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## Hetero-oligomerization of the P2Y<sub>11</sub> with the P2Y<sub>1</sub> receptor controls receptor internalization and ligand selectivity of the P2Y<sub>11</sub> receptor

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Short title:  $P2Y_1 - P2Y_{11}$  receptor hetero-oligomers

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### Abstract

Nucleotides signal through purinergic receptors. The nucleotide P2 receptors are subdivided into ionotropic P2X receptors and metabotropic P2Y receptors. The tissue-specificity of nucleotide signaling requires diversity of function within the family of the purinergic receptors. In the present study, hetero-oligomerization between two metabotropic P2Y receptor subtypes is established. The P2Y<sub>1</sub> and P2Y<sub>11</sub> receptors were found to associate physically, when co-expressed in HEK293 cells. The association was detected by copulldown, immunoprecipitation and Fluorescence Resonance Energy Transfer (FRET) experiments. We found a striking functional consequence of the interaction between the P2Y<sub>11</sub> receptor and the P2Y<sub>1</sub> receptor. The interaction promotes agonist-induced internalization of the  $P2Y_{11}$  receptor. This is remarkable because the  $P2Y_{11}$  receptor by itself is completely endocytosis-reluctant. Co-internalization of these receptors was also seen in 1321N1 astrocytoma cells, co-expressing both  $P2Y_{11}$  and  $P2Y_1$  receptors, upon stimulation with ATP or the P2Y<sub>1</sub> receptor specific agonist 2-MeS-ADP. 1321N1 astrocytoma cells do not express endogenous P2Y receptors. Moreover, in HEK293 cells the P2Y<sub>11</sub> receptor was found to associate functionally with the endogenous P2Y<sub>1</sub> receptors. Treatment of HEK293 cells with siRNA directed against the P2Y<sub>1</sub> receptor diminished the agonist-induced endocytosis of the heterologously expressed P2Y<sub>11</sub>GFP receptor. Pharmacological characteristics of the P2Y<sub>11</sub> receptor expressed in HEK293 cells were determined by recording Ca<sup>2+</sup> responses after nucleotide stimulation. This analysis revealed a ligandspecificity, which was different from the agonist profile established in cells expressing the P2Y<sub>11</sub> receptor as the only metabotropic nucleotide receptor. Thus, the hetero-oligomerization of the P2Y<sub>1</sub> and P2Y<sub>11</sub> receptors allows novel receptor functions of the P2Y<sub>11</sub> receptor in response to extracellular nucleotides.

**Key words**: Purinergic receptor, hetero-oligomerization, pulldown experiment, Fluorescence Resonance Energy Transfer (FRET), endocytosis, calcium rise

### Abbreviations

1321N1, human astrocytoma cell line; 2-MeS-ADP, 2-methylthio adenosine 5'-diphosphate; adenosine-5'-triphosphate; 2'(3')-O-(4-Benzoylbenzoyl)adenosine 5'-ATP, BzATP, triphosphate; [Ca<sup>2+</sup>]<sub>i</sub>, intracellular free calcium concentration; DMEM, Dulbecco's modified Eagles' medium; FCS, fetal calf serum; G418, Geneticine; GFP, green fluorescent protein; GPCR, G protein-coupled receptor; G protein, guaninenucleotide-binding protein; GSH, glutathion sepharose; IP, immunoprecipitation; MATra, Magnet Assisted Transfection; MRS2179, N<sup>6</sup>-methyl-2'-deoxyadenosine-3',5'-bisphosphate; NaHBS, HEPES buffered saline NF157, 8,8'-[Carbonylbis[imino-3,1-phenylenecarbonylimino solution; (4-fluoro-3,1phenylene) carbonylimino]] bis-1,3,5-naphthalenetrisulfonic acid; PAR-2-AP, PAR-2activating peptide; ROI, region of interest; siRNA, small interfering RNA; UDP, uridine-5'diphosphate;

### Introduction

The currently accepted notions of G protein-coupled receptor (GPCR) organization in the plasma membrane include the property that these proteins can oligomerize among each other. These oligomers link distinct signaling pathways and integrate receptor functions [1]. Visualization of rhodopsin dimers and higher order oligomers in native murine disc membranes by transmission electron and atomic force microscopy is an outstanding example of receptor oligomerization [2]. The possible functional consequences of hetero-oligomerization among GPCRs are manifold, including enhanced surface expression [3, 4], change in selectivity towards different G proteins [5] and alteration in the potency of receptor agonists [6]. Cross-internalization as a consequence of GPCR hetero-oligomerization may play an important role in regulating the desensitization of inter-connected signalling pathways. An influence on receptor endocytosis has been reported for several known GPCR hetero-oligomers [7-9]. In Parkinson's disease the hetero-dimerization of dopamine D<sub>2</sub> and adenosine  $A_{2A}$  receptors seems to accelerate the development of tolerance against L-DOPA, as increased adenosine levels in patients treated with L-DOPA lead to chronic activation of both receptors and thus to a greater extent of desensitization [1].

Hetero-oligomerization among metabotropic purinergic receptors originating from different subgroups has already been described. The purinergic receptor family consists of two different subgroups, the P1 and P2 receptors. P1 receptors (A<sub>1</sub>, A<sub>2A</sub>, A<sub>2B</sub>, A<sub>3</sub>) are responsive to adenosine, whereas the metabotropic P2 receptors (P2Y<sub>1,2,4,6,11-14</sub>) are activated by nucleotides as natural agonists [10]. An interaction between members of both subgroups was found for the A<sub>1</sub> receptor in a heteromeric association with the P2Y<sub>1</sub> or P2Y<sub>2</sub> receptor, respectively [11, 12]. The A<sub>1</sub>-P2Y<sub>1</sub> receptor hetero-oligomer was shown to create a P2Y-like adenosine receptor by creating ligand-binding sites with a A<sub>1</sub>-P2Y<sub>1</sub> receptor hybrid selectivity. In contrast, the association of the A<sub>1</sub> receptor with the P2Y<sub>2</sub> receptor did not seem to affect the ligand selectivity at the receptors, but stimulation of the P2Y<sub>2</sub> receptor with UTP in the A<sub>1</sub>-P2Y<sub>2</sub> receptor complex caused uncoupling of the A<sub>1</sub> receptor from G<sub>i</sub> proteins [11].

Oligomerization between different P2Y nucleotide receptor subtypes has not yet been reported. However, this is an important issue, since nearly all tissues express more than one subtype of purinergic receptors which enables a functional crosstalk between them in response to nucleotide release and subsequent degradation. The P2Y<sub>1</sub> and P2Y<sub>11</sub> receptors are found to be co-expressed by several tissues, including specific parts of the brain [13], smooth muscle cells and cells of the immune system [14]. Both receptors respond to adenine nucleotides but prefer different derivatives [15]. Slight variations in the nucleotide binding

pocket of the  $P2Y_{11}$  receptor as compared to that of the  $P2Y_1$  receptor, might account for the difference in agonist selectivities [16].

