Bioluminescence Resonance Energy Transfer Based G Protein-Activation Assay to Probe Duration of Antagonism at the Histamine H₃ Receptor

Tamara A. M. Mocking, Maurice C. M. L. Buzink, Rob Leurs and Henry F. Vischer

Amsterdam Institute for Molecules, Medicines and Systems (AIMMS), Division of Medicinal Chemistry, Faculty of Science, Vrije Universiteit Amsterdam, De Boelelaan 1108, 1081 HZ Amsterdam, The Netherlands
* Correspondence: h.f.vischer@vu.nl

Received: 8 July 2019; Accepted: 28 July 2019; Published: 30 July 2019

Abstract: Duration of receptor antagonism, measured as the recovery of agonist responsiveness, is gaining attention as a method to evaluate the ‘effective’ target-residence for antagonists. These functional assays might be a good alternative for kinetic binding assays in competition with radiolabeled or fluorescent ligands, as they are performed on intact cells and better reflect consequences of dynamic cellular processes on duration of receptor antagonism. Here, we used a bioluminescence resonance energy transfer (BRET)-based assay that monitors heterotrimeric G protein activation via scavenging of released Venus-Gᵦ₁γ₂ by NanoLuc (Nluc)-tagged membrane-associated-C-terminal fragment of G protein-coupled receptor kinase 3 (masGRK3ct-Nluc) as a tool to probe duration of G protein-coupled receptor (GPCR) antagonism. The Gᵦᵢ-coupled histamine H₃ receptor (H₃R) was used in this study as prolonged antagonism is associated with adverse events (e.g., insomnia) and consequently, short-residence time ligands might be preferred. Due to its fast and prolonged response, this assay can be used to determine the duration of functional antagonism by measuring the recovery of agonist responsiveness upon washout of pre-bound antagonist, and to assess antagonist re-equilibration time via Schild-plot analysis. Re-equilibration of pre-incubated antagonist with agonist and receptor could be followed in time to monitor the transition from insurmountable to surmountable antagonism. The BRET-based G protein activation assay can detect differences in the recovery of H₃R responsiveness and re-equilibration of pre-bound antagonists between the tested H₃R antagonists. Fast dissociation kinetics were observed for marketed drug pitolisant (Wakix®) in this assay, which suggests that short residence time might be beneficial for therapeutic targeting of the H₃R.

Keywords: histamine H₃ receptor (H₃R); G protein-coupled receptor (GPCR); re-equilibration; ligand binding kinetics; residence time

1. Introduction

Drug-target binding kinetics has established its relevance in drug discovery as target residence time of a drug might better predict its in vivo effect, as compared to its equilibrium binding affinity (pKᵢ) for the target [1]. Residence time is the reciprocal of the dissociation rate constant (kₒff) of a receptor-bound drug, which has mainly been derived from competition binding assays using the Motulsky and Mahan method [2]. Recently, a bioluminescent resonance energy transfer (BRET)-based binding assay on living cells using a Nanoluc (Nluc)-tagged receptor and fluorescent tracer has been introduced for determination of kinetics parameter of unlabeled ligands [3–5]. However, the Motulsky and Mahan analysis to estimate kinetic binding parameters for unlabeled ligands in these competitive binding assays was found to be sensitive to the used labeled probe [6]. Alternatively, functional assays
can be employed to estimate the duration of receptor occupancy by antagonists [7,8]. In contrast to binding assays that are routinely performed with cell homogenates or isolated membranes, functional assays are performed on intact living cells and, consequently also account for dynamic cellular processes such as effector pre-coupling, receptor reserve and receptor regulation. Fast and/or real time cellular responses such as Ca\(^{2+}\) influx, cyclic adenosine-monophosphate (cAMP) production and cellular impedance-based assays, have recently been used to measure the recovery of receptor responsiveness as measure of pre-bound antagonist dissociation upon washout of unbound antagonist [3,7,9–12]. Indeed, recovery time of G\(\alpha_{q}\)-coupled histamine H\(_1\) receptor (H\(_1\)R) responsiveness in a Ca\(^{2+}\) mobilization and label free dynamic mass redistribution (DMR) assay after antagonist washout highly correlated to residence times of these antihistamines, as determined in competitive radioligand binding assays [7]. Moreover, response recovery determined utilizing washout experiments corresponds well to the in vivo duration of neurokinin 1 receptor antagonism [12]. Alternatively, estimation of duration of antagonism can be monitored in real-time, utilizing a BRET-based \(\beta\)-arrestin 2 recruitment assay, following co-addition of agonist and antagonist resulting in overshoot-patterns similar to competition association binding assays [13]. In addition, a BRET sensor that monitors cAMP accumulation could effectively determine receptor recovery time (RecT) after washout of pre-incubated antagonist and RecT correlated to residence times from competition binding assays [3]. However, the delayed response of this biosensor upon agonist stimulation of the G\(\alpha_{i}\)-coupled receptor hampers quantification of kinetic parameters for relatively fast dissociating ligands.

