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5-HT₄ Receptors Constitutively Promote the Non-Amyloidogenic Pathway of APP Cleavage and Interact with ADAM10

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ABSTRACT:

In addition to the amyloidogenic pathway, Amyloid Precursor Protein (APP) can be cleaved by α secretases, producing soluble and neuroprotective APP alpha (sAPP α) (non-amyloidogenic pathway) and thus preventing the generation of pathogenic Amyloid- β . However, the mechanisms regulating APP cleavage by α -secretases remain poorly understood. Here, we showed that expression of serotonin type 4 receptors (5-HT₄Rs) constitutively (without agonist stimulation) induced APP cleavage by the α -secretase ADAM10 and the release of neuroprotective sAPP α in HEK-293 cells and cortical neurons. This effect was independent of cAMP production. Interestingly, we demonstrated that 5-HT₄ receptors physically interacted with the mature form of ADAM10. Stimulation of 5-HT₄ receptors by an agonist further increased sAPP α secretion and this effect was mediated by cAMP/Epac signalling. These findings describe a new mechanism whereby a GPCR constitutively stimulates the cleavage of APP by α -secretase and promotes the non-amyloidogenic pathway of APP processing.

KEYWORDS: Alpha-secretase; Alzheimer's disease; sAPP alpha; serotonin

INTRODUCTION

According to the amyloid hypothesis, alteration in synaptic transmission and neuronal loss observed in Alzheimer's disease (AD) mainly result from the formation of toxic Amyloid- β (A β) oligomers followed by the extracellular accumulation of A β aggregates. A β is produced by the successive cleavage of a transmembrane Amyloid Precursor Protein (APP) by β -secretase and γ -secretase (1). In addition to this amyloidogenic pathway, APP can be cleaved by α -secretases, a set of membrane-bound proteases of the ADAM (A Disintegrin And Metalloprotease) family, generating the soluble APP ectodomain (sAPP α) and a membrane-bound carboxy-terminal fragment (non-amyloidogenic pathway). As α -secretases cleave APP within the A β sequence, APP shedding by α -secretase prevents the generation of the pathogenic A β peptide (2, 3). Therefore, enhancing α -secretase expression or activity has been considered as a valuable strategy for inhibiting A β formation. For instance, it has recently been shown that activation of the transcription of the gene encoding the α -secretase ADAM10, by retinoic acid and sirtuin 1 (SIRT1), reduces A β production (4-6).

Previous reports have shown that G protein-coupled receptors (GPCRs) can differentially affect A β peptide production by either modulating the cellular trafficking of APP or by influencing the activity and trafficking of α -, β - and γ -secretases. Moreover, both the expression and the stimulation of GPCRs can affect APP metabolism (7). GPCRs that enhance sAPP α production by stimulating α -secretase activities include the muscarinic M₁-M₃ receptors, mGlu2 metabotropic glutamate receptor, serotonin 2A (5-HT_{2A}) and 5-HT_{2C} receptors, Corticotropin-Releasing Factor (CRF) receptor 1, purinergic receptor P2X7 and Pituitary Adenylate Cyclase-Activating Polypeptide (PACAP) type 1 (PAC₁) receptor (7). These GPCRs are supposed to have a beneficial effect, because sAPP α exerts neuroprotective and neurotrophic effects and enhances long-term potentiation (*8, 9*). In addition, promoting sAPP α formation generally decreases the production of toxic A β peptides. Several signalling cascades, such as the PKC, cAMP/PKA, MAP-Kinase and PI3K pathways, have been involved in the stimulation of sAPP α release mediated by these GPCRs (*10*). However, the mechanisms by which they activate α -secretases are not always well understood.

The 5-HT₄ receptor (5-HT₄R) is another GPCR raising particular interest in view of its possible influence at different levels in the pathogenesis of AD. Activation of 5-HT₄Rs stimulates acetylcholine release in prefrontal cortex and hippocampus and improves learning and memory in various pre-clinical paradigms of memory acquisition and retention (*11, 12*). These findings suggest that 5-HT₄R agonists might be used for improving cholinergic function and cognition, which are both compromised in AD. Stimulation of 5-HT₄Rs also enhances sAPP α release and reduces A β formation in neuronal cultures (*13, 14*). Whether the effect of 5-HT₄Rs on APP metabolism reflects an increase in α -secretase activity or is the result of a modification of cellular trafficking of α -secretases and APP remains to be elucidated. Like many GPCRs, 5-HT₄R interacts with a large set of intracellular proteins. Some of them, such as

p11 and SNX27, are known to affect 5-HT₄R transport and might also influence the trafficking of proteins physically and functionally connected with the receptor (*15, 16*). Further supporting the implication of 5-HT₄R in AD, the number of receptor binding sites is reduced in the hippocampus of patients with AD (*17*). However, the specific effects of 5-HT₄R expression and activation on APP metabolism remain to be elucidated.

In this study, we show that 5-HT₄R expression at physiological levels (around 120-550 fmol/mg of protein) (*17-21*) enhances sAPP α release in transfected HEK-293 cells and cortical neurons in primary culture *via* two different processes, a "constitutive" release which has never been reported and a previously described agonist-regulated one (*13, 14*). We also show that 5-HT₄ receptors physically associate with ADAM10, a mechanism that may be involved in the constitutive release of sAPP α .

RESULTS AND DISCUSSION

Expression of 5-HT₄ receptors enhances sAPP release from HEK-293 cells and cortical neurons. To investigate the impact of 5-HT₄R expression on APP metabolism, we first co-transfected HEK-293 cells with the SEAP-tagged APP plasmid (Figure 1A) and increasing amounts of the Myctagged 5-HT₄R construct. 5-HT₄R density at the cell surface was assessed by ELISA (Figure 1B) and ³H]GR 113808 saturation binding (from 120 to 750 fmol/mg protein corresponding to 25 to 500 ng cDNA/10⁷ cells, respectively). Expression of increasing amounts of 5-HT₄R did not significantly affect total APP (APP band intensity/actin band intensity = $97 \pm 4\%$ of control, $96 \pm 7\%$, $108 \pm 13\%$ or $89 \pm$ 18% for 25, 100, 250 or 500 ng of 5-HT_R cDNA respectively, n≥3 per each group) or ADAM10 expression (intensity of ADAM10 bands/actin band intensity = $104 \pm 5\%$ of control, $96 \pm 11\%$, $113 \pm$ 18% or 87 \pm 21% for 25, 100, 250 or 500 ng of 5-HT₄R cDNA respectively, n \geq 3 per each group) (Figure 1B), while it potently enhanced the release of soluble APP (sAPP) already at low receptor expression (Figure 1B). This effect was independent of the 5-HT₄R variant used: the mouse (a, b, e and f) and human (a, g and i) C-terminal variants as well as further truncations of the 5-HT₄R C-terminal domain (Δ 358 and Δ 329) could all promote sAPP release (Supplemental Figure 1, A-C). Overexpression of 5-HT,R in primary cultures of cortical neurons produced a comparable increase in sAPP release (Figure 1C) without affecting APP and ADAM10 levels (APP band intensity/actin band intensity = $88 \pm 19\%$ of control, intensity of ADAM10 bands/actin band intensity = $111 \pm 14\%$ for 2.500 ng of 5-HT₄R cDNA n \geq 3 per each group). Expression of similar amounts of PAC₁ or muscarinic M₃ receptor, which both increase sAPP α release upon activation by their respective agonists (22, 23), failed to enhance sAPP release in HEK-293 cells in the absence of agonist (Figure 1D). Moreover, comparative levels of over-expressed 5-HT₆ receptor, another serotonin receptor positively coupled to the G_s/cAMP pathway, did not induce sAPP production (Figure 1D). Similarly, only over-expression of 5-HT₄R, but

not of PAC₁, M_3 or 5-HT₆ receptors, in cortical neurons enhanced the release of sAPP from neurons (Figure 1E).