We suspected that an interaction between the  $P2Y_{11}$  receptor and other P2Y receptors might exist, as the ligand specificity of the endogenously expressed  $P2Y_{11}$  receptors does not always agree with the profile found at the cloned and heterologously expressed  $P2Y_{11}$ receptor. In the present study we provide evidence showing that the observed discrepancies might be due to the formation of hetero-oligomers between the  $P2Y_1$  and  $P2Y_{11}$  receptor. In addition, we found that the hetero-oligomer formation induces internalization of  $P2Y_{11}$ receptor, which by itself is endocytosis-reluctant. This has an important impact on  $P2Y_{11}$ receptor desensitization.

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### **Material and Methods**

### Materials

Geneticine (G418 sulphate; Calbiochem, Darmstadt); 2-MeS-ADP, ATP, BzATP, MRS2179, UDP, UTP (Sigma, Deisenhofen); Rp/Sp-ATP- $\alpha$ -S (Biolog, Bremen); Dulbecco's modified Eagle's medium (DMEM), DMEM/Ham's medium, Penicillin/Streptomycin (10 000/10 000 U/ml), Trypsin/EDTA (0.05%/0.02%), fetal calf serum (FCS), (Seromed, Biochrom, Berlin); cell culture dishes (Nunc, Wiesbaden); cover slips (22 mm) (OmniLab); fura 2-AM (Biomol, Hamburg/Molecular Probes); Complete protease inhibitor cocktail, DOTAP and Fugene 6 transfection reagents (Roche, Mannheim, Germany); Glutathione sepharose (GE Healthcare, München, Germany); cDNA plasmid for Cerulean Fluorescent Protein was a kind gift of Dr. David W. Piston (Vanderbilt University Medical Center, Nashville, TN, USA). NF157, a P2Y<sub>11</sub> receptor antagonists with nanomolar potency, was synthesized by M. U. K. by methods published previously [17]. The DNA sequence of the human P2Y<sub>11</sub> receptor (GenBankTM/EBI accession number AF030335) was kindly provided by Dr. D. Communi (Brussels, Belgium).

### **Receptor constructs**

The P2Y<sub>1</sub> receptor was constructed bearing a myc-epitope or a Cerulean fluorescent protein tag at its C-terminus. The P2Y<sub>1</sub> receptor DNA sequence was placed between the EcoRI/SmaI restriction sites of the pcDNA3.1/mycHis (A) vector (Invitrogen, Karlsruhe, Germany) or peGFPN1 vector (Clontech, Heidelberg, Germany). For generating the P2Y<sub>1</sub> receptor tagged with the Cerulean fluorescent protein, the GFP tag was replaced by the cDNA of Cerulean using PCR cloning. The P2Y<sub>11</sub> receptor tags (GST, GFP, Citrine fluorescent protein and the Cerulean-Citrine double tag) were all present at the C-terminus of the receptor. The GST tag was amplified from pGEX vector (GE Healthcare, München, Germany) using PCR cloning and placed downstream of the P2Y<sub>11</sub> receptor DNA sequence following a FXa cleavage site insert in a pcDNA3.1(+) vector. The fluorescent tagged receptor was generated by placing the P2Y<sub>11</sub> receptor sequence between the EcoRI/BamHI restriction sites of the peGFPN1 or peYFPN1 vector. The Q69M mutation in the YFP-tag to produce the Citrine-tag and the R268A mutation in the P2Y<sub>11</sub> receptor sequence were introduced using the site-directed mutagenesis QuikChange kit (Stratagene, La Jolla, CA, USA). For construction of a FRET positive control, DNA encoding for the Cerulean protein was ligated into the BamHI-digested  $P2Y_{11}$ Citrine vector by PCR cloning. The  $P2Y_4$  and protease-activated receptor (PAR)-2 were present as HA fusion proteins. The N-terminal HA-tagged P2Y<sub>4</sub> receptor was cloned into the pcDNA3.1(+) vector. The PAR-2 receptor was present in the pCMV-HA vector (Clontech) bearing the HA-epitope at the C-terminus of the receptor. All constructs were checked by sequencing analysis.

### Cell culture and transfection

HEK293 cells were grown in DMEM/HAM's F12 medium supplemented with 10% FCS, 100 U/ml penicillin, and 100 IU/ml streptomycin in a 5% CO<sub>2</sub>/ 95% air, humidified atmosphere at 37°C. Transient transfection of receptor constructs was performed using MATra-A (magnet-assisted transfection for adherent cells) reagent (IBA, Göttingen, Germany) as per manufacturer's protocol. Stable transfection of the cells was done using DOTAP transfection reagent as described previously [18]. 1321N1 human astrocytoma cells were cultured in high-glucose DMEM supplemented with 5% FCS, 100 U/ml penicillin, and 100 IU/ml streptomycin at 37°C in 10% CO<sub>2</sub>. Transfection of the cells was done using Fugene 6 as described before [15]. A stable expression of the receptor constructs was achieved by keeping G418 throughout the culture period in the medium of both cell lines.

### **Co-Pulldown and Co-Immunoprecipitation**

Transiently transfected (48 h) HEK293 cells were lysed in RIPA buffer (50 mM Tris, pH 8.1; 1% Igepal CA630; 0.25% Na-deoxycholate; 150 mM NaCl; 1 mM EDTA; 1 mM NaF; protease inhibitor cocktail), harvested by scraping, briefly sonicated and centrifuged at 14.000 x g for 15 min at 4°C. The whole cell lysate was collected from the resulting supernatant. The protein content was determined by the Bradford method [19]. For Co-Pulldown experiments whole cell lysates were diluted with RIPA-buffer to a final protein concentration of 500 µg/ml and incubated overnight on a rocker at 4°C with Glutathione Sepharose (GSH-beads). For Co-IP whole cell lysates were diluted with RIPA-buffer to a final protein concentration of 250 µg/ml and incubated with 1.5 µg anti-myc antibody (mouse monoclonal, Invitrogen) overnight as described above. Protein A/G Plus Agarose (Santa Cruz, Heidelberg, Germany) was added and mixed with the lysates for at least 2 h at 4°C. Pulldown fractions or immunoadsorbents were recovered by centrifugation and washed three times in lysis buffer. Samples were eluted in 10 µl 2X Laemmli buffer to prepare for loading on a SDS-PAGE.

### Western Blot analysis

Samples were resolved by SDS-PAGE, followed by electro-transfer to nitrocellulose membrane. The membrane was blocked and probed with antibodies against the myc-epitope (Invitrogen, 1:5000), GST-tag (Santa Cruz, 1:20.000) and HA-epitope (Cell Signalling, 1:1000). After washing, the membrane was further incubated with peroxidase-conjugated anti-rabbit IgG or anti-mouse IgG (1:20.000 or 1:10.000, resp.) and visualized using the

SuperSignal West Pico chemiluminescent substrate (Pierce, Rockford, IL, USA). Membranes were stripped before reprobing.

