. Like dimaprit (**10**), VUF8430 (**11**) originated from an  $H_2R$  medicinal chemistry program [25]. A convenient microwave-assisted synthesis route for the preparation of this compound was also recently reported, making this ligand readily accessible [25].

The quinoxalinone class of  $H_4R$  ligands represented by compound **12** (Fig. 2) has been developed by scientists from J&J Pharmaceutical and was also discovered using a fragment-based drug design approach at VU University Amsterdam [22,23]. Originally developed as 5-HT $_3$  ligands in 1981 [26], the quinoxalin-(1H)2-ones have high affinity for the  $H_4R$ , with compound **12** as the most potent example in the series (hH $_4R$   $K_i$  = 4 nM as determined in our labs and 32 nM as reported in Ref. [27]). Although literature reports no functional data, it was found that quinoxaline **12** behaves as a partial agonist in a  $H_4R$ -driven CRE-beta-galactosidase reporter gene assay (pEC $_{50}$  = 8.99  $\pm$  0.14,  $\alpha$  = 0.55, unpublished data).

# H<sub>4</sub>R antagonists

In the search for novel  $H_4R$  antagonists or inverse agonists, a variety of ligand classes have been identified. An antagonist that was discovered shortly after cloning of the  $H_4R$  gene in 2000 is the imidazole-containing ligand, thioperamide (13). Originally developed as an  $H_3R$  antagonist, thioperamide (13) has an  $H_4R$  affinity that is two to threefold lower than that for the  $H_3R$  while acting as

an H<sub>4</sub>R inverse agonist [20]. The early availability of recombinant systems to measure ligand affinity for H<sub>4</sub>R has had a great impact on the identification of novel classes of compounds. In the pregenomic era, medicinal chemistry programs for the previously discovered H<sub>1</sub>-H<sub>3</sub> receptors were limited with respect to the throughput of ligand screening campaigns and many of the initial reference compounds were based on the endogenous ligand. For the H<sub>4</sub>R, high-throughput screening (HTS) was set up at the very beginning of the discovery efforts. This led to the early identification of the non-imidazole antagonist JNJ7777120 (14, Fig. 3) by Johnson & Johnson (J&J), undoubtedly one of the most active players in the field [28]. This highly potent indole carboxamide has a  $K_i$  of 4 nm and behaves as a neutral antagonist, although a debate on the exact functional behavior of this compound is ongoing. Recently it was shown that 14 can also behave as a partial inverse agonist (EC<sub>50</sub> = 37.7  $\pm$  8.5 nm,  $\alpha$  = -0.31  $\pm$  0.07) in a steady-state GTPase assay [29]. Nevertheless, the compound has been used effectively in a variety of animal models of inflammatory disease and is so far considered the reference antagonist of choice to study the H<sub>4</sub>R. Although **14** has reasonable oral bioavailability (22%), the compound suffers from a short in vivo half life in rats (0.8 hours) [30]. Therefore, its use in prolonged in vivo studies (e.g. models of chronic inflammation and asthma) will require either frequent administration or high doses to maintain a therapeutically effective plasma concentration. Such high doses might compromise the relatively good selectivity profile of this compound, in particular its selectivity over the  $H_3R$  ( $hH_3R$   $pK_i = 5.3$ ) [31].

#### FIGURE 3

Selected H<sub>4</sub>R ligands developed by J&J together with thioperamide (13) and benzimidazole 17 from Pfizer Inc.

Based on JNJ7777120 (14), J&J developed a variety of structurally diverse compound classes. Benzimidazole 15 and thienopyrrole 16 (Fig. 3) analogs were prepared as bioisosteres of the indole carboxamides [30,32]. Both 15 and 16 have affinities close to that of indole 14 and showed somewhat improved metabolic stability *in vitro*. *In vivo* studies of 15, however, revealed similar stability problems and compound 16 lacked oral bioavailability altogether. Scientists from Pfizer Inc. have successfully replaced the *N*-methylpiperazine moiety of 15 with an octahydropyrrolo[3,4-c]pyrrole bioisostere (compound 17, Fig. 3) [33]. In addition, this series contains several potent analogs that contain an amidine group that connects the benzimidazole group with the octahydropyrrolo[3,4-c]pyrrole moiety (e.g. compound 17).