Expression of 5-HT₄ receptors specifically enhances sAPPα release. To identify the nature of the sAPP form (sAPPα or sAPPβ) secreted upon 5-HT₄R expression, we used a sandwich ELISA kit based on two antibodies, one directed against the N-terminal sequence of APP and the other recognizing the neo-sAPPα epitope. Using this kit, we showed that over-expression of 5-HT₄R in both HEK-239 cells and in cortical neurons led to an increase in the release of sAPPα (Figure 2A). Furthermore, SEAP-tagged APP in supernatants from transfected HEK-293 cells was immunoprecipitated with the 22C11 antibody and immunodetected using a site-specific antibody (7A6) that specifically recognizes sAPPα, but not shorter sAPP species, such as sAPPβ' and sAPPβ (Figures. 1A and 2B) (24). The immunoreactive signals detected by 7A6 antibody increased concomitantly with 5-HT₄R expression, while no significant change of APP and ADAM10 expression was observed (Figure 2B). Mutation of the residues (R⁶⁰⁹D and K⁶¹²E) surrounding the α-secretase cleavage site (α-site APP mutant) can impair about 50% of the α-cleavage of APP (25). Co-expression of this mutant with 5-HT₄R in HEK-293 cells strongly decreased 5-HT₄R-induced sAPP release (Figure 2C), indicating that 5-HT₄R expression stimulates the non-amyloidogenic α-cleavage of APP.

Several metalloproteinases, including ADAM9, ADAM10 and ADAM17 (also called TACE, Tumour Necrosis Factor α -Converting Enzyme) have been proposed as potential α -secretases that promote the shedding of APP ectodomain. TAPI-1, a metalloproteinase inhibitor, almost completely abolished 5-HT₄R-induced sAPP α release from HEK-293 cells (Figure 2D), suggesting the involvement of a member of the ADAM family of metalloproteinases. Moreover, reduction of ADAM10 expression by transfection of *ADAM10*-specific siRNA (Figure 2E) diminished basal processing of APP in absence of 5-HT₄ receptor and also markedly decreased sAPP α release following 5-HT₄R expression (Figure 2F). This indicates that ADAM10 contributes significantly to 5-HT₄R-induced sAPP release. In contrast, specific inhibitors of β -secretase (KTEEISEVN-statine-VAEF) or γ -secretase (L-685,458) did not affect the release of sAPP α induced by 5-HT₄R (Supplemental Figure 2, A and B).

5-HT₄R stimulation further enhances sAPP\alpha release. Previous studies have shown that 5-HT₄R stimulation by agonists increases sAPP α production (*26*). Consistent with these findings, treating HEK-293 cells that transiently express 5-HT₄R and SEAP-tagged APP with prucalopride, a 5-HT₄R agonist, for 2 hrs increased by ~ 50% the release of sAPP α induced by 5-HT₄R expression (Figure 3A and B). This effect was blocked by GR 113808 and RO 116-0086, a neutral antagonist and an inverse agonist of 5-HT₄R at G_s signalling, respectively. Neither GR 113808 nor RO 116-0086 alone altered the secretion of sAPP in the absence of the agonist (Figure 3B).

5-HT₄R stimulation by prucalopride increased sAPP α release by cultured cortical neurons that transiently expressed 5-HT₄R and SEAP-tagged APP in a concentration-dependent manner (pEC₅₀ = 0.85 ± 0.22 μ M, n = 3), as assessed by monitoring the SEAP activity (Figure 3C). Similarly, prucalopride

stimulated sAPP α release from endogenous APP, as assessed using a sAPP α specific ELISA kit (Figure 3D). Like in HEK-293 cells, the prucalopride effect was blocked by GR 113808 and RO 116-0086 (Figure 3E). Treatment of neurons with maximally effective concentrations of various synthetic 5-HT₄R agonists (10 μ M RS 67333, 3 μ M prucalopride, 10 μ M BIMU8, 1 μ M cisapride) (*27*) or with 10 μ M 5-HT stimulated sAPP secretion to a similar extent as prucalopride. In contrast, 5-HT₄R ligands exhibiting either antagonist (GR 113808, ML 10375) or inverse agonist (RO 116-1148, SB 207266) properties on 5-HT₄ receptors coupled to G_s (*27*) did not significantly affect sAPP release from neurons (Figure 3F). However, RO 116-0086 and RO 116-2617, two inverse agonists at G_s signalling decreased basal sAPP release. GPCRs proteins are adapting many conformations. The constitutive 5-HT₄R conformation which induces cAMP production (via G protein) is unlikely to be similar to the constitutive conformation of the same 5-HT₄ receptor which is producing sAPP α (without a need of G protein). The pharmacology of these two different constitutive conformations is likely to be different. Thus, it is not surprising that some "classical" inverse agonists on the 5-HT₄ receptor inducing constitutive cAMP production are not inverse agonists on the constitutive conformation required for sAPP α production.

The constitutive sAPP α release induced by 5-HT₄R expression is independent of the cAMP/Epac pathway. As cAMP and Epac (cAMP-responsive Rap1 Guanine Nucleotide Exchange Factor) regulate sAPP α release elicited by 5-HT₄R stimulation (*13*), we investigated whether receptor coupling to G_s and recruitment of the cAMP/Epac pathway were also necessary for enhancing constitutive sAPP α secretion upon 5-HT₄R expression. Expression of a dominant-negative form of G_s strongly reduced agonist-independent cAMP accumulation in HEK-293 cells expressing 5-HT₄Rs, but did not affect sAPP release (Figure 4A). Moreover, expression of 5-HT₄R mutants with high level of constitutive G_s signalling activity (Y³⁰²F, N³⁰⁸D or A²⁵⁸L) (*28, 29*) increased basal cAMP production but were less active to increase sAPP release in comparison to wild type 5-HT₄R (Figure 4B). Similarly, over-expression of a dominant negative form of Epac1 did not modify the constitutive sAPP secretion induced by 5-HT₄R expression (Figure 4C), whereas the agonist (prucalopride)-dependent sAPP release was reduced, as previously reported (*13*) (Figure 4C). As expected, the Epac1 dominant negative form did not affect cAMP production elicited by 5-HT₄R activation (Figure 4D). Collectively, these results indicate that only agonist-elicited sAPP secretion is dependent on 5-HT₄R coupling to G_s protein and engagement of the cAMP/Epac pathway.

