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The P2Y12 receptor regulates microglial activation by extracellular nucleotides

by

Sharon E. Haynes

DISSERTATION

Submitted in partial satisfaction of the requirements for the degree of

DOCTOR OF PHILOSOPHY

in

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in the

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of the

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by

Sharon Haynes

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Abstract

The P2Y₁₂ receptor regulates microglial activation by extracellular nucleotides

Sharon Haynes

Microglia are primary immune sentinels of the central nervous system (CNS). Following injury, these cells migrate or extend processes towards sites of tissue damage. CNS injury is accompanied by the release of nucleotides, serving as signals for microglial activation or chemotaxis. Microglia express several purinoceptors, including a Gi-coupled subtype that has been implicated in ATP/ADP-mediated migration in vitro. Here we show that microglia from mice lacking Gi-coupled P2Y12 receptors exhibit normal baseline motility but are unable to polarize, migrate, or extend processes towards nucleotides in vitro or in Microglia in $P2ry12^{-1}$ mice display significantly diminished directional vivo. branch extension towards sites of cortical damage in the living mouse. Moreover, $P2Y_{12}$ expression is robust in the "resting" state, but dramatically reduced after microglial activation. These results suggest that P2Y₁₂ is a primary site at which nucleotides act to induce microglial chemotaxis at early stages of the response to local CNS injury.

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CHAPTER 1

INTRODUCTION

INTRODUCTION

Microglia are the primary mediators of the immune response that occurs following central nervous system (CNS) injury or disease. They are derived from blood monocytes, sharing a developmental lineage with peripheral macrophages, and invade the central nervous system during embryonic development¹. In the healthy adult brain, microglia comprise 10% of total brain cells² and are found in a resting, highly ramified state in which they keep a close surveillance over their local environment. In this "un-activated" state, microglia are rapidly projecting and retracting their elaborate ramifications, possibly as a means of surveying their local environment^{3,4}. When injury or disease strikes, microglia become "activated" and migrate or send cellular projections towards the site of damage³⁻⁵. In this activated state, they take on a macrophage-like phenotype⁵ and are able to phagocytose debris, release cytokines or inflammatory factors, and present antigens to infiltrating T cells^{6,7}. Microglia are thought to be both protective and deleterious to the CNS because the neuromodulating factors they secrete can promote either cell survival or cell death⁸. It has also been suggested that microglia facilitate synapse formation and reconstruction⁹ and modulate pain sensitivity following nerve damage¹⁰.

Microglia use numerous signaling mechanisms to monitor the physical condition of their local environment². For example, extracellular nucleotides released at an

injury site may activate purinoceptors on microglia. Some aspects of microglial response to injury can be mimicked by injection of ATP, or attenuated by broadspectrum inhibitors of purinergic signalling³, suggesting that nucleotides function as regulators of microglial behavior *in vivo*. Specifically, it has been shown that cortical tissue damage inflicted by a two-photon laser induces local microglia to extend cellular projections towards the injury. This response can be abrogated by a saturating concentration of extracellular ATP³, implicating that a nucleotide gradient is important to elicit this behavior.

Purinergic receptors, designated P, are subdivided into classes depending on their mode of action. P1 receptors are activated by adenine and uridine, whereas P2 receptors are activated by ATP, UTP and their derivatives. The postscript X designates those that are ionotrophic, or ligand-gated ion-channels, and Y designates those that are metabotrophic, or G-protein coupled receptors^{11,12}. Nucleotides activate microglia via a number of purinoceptors on the cell surface such as P2X₄, P2X₇, P2Y₁, P2Y₂, and P2Y₁₂¹³. Although the roles of these purinoceptors are poorly understood, it has been shown that P2X₄ may play a role in tactile allodynia after nerve injury through ATP-evoked BDNF release^{14,15} and P2X₇ may induce the release of TNF- α and IL-1 β ¹⁶. Low levels of ATP or ADP exposure induce microglia to release neuroprotective factors, such as plasminogen, whereas high levels induce neurotoxic factors, such as nitric oxide, although the receptor(s) mediating this response is unknown¹³.