### Immunocytochemistry

Transiently transfected 1321N1 cells cultured on 22 mm coverglasses were fixed in 4% paraformaldehyde, blocked and permeabilised in FSBB buffer and incubated in mouse monoclonal anti-myc antibody (Invitrogen; 0.8  $\mu$ g/ $\mu$ l) overnight at 4°C. After washing, the secondary antibody was applied (Alexa IgG Goat-anti-mouse A<sub>555</sub>-1:200: Molecular Probes) for 2 h at room temperature. Cells were washed again and mounted in Vectashield mounting medium (Axxora, Grünberg, Germany). Visualization of the GFP fluorescence and immunostaining was done with a Zeiss LSM510 Confocal laser scanning microscope equipped with a Plan-Apochromat x63 objective. GFP was excited using a 488-nm argon/krypton laser, and the emitted fluorescence was detected with a 505-530-nm band pass filter. Alexa Fluor 555 was detected using a 543-nm helium/neon laser and a 560-nm long pass filter.

### **FRET** assay

HEK293 cells transiently transfected (48 h) with the fluorescent tag receptors (P2Y<sub>1</sub>Cerulean, P2Y<sub>11</sub>Citrine or P2Y<sub>11</sub>Cerulean-Citrine) or the fluorescent tags only were imaged on a Zeiss LSM510 confocal laser scanning microscope equipped with a Plan-Apochromat x63 objective. For FRET analysis, emission spectra were measured at 458 nm excitation employing an argon/krypton laser and a beamsplitter NT80/20. The emitted fluorescence was detected from 473 nm to 580 nm in 10.6 nm steps using the lambda modus. The Cerulean protein showed an emission maximum at 483 nm and the emission maximum of the Citrine protein was at 526 nm. HEK293 cells co-transfected with Cerulean and Citrine fluorescent proteins served as negative control. The double tagged P2Y<sub>11</sub> receptor bearing a Cerulean tag that was connected to a Citrine tag by a 16 amino acid long linker was used as a FRET positive control. Fluorescence intensities (I) were normalized to the emission maximum of the Cerulean tag (donor) at 483 nm to display sample spectra in the graph. FRET ratios were calculated using the following equation:  $ERET = (R) = \frac{I_{526mn}}{R}$ 

calculated using the following equation:  $FRET_{ratio}(R) = \frac{I_{526nm}}{I_{483nm}}$ .

### Agonist-induced internalization

Agonist-induced internalization was measured as described previously [20]. Briefly, HEK293 cells stably expressing the P2Y<sub>11</sub>GFP receptor were seeded on coverslips (30 mm diameter). Coverslips were placed in a Perfusion, Open and Closed (POC)-chamber (PeCon GmbH, Erbach, Germany) and incubated on stage in 5% CO<sub>2</sub> at 37°C in complete culture medium.

Images were taken on a Zeiss inverted LSM 510 META laser scanning confocal microscope. Cells were stimulated with various agonists and live-imaging was done for 60 min. Fluorescence intensities of confocal images were analyzed using the Zeiss LSM 510meta software histo macro. Regions of interest (ROI = ellipse 30-39  $\mu$ m<sup>2</sup>) were set in the cytosol of single cells and the average fluorescence intensity in the ROI was determined over time. The intensity values for 10-60 min were normalized to the starting value at 0 min.

### siRNA

 $P2Y_1$  receptor specific siRNA [sense r(CUC UCC UCU GAG GAG AAA A)dTdT] and non-silencing control siRNA were from Qiagen (Hilden, Germany). HEK-P2Y<sub>11</sub>GFP cells were seeded on coverslips (30 mm diameter) pre-coated with 0.01% poly-L-lysine and transfected at 30-40% density with siRNA using MATra-A reagent as described above. Experiments and analysis of the P2Y<sub>1</sub> receptor knockdown were done 48 h post-transfection. For western blot analysis of the P2Y<sub>1</sub> receptor knock-down the nitrocellulose membrane was incubated with an anti-P2Y<sub>1</sub> antibody (Alomone, München, Germany) at a 1:500 dilution.

### **Calcium measurements**

The cells were plated on coverslips (diameter = 22 mm), and single cell measument was done after 3 days, when the cells were 30-50% confluent. The changes of free intracellular Ca<sup>2+</sup> concentration ( $[Ca^{2+}]_i$ ) were measured, as described before [21]. Briefly, cells were preincubated with 2 µM fura-2AM (for 1321N1cells : plus 0.02% pluronic acid) at 37°C for 30 min in NaHBS (HEPES buffered saline solution: 145 mM NaCl, 5.4 mM KCl, 1.8 mM CaCl<sub>2</sub>, 1 mM MgCl<sub>2</sub>, 25 mM glucose and 20 mM HEPES/ Tris pH 7.4). Stimulation of the cells with different concentrations of various agonists was done under continuous superfusion with a flow rate of 1 ml/min at 37°C [18]. Fluorescence intensity was recorded alternately at 340 and 380 nm excitation and 520 mm emission.

### Data analysis

Statistical analysis of the data was done using Graph pad prism. Data were checked for gaussian distribution. In the case of gaussian distribution of data, further analysis was done using the one-way ANOVA test and the Tukey or Dunett test as a post test. Significant difference was taken at a P value < 0.05.

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### **Results and Discussion**

### Physical association of the P2Y<sub>1</sub> and P2Y<sub>11</sub> receptors in HEK293 cells

In the current study we investigated the possible interaction of two members of the P2Y receptor family. The P2Y<sub>1</sub> receptor and P2Y<sub>11</sub> receptor are close homologues in their receptor family. So far a crosstalk between both receptors has not been hypothesized. For the P2Y<sub>1</sub> receptor the ability to interact with other GPCRs became obvious in a study exploring the hetero-dimerization with the A<sub>1</sub> adenosine receptor [12]. In the present study, we analysed whether there is a physical interaction of the P2Y<sub>11</sub> receptor with the P2Y<sub>1</sub> receptor. For that purpose we carried out co-pulldown, co-immunoprecipitation, and FRET experiments.

The P2Y<sub>11</sub> receptor expressed as a GST fusion protein and the P2Y<sub>1</sub> receptor bearing a mycHis tag were co-expressed in HEK293 cells. This allowed us to check for their physical interaction by pulldown and immunoprecipitation experiments using whole cell lysates (Figs. 1 and 2). Western blot analysis of lysates from co-transfected cells revealed smearlike antimyc reactive bands between 35 kDa and 75 to 100 kDa (Fig. 1A, lane 1). The smearlike staining probably represents several glycosylated forms of the receptor, as reported previously by Nakata and co-workers [12]. In addition, the homogenate may also contain receptor proteins, which are not fully processed, and higher molecular mass bands may correspond to oligomerized receptor. The main band of the P2Y<sub>1</sub> receptor protein is indicated by arrowhead in Fig. 1A.Western blot analysis applying a GST-antibody resulted in anti-GST reactive bands of the predicted molecular mass (~65 kDa) for the P2Y<sub>1</sub>IGST receptor (Fig. 1B, lane 1).

Pulldown experiments with lysates of cells co-expressing both receptors (Fig. 1A,B, lane 5) showed that the P2Y<sub>1</sub>mycHis receptor (Fig. 1A, lane 5) was caught together with the P2Y<sub>11</sub>GST receptor (Fig. 1B, lane 5) at the GSH beads. The prominent band marked by arrowheads, as well as a band below that, was greatly enriched at the GSH beads. The interaction of the P2Y<sub>1</sub> and P2Y<sub>11</sub> receptor was strong, as seen from the myc-staining in the pulldown sample. There the staining was much more intense than in the 10% input (Fig. 1A, compare lanes 5 and 1, respectively). The association seems to be constitutive because agonist addition did not alter the interaction of both receptors (data not shown).