5-HT₄ receptors physically interact with the α -secretase ADAM10. To further explore the possible mechanism by which 5-HT₄R constitutively activates ADAM10-dependent APP cleavage, we examined whether 5-HT₄R physically interacts with this α -secretase. ADAM10 co-immunoprecipited with HA-tagged-5-HT₄R in HEK-293 cells (Figure 5A). Although two immunoreactive ADAM10 species with apparent molecular masses of 95 (immature pro-enzyme) and 69 kDa (mature, active form) were detected in whole cell lysates by Western blotting, only the 69 kDa mature form, which originates from the immature form by removal of a 194 amino acid pro-domain and is localized at the plasma

membrane (*30*), was co-immunoprecipitated with 5-HT₄R. Prucalopride stimulation of 5-HT₄R did not modify the ability of the receptor to interact with ADAM10, thus demonstrating the existence of a "constitutive" specific complex composed of 5-HT₄R and active ADAM10 (Figure 5A).

Active ADAM10 did not co-immunoprecipitate with over-expressed 5-HT₆ (Figure 5B), PAC₁ (Figure 5C) or M₃ receptors (Figure 5D). Furthermore, different 5-HT₄R mutants ($A^{258}L$, $N^{308}D$ and $Y^{302}F$) and truncated forms devoid of the C-terminal domain (Δ 327 and Δ 329) showed the same ability to interact with ADAM10 as the wild type receptor (Supplemental Figure 1, D-G).

Promotion of non-amyloidogenic pathway by 5-HT₄ receptors. The nature of APP processing is dependent on its cellular localization and on the cellular proteases involved in its cleavage. The present study shows that 5-HT₄R expression, at physiological levels (120-550 fmol/mg proteins) (17-21), constitutively promotes the non-amyloidogenic processing of APP and that 5-HT₄ receptors physically interact with the mature form of the α -secretase ADAM10. Several reports described the overlapping expression of APP and ADAM10 (but not ADAM17) in adult brain neurons (31) and suggested that ADAM10 could be the α -secretase involved in the non-amyloidogenic pathway of APP, particularly in neurons (30). This assumption was confirmed by RNAi-mediated silencing of ADAM10 and by using conditional ADAM10 knock-out mice (24, 32). These findings were extended by the present study, which revealed association of the 5-HT₄R with ADAM10. The ability of 5-HT₄R to physically interact with mature ADAM10 was not shared by other GPCRs known to stimulate sAPPa secretion upon stimulation by their respective agonists. Indeed, expression of the PAC₁ and M₃ receptors did not trigger the formation of such a protein complex. Moreover, ADAM10 is retained in the ER by an arginine stretch (RRR) (33). Future work should test the possibility that association of ADAM10 with 5-HT₄R masks this sequence, allowing the translocation of both proteins to the plasma membrane, a process associated with ADAM10 maturation.

It has been previously reported that stimulation of 5-HT₄R promotes sAPP α release in a cAMP/Epac-dependent manner (*13*) and these findings were confirmed in the present study. Although 5-HT₄R displays high level of constitutive activity at the G_s-adenylyl cyclase pathway (*28*), several lines of evidence argued against a role of the cAMP/Epac pathway in the constitutive effect of 5-HT₄R on APP processing: 1) inhibition of cAMP formation by an inverse agonist or by over-expression of the inactive form of G_s had no effect on the level of sAPP α release elicited by 5-HT₄R expression; 2) expression of a dominant negative Epac mutant did not affect constitutive induction of sAPP α release by 5-HT₄R; and 3) mutants of 5-HT₄R that constitutively increase adenylyl cyclase activity did not stimulate sAPP α secretion more than the wild type receptor. Collectively, these findings demonstrate that 5-HT₄R promotes the non-amyloidogenic processing of APP *via* different mechanisms implicating either the canonical G_s protein-dependent signalling (agonist-elicited response) or the formation of a protein complex including the receptor, the ADAM10 α -secretase and, probably, additional protein partners that remain to be identified (agonist-independent response).

One promising strategy to decrease the accumulation of $A\beta$ peptides in the brain has been to inhibit β - and γ - secretase activities. However, the development of γ -secretase inhibitors has been challenging because this enzyme has many endogenous substrates. Several γ -secretase inhibitors have been synthesized and have shown unequivocal efficiency in reducing $A\beta$ plaque deposition in preclinical models of AD. However, clinical trials have revealed severe adverse side effects associated with the blockade of Notch processing (*34*). The recent failure of a γ -secretase inhibitor to slow down the progression of the disease, accompanied by worsening of cognitive performances has halted its development (*35*). However, the recent discovery that down-regulation of GSAP (γ -secretase activating protein) selectively prevents $A\beta$ peptide production without affecting Notch signalling (*36*) opens new perspectives.

Strategies for increasing the production of the neuroprotective sAPP α are also actively investigated. 5-HT₄R expression clearly increased sAPP α release and it was previously demonstrated that 5-HT₄R activation concomitantly decreased A β formation while increasing sAPP α release (13, 14). However, α -cleavage is not always coupled to a reduction in A β level (3). The role of ADAM10, the physiological α -secretase in neurons, in the regulation of late-onset AD has been suggested by the discovery of non-synonymous mutations in the ADAM10 pro-domain in seven late-onset AD families (37). In line with these findings, activation of ADAM10 expression by vitamin A derivatives (4) or the transcription factor SIRT1 (6) decrease A β peptides production and amyloid plaque formation in APP/PSEN1 transgenic mice. In addition, SIRT1-induced ADAM10 expression increases Notch processing and the release of Notch/intracellular domain (NICD), which activates genes involved in neuronal repair in adult brain (38). The efficacy of ligands of some GPCRs, such as M₁R, 5-HT₄R and PAC₁R, in reducing A β production and/or improving cognition in pre-clinical models of AD is now well documented (39). Here, we describe a new mechanism whereby 5-HT₄Rs constitutively activate the α -secretase ADAM10 and sAPP α release. Together with the ability of 5-HT_xR agonists to increase learning and memory, via acetylcholine release (12), the present findings suggest that this GPCR might represent an interesting target for AD treatment. Several studies reported a significant drop of 5-HT₄R binding sites in post-mortem brains of AD patients (17, 40, 41). In this context, the identification of 5-HT₄R-interacting proteins capable of modulating the trafficking of the 5-HT₄R/ADAM10 complex will be of high relevance.