The HTS efforts of J&J also resulted in another, structurally distinct, benzimidazole-containing hit (**18**) [34,35]. A very extensive series of compounds was synthesized and described in six patents that cover a wide variety of alterations of the structures illustrated in Fig. 3 (e.g. structures **18** and **19**) [36–39]. In these compounds, the benzimidazole scaffold is attached to a substituted phenyl or heterocycle which, in turn, is connected by a flexible alkyloxy or alkyl amine chain to a piperidine or (homo)-piperazine moiety. These compounds are claimed to have high affinity for the  $H_4R$ , in our hands a potent analog was found to behave as a full agonist in a functional  $H_4R$  assay. A closely related series of compounds in which the benzimidazole group could be replaced by a di-phenyl substituted imidazole (compound **20**) has

#### FIGURE 4

Quinoxaline and quinazoline compounds developed at VU University Amsterdam.

also been discovered [40]. The potencies of these ligands are, however, a bit lower than those found for the benzimidazolebased series.

Using a pharmacophore model constructed from the antipsychotic drug clozapine and the reference H<sub>4</sub>R antagonist JNJ7777120 (14), the benzyl substituted quinoxaline 21 (Fig. 4) was developed [23]. By combining the SAR of the quinoxaline series and the pharmacophore model, a series of quinazoline compounds (e.g. compound 22) was designed in a successful scaffold hopping exercise [41]. The thiophene moiety of compound 22 was replaced by a sulfonamide group, as illustrated by compound 23. This replacement gave a series of compounds that tolerates a variety of substituents on the sulfonamide moiety while retaining excellent affinity for the H<sub>4</sub>R (R. Smits, Design and synthesis of new histamine H4 receptor ligands, PhD thesis, VU University Amsterdam, 2009, ISBN: 987-90-8891-091-3).

Bayer Healthcare AG was the first to patent two series of aminopyrimidines as  $H_4R$  antagonists (e.g. **24** and **25**, Fig. 5) [42,43]. Exact IC<sub>50</sub> values of these new compounds are not mentioned in the patents, but it was indicated that the affinities of the most active compounds are below 20 nm. Since the appearance of the first aminopyrimidine patents from Bayer Healthcare AG, other companies have eagerly used the 2-aminopyrimidine scaffold to develop their own H<sub>4</sub>R ligand series. Tolerance to the introduction of a wide variety of substituents on the aminopyrimidine scaffold is generally found on the 5- and 6-positions (see also Fig. 5). In most patents the 2-amino group is left unsubstituted and the Nmethylpiperazine moiety is usually among the most potent analogs. The N-methylpiperazine group can, however, be replaced with other amines with the retention of good affinity. The tolerated chemical variation of this basic side-chain is somewhat limited and the most important reason appears to be a restriction of size. On other scaffolds, the piperazine group is very intolerant