Extracellular concentrations can rise rapidly and dramatically when injured cells lose membrane integrity and release their cytosolic contents¹⁷. ATP is released from neurons, astroalia, and microalia¹⁸, and functions in cell-cell communication. Regulated nucleotide release from neurons and astrocytes may occur by exocytosis or through various ATP pumps such as the multi-drug resistance (MDR) protein or cystic fibrosis transmembrane conductance regulator (CFTR)^{19,20}. ATP may be released from microglia following activation and may serve as a positive feedback signal²¹. The role of nucleotides as autocrine and paracrine cell modulators has been well documented in the immune, nervous, and cardiovascular systems. In the immune system, extracellular ATP activates mast cell degranulation as well as induces the formation of a nonselective membrane pore permeable to molecules up to 900Da in lymphocytes and macrophages²⁰. There is evidence that suggests ATP may also be utilized as a chemotactic agent in dendritic cells, hematopoietic cells that share a developmental lineage with microglia and macrophages²². In the nervous system, purinoceptors on neurons mediate ATP-dependent membrane depolarization and action potential firing²³.

It has recently been shown that ATP or ADP can evoke membrane ruffling (actin polymerization) and directed migration of cultured microglia through a pertussis toxin-sensitive signalling pathway, thus implicating G_i -coupled receptor(s)-mediated inhibition of cyclic AMP synthesis²⁴. The vast majority of metabotropic P2Y receptors transduce their signals through G_q /phospholipase C pathways²⁵.

However, in 2001 the first G_i -coupled purinergic receptor was identified, termed P2Y₁₂, and was shown to be expressed in glial cells of the brain^{26,27}. As such, P2Y₁₂ is an attractive candidate for mediating morphological responses of microglia to extracellular nucleotides.

Although ADP is the primary agonist of P2Y₁₂, ATP is a partial agonist; thus ATP acts as an antagonist when receptor reserves are low (perhaps in platelets) and a weak agonist when receptor reserves are high²⁸. Activation of P2Y₁₂ induces the dissociation of heterotrimeric G-protein subunits so that G_{α} may inhibit adenylyl cyclase and $G_{\beta\gamma}$ may activate inwardly rectifying K⁺ (K_{IR}) channels²⁶. In fact, there has been evidence that microglia express K_{IR} channels²⁹, although the role of this channel in microglial physiology is not understood. In addition, P2Y₁₂ has been shown to activate Rac GTPase, which in turn activates actin polymerization at the leading edge of a migrating cell²⁴. Recent evidence suggests that P2Y₁₂ desensitizes after activation via a kinase-dependent mechanism in platelets³⁰ and is localized to lipid rafts in the membrane³¹.

The P2Y₁₂ receptor was initially identified on platelets, where it regulates their conversion from the inactive to active state during thrombosis^{26,32,33}. Thus, growth of a nascent clot is dependent on the release of ADP (and other factors such as thromboxane A_2)³⁴ from activated platelets or damaged red blood cells, thereby promoting a feed-forward mechanism to recruit platelets into the active, clotting competent state. Indeed, peripherally active P2Y₁₂ antagonists, such as

ticlopidine (Ticlid) and clopidogrel (Plavix), block platelet activation and reduce the risk of recurrent heart attack or stroke³⁵. Moreover, mutations in the P2Y₁₂ receptor (in humans or mice) result in prolonged bleeding times due to the reduced ability of P2Y₁₂-deficient platelets to change shape and undergo coagulation in response to ADP released after vascular injury³³.

To determine the physiological role of P2Y₁₂ in microglia, we, in collaboration with researchers formerly of COR Therapeutics, generated mice genetically engineered to lack P2Y₁₂ receptors. In order to facilitate imaging studies of microglia, we crossed $P2ry12^{-r}$ mice with a line of mice genetically engineered to express green fluorescent protein (GFP) specifically in microglia and their precursor cells. These mice, a gift from Dr. Dan Littman at New York University, were made by replacing the gene for fractalkine receptor CX₃CR1 with the gene encoding GFP. $Cx3cr1^{GFP/GFP}$ mice exhibit a normal microglial response when challenged with sciatic nerve lesion³⁶, allowing us to use mice that are heterozygous for CX₃CR1 in some of our migration studies knowing a deficiency in microglial response will not be due to the genetic disruption of this receptor.

In light of the importance of this receptor in platelet activation, we asked whether P2Y₁₂ might play a similar role in microglia by contributing to their ability to respond to injury through their capacity to detect nucleotide release. The paucity of subtype-selective agonists or antagonists has hindered efforts to rigorously examine physiological roles for specific P2 receptor subtypes *in vivo*, and we

have therefore taken a genetic approach to address this question. By examining microglial behavior in a variety of *in vitro* and *in vivo* systems, we now show that $P2Y_{12}$ is, indeed, an essential component of the signalling pathway through which extracellular purines promote directed microglial movement following CNS injury. Moreover, our findings suggest that, as in platelets, signalling through $P2Y_{12}$ is most relevant during early stages of the microglial activation process.