METHODS

Plasmids. The plasmid encoding the inactive form of the G_s protein was kindly given by Dr. Catherine H. Berlot (Weis Center for Research Geisinger Clinic, Danville, Canada). The plasmids encoding c-Myc-tagged 5-HT₄R, HA-tagged 5-HT₄R, Flag-tagged 5-HT₄R, HA-tagged 5-HT₆R and the dominant-negative form of Epac2 were previously described (*28, 42, 43*). The SEAP-tagged APP695 cDNA in pRK5 was generated by adding the secreted embryonic alkaline phosphatase (SEAP)

sequence, flanked by glycine linkers and the PspX I (5') or BsrG I (3') restriction sites, to the N-terminus of mouse APP695, after L³⁹⁸ (Figure 1A). The plasmids encoding the HA-tagged M₃ and Myc-tagged PAC₁ receptors were kindly provided by Dr. Arnaud Monteil and Dr. Laurent Journot respectively (Institut de Génomique Fonctionnelle, Montpellier, France).

Antibodies. The mouse anti-sAPP antibody (22C11, directed against the N-terminal part of APP; 1:80 dilution) (44) was purchased from Millipore (Billerica, USA). The rat anti-sAPPα antibody (7A6, murine sAPPα-specific, 1:2 dilution) was kindly given by Dr. Stefan F Lichtenthaler (Adolf Butenandt Institute - Ludwig Maximilians University, Munich, Germany) (24). The rabbit anti-ADAM10 antibody (directed against the C-terminal domain, 1:500 dilution), the mouse anti-c-Myc antibody (9E10) and the rabbit anti-Flag antibody were purchased from Sigma-Aldrich (Saint Quentin Fallavier, France). The mouse anti-HA antibody was purchased from Life Technologies (Saint Aubin, France). The rat anti-ADAM10 antibody (directed against the ectodomain domain, 1:500 dilution) was purchased from R &D Systems (Minneapolis,USA). Mouse anti-HA and rabbit anti-GFP antibodies were purchased from Life Technologies Invitrogen (Carlsbad, USA). The mouse anti-SEAP antibody was purchased from Life anti-seapone (Souffelweyershein, France). Horseradish peroxidase (HRP)-conjugated anti-rabbit anti-rabbit anti-mouse anti-mouse anti-rabbit anti-rabbit anti-mouse anti-rabbit anti-rabbit anti-rabbit anti-rabbit anti-mouse anti-rabbit anti-rabbit anti-seapone (Souffelweyershein, France).

Materials. All media, sera and antibiotics used for cell culture were purchased from Lonza (Basel, Switzerland). GR 113808 and GR 125487 were synthesized and generously provided by GlaxoSmithKline (Brentford, UK). 5-HT (serotonin creatine sulfate), Phorbol 12-myristate 13-acetate (PMA), the β-secretase inhibitor KTEEISEVN-statine-VAEF, the γ-secretase inhibitor L-685458, (-)isoproterenol hydrochloride and metoclopramide (*N*-(2-dimethylamino)-ethyl)-4-amino-5-chloro-2methoxybenzamide) were from Sigma. The TNF-α Protease Inhibitor-1 (TAPI-1) was purchased from Merck (Darmstadt, Germany). Prucalopride was synthesized and kindly provided by Dr. Jan Schuurkes (Shire-Movetis n.v., Turnhout, Belgium). RO 116-0086, RO 116-1148 and RO 116-2617 were generously provided by Roche Bioscience (Palo Alto, USA). RS 67333 was purchased from Tocris Bioscience (Bristol, UK). BIMU8 was kindly provided by Boehringer Ingelheim (Rhein, Germany). Cisapride was synthesized and obtained from Janssen Pharmaceutica (Beerse, Belgium). ML 10375 was a generous gift from Dr. Langlois (CNRS-BIOCIS, Châtenay-Malabry, France). SB 207266 was obtained from Fournier-Debat Laboratories (Daix, France).

Cell culture and transfections. Human Embryonic Kidney 293 (HEK-293) cells were grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% dialyzed foetal calf serum (FCS). Cells were transfected at 60-70% confluence by electroporation, as previously described (*28*). 24 hrs after transfection, cells were serum starved for 4 hrs before assays.

Primary cultures of cortical neurons were prepared as previously described (45). Briefly, cells dissociated from the cortex of 15-16 day/old Swiss mouse embryos were plated in serum-free medium in 24-well culture dishes (0.5 x 10^6 cells/well). Cultures were maintained at 37° C in a humidified

atmosphere in 5% CO₂ and 95% H₂O, in DMEM/F12 supplemented with 10% of hormone mixture for 4 days. Cortical neurons were transfected by electroporation using AMAXA (Lonza, Basel, Switzerland) according to the manufacturer's instructions. Media were collected for 12 hrs at DIV4.

Small interfering RNA (siRNA) transfection. A mix of double-stranded siRNA sequences 5'-AAUGAAGAGGGACACUUCCCUdTdT-3' and 5'-AAGUUGCCUCCUCCUAAACCAdTdT-3' (Eurogentec, Liege, Belgium) was used to selectively target *ADAM10*. A small RNA duplex, which has no silencing effect, was used as control (negative control siRNA duplex from Eurogentec). All the siRNAs used in this study have been described earlier and validated (*46*). Early passage HEK-293 cells at 40% confluence in 24-well plates were transiently transfected with 400 ng of plasmid encoding wild type 5-HT₄R and 500 ng of plasmid encoding SEAP-tagged-APP. siRNAs were transfected at 150 nM using Lipofectamine 2000, according to the manufacturer's recommendations (Life Technologies-Invitrogen). Cells were lysed 2 days after siRNA transfection in order to quantify APP expression.

Measurement of sAPP. To determine the total and intracellular concentration of APP and of secreted APP, cells were washed with PBS at the beginning of the assay and pre-incubated for 2 hrs in serum-free medium supplemented with 5-HT₄R ligands and drugs were added for the indicated times. Cells were then lysed after collection of conditioned supernatants containing secreted proteins according to the two following methods: (i) collected media were centrifuged at 300 x g for 10 min and then at 20,000 x g for 20 min to remove cells and cell debris. Proteins were concentrated by precipitation with 10% ice-cold tricholroacetic acid (TCA) (at 4°C for 2 hrs), centrifuged at 38,000 x g at 4°C for 25 min and pellets were washed with diethyl ether three times. TCA precipitates were reconstituted in solubilization buffer (20 mM HEPES buffer; 250 mM Tris buffer pH6.8; 40% SDS; 40% glycerol; 40% 2-mercaptoethanol and traces of bromophenol blue). The protein amount of each sample was determined with the bicinchoninic acid assay (Sigma). Following precipitation, samples were separated by electrophoresis on 7.5% SDS-PAGE gels and transferred onto nitrocellulose membranes; (ii) in order to measure sAPP release, HEK-293 cells were transiently transfected with APP-SEAP and plated in 6-well culture dishes. Spent media were incubated overnight at 4°C with Dynabeads® protein G (Life Technologies-Invitrogen, Carlsbad, USA) that had been pre-coupled to 4 µg of 22C11 polyclonal antibody. After 5 washes in lysis buffer (20 mM HEPES; 150 mM NaCl; 1% NP40; 10% glycerol; 4 mg/ml dodecylmaltoside) supplemented with protease inhibitors and phosphatase inhibitors (10 mM NaF; 2 mM Na⁺-vanadate and 1 mM Na⁺-pyrophosphate), immunoprecipitated proteins were eluted in solubilization buffer and separated by electrophoresis (see above). To detect intracellular APP, cells were lysed in 100 μ L lysis buffer (see above), reconstituted in solubilization buffer and separated by electrophoresis (see above).