In this study, we employed a BRET-based assay to measure human histamine H\(_3\) receptor (hH\(_3\)R)-induced G\(\alpha_{i2}\)-protein activation by measuring the very rapid recruitment of Venus-G\(\beta_{1}\)G\(\gamma_{2}\) to a membrane-associated-C-terminal fragment of G protein-coupled receptor kinase 3 (i.e., masGRK3ct-Nluc) that scavenges released G\(\beta\)\(\gamma\) complexes with high affinity [14,15].

The G protein-coupled receptor (GPCR) H\(_3\)R is a G\(\alpha_{i}\)-coupled receptor that functions as pre-synaptic auto-and heteroreceptor and thereby regulates the release of histamine and various other neurotransmitters in the brain [16]. Due to this prominent role in the central nervous system the H\(_3\)R has been associated with a variety of neuropsychiatric disorders such as Parkinson’s disease, epilepsy, learning and sleeping disorders. Although several compounds targeting this receptor have entered clinical trials for different indications, often they are withdrawn in early stages of clinical trials because they are either ineffectual or induce side effects like insomnia [17–19]. Many compounds already fail in pre-clinical stages, e.g., ABT-239 due to hERG mediated cardiac toxicity and imidazole containing ligands, like iodophenpropit, due to poor penetration of the blood brain barrier and interference with CYP enzymes. A promising drug candidate as PF03654746 has been tested for several indications (a.o. attention deficit hyperactivity disorder (ADHD), Alzheimer’s disease, allergic rhinitis and schizophrenia), but was discontinued without disclosure of results [20]. Yet, recently pitolisant (Wakix®) has been approved [21] by the European Medicines Agency in 2016 for the treatment of narcolepsy to improve wakefulness in patients suffer from excessive daytime sleepiness [22]. In the US, pitolisant has received Breakthrough Therapy and Fast Track designations from the Food and Drug Administration and approval to market this new medication in the United States in 2019 has been requested [23].

Ligands with a short duration of action might be beneficial for the H\(_3\)R to minimize the side effects associated with prolonged inhibition of H\(_3\)R signaling due to its prominent role in neurotransmission. H\(_3\)R antagonists PF03654746, pitolisant and iodophenpropit were assessed for their duration of H\(_3\)R antagonism in a newly developed functional assay format, using living cells. Differences in the recovery of receptor responsiveness and pre-bound antagonists re-equilibration could be observed between the tested H\(_3\)R antagonists.
2. Results

2.1. BRET between Venus-Gβ1γ2 and masGRK3ct-Nluc to Measure hH3R-Induced Gαi2 Activation

Heterotrimeric G protein activation by GPCRs can be monitored as the release of Venus-tagged Gβ1γ2 subunits and their subsequent recruitment toward the membrane-associated Gβγ-scavenger masGRK3ct-Nluc, which increases the BRET signal [14]. Stimulation of adherent HEK293T cells that were co-transfected with Venus-Gβ1γ2, masGRK3ct-Nluc, hH3R and Gαi2 cDNA with 1 µM histamine induced a rapid increase in BRET between Venus-Gβ1γ2 and masGRK3ct-Nluc (Figure 1A), whereas no significant BRET change was observed in cells that were not co-transfected with Gαi2 cDNA (p = 0.5193, one-way ANOVA). This confirms previous observations that co-expression of exogenous Gα proteins is required for appropriate localization of Venus-Gβ1γ2 [14]. Histamine did also not change BRET between Venus-Gβ1γ2 and masGRK3ct-Nluc in cells that were not transfected with hH3R cDNA (p = 0.8884, one-way ANOVA), demonstrating that the observed Gβ1γ2 release is indeed mediated via the hH3R (Figure 1A).