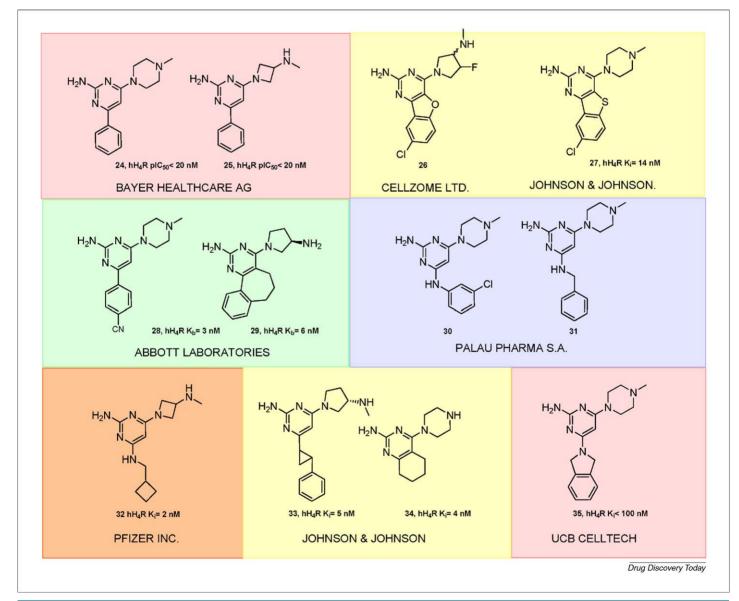


FIGURE 5

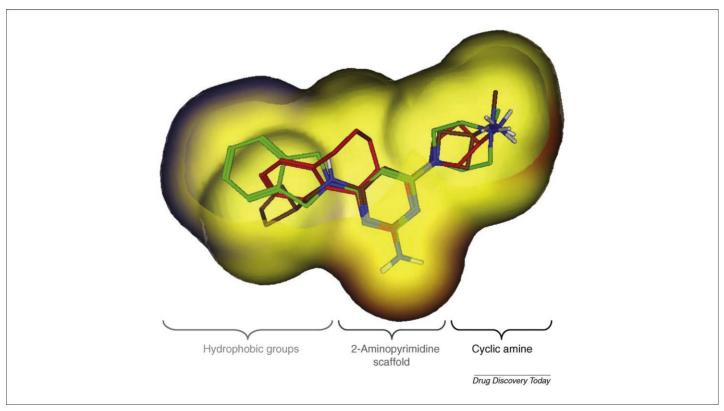
Selected patent-protected compounds that share the privileged aminopyrimidine scaffold as a common element for the H<sub>4</sub>R.

to substitution with anything much larger than a methyl or ethyl group, suggesting a similar restriction of the size of the basic sidechain [22,28]. Initially, *N*-methylpiperazine replacements consisted of azetidine (e.g. compound **25**), aminopyrrolidine or diazepane rings substituted with small aliphatic groups.

An interesting series of derivatives that has appeared in patent literature contains a benzofused aminopyrimidine scaffold (e.g. compound 26) [44]. This series comprises a phenyl ring that is directly connected to a 2-aminopyrimidine ring. Such a scaffold is very similar to the first aminopyrimidine series claimed by Bayer Healthcare AG (compare compound number 24 with compound number 26) [42,43]. To gain a novel IP position, however, the aromatic phenyl and pyrimidine rings were fused with an oxygen atom to give the conformationally constrained analog 26. After the initial discovery of the benzofuropyrimidines by Argenta Discovery [44], this series has now been taken into development by Cellzome Ltd. Much attention has been paid to alteration of the Nmethylpiperazine moiety, resulting in a large number of azetidine and aminomethylpyrrolidine analogs of compound 26 [45-51]. Cellzome Ltd. has announced that they have developed a very potent hH<sub>4</sub>R inverse agonist (CZC-13788, undisclosed structure) that had been planned to enter phase I clinical trials in 2008 for the treatment of allergic rhinitis [52]. This compound has a  $K_i$  of 0.17 nM for the human H<sub>4</sub>R and is claimed to be at least 10,000-fold selective over the other HR subtypes [52].

J&J has also directed some effort toward the aminopyrimidine scaffold after the initial development of several of the compound classes described earlier (Fig. 3). The aminopyrimidines claimed are based on the fused aminopyrimidines (e.g. compound **27**) and are very similar to the compounds claimed by Cellzome Ltd. [53].

Following an HTS campaign, medicinal chemistry efforts by Abbott Laboratories gave compound **28** as a high affinity  $H_4R$  ligand with good oral availability ( $F_{po}/_{iv} = 31\%$  in rats) and active in various animal models (zymosan-induced peritonitis, carrageenan-induced hyperalgesia and murine itch), despite its decreased potency at the rat  $H_4R$  (rH4R p $K_i = 6.56 \pm 0.06$ ) [54]. A SAR study of **28** that focused on the replacement of the 4-cyanophenyl group failed to yield a compound with improved potency at the rat  $H_4R$ . By using a strategy that has also been used by scientists from J&J and Cellzome Ltd. (compounds **26** and **27** Fig. 5), a series of conformationally constrained aminopyrimidines was prepared that retain high  $H_4R$  potency across the species [55]. It has been reported that some fused cyclohepta[1,2-d]pyrimidine compounds (e.g. A-943931 (**29**)) bind the  $H_4R$  in the low nanomolar