sAPP release was also estimated by measuring the extracellular SEAP activity released by HEK-293 cells transiently transfected with the SEAP-APP plasmid. One day post-transfection, drugs were added for the indicated times (see legends), then cell culture media were collected and SEAP activity measured by adding the chromogenic substrate p-Nitrophenyl phosphate disodium hexahydrate (Sigma) according to the manufacturer's instructions. The reaction was read at 405 nm using an Infinite 2000 luminescence counter (Tecan, Männedorf, Switzerland). Specific quantification of sAPP α release was performed by using the sAPP α ELISA kit from IBL International (Hamburg, Germany) according to the manufacturer's instructions. The reaction was read at 620 nm and 450 nm using an Infinite 2000 luminescence counter (Tecan).

Immunoblotting. Proteins were separated on 7.5% SDS-PAGE gels and transferred by semi-dry blotting onto nitrocellulose membranes (Hybond-C, Amersham Biosciences, Orsay, France). Membranes were blocked at room temperature in Tris-buffered saline blocking solution containing 5% (w/v) milk powder, 0.25% v/v Tween 20 for 1 hr. Membranes were then incubated with the primary antibodies at 4 °C overnight, washed extensively and incubated with the secondary antibodies at room temperature for 1 hr. Immunoreactivity was detected using the Chemiluminescence Reagent Plus kit (Perkin Elmer, Courtaboeuf, France). Immunoreactive bands were quantified by densitometry using Image J and GraphPad PRISM (GraphPad Software, San Diego, USA).

Co-immunoprecipitation. Cells were seeded in 150 mm-plates (10^6 cells/plate) 24 hrs before the experiment. Cross-linking was carried out in PBS completed with 1.25 mM dithiobissuccinimidyl propionate (DSP) (Pierce, Perbio-Brebières, France) (a membrane-permeable, hydrolysable, covalent cross-linker) for 30 min. The cross-linking reaction was stopped with 50 mM Tris pH 7.4. After one wash in PBS, cells were incubated with lysis buffer (see above) at 4°C for 1 hr. Samples were centrifuged at 20,000 x g for 15 min. Solubilized proteins were incubated at 4°C with 20 µl of Dynabeads® protein G (Life Technologies-Invitrogen) pre-coupled with 4 µg of antibody overnight. Immunoprecipitated proteins were eluted in Laemmli sample buffer, resolved by SDS-PAGE gel electrophoresis and detected by Western blotting.

Determination of cAMP production. HEK-293 cells were transiently transfected with plasmids encoding the indicated receptors and/or SEAP-APP, then seeded in 24-well plates (200,000 cells/well). 24 hrs after transfection, cells were exposed to the indicated concentrations of 5-HT₄R ligands in the presence of 0.1 mM L-ascorbic acid and 0.1 mM of the phosphodiesterase inhibitor RO-20-1724, at 37°C in 250 µl of HBS (20 mM HEPES; 150 mM NaCl; 4.2 mM KCl; 0.9 mM CaCl₂; 0.5 mM MgCl₂; 0.1% glucose; 0.1% BSA). Cells were then lysed by addition of the same volume of Triton-X100 (0.1%). Quantification of cAMP production was performed by HTRF by using the cAMP Dynamic kit (Cisbio Bioassays) according to the manufacturer's instructions.

Cell surface ELISA (Enzyme-Linked Immunosorbent Assay). HEK-293 cells transiently transfected with the appropriate cDNAs were seeded in 96-well plates (40,000 cells/well). 24 hrs after transfection, cells were treated with drugs, washed, fixed with 4% paraformaldehyde at room temperature for 10 min and blocked with PBS + 1% FCS. Cells were then incubated with primary antibodies in the same buffer for 1 hr followed by HRP conjugated secondary antibodies at 1:8000 for 1hr. After extensive washes, the chromogenic substrate was added (Supersignal® ELISA Femto

Maximum Sensitivity, Pierce). Chemiluminescence was detected and quantified with an Infinite 2000 luminescence counter (Tecan).

Statistical Analysis. All represented experiments were performed in triplicate. Statistical significance was determined with the Student-Newman-Keuls test using GraphPad Prism (GraphPad Software). P values <0.05 were considered as statistically significant. Data corresponded to the mean \pm SD of three independent experiments.

ASSOCIATED CONTENT

Supporting Information

Supplemental Figure 1: 5-HT₄R variants (mouse a, b, e and f; human a, g and i), truncated forms (Δ 358 and Δ 329) and mutants ($A^{258}L$, $N^{308}D$ and $Y^{302}F$) promote sAPP release and physically interact with ADAM10; Supplemental Figure 2: sAPP release induced by 5-HT₄R expression is not affected by β - or γ -secretase inhibitors; Supplemental figure legends. This material is available free of charge via the Internet at http://pubs.acs.org.

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Author contributions

[¶]These authors contributed equally to this work.

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Notes

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ABBREVIATIONS

APP, Amyloid Precursor Protein; sAPP α , soluble APP alpha; sAPP β , soluble APP beta; SEAP: Secreted Embryonic Alkaline Phosphatase, 5-HT₄R, serotonin type 4 receptor; AD, Alzheimer's Disease; ADAM, A Disintegrin And Metalloprotease; A β , Amyloid- β peptide; GPCR, G Protein-Coupled Receptor.