To test for the specificity of the pulldown experiment, several control conditions were applied. The GST pulldown experiment with single transfected HEK293 cells, expressing either the P2Y<sub>1</sub>mycHis or the P2Y<sub>11</sub>GST receptor showed no myc-immunoreaction in the Western blot analysis (data not shown). Also, the P2Y<sub>1</sub>mycHis receptor did not interact with the GST tag, which was examined using GST-preloaded GSH beads in a pulldown test (Fig. 1A, lane 6). This was further confirmed by cleaving the P2Y<sub>11</sub> receptor from the GST fusion

protein at the Factor Xa (FXa) cleavage site using this protease. The FXa treatment led to the disappearance of the myc-immunoreaction in the pulldown sample (Fig. 1A, lane 7). When we used a mixture of lysates from HEK293 cells transfected with a single receptor, HEK-P2Y<sub>1</sub>mycHis or HEK-P2Y<sub>11</sub>GST, the GST-pulldown fraction did not show any myc-immunoreaction (Fig. 1A, lane 4). This excludes that the formation of aggregates during the pulldown procedure might be a possible cause of the positive co-pulldown result [22].

We further investigated the specificity of the interaction between both receptors. The P2Y<sub>11</sub>GST receptor was co-transfected with a non-related GPCR. Here we used the protease-activated receptor-2 (PAR-2) (bearing a HA tag). The PAR-2-HA protein showed a smear band (~37-70 kDa, Fig. 1C, lane 2) which is consistent with previous results [23]. The pulldown experiment in Fig. 1C, lane 8, revealed no HA-immunoreactivity, thereby excluding an interaction of the P2Y<sub>11</sub> receptor with the PAR-2 receptor. However, co-transfection of the P2Y<sub>11</sub>GST receptor with the P2Y<sub>4</sub>-HA nucleotide receptor (~37 kDa) resulted in a physical interaction of both receptors, as demonstrated by GST pulldown experiment and subsequent Western blot with an anti-HA antibody (Fig. 1C, lane 9). This implies the interesting possibility of P2Y receptor pooling in HEK293 cells.

The stability of the P2Y<sub>1</sub> and P2Y<sub>11</sub> receptor interaction was further examined by coimmunoprecipitation of the P2Y<sub>11</sub>GST receptor by an anti-myc antibody from P2Y<sub>1</sub> myc / P2Y<sub>11</sub> GFP co-transfected cells. The P2Y<sub>11</sub>GST receptor clearly co-immunoprecipitated together with the myc-tagged P2Y<sub>1</sub> receptor, as detected by Western blot analysis using a GST-antibody (Fig. 2A, lane 3). The signal was specific, as lysates precipitated with an unspecific IgG showed no anti-GST-reactive bands (Fig. 2A, lane 4). Moreover, like in the pulldown test, a mixture of lysates from single receptor-transfected HEK293 cells did not result in co-precipitation of the P2Y<sub>11</sub>GST receptor (Fig. 2A, lane 2).

Furthermore, we analyzed the probability that the transiently overexpressed receptors were present in small membrane patches of HEK293 cells without direct physical interaction [22]. Therefore, the total lysates of double-transfected cells were extracted for 2 h at 4°C in lysis buffer and then centrifuged at 50.000 x g for 30 min to remove undissolved membrane patches [24]. After that procedure, the P2Y<sub>1</sub>mycHis receptor (Fig. 3A, lane 1 and 2) and the P2Y<sub>11</sub>GST receptor (Fig. 3B, lane 2) were present in the extract. When we performed a pulldown experiment with these extracts the extracted P2Y<sub>1</sub>mycHis receptor (Fig. 3A, lane 4) could be found together with the P2Y<sub>11</sub>GST receptor (Fig. 3B, lane 4) at the GSH beads. This confirmed the direct physical interaction of the two receptors.

Moreover, in a FRET assay we investigated the physical association of the P2Y<sub>1</sub> and P2Y<sub>11</sub> receptor in living HEK293 cells. As donor molecule we employed an improved variant of the cyan fluorescent protein named Cerulean which is especially useful for FRET [25]. Citrine fluorescent protein was used as acceptor molecule. This version of the yellow fluorescent protein shows enhanced pH and chloride resistance, thereby improving its usefulness in a FRET assay [26]. The P2Y<sub>1</sub> receptor bearing the Cerulean tag and the P2Y<sub>11</sub> receptor bearing the Citrine tag were transiently co-expressed in HEK293 cells and the fluorescence spectra were determined, as described in methods. Negative control conditions were mimicked by co-expressing the fluorescent tags only. A representative spectrum of the negative control is shown in Fig. 4A (dotted line) and the FRET ratio (n=6) was calculated as described in methods. Maximum FRET ratio (n=6) in our experiments was obtained by expressing the  $P2Y_{11}$  receptor double-tagged with a Cerulean tag linked to a Citrine tag by a 16 amino acid linker (positive control) (Fig. 4B). Excitation of Cerulean at 458 nm resulted in an energy transfer to the linked Citrine protein leading to an emission maximum at the appropriate wavelength of 526 nm (Fig. 4A, line with dash-dots). The emission maximum of the positive control correlated with the fluorescence spectrum of Citrine alone (data not shown). Confocal imaging of HEK293 cells co-expressing the P2Y<sub>1</sub>Cerulean and P2Y<sub>11</sub>Citrine receptor also showed an emission maximum at 526 nm (Fig. 4A, continous line). The FRET ratio (n=9) was smaller than in the positive control but clearly increased as compared to the negative control (Fig. 4B). This suggests a greater distance of the fluorophores in case of the hetero-oligomerized receptors than in case of the positive control, where both fluorescent proteins were closely connected by a linker of 16 amino acids.

# Interaction of P2Y<sub>11</sub> and P2Y<sub>1</sub> receptors promotes agonist-induced internalization of the endocytosis-resistant P2Y<sub>11</sub> receptor

The functional consequence of the formation of  $P2Y_1-P2Y_{11}$  receptor hetero-oligomers was studied by checking the impact on receptor endocytosis of the heterologously expressed receptors. For these studies we used 1321N1 astrocytoma cells, because this cell line lacks endogenous P2Y receptors [15, 27]. 1321N1 cells were firstly single transfected with the GFP-tagged P2Y<sub>1</sub> or P2Y<sub>11</sub> receptor to check for endocytosis properties of the respective individual receptor. Cells were imaged on a Zeiss inverted LSM 510 META laser scanning confocal microscope before and during agonist stimulation, as described in the methods section.