# FIGURE 6

Pharmacophore for 2-aminopyrimidine H<sub>4</sub>R ligands. The affinity H<sub>4</sub>R ligands that use the 2-aminopyrimidine template is based on three structural elements. The first is the 2-aminopyrimidine moiety that connects the other two elements by the substitution of the aminopyrimidine 4- and 6-positions. The second element consists of hydrophobic aromatic or aliphatic groups. These hydrophobic groups can be branched, straight, or fused to the pyrimidine core (e.g. compound 29, in red). There appears to be great tolerance for structurally diverse substituents as illustrated by the large number of patent applications covering the 2-aminopyrimidine scaffold. The third element is a cyclic amine of which the most commonly encountered example is the *N*-methylpiperazine (NMP) group (e.g. compound 35 in green). On the 2-aminopyrimidine scaffold, the NMP moiety can be substituted with small heterocyclic aliphatic rings that contain a basic nitrogen atom (e.g. azetidine 32 in brown). Substituents in this area are restricted by size and can usually not be much larger than one six-membered or two smaller fused rings. (The model was generated with MOE2007.0902. Compounds 29, 32 and 25 were superimposed with the flexible alignment module. Then the 4.5 ÅvdW interaction surface was calculated as a representation of the H<sub>4</sub>R binding site. Hydrophobic surface is shown in yellow, hydrogen bonding surface in red and mild polar surface in blue).

range and are equipotent at rat and mouse  $\rm H_4Rs$ . In the rat, A-943931 (**29**) combines good oral bioavailability ( $F_{\rm po/iv} = 34\%$ ) with an improved *in vivo* half-life ( $t_{1/2} = 2.6$  hours compared to  $t_{1/2} = 0.8$  hours for the widely used antagonist JNJ7777120 (**14**)). Although the selectivity of JNJ7777120 (**14**, 1250-fold) for  $\rm H_4R$  over  $\rm H_3R$  is slightly better than that of A-943931 (**29**, 640-fold), the improved pharmacokinetic profile of A-943931 (**29**) allows for less frequent administration and lower dosing in animal models that require chronic dosing.

An aminopyrimidine patent filed by Palau Pharma covers several compounds that are closely related to the initial 2-aminopyrimidines from Bayer Healthcare AG [56]. In these compounds (e.g. **30** and **31**), the phenyl group has not been directly attached to the pyrimidine heterocycle but instead an amino group is used as a linker between both aromatic groups. An elaborate series was claimed with various aromatic substituents, consisting primarily of substituted phenyl groups, but thiophene, naphthalene and benzyl groups were also included. In the patent application it was noted that a 10 µM concentration of the ligands displaced at least 50% of [<sup>3</sup>H]histamine binding [56]. Another patent application covering the aminopyrimidine scaffold has been filed by Pfizer Inc. [57]. Although these compounds are very closely related to the diaminopyrimidines from Palau Pharma, the focus in this series lies on aliphatic substituents (e.g. compound 32). Where most of the previously patented aminopyrimidines contained a second aromatic group, most Pfizer Inc. compounds have straight or branched aliphatic groups. Apparently, the H<sub>4</sub>R affinity can be increased by the introduction of hydrophobic groups that are either aromatic or aliphatic in nature, both leading to very potent compounds in the low nanomolar range (Fig. 6).

A recent addition to the  $H_4R$  patent literature is a series of structurally diverse aminopyrimidine compounds from J&J [58]. This includes cyclopropylphenyl substituted pyrimidine (**33**) and a tetrahydroquinazoline (**34**). Together with the previously discussed aminopyrimide ligands, compounds **33** and **34** yet again demonstrate the great tolerance of the  $H_4R$  to a wide variety of hydrophobic substituents. Exploiting the tolerance to chemical diversity, UCB Celltech has prepared a series of aminopyrimidines that includes several compounds with (partially)saturated heterocyclic rings attached to the 6-position (e.g. compound **35**) [59]. The point of attachment can not only be the nitrogen atom as exemplified by compound **35**, but also be a carbon atom of a cyclic substructure, such as a pyrrole or piperidine ring. These compounds have been reported to bind the  $H_4R$  in the nanomolar range.

# Clinical applications for H<sub>4</sub>R antihistamines

 $\rm H_4R$  agonists and antagonists have an important role to play in the elucidation of the (patho)physiological role of the  $\rm H_4R$  and to explore the therapeutic potential of this receptor in human disease. The use of  $\rm H_4R$  ligands in animal models, so far mostly done with reference antagonist JNJ7777120 (**14**), has yielded a variety of interesting results.

# **Pruritis**

Recently the role of the  $H_1R$  and  $H_4R$  in the attenuation of pruritis (itch) has been reported in several papers [60–62]. In mice, the selective  $H_4R$  agonist 4-MeHA (7) was shown to induce itch in a

dose-dependent way and this effect could be blocked by the pretreatment with JNJ7777120 (**14**) [61]. The antipruritic effect of **14** was far superior to other antihistamines and only the centrally acting  $H_1R$  antagonist diphenhydramine showed some antipruritic activity. This pruritic response could not be abolished completely by either an  $H_1R$  or  $H_4R$  antagonist alone, but when both were administered simultaneously, almost no scratching behavior was observed. This observation clearly implicates both HR subtypes in the pruritic response and suggests that in pruritis, simultaneously antagonizing both the  $H_1R$  and  $H_4R$  might have added clinical benefit over  $H_1R$  or  $H_4R$  monotherapy.