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FIGURE LEGENDS

Figure 1. 5-HT₄R expression constitutively enhances sAPP release. (A) Schematic representation of APP. The position of the SEAP-tag is depicted. The epitopes of the anti-sAPP (22C11) and -sAPPa antibodies (7A6) and the β -, β' -, α - and γ -secretase cleavage sites are also indicated. Mb, membrane. (B, C) HEK-293 cells (B) or embryonic cortical neurons (C) were transiently transfected with plasmids encoding Myc-tagged 5-HT₄R and SEAP-tagged APP (500 ng/10⁷ cells or 2,500 ng/6.10⁶ cells, respectively). 24 hrs (B) or four days (C) post-transfection, cell surface expression of 5-HT₄R was assessed by ELISA in non-permeabilized cells using the anti-Myc antibody (B and C, white bars). In parallel, sAPP release during a 2 hr-period (B) or 12 hr-period (C) was evaluated by measuring the alkaline phosphatase activity (colorimetric assay) of cell supernatants (B and C, grey bars). (B, C) Total APP expression was detected in cell lysates in the corresponding transfection conditions, using the 22C11 antibody. Expression levels of ADAM10, 5-HT,R and actin were provided as controls. (D) HEK-293 cells were transiently transfected with plasmids encoding epitope-tagged 5-HT₄R ($300 \text{ ng}/10^7 \text{ cells}$), PAC₁R (700 ng/10⁷ cells), M₃R (300 ng/10⁷ cells) or 5-HT₆R (75 ng/10⁷ cells) to ensure equivalent cellsurface expression of the different receptors, and SEAP-tagged APP (500 ng/ 10^7 cells). Cell surface expression of the receptors (white bars) and quantification of sAPP release (grey bars) were measured as described in (B) using appropriate primary antibodies. (E) Embryonic cortical neurons were transiently co-transfected with a plasmid encoding epitope-tagged 5-HT₄R (800 ng/6.10⁶ cells), PAC₁R $(1,500 \text{ ng}/6.10^6 \text{ cells}), M_3R (1,000 \text{ ng}/6.10^6 \text{ cells}) \text{ or } 5\text{-HT}_6R (800 \text{ ng}/10^7 \text{ cells}) \text{ and SEAP-tagged APP}$ (2,500 ng/6.10⁶ cells). Cell surface expression of the receptors (white bars) and quantification of sAPP release during a 4 day-period was performed as described in (C). CTL, Condition transfected with only SEAP-tagged APP. Data are the means \pm SD of at least three independent experiments. * p<0.05, ** p< 0.01 vs. CTL corresponding value.

Figure 2. 5-HT₄**R specifically enhances sAPPα release.** (A) HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged 5-HT₄R (250 ng/10⁷ cells), SEAP-tagged APP (500 ng/10⁷ cells). 2hrs-sAPPα release in the medium was detected using an ELISA kit designed to quantify mouse/rat sAPPα. Cortical neurons were transfected with empty plasmid (CTL) or plasmid encoding Flag-tagged 5-HT₄R (250 ng/6.10⁶ cells). Four days after transfection, sAPPα released by neurons during 12 hrs was quantified using the same kit. ** p<0.01 vs. corresponding CTL value. (B) Proteins in supernatants from HEK-293 cells transiently transfected with plasmids encoding Myc-tagged 5-HT₄R and SEAP-tagged APP (500 ng/10⁷ cells) were immunoprecipitated with the 22C11antibody. Immunoprecipitated complexes were analysed by western blotting with the 7A6 (anti-sAPPα) antibody (Upper panel). Expression levels of APP, ADAM10, 5-HT₄R and actin in cell lysates were provided as controls (Lower panel). (C) HEK-293 cells were transiently co-transfected with plasmids encoding

Myc-tagged 5-HT₄R (250 ng/10⁷ cells) and either wild type SEAP-tagged APP or the α site APP mutant (R⁶⁰⁹D, K⁶¹²E) (500 ng/10⁷ cells). sAPP release during 2 hrs was evaluated by measuring the alkaline phosphatase activity in the cell medium, ** p<0.01 vs. corresponding value in cells expressing the wild type SEAP-tagged APP. (D) HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged 5-HT₄R (250 ng/10⁷ cells) and SEAP-tagged APP (500 ng/10⁷ cells). 24 hrs post-transfection, cells were treated with vehicle alone or 50 μ M TAPI-1 and SEAP activity during 30, 60 and 120 minutes was measured in the spent medium. * p<0.05, ** p<0.01 vs. corresponding Value in absence of TAPI-1. (E, F) HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged 5-HT₄R (250 ng/10⁷ cells), SEAP-tagged APP (500 ng/10⁷ cells) and either control (CTL) or ADAM10 siRNA. ADAM10 expression was assessed by Western blotting (E) and sAPP α release (F) as described in A, ** p<0.01 vs. 5-HT₄R-mediated release in cells transfected with control siRNA. CTL, Condition transfected with only SEAP-tagged APP. Data are the means ± SD of at least three independent experiments.

Figure 3. 5-HT₄**R stimulation further enhances sAPPα release induced by 5-HT**₄**R expression. (**A, B) HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged 5-HT₄**R** (250 ng/10⁷ cells) and SEAP-tagged APP (500 ng/10⁷ cells). 24 hrs post-transfection, cells were treated with vehicle alone (CTL) or 1 µM prucalopride in the presence or not of 5 µM GR 113803 or 0.1 µM RO 116-0086 for 2 hrs and then sAPP release was evaluated by measuring the alkaline phosphatase activity in the medium. (C, D) Embryonic cortical neurons were transiently transfected with plasmids encoding Flag-tagged 5-HT₄R (800 ng /6.10⁶ cells) with (C) or without (D) SEAP-tagged APP (2,500 ng/6.10⁶ cells). Four days after transfection, neurons were treated with increasing concentrations of prucalopride and sAPP release during 12 hrs was evaluated by measuring the alkaline phosphate activity (C) or by using the sAPPα ELISA kit (D). (E, F) Cortical neurons co-transfected with plasmids encoding Flag-tagged 5-HT₄R (250 ng/6.10⁶ cells) and SEAP-tagged APP (2,500 ng/6.10⁶ cells) were exposed to the indicated 5-HT₄R agonists/antagonists for 12 hrs and sAPP release was evaluated by measuring the alkaline phosphata activity. CTL, Condition transfected with only SEAP-tagged APP. Data are the means ± SD of at least three independent experiments. In (B), (E) and (F), * p<0.05 vs. the corresponding value in cells not treated with agonist.

Figure 4. Role of the G_s/cAMP/Epac pathway in 5-HT₄R-induced sAPP α release. HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged 5-HT₄R (250 ng/10⁷ cells), SEAP-tagged APP (500 ng/10⁷ cells) and a dominant negative G_s protein (A) or a dominant-negative form of Epac1 (500 ng/10⁷ cells) (C, D). (B) HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged wild type 5-HT₄R or mutants (Y³⁰²F, N³⁰⁸D or A²⁵⁸L) with high level of constitutive G_s signalling activity and SEAP-tagged APP (500 ng/10⁷ cells). 24 hrs after transfection, cAMP

accumulation and sAPP release during 2 hrs were measured. CTL, Condition transfected with only SEAP-tagged APP. Data are the means \pm SD of at least three independent experiments. In (D) and (F), * p<0.05 vs. the corresponding value in cells not treated with agonist.