1321N1-P2Y<sub>1</sub>GFP cells were stimulated with 10  $\mu$ M 2-MeS-ADP and the receptor trafficking was observed for 60 min. After 30 min of stimulation, receptor endocytosis could be clearly detected (data not shown) and was complete at the end of 60 min (Fig. 5, A2). This resembles the findings of a recent study [28], where the P2Y<sub>1</sub> receptor was found to be internalized with a similar kinetics. In contrast, 1321N1-P2Y<sub>11</sub>GFP expressing cells did not show any sign of receptor endocytosis even after 60 min of stimulation with 100  $\mu$ M ATP (Fig. 5, B2). This substantiates the lack of agonist-induced receptor internalization of the P2Y<sub>11</sub> receptor. The P2Y<sub>6</sub> receptor behaves similarly and shows only slight reduction in cell surface expression after long-time exposure to agonists [29]. In contrast, the other members of the G<sub>q</sub>-coupled P2Y receptor subgroup are able to undergo agonist-induced internalization, as shown in studies investigating the P2Y<sub>1</sub> receptor [30], the P2Y<sub>2</sub> receptor [20] or the P2Y<sub>4</sub> receptor [29].

However, co-transfected 1321N1 cells, expressing the P2Y<sub>1</sub>mycHis and P2Y<sub>11</sub>GFP receptor, yielded different results. Cells were stimulated with either 100 µM ATP or 10 µM 2-MeS-ADP and fixed after 60 min of stimulation. Fixed cells were stained with anti-myc antibody and visualized on a Zeiss inverted LSM 510 META laser scanning confocal microscope. The green fluorescence of the  $P2Y_{11}GFP$  receptor (Fig. 5, C1) and the red fluorescence of the Alexa<sub>555</sub> dye visualized myc-staining (Fig. 5, C2) co-localized nicely at the plasma membrane as shown in the merged image (Fig. 5, C3). Upon addition of 100 µM ATP for 60 min, endocytosis of the  $P2Y_1$  mycHis receptor (Fig. 5, D2), and now also of the P2Y<sub>11</sub>GFP receptor (Fig. 5, D1), was observed. Fig. 5, D3 shows that the receptors cointernalized to the same compartment (yellow colour) as well as to different endocytotic compartments. The zoomed pictures in Fig. 5, D3 emphasize the co-localization of the  $P2Y_1mycHis$  and  $P2Y_{11}GFP$  receptor in intracellular compartments. Similar observations were made after stimulating the co-transfected cells with the P2Y<sub>1</sub> receptor agonist 2-MeS-ADP at 10 µM (Fig. 5, E1-E3). Again the myc-staining of the P2Y<sub>1</sub>mycHis receptor and the green fluorescence of the P2Y<sub>11</sub>GFP receptor co-localized in the same intracellular compartments (Fig. 5, E3). This was an important finding, as Communi et al. reported that 2-MeS-ADP has no activity at the  $P2Y_{11}$  receptor [31]. Therefore, we can conclude that stimulation of P2Y<sub>1</sub> receptors by the selective agonist 2-MeS-ADP caused co-internalization of the  $P2Y_1$ - $P2Y_{11}$  receptor hetero-oligomers.

To further validate this conclusion, the interaction of the  $P2Y_{11}$  receptor with an endogenously expressed  $P2Y_1$  receptor was investigated. HEK293 cells endogenously express the  $P2Y_1$  receptor [32]. Therefore, we stably transfected these cells with the  $P2Y_{11}$  receptor

GFP fusion protein and observed receptor endocytosis by live imaging. Cells were stimulated on stage and imaged on a Zeiss inverted LSM 510 META confocal microscope. Stimulation of the HEK-P2Y<sub>11</sub>GFP cells with ATP clearly induced receptor internalization (Fig. 6, A2-3). This endocytosis was blocked by the specific P2Y<sub>1</sub> receptor antagonist MRS2179 (Fig. 6, B2-3). Endocytosis of the P2Y<sub>11</sub>GFP receptor could also be induced by the P2Y<sub>1</sub> receptorspecific agonist 2-MeS-ADP (Fig. 6, C2-3), whereas the potent P2Y<sub>11</sub> receptor agonist BzATP turned out to be completely ineffective (Fig. 6, D2-3). In addition, preincubation with a high concentration of NF157, a specific and highly potent P2Y<sub>11</sub> receptor antagonist [17], did not inhibit ATP-induced receptor internalization (Fig. 6, E2-3).

As 2-MeS-ADP was able to induce the endocytosis of the P2Y<sub>11</sub>GFP receptor in HEK293 cells, whereas specific ligands of the P2Y<sub>11</sub> receptor were ineffective, we investigated whether an unresponsive P2Y<sub>11</sub> receptor was able to be co-internalized by P2Y<sub>1</sub> receptor activation. We have previously shown that the R268A-P2Y<sub>11</sub> receptor mutant ( $_{A268}P2Y_{11}$ -R) shows a 1000-fold reduced potency for ATP [16]. Therefore, we stably expressed the  $_{A268}P2Y_{11}$  receptor in HEK293 cells. For detection of receptor endocytosis, the HEK $_{A268}P2Y_{11}$ GFP cells were stimulated on stage with an ATP concentration (100 µM) that was not able to activate the mutant receptor, and cells were imaged by confocal microscopy. The results depicted in Fig. 6, F2-3 demonstrate that during the total time period of imaging (60 min) the GFP tagged  $_{A268}P2Y_{11}$  receptor did not show any sign of internalization. The receptor remained at the plasma membrane. The same was true when we stimulated the cells with 2-MeS-ADP (data not shown). Real time PCR showed that HEK<sub>A268</sub>P2Y<sub>11</sub>GFP, mock transfected as well as HEK-P2Y<sub>11</sub>GFP transfected cells all had comparable levels of P2Y<sub>1</sub> receptor mRNA (supplementary material). Therefore, the lack of internalization was not due to a lower P2Y<sub>1</sub> receptor level.

However, the  $_{A268}P2Y_{11}GST$  receptor expressed in HEK293 cells still displayed physical association with the co-transfected P2Y<sub>1</sub>mycHis receptor, as analysed by pulldown experiments from cellular extract. The data from the co-pulldown experiment are shown in Fig. 3A, lane 3. Therefore, we conclude that a mutant P2Y<sub>11</sub> receptor with greatly reduced agonist affinity was not able to allow the functional interaction with the P2Y<sub>1</sub> receptor.

Interestingly, the induction of the endocytosis of the wild type  $P2Y_{11}GFP$  receptor in HEK293 cells was specific for  $P2Y_1$  receptor agonists, because stimulation of the cells with UTP acting on the endogenous  $P2Y_2$  receptor (Fig. 6, G2-3) was not able to induce endocytosis of the  $P2Y_{11}GFP$  receptor. Also, stimulation with PAR-2-activating peptide, which can specifically activate the PAR-2 receptor without proteolysis [33, 34], did not result

in endocytosis of the  $P2Y_{11}GFP$  receptor (Fig. 6, H2-3). Both stimuli elicited a clear calcium response in the cells (data not shown) and real time PCR analysis confirmed the expression of the  $P2Y_2$  receptor and PAR-2 receptor in HEK- $P2Y_{11}GFP$  cells (supplementary material).