![Figure 1](https://example.com/figure1.png)

**Figure 1.** Human H3R-mediated Gαi2 activation in HEK293T cells measured as bioluminescence resonance energy transfer (BRET) ratio between Venus-Gβ1γ2 and masGRK3ct-Nluc. (A) Real time BRET ratio in HEK293T cells expressing hH3R and/or Gαi2 upon stimulation with 1 µM histamine at t = 0. (B) Concentration-dependent increase in BRET ratio in cells co-expressing hH3R and Gαi2 upon stimulation with H3R agonists. (C) BRET ratio in cells co-expressing hH3R and Gαi2 upon stimulation with 1 µM histamine in the presence of 10 µM histamine receptor subtype-selective antagonists. (D) Decrease in histamine-induced (1 µM) BRET ratio in HEK293T cells co-expressing Gαi2 in the presence of increasing concentrations H3R antagonists. Representative graphs of 3 experiments performed in triplicate are shown and data are mean ± SD. BRET ratio was measured after 10 min incubation (B–D). BRET ratios are corrected for vehicle. Significance was determined by a Student’s t-test *** (p < 0.001).

The known H3R agonists histamine, imetit, immepip and VUF8328 all induced a concentration-dependent increase in BRET ratio between Venus-Gβ1γ2 and masGRK3ct-Nluc in cells that co-express hH3R and Gαi2 (Figure 1B) with potencies (pEC50) and intrinsic activities (α) that are comparable to those reported in a cAMP-responsive element (CRE)-driven reporter gene assay or [3H]-cAMP accumulation assay (Table 1) [24,25]. Histamine, imetit, and immepip acted as full agonist in the Gβ1γ2 release assays, whereas VUF8328 acted as partial agonist (Figure 1B).
Table 1. Potencies, intrinsic activities ($\alpha$) of ligands in hH$_3$R-mediated G$_{\alpha i2}$ activation in HEK293T cells measured as recruitment of Venus-G$\beta_1 \gamma_2$ to masGRK3ct-Nluc. Data shown are mean ± S.E.M. of three experiments performed in triplicate.

| Compound    | pEC50 ± S.E.M. | $\alpha$ ± S.E.M. | pKi $^a$  |
|-------------|----------------|-------------------|-----------|
| Histamine   | 8.3 ± 0.0      | 1.0 ± 0.0         | 6.7       |
| Imetit      | 9.6 ± 0.1      | 1.1 ± 0.1         | 8.6       |
| immepip     | 9.9 ± 0.1      | 1.0 ± 0.1         | 8.8       |
| VUF8328     | 9.4 ± 0.1      | 0.7 ± 0.2         | 8.5       |

$^a$ pKi values obtained from previously reported competition binding studies [3,24].

The H$_3$R antagonist pitolisant (10 $\mu$M) fully inhibited Venus-G$\beta_1 \gamma_2$ recruitment to masGRK3ct-Nluc in response to 1 $\mu$M histamine, while the H$_1$R, H$_2$R and H$_4$R antagonists (10 $\mu$M) mepyramine, tiotidine and JNJ7777120, respectively, were ineffective, confirming that the observed BRET change is indeed hH$_3$R dependent (Figure 1C). Moreover, the H$_3$R antagonists iodophenpropit, PF03654746, and pitolisant, inhibited histamine-induced G$\beta_1 \gamma_2$ release in a concentration-dependent manner and yielding pK$_B$ values that are comparable to their pK$_i$ values in a previously reported NanoBRET-based competition binding assay on intact cells that express Nluc-hH$_3$R (Figure 1D and Table 2) [3].