Figure 5. 5-HT₄**R interacts with the** α-secretase ADAM10. (A) HEK-293 cells were transiently transfected with plasmids encoding HA-tagged 5-HT₄R (250 ng/10⁷ cells) and SEAP-tagged APP (500 ng/10⁷ cells). 24 hrs after transfection, Cells were treated 2 hours with vehicle alone or with 1 μ M prucalopride (Prucalo.), then protein lysates were immunoprecipitated with the anti-HA antibody. Whole cell extracts and immunoprecipitated material were analysed by Western blotting using antibodies against ADAM10 (mature (m) and immature (im) forms) and against HA-tag. (B, C, D) HEK-293 cells were transiently co-transfected with plasmids encoding Myc- (C) or HA-tagged 5-HT₄R (B, D), HA-5-HT₆ R (B), Myc-PAC₁R (C) or HA-M₃R (D) and SEAP-tagged APP. 24 hrs after transfection, protein lysates were immunoprecipitated as described in A. (E) Data correspond to the ratio of ADAM10 co-immunoprecipitation to 5-HT₄R immunoprecipitation signals. ND, not detectable. CTL, Condition transfected with only SEAP-tagged APP. The data illustrated are representative of three independent experiments performed with different sets of cultured cells.

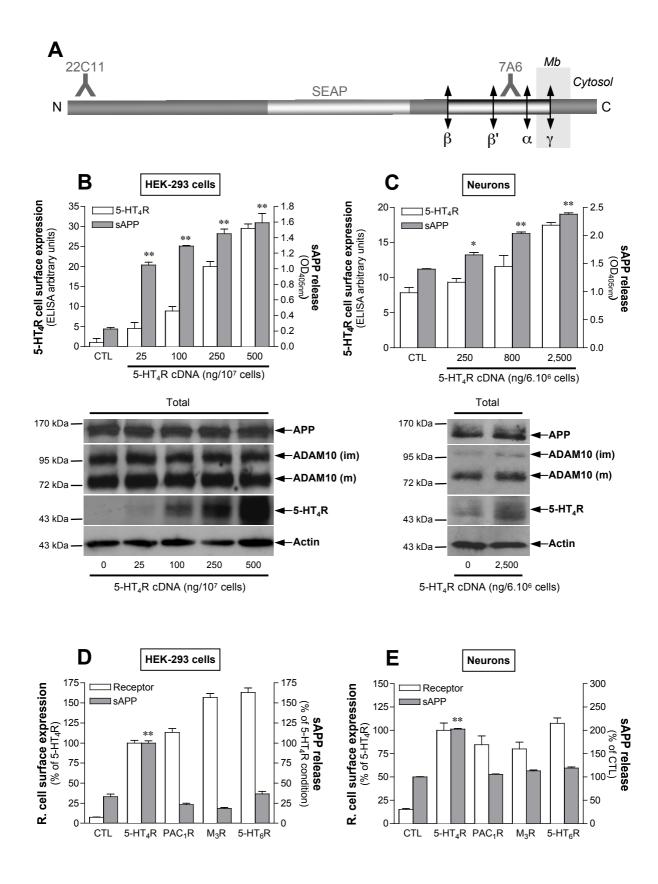
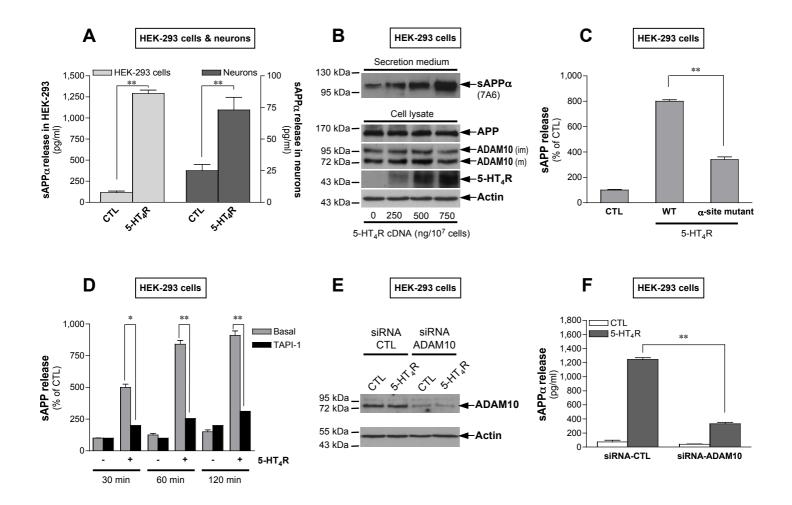


Figure 1



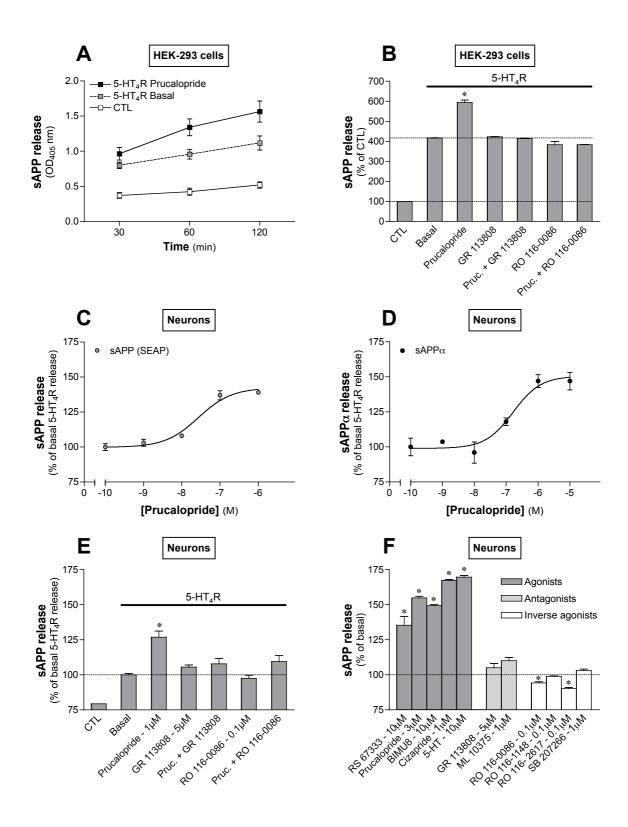
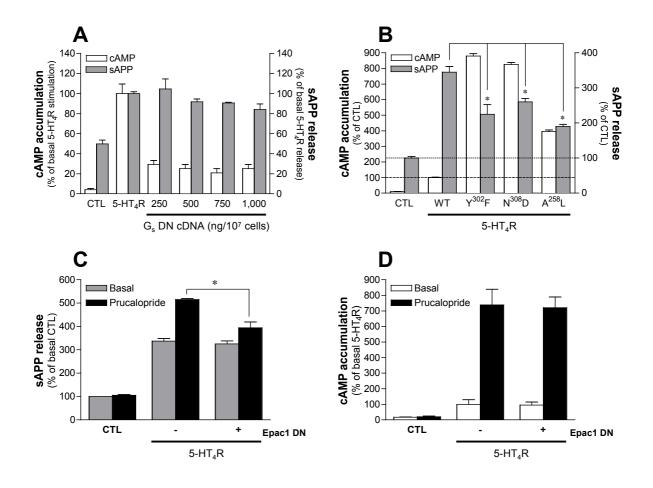