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CHAPTER 2

Gi-COUPLED PURINERGIC RECEPTORS ON MICROGLIA

GI-COUPLED PURINERGIC RECEPTORS ON MICROGLIA

The G₁-coupled P2Y₁₂ receptor is expressed on brain microglia but not peripheral or infiltrating macrophages

We, and others, have previously shown that P2Y₁₂ transcripts are expressed by glial cells of the spinal cord and brain, specifically microglia^{1,2}. To determine whether receptor protein is located on the microglial cell surface, we generated an antibody that recognizes the C-terminus of mouse P2Y12 and used this polyclonal antiserum to stain sections from spinal cord and brain. Indeed, robust signals were clearly observed in a population of glial cells evenly distributed throughout both white and gray matter. Identification of these cells as microglia was confirmed by co-localization with two independent microglial markers, including enhanced green fluorescent protein (GFP) expressed from the CX₃CR1 fractalkine receptor promoter³ and the integrin CD11b (Fig. 1a, b). Moreover, P2Y₁₂ immunoreactivity was not observed in GFAP-expressing astrocytes (Fig. Higher magnification images showed that P2Y12 protein was localized 1a). predominantly to the cell surface, including the elaborate ramified processes emanating from the cell body (Fig. 1c). Thus P2Y₁₂ is poised to enable microglia to detect changes in extracellular nucleotide concentration as they survey their local environment. Importantly, P2Y₁₂-deficient mice were devoid of receptor immunoreactivity (Fig. 1a), but showed normal prevalence, distribution, and

morphology of "resting" microglia (Fig. 2), suggesting that the receptor is not required for proper development or CNS localization of these cells.

Few markers exist that can differentiate between microglia of the central nervous system and macrophages that reside in peripheral tissues or infiltrate the CNS after injury. Previous *in situ* hybridization studies have shown that P2Y₁₂ transcripts are observed in microglia but not spleenic macrophages¹. Here, we confirm this observation at the protein level by showing that peripheral macrophages are devoid of P2Y₁₂ immunoreactivity (**Fig. 1b**), demonstrating that this receptor is, indeed, an excellent molecular marker for differentiating between these two closely related phagocytic cell types.

P2Y₁₂ is down regulated upon microglial activation

Microglia are believed to mediate a number of physiological responses to neural injury that include both short and long term actions⁴. Expression of various cell surface markers and cellular phenotypes are known to change following CNS injury⁵ and we therefore asked whether P2Y₁₂ expression varies with the extent of microglial activation. To address this question, we first used an *ex vivo* tissue preparation to examine the time-course of P2Y₁₂ expression following neural injury. To induce neural injury and facilitate microscopic examination over an extended period, thick (400 µm) tissue slices were cut from hippocampi of *Cx3cr1*^{+/GFP} expressing neonatal (P4–7) mice. Over a 24 h time-course, we observed trauma-induced activation of microglia as typically characterized by

their transformation from a highly ramified to amoeboid morphology⁶. Interestingly, $P2Y_{12}$ expression showed a dramatic and continuous decrease such that by 24 h, expression was barely observable (**Fig. 3a**). This correlated with a decrement in the number of primary projections emanating from the cell body as microglia progressed towards a more amoeboid state (**Fig. 3b**). Indeed, when $P2Y_{12}$ expression was quantified on a single cell level, a linear correlation was revealed between expression level and the morphological state of the cell (**Fig. 2c-e**).

We next asked whether $P2Y_{12}$ expression changes as a function of microglial activation *in vivo*. To accomplish this, we examined $P2Y_{12}$ immunoreactivity following a bolus injection of lipopolysaccharide (LPS) into the striatum, a treatment that activates microglia within the injected area⁷ and which has been shown to decrease 2MeSADP-evoked calcium increases in cultured mouse microglia⁸. Four days after LPS injection, virtually all microglia within proximity of the injection site exhibited an amoeboid-like morphology, and $P2Y_{12}$ immunoreactivity was virtually undetectable in these cells (**Fig. 4a**). Reduced $P2Y_{12}$ expression was confirmed at the transcriptional level by *in situ* hybridization histochemistry and northern blotting, where a substantial decrease in messenger RNA levels was observed (**Fig. 4b, c**). This was further validated by quantitative RT-PCR performed with total RNA isolated from a region of brain containing the injection site, which showed a 60% reduction in P2Y₁₂ transcripts (data not shown).