Table 2. Equilibrium dissociation constants (pK$_B$), pA$_2$-values and slopes from Schild-regression and recovery time of ligands in hH$_3$R-mediated G$_{\alpha i2}$ activation in HEK293T cells measured as recruitment of Venus-G$\beta_1 \gamma_2$ to masGRK3ct-Nluc. Data shown are mean ± S.E.M. of three experiments performed in triplicate except for RecT, which was measured in singlo.

| Compound      | pK$_B$ $^a$ | pK$_i$ $^b$ | Imetit 0 min | VUF8328 0 min | VUF8328 10 min | RecT $^c$ (min) |
|---------------|------------|------------|-------------|--------------|---------------|-----------------|
|               | pA$_2$ | Slope     | pA$_2$ | Slope | pA$_2$ | Slope | pA$_2$ | Slope |
| Iodophenpropit| 9.5 ± 0.1 | 9.4       | 9.6 ± 0.0 | 1.1 ± 0.0 | 9.3 ± 0.2 | 1.1 ± 0.1 | 9.5 ± 0.3 | 1.1 ± 0.1 | 1.4 ± 0.2 |
| Pitolisant    | 9.1 ± 0.0 | 8.9       | 8.9 ± 0.1 | 1.1 ± 0.0 | 8.4 ± 0.0 | 1.4 ± 0.0 | 8.7 ± 0.2 | 1.1 ± 0.1 | 1.7 ± 0.3 |
| PF03654746   | 9.9 ± 0.1 | 9.3       | 9.8 ± 0.1 | 1.0 ± 0.1 | 9.3 ± 0.2 | 1.5 ± 0.2 | 9.6 ± 0.2 | 1.2 ± 0.1 | 3.6 ± 0.3 |

$^a$ pK$_B$ values are calculated from IC$_{50}$ values, $^b$ pK$_i$ values obtained from previously reported competition binding studies [3], $^c$ RecT values are calculated as 1/k$_{rec}$.

2.2. Duration of Functional hH$_3$R Antagonism

The very short time span to achieve maximal Venus-G$\beta_1 \gamma_2$ recruitment to masGRK3ct-Nluc upon agonist H$_3$R stimulation allows the detection of duration of antagonism by measuring the recovery of hH$_3$R responsiveness to agonist stimulation over time. To this end, hH$_3$R was blocked by pre-incubating the cells with antagonist (10 × IC$_{50}$ concentration) for 1 h, followed by the removal of unbound antagonist to initiate dissociation from the hH$_3$R. Cells were then stimulated with a saturating concentration of imetit (10 $\mu$M) after different time intervals and BRET changes between Venus-G$\beta_1 \gamma_2$ and masGRK3ct-Nluc were measured directly and BRET ratio 10 s after agonist stimulation were plotted (Figure 2A,B). The imetit-induced response recovered to a similar steady-state level upon washout of all three antagonists, which was slightly lower as compared with control cells that were pre-incubated with vehicle (Figure 2B). The hH$_3$R recovery time was comparable upon washout of iodophenpropit and pitolisant, while hH$_3$R recovery was 2 to 2.5-fold slower for PF03654746 (Table 2).
However, allowing pre-bound antagonists to re-equilibrate with VUF8328 for 10 min, the Schild regression slopes were not significantly different from unity for iodophenpropit-mediated inhibition of imetit- and VUF8328-induced hH3R signaling, indicating that iodophenpropit acted as competitive antagonist (Figure 3D). In contrast, slopes of Schild-regression for PF03654746 and pitolisant were larger than 1 (slope (0 min): 1.5 ± 0.2 and slope (0 min): 1.4 ± 0.0, respectively) if hH3R antagonism for pitolisant and PF03654746, respectively (Figure 3H,L). In contrast, iodophenpropit displayed surmountable antagonism by shifting the imetit and VUF8328 concentration-response curves (Figure 3F,G,J,K). However, allowing pre-bound antagonists to re-equilibrate with VUF8328, as revealed by the reduced maximum and rightward-shift of the concentration-response curves (Figure 3F,G,J,K). However, allowing pre-bound antagonists to re-equilibrate with VUF8328 for 10 min before measuring receptor response resulted in surmountable and near-surmountable antagonism for pitolisant and PF03654746, respectively (Figure 3H,L). In contrast, iodophenpropit displayed surmountable antagonism by shifting the imetit and VUF8328 concentration-response curves parallel rightward without attenuating the maximal response if measured either directly or 10 min after agonist stimulation (Figure 3B,C). The observed difference in re-equilibration time confirmed the slightly longer receptor recovery time (RecT) for PF03654746 in comparison with iodophenpropit and pitolisant. Double logarithmic Schild plot analysis revealed a linear relationship between the equiactive dose ratios (DR-1) and antagonist concentration for all three antagonists (Figure 3E,IM). The Schild regression slopes were not significantly different from unity for iodophenpropit- mediated inhibition of imetit- and VUF8328-induced hH3R signaling, indicating that iodophenpropit acted as competitive antagonist (Figure 3D). In contrast, slopes of Schild-regression for PF03654746 and pitolisant were larger than 1 (slope(0 min): 1.5 ± 0.2 and slope(0 min): 1.4 ± 0.0, respectively) if hH3R antagonism was assessed immediately after agonist addition or after 10 min re-equilibration (Figure 3A).