Figure 3



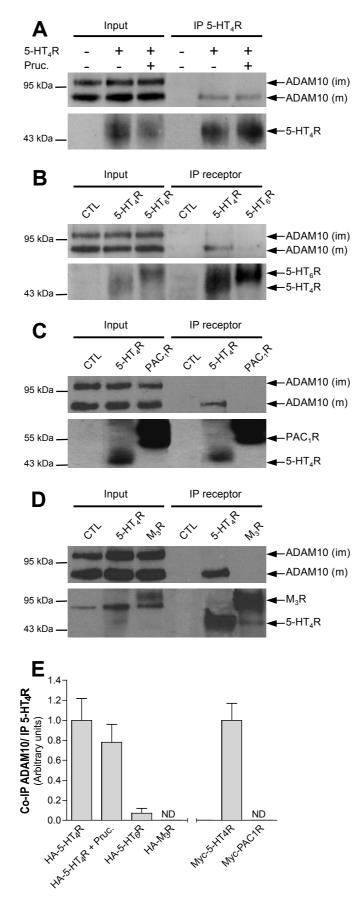


Figure 5

5-HT₄ RECEPTORS CONSTITUTIVELY PROMOTE THE NON-AMYLOIDOGENIC PATHWAY OF APP CLEAVAGE AND INTERACT WITH ADAM10

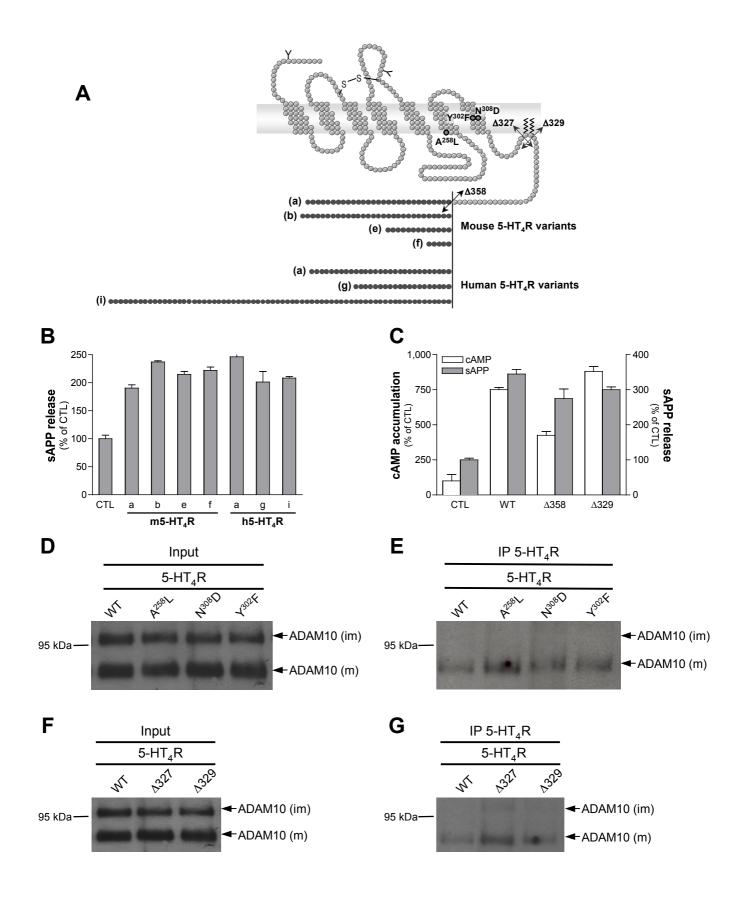
Maud Cochet, Romain Donneger, Elisabeth Cassier, Florence Gaven, Stefan F. Lichtenthaler, Philippe Marin, Joël Bockaert, Aline Dumuis^{*} and Sylvie Claeysen^{*}

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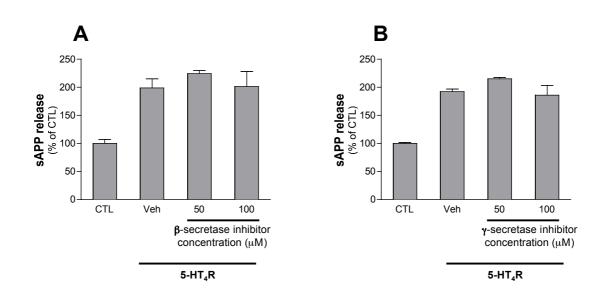
SUPPLEMENTAL FIGURES

- **Supplemental Figure 1:** 5-HT₄R variants (mouse a, b, e and f; human a, g and i), truncated forms (Δ 358 and Δ 329) and mutants ($A^{258}L$, $N^{308}D$ and $Y^{302}F$) promote sAPP release and physically interact with ADAM10.
- **Supplemental Figure 2:** sAPP release induced by 5-HT₄R expression is not affected by β or γ -secretase inhibitors.

SUPPLEMENTAL FIGURE LEGENDS



Supplemental Figure 1



SUPPLEMENTAL FIGURE LEGENDS

Supplemental Figure 1: 5-HT₄R variants (mouse a, b, e and f; human a, g and i), truncated forms (Δ 358 and Δ 329) and mutants ($A^{258}L$, $N^{308}D$ and $Y^{302}F$) promote sAPP release and physically interact with ADAM10.

A, Schematic representation of 5-HT₄R variants and mutants. (*B-G*) HEK-293 cells were transiently transfected with the indicated plasmids encoding HA-tagged 5-HT₄R variants, mutants or truncated forms (200 ng/10⁷ cells). *B*, *C*, 24 hrs after transfection, cAMP accumulation and sAPP release (alkaline phosphate activity) were measured. Results are the means \pm SD of at least three independent experiments. *D-G*, Protein lysates were immunoprecipitated with the anti-HA antibody. Whole cell extracts and immunoprecipitated material were analysed by Western blotting using the anti-ADAM10 C-terminal antibody. The results are representative of three independent experiments performed with different sets of cultured cells.

Supplemental Figure 2: sAPP release induced by 5-HT₄R expression is not affected by β - or γ -secretase inhibitors.

HEK-293 cells were transiently transfected with plasmids encoding Myc-tagged 5-HT₄R (250 ng/10⁷ cells) and SEAP-tagged APP (500 ng/10⁷ cells). 24 hrs after transfection, cells were treated with vehicle alone or the indicated concentrations of a β -secretase (KTEEISEVN-statine-VAEF) (*A*) or a γ -secrease (L-685,458) (*B*) inhibitor for 2 hrs and sAPP release was evaluated by measuring the alkaline phosphatase activity. Data are the means \pm SD of at least three independent experiments.

Non-amyloidogenic pathway