To further validate the influence of the  $P2Y_1$  receptor in the process of  $P2Y_{11}$  receptor internalization, we attempted to knock-down P2Y<sub>1</sub> receptor expression in HEK293 cells. Oligonucleotides selected against a specific region in the coding sequence of the receptor were applied to HEK-P2Y<sub>11</sub>GFP cells using magnet-assisted transfection (MATra) reagent. Cells treated with the P2Y<sub>1</sub> receptor siRNA showed approximately 50% knock-down in the mRNA level of the  $P2Y_1$  receptor (data not shown). The decrease in  $P2Y_1$  receptor expression, as confirmed on the protein level, was about 50% (Fig. 7, inset). Receptor expression remained unaffected when cells were treated with MATra reagent only or with control siRNA (data not shown). Treatment of the HEK-P2Y<sub>11</sub>GFP cells with P2Y<sub>1</sub> receptor siRNA induced a significantly diminished P2Y<sub>11</sub>GFP receptor endocytosis. The endocytosis was quantified by measuring the time course of the increase in GFP fluorescence in the cytosol of transfected cells upon agonist stimulation (Fig. 7). The time-dependent increase in GFP fluorescence intensity in the cytosol of ATP-stimulated cells was significantly smaller in P2Y<sub>1</sub>-siRNA treated cells than in cells, which were not treated with P2Y<sub>1</sub>-siRNA. The extent of reduction in P2Y<sub>1</sub> receptor mRNA transcripts correlated with the reduction in endocytosis of the P2Y<sub>11</sub> receptor. Cells treated with control siRNA or only transfection reagent behaved like control cells (Fig. 7). This verifies the specificity of the interaction of  $P2Y_1$  and  $P2Y_{11}$ receptors in the receptor endocytosis of the P2Y<sub>11</sub>GFP receptor in HEK293 cells.

In conclusion, the phenomenon of co-internalization may be a very important functional consequence of GPCR hetero-oligomerization. This controls desensitization and resensitization of GPCRs [35]. Hetero-oligomerization among GPCRs can induce internalization of receptors which were previously considered reluctant to endocytosis. The somatostatin receptor (SSTR) SSTR1 failed to internalize upon agonist stimulation in several cell lines. However, the SSTR1 present as a heterodimer with the SSTR5 was found to display sufficient agonist-induced internalization [36].

Similar findings are presented here for the  $P2Y_1$  and  $P2Y_{11}$  receptor. The  $P2Y_{11}$  receptor, when expressed alone, lacks any desensitization, as shown here and found previously by others [37]. This property of the the  $P2Y_{11}$  receptor seems to be important for normal receptor function in the maturation process of dendritic cells (DC) which express  $P2Y_{11}$  receptors. These antigen-presenting cells are trapped in the epicenter of inflammation due to activation of the  $P2Y_{11}$  receptor by high extracellular ATP concentrations. This

prolongs exposure to maturation-inducing factors [38]. Thereby, the inability of the P2Y<sub>11</sub> receptor to desensitize seems to be important in overriding the chemotactic effects of nucleotides acting on other P2Y receptors. However, distinct DC subsets respond differently to ATP in terms of migration. DCs directly isolated from human blood show no arrest in migration in response to ATP treatment. Interestingly, these DC subpopulations display significant mRNA expression levels for the P2Y<sub>1</sub> receptor, opposed to a negligible PCR signal usually found in DCs (Langerhans cells, monocyte-derived DCs) which do not migrate in the presence of high ATP concentrations [38]. We hypothesize that the presence of the P2Y<sub>1</sub> receptor in DCs isolated from human blood enables the desensitization of the P2Y<sub>11</sub> receptor in response to ATP, and this controlls the chemotactic behaviour and maturation of the cells.

### Influence of the hetero-oligomerization on the ligand selectivity of the P2Y<sub>11</sub> receptor

The hetero-oligomerization of the P2Y<sub>11</sub>GFP receptor with the endogenous P2Y<sub>1</sub> receptor seems to modify the activity of known agonists and antagonists at the receptor, as observed in the endocytosis experiments described above. BzATP was ineffective in inducing receptorendocytosis and NF157 could not inhibit the ATP-induced internalization of the P2Y<sub>11</sub>GFP receptor. So far there is already evidence that formation of hetero-oligomers can result in a change in pharmacology of GPCRs [39]. The A<sub>1</sub>-P2Y<sub>1</sub> receptor hetero-dimer was found to have no significant affinity for MRS2179, whereas A<sub>1</sub> receptor antagonists did still bind. In contrast, the modified ligand binding pocket in the A<sub>1</sub>-P2Y<sub>1</sub> heterodimer appears to bind the P2Y<sub>1</sub> receptor agonist ADP $\beta$ S, but slightly less the A<sub>1</sub> receptor ligands [12]. Therefore, we also investigated receptor ligand selectivities. This was done by measuring the intracellular calcium rise in HEK293 cells expressing the P2Y<sub>11</sub>GFP receptor. As described in methods, cells were subjected to single-cell calcium measurements using the calcium indicator fura-2.

P2Y<sub>11</sub>GFP receptor-expressing HEK293 cells responded to the potent P2Y<sub>11</sub> receptor agonist BzATP, but the recently developed, highly potent P2Y<sub>11</sub> receptor antagonist NF157 [17] was not able to inhibit this response (Fig. 8A). However, when HEK-P2Y<sub>11</sub>GFP cells were preincubated with the specific P2Y<sub>1</sub> receptor antagonist MRS2179, surprisingly the response to BzATP was inhibited (Fig. 8A). This was different when the P2Y<sub>11</sub>GFP receptor was expressed singly in 1321N1 cells. There MRS2179 showed no effect on the activity of BzATP, but preincubation with NF157 totally abolished the activity of BzATP at the P2Y<sub>11</sub> receptor (Fig. 8C). MRS2179 was able to interfere with the BzATP-stimulated [Ca<sup>2+</sup>]<sub>i</sub> rise and the ATP-induced internalization of the P2Y<sub>11</sub>GFP receptor, whereas NF157 was not able to inhibit any of these effects. Thus, the interaction of the  $P2Y_{11}GFP$  receptor with the endogenous  $P2Y_1$  receptor in HEK293 cells apparently influences the ligand selectivity of the receptor, as measured by the induction of a  $[Ca^{2+}]_i$  rise.

BzATP normally shows no activity at the P2Y<sub>1</sub> receptor [40]. Therefore, it can be concluded that BzATP and MRS2179 both bind to the hetero-dimer and that the ligand selectivity of both receptors is changed due to dimerization. However, there are some surprises regarding the activity of BzATP. This ligand could induce a  $[Ca^{2+}]_i$  increase in HEK-P2Y<sub>11</sub>GFP cells but was not able to stimulate receptor endocytosis. Such inconsistencies in the action of BzATP have also been found by other groups [41, 42]. Taken together, this indicates that BzATP can only induce specific receptor conformations in accordance with the probabilistic model of GPCR function [43].