### 2.3. Re-Equilibration Time Determines Whether hH3R Antagonism is Insurmountable or Surmountable

Receptor-induced Venus-Gβ1γ2 recruitment to masGRK3ct-Nluc can be detected after different incubation times following stimulation with agonist, which offers the opportunity to evaluate the re-equilibration time of pre-bound antagonist within the same agonist-induced response. Cells were pre-incubated for 1 h with multiple concentrations of antagonist and subsequently stimulated with increasing concentrations agonist and BRET ratio was calculated either immediately after agonist addition or after 10 min re-equilibration (Figure 3A).

Pitolisant and PF03654746 acted as insurmountable antagonists if hH3R-mediated Venus-Gβ1γ2 release was measured directly following stimulation with full agonist imetit or partial agonist VUF8328, as revealed by the reduced maximum and rightward-shift of the concentration-response curves (Figure 3E,G,J,K). However, allowing pre-bound antagonists to re-equilibrate with VUF8328 for 10 min before measuring receptor response resulted in surmountable and near-surmountable antagonism for pitolisant and PF03654746, respectively (Figure 3H,L). In contrast, iodophenpropit displayed surmountable antagonism by shifting the imetit and VUF8328 concentration-response curves parallel rightward without attenuating the maximal response if measured either directly or 10 min after agonist stimulation (Figure 3B,C). The observed difference in re-equilibration time confirmed the slightly longer receptor recovery time (RecT) for PF03654746 in comparison with iodophenpropit and pitolisant. Double logarithmic Schild plot analysis revealed a linear relationship between the equiactive dose ratios (DR-1) and antagonist concentration for all three antagonists (Figure 3E,IM). The Schild regression slopes were not significantly different from unity for iodophenpropit- mediated inhibition of imetit- and VUF8328-induced hH3R signaling, indicating that iodophenpropit acted as competitive antagonist (Figure 3D). In contrast, slopes of Schild-regression for PF03654746 and pitolisant were larger than 1 (slope(0 min): 1.5 ± 0.2 and slope(0 min): 1.4 ± 0.0, respectively) if hH3R antagonism was assessed immediately after agonist addition or after 10 min re-equilibration (Figure 3A).

![Figure 2](image-url) **Figure 2.** Recovery of hH3R-mediated Gα12 activation measured in time as BRET ratio between Venus-Gβγ and masGRK3-Nluc after inhibition with antagonists was monitored for 10 s after agonist addition with 60 s time intervals. (A) hH3R-expressing HEK293T cells were pre-incubated for 1 h with antagonist (10 × IC50 concentration) followed by rapid washout of unbound antagonist and stimulation with 10 μM imetit after different incubation times. (B) Recovery rate of hH3R responsiveness after washout of pre-bound iodophenpropit, pitolisant and PF03654746 BRET ratio’s plotted are calculated 10 s after agonist stimulation. Representative graph of 3 independent experiments performed in singular is shown.
signaling is measured directly upon VUF8328 stimulation, indicating non-competitive antagonism due to insufficient re-equilibration time [26,27]. After 10 min re-equilibration with VUF8328, the Schild regression slopes for both pitolisant and PF03654746 were not significantly different from unity ($p = 0.15$ and 0.18 for Pitolisant and PF03654746, respectively, Student’s t-test) (Table 2, Figure 3I,M). The fitted $pA_2$ values for all three antagonists were comparable between the responses that were measured directly or after 10 min upon agonist addition, and in the same order as their $pK_B$ and $pK_i$ values (Table 2).