#### **Asthma**

A role for the  $H_4R$  in a murine ovalbumin-induced inflammation model of asthma was recently found by Dunford *et al.* [63]. It was demonstrated that the  $H_4R$  is involved in the activation of  $CD4^+$  cells by dendritic cells. The administration of JNJ7777120 (**14**) showed significant anti-inflammatory responses during both the sensitization and effector phases. This finding indicates that the  $H_4R$  is involved in the initial priming of the immune system after allergen challenge. More importantly, the administration of a  $H_4R$  antagonist is also efficacious after the animals had been sensitized for ovalbumin. This discovery is of great interest because, a therapeutic intervention will only be done after a disease has become manifest [63].

# **Allergic rhinitis**

The presence of the H<sub>4</sub>R in nasal tissue was first discovered by Nakaya et al. [64]. In addition, a more recent finding showed that there is a significant increase in the level of H<sub>4</sub>R in human nasal polyp tissue taken from patients with chronic rhinosinusitis (infection of the nose and nasal cavities) when compared to normal nasal mucosa [65]. Jókúti et al. suggest that the administration of H<sub>4</sub>R antagonists might be a new way to treat nasal polyps and chronic rhinosinusitis. The administration of H<sub>4</sub>R antagonists may prevent the accumulation of eosinophils as a result of impaired cell chemotaxis toward polypous tissue [65]. Although scientific data on the role of the H<sub>4</sub>R in rhinitis is limited, at present, it is the only indication for which an H<sub>4</sub>R inverse agonist (CZC-13788) is reported to be in preclinical development [52]. Apparently, histamine is implicated in allergic rhinitis by acting on three HR subtypes, the H<sub>1</sub>R, H<sub>3</sub>R and H<sub>4</sub>R [66]. For many years, the classical application of H<sub>1</sub>R antagonists (antihistamines) has been the treatment of allergic rhinitis. H<sub>1</sub>R antagonists relieve edema and vasoconstriction, both important symptoms of the disease, but these drugs do not affect the underlying inflammatory responses [66]. After the discovery of the H<sub>3</sub>R and H<sub>4</sub>R subtypes, the traditional role for H<sub>1</sub>R antagonists in rhinitis has been reappraised [66]. It has been shown that the  $H_3R$  agonist (R)- $\alpha$ -methylhistamine (2) can induce the dilatation of nasal blood vessels and that this effect can be counteracted by the H<sub>3</sub>R antagonist/H<sub>4</sub>R agonist clobenpropit (5) [66]. Although a role for the H<sub>4</sub>R cannot be ruled out, this H<sub>3</sub>R antagonist-mediated mechanism in nasal decongestion has certainly caught the attention of scientists from Pfizer Inc. Recently, patient recruitment started for a Phase II clinical trial to test a H<sub>3</sub>R antagonist (PF-03654746, unpublished structure) as a novel nasal decongestant in patients with seasonal allergic rhinitis (http://www.clinicaltrials.gov). A dual target

approach is being pursued by GSK that is currently recruiting patients to test a systemic H<sub>1</sub>/H<sub>3</sub> antagonist (GSK835726, unpublished structure) for seasonal allergic rhinitis in a Phase I clinical trial (http://www.clinicaltrials.gov). A second Phase I trial with another H<sub>1</sub>/H<sub>3</sub> antagonist (GSK1004723, unpublished structure) for intranasal administration to treat rhinitis has recently been completed (http://www.clinicaltrials.gov). With these compounds, the mode of action of the classical H<sub>1</sub>R antagonist is combined with the potential clinical benefit of added nasal decongestion by H<sub>3</sub>R blockade. The synergistic role of the H<sub>1</sub>R and H<sub>3</sub>R has been demonstrated in vivo in experiments performed at Schering-Plough [67]. In view of the role of the H<sub>4</sub>R in allergic rhinitis, other potential treatment paradigms may also be considered, such as combining H<sub>1</sub>/H<sub>4</sub>, H<sub>3</sub>/H<sub>4</sub> or even H<sub>1</sub>/H<sub>3</sub>/H<sub>4</sub> antagonists/inverse agonist activity in the same molecule.