The A268P2Y<sub>11</sub>GFP receptor did not functionally interact with the endogenous P2Y<sub>1</sub> receptor in HEK293 cells with regard to internalization. However, there was a physical interaction (Fig. 3). Therefore, we examined whether the ligand selectivity of this P2Y<sub>11</sub> receptor mutant was influenced by the interaction with the P2Y<sub>1</sub> receptor. We measured the intracellular calcium rise in HEK<sub>A268</sub>P2Y<sub>11</sub>GFP cells. The cells responded to 100  $\mu$ M ATP in the same way as mock-transfected cells, confirming the decreased potency of ATP at the P2Y<sub>11</sub> receptor mutant (Fig. 8B). However, the more potent P2Y<sub>11</sub> receptor agonist BzATP elicited a significant response at a concentration of 100  $\mu$ M at the R268A mutant (Fig. 8A). Pretreatment of the HEK<sub>A268</sub>P2Y11GFP cells with MRS2179 did not affect the action of BzATP at the receptor, in contrast to the observation made at the unmutated receptor. Therefore, this arginine seems to be a critical residue in the newly formed nucleotide binding pocket of the hetero-oligomer, as the physical interaction of both receptors was unaffected after mutation of this residue.

Two theories are currently discussed as to how GPCR oligomers are formed. One is the ,contact dimerization' and the other is the ,domain swapping' theory [1]. In the case of the  $P2Y_1-P2Y_{11}$  receptor hetero-dimer our findings suggest that it is formed as a ,domain swapped' dimer because the change in the pharmacological profile of P2Y receptor ligands implies the creation of a newly formed binding pocket. In agreement with this suggestion is our finding that UDP induced a drastic  $[Ca^{2+}]_i$  increase in HEK-P2Y<sub>11</sub>GFP cells (data not shown). UDP is known to activate the P2Y<sub>6</sub> receptor, but this receptor was not expressed in our HEK293 cells. Interestingly, previous studies have reported that uridine nucleotides can activate the P2Y<sub>11</sub> receptor [44]. However, at the P2Y<sub>11</sub> receptor expressed in our batch of 1321N1 cells we could not confirm this finding. Our new findings in HEK293 cells suggest that the activity of uridine nucleotides might depend on the presence of the  $P2Y_1$  receptor.

### Conclusion

The present study provides convincing evidence for a physical interaction of the P2Y<sub>1</sub> and the P2Y<sub>11</sub> receptors in hetero-oligomers. This hetero-oligomerization enables the agonist-induced internalization of the P2Y<sub>11</sub> receptor. The P2Y<sub>11</sub> receptor expressed alone is reluctant to endocytosis and lacks desensitization. This property of the P2Y<sub>11</sub> receptor seems to be important for normal P2Y<sub>11</sub> receptor function. We hypothesize that the presence of P2Y<sub>1</sub> receptors in dendritic cells isolated from human blood enables the desensitization of the P2Y<sub>11</sub> receptor in response to ATP, whereby chemotactic behaviour and maturation of these cells are controlled. This is an important functional consequence of the heteromeric association of the P2Y<sub>11</sub> receptor leading to the alteration of its behaviour.

We have previously shown that the R268A-P2Y<sub>11</sub> receptor mutant ( $_{A268}P2Y_{11}$ ) shows a 1000-fold reduced potency for ATP [16]. The  $_{A268}P2Y_{11}$ GFP receptor mutant did not functionally interact with the endogenous P2Y<sub>1</sub> receptor in HEK293 cells, since it did not co-internalize with the activated P2Y<sub>1</sub> receptor. Nevertheless, we found physical interaction between the  $_{A268}P2Y_{11}$  receptor and the P2Y<sub>1</sub> receptor by co-pulldown.

In addition, we also found that the association of the  $P2Y_1$  and the  $P2Y_{11}$  receptor influences the ligand selectivity of the  $P2Y_{11}$  receptor. Therefore we suggest that some discrepancies reported in the literature concerning the responsiveness of the  $P2Y_{11}$  receptor to certain nucleotides could be due to a varying degree of co-expression of the  $P2Y_1$  receptor.

The existence of GPCR homo- or hetero-oligomers is now largely accepted. The functional relevance of GPCR oligomers includes modulation of internalization processes or alteration of their agonist/antagonist potency. Both have to be considered regarding the development of subtype-specific drugs [6]. The importance of GPCR oligomerization becomes apparent, when we consider the organization of receptors and their signaling elements in specific microdomains. This clearly affects cellular communication processes [45]. The homo- or hetero-dimerization of purinergic GPCRs thereby allows fine tuning of the responses of cells to extracellular nucleotides.

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### **Figure legends**

Fig. 1: Co-Pulldown (PD) of the P2Y<sub>1</sub>mycHis and P2Y<sub>11</sub>GST receptors from HEK293 cells. For the pulldown, total lysates of transfected HEK293 cells expressing the indicated receptors were incubated overnight with GSH beads at 4°C, as described in methods. Western blot (WB) analysis was performed using anti-myc (A), anti-GST (B) or anti-HA (C) antibodies. Lane 1-3: Input signals from total lysates (10% of PD samples) of co-transfected HEK293 cells. Lane 4: Mixture of lysates from P2Y<sub>1</sub>mycHis and P2Y<sub>11</sub>GST single transfected cells and subsequent incubation with GSH beads. Lane 5: Co-pulldown of the P2Y1mycHis and  $P2Y_{11}GST$  receptor from lysates of co-transfected cells (compare with lane 1). Lane 6: Lysate of P2Y<sub>1</sub>mycHis single transfected cells incubated with GST-preloaded GSH beads. Lane 7: P2Y<sub>1</sub>mycHis/P2Y<sub>11</sub>GST receptor co-pulldown, beads after incubation with FXa. Lane 8: Lysates of PAR-2-HA/P2Y<sub>11</sub>GST co-transfected cells incubated with GSH beads. Lane 9: Co-pulldown of the P2Y<sub>4</sub>HA and P2Y<sub>11</sub>GST receptor from lysates of co-transfected cells. Approximate molecular mass: P2Y<sub>1</sub>mycHis ~ smearlike staining between 35 and 100 kDa; P2Y<sub>11</sub>GST ~ 65 kDa; PAR-2-HA ~ smearlike bands from 37-70 kDa; P2Y<sub>4</sub>HA ~ 37 kDa. Here and in Figs. 2 and 3, Western blots shown are representative of at least two independent experiments analyzing the various receptors and testing the different conditions. \* indicates positive pulldown signal. The arrowheads point to the main band of the P2Y<sub>1</sub>mycHis protein.

**Fig. 2:** Co-Immunoprecipitation (IP) of the P2Y<sub>1</sub>mycHis and P2Y<sub>11</sub>GST receptor from HEK293 cells. Total lysates of differently transfected HEK293 cells were incubated first overnight with anti-myc antibody and then with Agarose Plus A/G beads at 4°C, as described in methods. Western blot (WB) analysis was performed using anti-GST (A) or anti-myc (B) antibodies. **Lane 1:** Input signal from total lysate (10% of IP samples). **Lane 2:** Mixture of lysates from P2Y<sub>1</sub>mycHis and P2Y<sub>11</sub>GST single transfected cells immunoprecipitated with anti-myc antibody. **Lane 3:** Co-IP of P2Y<sub>1</sub>mycHis and P2Y<sub>11</sub>GST receptor from lysates of co-transfected cells (compare with lane 1) using anti-myc antibody (arrow points to P2Y<sub>11</sub>GST protein). **Lane 4:** Precipitation of lysates from P2Y<sub>1</sub>mycHis/P2Y<sub>11</sub>GST cells using unspecific mouse IgG (negative control). Molecular mass for receptors is given in legend to Fig. 1. The arrowhead indicates a signal corresponding to Protein A. \* the indicated band represents non-reduced IgG due to unboiled samples.