![Figure 3](image-url)  
**Figure 3.** Concentration response curves of imetit in the presence of increasing concentrations of antagonists iodophenpropit (B–D), pitolisant (F–H) or PF03654746 (J–L) measured as BRET ratio between Venus-GRK3ct-Nluc and Gβ1γ2 in HEK293T cells co-expressing hH3R and Gαi2. (A) Cells were pre-incubated with antagonists for 1 h prior to stimulation with imetit (B,F,I) or VUF8328 (C,D,G,H,K,L) and BRET ratios were measured immediately (B,C,F,G,J,K) or 10 min (D,H,L) after agonist stimulation. Schild-plots for iodophenpropit (E), pitolisant (I), and PF03654746 (M) contain three regression-lines for the three different DRC panels. Representative graphs of 3 experiments performed in duplicate are shown and data are mean ± SD.

3. Discussion

The preferred residence time and consequent duration of drug action might be highly target dependent [28,29]. For example, antihistamines should preferentially occupy the H1R for a long duration to prevent an allergic reaction in the case of hay fever. On the other hand, prolonged blockade of the H3R leads to undesired on-target side effect insomnia, and consequently relatively short residence time antagonists might be preferred for this receptor [19,30]. Early withdrawal of drug candidates
from clinical trials might be reduced if duration of drug action is already taken into account at a very early stage in the drug discovery process. Recently, we evaluated the duration of functional H₃R antagonism using the BRET-based cAMP biosensor CAMYEL, and indeed observed a correlation between the target recovery time upon antagonist washout and target residence time in competitive association binding assays [3]. However, detection of H₃R-mediated reduction in cAMP levels using this CAMYEL biosensor was delayed in time upon agonist stimulation, due to the required activation of adenylyl cyclase by forskolin. Therefore, prompt detection of Gαᵢ protein activation via Gβ₁γ₂ release rather than the downstream effect on cAMP levels might be more suitable to evaluate antagonist binding kinetics. Indeed, fast recovery (<20 min) of H₃R-induced Gβ₁γ₂ release to steady-state level in response to agonist stimulation upon washout of pre-bound iodophenpropit, pitolisant, and PF03654746 was observed. However, steady-state levels were slightly lower as compared with the agonist-induced response in cells that were pre-treated with vehicle, which might be the consequence of re-binding of antagonist that has dissociated from the receptor or partitioned in the cell membrane after washout of unbound antagonist [31]. In contrast, a full recovery of the hH₃R responsiveness was previously observed after >30 min following washout of pre-bound pitolisant in the CAMYEL biosensor assay [3]. The recovery of agonist-induced Gβ₁γ₂ release was approximately two-fold faster upon iodophenpropit and pitolisant washout as compared with PF03654746, suggesting that the latter antagonist has a longer H₃R residence time. Competitive association binding on intact HEK293T cells, however, previously indicated that pitolisant and PF03654746 have a comparable residence time of approximately 15 min [3,32]. The incomplete re-equilibration of pre-bound PF03654746 within the 75 s time-course of a transient Ca²⁺ peak response leads to a reduced maximum of the R-α-methyl-histamine concentration response curves as unoccupied receptors are not available for agonist stimulation [32]. Similar insurmountable antagonism by pre-bound PF03654746 was observed when BRET between Venus-Gβ₁γ₂ and masGRK3ct-Nluc was measured within the same short time frame after agonist-stimulation, whereas PF03654746 acted as (nearly) surmountable antagonist (i.e., parallel dextral shift of agonist response curve without attenuation of maximum response) if this response was measured after a 10 min re-equilibration period. Hence, the required re-equilibration time for a pre-bound insurmountable antagonist (i.e., parallel dextral shift combined with depression of maximum agonist response) to become a surmountable antagonist (i.e., parallel dextral shift of agonist response curve without attenuation of maximum response) is indicative for the target residence time of the antagonist [27,33]. Indeed, short residence time antagonist iodophenpropit caused a parallel rightward shift of these curves directly upon agonist stimulation, whereas longer residence time antagonists pitolisant and PF03654746 decreased the maximum response directly following VUF8328 stimulation but not after 10 min incubation period, suggesting that these pre-bound antagonists re-equilibrate within this time-frame. In conclusion, knowledge on how long an antagonist occupies its target is considered to be an important parameter in drug discovery. The very fast kinetics of the Gβ₁γ₂ release response makes this BRET-based assay a valuable tool to measure duration of target occupancy by washout of pre-bound antagonist and detection of the recovery of agonist responsiveness. Moreover, in contrast to frequently used, but very transient calcium responses, this Gβ₁γ₂ release can be measured after increasing time intervals following agonist stimulation, which allows detection of antagonist re-equilibration by measuring Schild analysis experiments in time. The only marketed H₃R antagonist pitolisant (Wakix®) showed relative fast dissociation kinetics in both Gβ₁γ₂ release and hemi-equilibrium Ca²⁺ assays, suggesting that short-residence time antagonists might display best in vivo efficacy to therapeutically target hH₃R [32].
4. Materials and Methods