#### **Pain**

In 2007 it was suggested that the H<sub>4</sub>R might play a role in nociception [68]. The H<sub>4</sub>R antagonist JNJ7777120 (14) and its benzimidazole analog, VUF6002 (15), were both able to increase paw withdrawal latency in carrageenan-induced thermal hyperalgesia in rats [68]. This effect on the modulation of pain was recently confirmed by work from Abbott Laboratories that describes in vivo anti-nociceptive effects of the H<sub>4</sub>R antagonist A-943931 (39) [55]. This compounds is active against carrageenaninduced inflammatory pain and in a spinal nerve ligation model of neuropathic pain. The exact location of the H<sub>4</sub>Rs that are responsible for the observed antinociceptive effects remains unknown, although very recently new evidence for the presence of the H<sub>4</sub>R in the spinal cord and dorsal root ganglion has been reported [69]. The encouraging results seen with H<sub>4</sub>R antagonists in vivo may offer a new way to relieve pain that is unresponsive to existing therapies.

# Cancer

Although histamine has been known to play a role in the development of cancer, a role for the H<sub>4</sub>R has only recently been suggested on the basis of in vitro studies with HT29 and Caco-2 cells [70]. Histidine decarboxylase is often overexpressed in human colorectal cancer cells, resulting in an increased local production of histamine from the amino acid histidine. The selective H<sub>4</sub>R antagonist JNJ7777120 (14) was able to inhibit histamine-induced overexpression of cyclooxygenase-2 and subsequent production of PGE<sub>2</sub>. A similar finding was reported for H<sub>2</sub>R activation, which can be blocked by the H<sub>2</sub>R antagonist zolantidine [70]. PGE<sub>2</sub> is an

important mediator of proliferative and proangiogenic activity of cells. Activation of the H<sub>2</sub>R and H<sub>4</sub>R receptors may, therefore, explain the proliferative effect of histamine on HT29 and Caco-2 cells. A different relationship between the H<sub>4</sub>R and cancer has been proposed by Boer et al. [71]. It was found that human colon carcinoma cells had significantly reduced levels of H<sub>1</sub>R and H<sub>4</sub>R, but unchanged H<sub>2</sub>R levels, possibly causing a microenvironment that is favorable for tumor development. The low abundance of H<sub>1</sub>R and H<sub>4</sub>R leads to a more dominant role of H<sub>2</sub>R-mediated negative regulation of the Th<sub>1</sub> and Th<sub>2</sub> immune response against the carcinoma cells. In addition, H<sub>2</sub>R-mediated angiogenesis and metastatic spreading may also lead to malignant tumor formation [71]. Further studies in this direction will be conducted with H<sub>4</sub>R and H<sub>1</sub>R knock-out mice and results in this area are eagerly awaited.

# Inflammatory bowel disease

In an experimental model of colitis in the rat, the oral administration of a high dose of the H<sub>4</sub>R antagonists JNJ7777120 (14) or VUF6002 (15) was able to reduce (TBNS)-induced colon damage, the influx of neutrophils and levels of myeloperoxidase in colonic tissue [72]. Although several of the underlying physiological mechanisms need to be fully explored, these preliminary findings suggest a potential role for H<sub>4</sub>R antagonists in inflammatory bowel disease.

# **Conclusions and outlook**

Since the discovery of the H<sub>4</sub>R in 2000 the amount of publications and patent applications has shown a steady annual increase. Several useful selective H<sub>4</sub>R agonists and antagonists have been developed and now serve as important tools to delineate the role of the H<sub>4</sub>R in physiology and to establish its potential value as a drug target. The recent increase in patent applications from various players in the pharmaceutical industry clearly reflects the mounting interest in this new HR subtype as a drug target.

Several histamine-induced events are caused by the activation of multiple HR subtypes. In some diseases, such as allergic rhinitis and pruritis, these insights may offer new treatment paradigms with H<sub>4</sub>R antagonists that have affinity for multiple HR subtypes to provide added clinical benefit compared to single receptor selective histaminergic ligands. The development of compounds that display potency at the panel of HR subtypes can be considered an exciting opportunity to synergistically increase efficacy in vivo or for treating multiple histamine-induced symptoms of a single disease.

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