**Fig. 3:** Cellular extracts of differently transfected HEK293 cells were incubated overnight with GSH beads at 4°C, as described in methods. Western blot (WB) analysis was performed

using anti-myc (A) or anti-GST (B) antibodies. Lane 1-2: Input signals from extracts (10% of pulldown samples) of co-expressing cells. Lane 3-4: Co-pulldown of the  $P2Y_1mycHis$  receptor with the <sub>A268</sub>P2Y<sub>11</sub>GST receptor (lane 3) or the wild type  $P2Y_{11}GST$  receptor (lane 4) from lysates of co-transfected cells. Molecular mass of receptors is given in legend to Fig. 1. The arrowheads point to the main band of the  $P2Y_1mycHis$  protein.

**Fig. 4:** Analysis of the hetero-oligomerization of P2Y<sub>1</sub>Cerulean and P2Y<sub>11</sub>Citrine by FRET experiments. HEK293 cells transiently transfected with the fluorescent constructs were imaged by confocal microscopy to detect the fluorescence spectra at 458 nm excitation, as described in methods. (**A**) The graph displays representative emission spectra normalized to fluorescence intensity at 483 nm. (**B**) Analysis of FRET ratios. FRET ratios (R) were calculated using the following equation as explained in the methods section:  $FRET_{ratio}(R) = \frac{I_{526nm}}{I_{483nm}}$ . Statistical analysis was done using the one-way ANOVA test and the

Dunett test as a post test. \*\* p < 0.001

**Fig. 5:** Confocal imaging of internalisation of P2Y<sub>1</sub>GFP and P2Y<sub>11</sub>GFP receptors in transfected 1321N1 cells. (**A**) 1321N1 cells expressing the P2Y<sub>1</sub>GFP receptor were stimulated with 10  $\mu$ M 2-MeS-ADP on stage at 37°C (5% CO<sub>2</sub>) and living cells were imaged with a LSM510meta confocal microscope for up to 60 min. Pictures show unstimulated (US) cells (A1) and cells after 60 min of agonist sumulation (A2). Scale bar indicates 5  $\mu$ m. (**B**) 1321N1 cells stably expressing the P2Y<sub>11</sub>GFP receptor were stimulated with 100  $\mu$ M ATP and measured in the same way as P2Y<sub>1</sub>GFP expressing cells. (**C-E**) Transiently co-transfected 1321N1 cells expressing the P2Y<sub>11</sub>mycHis and the P2Y<sub>11</sub>GFP receptor were stimulated with nucleotides for 60 min, fixed with 4% PFA and stained with anti-myc antibody as described in methods. Images display the GFP fluorescence (C1, D1, E1) and myc-staining (C2, D2, E2) separately, as well as a merged picture (C3, D3, E3). White lines in the merged picture illustrate the x-z or y-z section-plane displayed in the zoomed pictures on top and besides the image. Scale bar indicates 10  $\mu$ m. Pictures shown are representative for at least three independent experiments.

**Fig. 6:** HEK-P2Y<sub>11</sub>GFP cells were visualized with a LSM510meta confocal microscope and stimulated with various agonists for up to 60 min at  $37^{\circ}$ C (5% CO<sub>2</sub>) as described in methods. Pictures show unstimulated (US) cells (A1-H1) and cells after 30 min (A2-H2) and 60 min

(A3-H3) of agonist stimulation. (A) Stimulation with 100  $\mu$ M ATP (n=5). (B) Cells preincubated with 100  $\mu$ M MRS2179 and stimulated with 100  $\mu$ M ATP (n=3). (C) Stimulation with 10  $\mu$ M 2-MeS-ADP (n=3). (D) Stimulation with 100  $\mu$ M BzATP (n=3). (E) Cells preincubated for 30 min with 10  $\mu$ M NF157 and stimulated with 100  $\mu$ M ATP (n=3). (F) Stimulation with 100  $\mu$ M UTP (n=4). (G) Stimulation with 100  $\mu$ M PAR-2-AP (n=3). (H) HEK<sub>A268</sub>P2Y<sub>11</sub>GFP cells stimulated with 100  $\mu$ M ATP (n=3). Results shown are representative for n independent experiments analysing 20-30 cells in each case. Scale bar represents 5  $\mu$ m.

**Fig. 7:** Kinetics of receptor endocytosis analyzed by the increase in fluorescence intensity of the P2Y<sub>11</sub>GFP receptor in the cytosol of siRNA transfected HEK293 cells. Images of HEK-P2Y<sub>11</sub>GFP cells transfected with siRNA and stimulated with 100  $\mu$ M ATP for up to 60 min were taken on a Zeiss LSM510 meta confocal microscope, 48 hours after transfection, and fluorescence intensities were analyzed using the Zeiss software. The average fluorescence intensity was normalised to the value at 0 min as described in methods. Displayed are curves for untreated cells (n=16), MATra transfection reagent treated cells (n=19), control siRNA treated cells (n=18) and P2Y<sub>1</sub> receptor siRNA (n=21) treated cells. Asterisks indicate significance (\* p<0.05) compared to untreated cells, analysed by one way ANOVA and Tukey's test. The inset shows a representative blot depicting the 50% decrease in P2Y<sub>1</sub> receptor antibody.

**Fig. 8**: Influence of hetero-oligomerization on the ligand selectivity of the receptors. Cells were stimulated with 100  $\mu$ M of agonists and preincubated with 100  $\mu$ M of antagonists, unless the subscript indicates treatment with the agent at 1  $\mu$ M concentration. Bar graphs show peak values ( $\Delta$ F340nm/F380nm) from [Ca<sup>2+</sup>]<sub>i</sub> measurements using fura-2. Data represent the mean±s e.m obtained in at least three independent experiments. (**A**) Measurements in HEK293 cells, which were mock transfected (blank bars), P2Y<sub>11</sub>GFP transfected (black bars) or <sub>A268</sub>P2Y<sub>11</sub>GFP transfected (cross-hatched bars). Cells were stimulated with BzATP. Preincubation of the cells with MRS2179 and NF157 are indicated. (**B**) Measurements in transfected HEK293 cells using ATP (100  $\mu$ M) as stimulating agonist. (**C**) Intracellular calcium rise measured in 1321N1 cells, stably expressing the P2Y<sub>11</sub>GFP receptor, evoked by the agonists indicated.

B



B



B

| Extracts<br>1 2<br>75 kDa –<br>A 50 kDa –<br>37 kDa –  | Pulldown<br>3 4          | WB: myc |
|--|--------------------------|---------|
| B 75 kDa   |                          | WB: GST |
| $\begin{array}{rcrcr} P2Y_{1}mycHis & + & + \\ P2Y_{11}GST & - & + \\ A268}P2Y_{11}GST & + & - \\ GSH \ beads & - & - \end{array}$ | + +<br>- +<br>+ -<br>+ - |         |
| Figure 3   |                          |         |
|  |                          |         |





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