4.1. Materials

PF03654746 was bought from Axon Medchem (Groningen, The Netherlands) and pitolisant was obtained from Griffin Discoveries (Amsterdam, The Netherlands). All other compounds were synthesized in-house as previously reported [34–36]. Human embryonic kidney 293T cells (HEK293T cells) were obtained from ATCC (Manassas, VA, USA). Fetal bovine serum (FBS) was obtained from Bodinco (Alkmaar, The Netherlands). Penicillin and streptomycin were purchased from GE healthcare (Uppsula, Sweden). Dulbecco’s Modified Eagles Medium (DMEM), trypsin-EDTA and Hanks’ Balanced Salt Solution (HBSS) were obtained from Gibco (Thermo Fisher Scientific, Waltham, MA, USA). Nano-Glo® (3.2 µL/mL) was obtained from Promega (Madison, WI, USA). Plasmids encoding for Venus155-239-Gβ1, Venus1-155-Gγ2 and masGRK3ct-Nluc were kindly provided by Dr. N. Lambert (Georgia Health Sciences University, Augusta, GA, USA). PcDNA3.1+ plasmid encoding human Gαi2 subunit was bought from cDNA Resource Center (Bloomsberg, PA, USA). Human H3R (Genbank accession no. AF140538) in pcDEF3 was previously described [37]. All other chemicals were of analytical grade and obtained from standard commercial sources.

4.2. Cell Culture and Transfection

Human embryonic kidney 293T cells (HEK293T cells) were cultured in DMEM supplemented with 10% FBS, penicillin (100 µg/mL) and streptomycin (50 µg/mL) at 37 °C with 5% CO2. Cells were transiently transfected in a 10 cm² dish with plasmids coding for Venus155-239-Gβ1 (0.4 µg), Venus1-155-Gγ2 (0.4 µg), masGRK3ct-Nluc (0.4 µg), hH3R (0.4 µg), Gαi2-protein (1.2 µg), and empty pcDEF3 (2.2 µg) using 20 µg 25 kDa linear polyethylenimine (Polysciences Inc, Warrington, PA, USA), as previously described [3]. The next day, 50,000 cells/well were transferred to a poly-L-lysine coated black 96 well plates (Greiner Bio-one GmbH, Frickenhausen, Germany) and grown for an additional 24 h.

4.3. BRET between Venus-Gβ1γ2 and masGRK3ct-Nluc to Measure H3R-Induced Gαi2 Activation

Agonist-induced BRET between Venus-Gβ1γ2 and masGRK3ct-Nluc was measured on adherent HEK293T cells co-expressing hH3R and Gαi2 in the presence of NanoGlo® (3.2 µL/mL) using a Mithras LB940 multimode microplate reader (Berthold, Germany). Cells were stimulated with histamine in the absence and presence of antagonist in HBSS buffer, and BRET (540–40 nm) and luminescence (480–20 nm) signals were monitored either immediately or after 10 min at 25 °C. Recovery of hH3R responsiveness was measured following 1 h pre-incubation with antagonist (10 × IC50 concentration) and two wash steps with HBSS to remove unbound antagonist. Next, cells were incubated in HBSS and stimulated with 10 µM imetit after various incubation times. BRET between Venus-G1β1γ2 and masGRK3ct-Nluc was immediately measured upon imetit stimulation and BRET ratio after 10 s were plotted. Re-equilibration of pre-bound antagonists was measured upon pre-incubation of the cells with increasing concentrations of antagonist for one hour prior to the addition of increasing concentrations imetit or VUF8328 and BRET ratio was calculated either immediately after agonist addition (0–3 min) or after 10 min re-equilibration.

4.4. Data Analysis

Ligand-induced BRET changes were calculated by dividing emission at 540–40 nm (Venus) by emission at 480–20 nm (Nluc) and baseline-corrected by subtracting BRET of vehicle. Representative graphs of at least three independent experiments are shown and analyzed by linear or non-linear regression using Prism 7.03 (GraphPad Software, San Diego, CA, USA). Agonist and antagonist concentration-response curves were fitted to three-parameter response models to obtain EC50 and
$\text{IC}_{50}$ values, respectively. The equilibrium dissociation constant of antagonist ($K_B$) for the hH$_3$R is calculated from $\text{IC}_{50}$ values using:

$$K_B = \frac{\text{IC}_{50}}{[A] \times \text{EC}_{50} + 1}$$  \hspace{1cm} (1)

where $[A]$ and $\text{EC}_{50}$ are the histamine concentration (10 $\mu$M) and potency, respectively. For receptor recovery the vehicle corrected BRET ratio at 10 s after agonist addition was plotted over time.

Recovery of receptor responsiveness ($Y$) was analyzed as function of time by a one-phase association model:

$$Y = Y_0 + (Y_{\text{max}} - Y_0) \times \left(1 - e^{-k_{\text{rec}} \times t}\right)$$  \hspace{1cm} (2)

where $Y_0$ is the hH$_3$R response induced by 10 $\mu$M imetit in the presence of (10 $\times$ $\text{IC}_{50}$ concentration) antagonist, $Y_{\text{max}}$ is the imetit-induced response upon reaching steady-state and $k_{\text{rec}}$ is the receptor-recovery rate in min$^{-1}$. The receptor recovery time (RecT) was calculated as the reciprocal of the $k_{\text{rec}}$.

Equiactive agonist concentrations for Schild analysis were determined from concentration response curves at 10 % max agonist response (vide supra) in the absence and presence of increasing antagonist concentrations. Dose ratios (DR) were calculated by dividing the equiactive agonist concentration in the presence of antagonist by the equiactive agonist concentration in the absence of antagonist. $DR$ minus 1 was plotted as function of antagonist concentration ($[B]$) in a double logarithmic graph and $pA_2$ and slope values were determined by linear regression:

$$\log(DR - 1) = \log[B] + pA_2$$  \hspace{1cm} (3)

Statistical analysis was performed using Graphpad Prism 7.03.

**Author Contributions:** Conceptualization, T.A.M.M., R.L. and H.F.V.; methodology, all authors; validation, all authors; formal analysis, M.C.M.L.B., T.A.M.M., H.F.V.; investigation, M.C.M.L.B., T.A.M.M.; resources R.L. and H.F.V.; writing—original draft preparation, T.A.M.M., R.L. and H.F.V.; writing—review and editing, all authors; visualization, T.A.M.M., R.L. and H.F.V.; supervision, R.L. and H.F.V.

**Funding:** This research was funded by The Netherlands Organization of Scientific research (NWO) TOPPUNT (“7 ways to 7TMR modulation (7-to-7)”) (grant 718.014.002) (R.L. and H.F.V.).

**Conflicts of Interest:** The authors declare no conflict of interest.

**Abbreviations**

- $\alpha$  
  Intrinsic activity
- ADHD  
  Attention deficit hyperactivity disorder
- ANOVA  
  Analysis of variance
- BRET  
  Bioluminescence resonance energy transfer
- cAMP  
  cyclic Adenosine-MonoPhosphate
- DMEM  
  Dulbecco’s Modified Eagles Medium
- DMR  
  Dynamic mass redistribution
- DR  
  Dose ratio
- GPCR  
  G protein-coupled receptor
- FBS  
  Fetal bovine serum
- H$_1$R  
  Histamine H$_1$ receptor
- H$_3$R  
  Histamine H$_3$ receptor
- HEK  
  Human embryonic kidney
- hERG  
  Human ether-a-go-go related gene
- hH$_3$R  
  Human H$_3$R
- $\text{IC}_{50}$  
  Concentration inhibitor needed to reduce response by 50%
- $k_{\text{off}}$  
  Dissociation rate constant
masGRK3ct membrane-associated-C-terminal fragment of G protein-coupled receptor kinase 3

Nluc NanoLuc luciferase

pEC$_{50}$ $-\log(\text{EC}_{50})$ where EC$_{50}$ is concentration required to obtain half maximal response

pKi $-\log K_i$, where $K_i$ is the equilibrium dissociation constant

RecT Recovery time

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