Taken together, our data support a relationship between microglial activation and reduced P2Y₁₂ expression in two different experimental systems, and suggest that P2Y₁₂ is an excellent molecular marker for visualizing brain microglia in the ramified state. Interestingly, recent studies suggest that P2Y₁₂ desensitizes upon platelet activation⁹, a phenomenon that may represent a functional correlate of the decreased receptor expression that we observe following microglial activation. Together, these findings suggest possible roles for P2Y₁₂ receptors at early rather than late stages of the microglial response to injury.

G_i-coupled purinergic receptors

Like platelets, microglia express a plethora of metabotropic and ionotropic purinergic receptor subtypes such as P2Y₁, P2Y₂, P2Y₁₂, P2X₄, and P2X₇¹⁰, suggesting that nucleotides play numerous roles in modulating immune responses to a range of pathophysiological insults of the CNS. For example, pharmacological studies have implicated P2X receptors on microglia in the development of neuropathic pain¹¹ or the release of neuroprotective or proinflammatory agents, such as tumor necrosis factor- α and interleukin-1 $\beta^{12,13}$. When studying the physiological role of P2Y₁₂, it is very important to consider other receptors that are activated by extracellular nucleotides. The presence of multiple receptors with similar pharmacological profiles may make it harder to determine the contribution of the P2Y₁₂ receptor to a particular cellular phenotype. Therefore, we decided to examine the pharmacology of a P2Y₁₂-

related receptor and determine whether this receptor contributes to microglial physiology.

In 2001 when the sequence of P2Y₁₂ was discovered², it was the only G_i-coupled purinergic receptor known to exist and actually was guite distantly related to the other known P2Y receptors (P2Y₁, P2Y₄, and P2Y₆ which couples to phospholipase C pathways, and P2Y₁₁ which activates the synthesis of cAMP). However, in recent years there have been a handful of other purported Gicoupled P2Y receptors found to lie just adjacent to the P2Y₁₂ locus on chromosome 3, and thus two distinct classes of P2Y receptors were classified¹⁴. This second class of receptors includes P2Y₁₂, P2Y₁₃, GPR 87, P2Y₁₄ (the UDPglucose receptor), and GPR 171 located within 114 kb of each other. While little is known about the orphan receptors GPR 87 and GPR 171, it is thought that P2Y₁₄ is involved in UDP-glucose induced migration of premature hematopoietic stem cells^{15,16}. A homology tree and the locus map for this class of receptors are shown in Fig. 5. The close proximity of these genes and striking similarity in sequence homology suggests that these receptors are the product of a recent evolutionary gene duplication event. Examining differences in pharmacology and sequence of these related proteins may help elucidate mechanisms of ligand activation.

 $P2Y_{13}$ is located only 4.5 kb away from $P2Y_{12}$ on chromosome 3 and shares 48% sequence homology^{17,18}. $P2Y_{13}$ is considered to be G_i-coupled because

pertussis toxin blocks ADP inhibition of cAMP production in CHO cells¹⁷. P2Y₁₂ and P2Y₁₃ share very similar pharmacological profiles; ADP acts as an agonist, ATP a partial agonist, and AR-C69931MX an antagonist¹⁴. We further examined the pharmacology of these receptors by two-electrode voltage clamp electrophysiology in Xenopus laevis oocytes. P2Y12 or P2Y13 mRNA was injected into oocytes along with the inwardly rectifying potassium channels Kir3.1 and Kir3.4, which allow K^{+} ions to pass through the membrane when activated by the G-protein. As shown in Fig. 6a, P2Y₁₂ and P2Y₁₃ are equally sensitive to ADP. Interestingly, 2MeSADP activates P2Y₁₂ at 100 fold less concentration than ADP, whereas 2MeSADP only activates P2Y₁₃ roughly 10 fold less concentration than ADP. In addition, the shape of the 2MeSADP-induced P2Y₁₃ current is different than the 2MeSADP-induced P2Y₁₂ current in that P2Y₁₃ inactivation is slower, or 2MeSADP is harder to wash out from P2Y₁₃ expressing oocytes. ATP activates both receptors to a similar extent, and ATP is about 50% less potent than ADP at both receptors. Indeed, it has previously been reported that P2Y₁₃ in 1321N1 cells bound radioactively labelled 2MeSADP more tightly than P2Y12¹⁴. There are several other differences in the pharmacology of these two receptors as shown in **Fig. 6a**. For example, both ADP-ribose and AP4A are weak agonists to P2Y₁₂ but approximately 2-4 fold stronger agonists to P2Y₁₃.

Some GPCRs are known to form multimers in their active state¹⁹, so we asked whether there is a synergism of the $P2Y_{12}$ and $P2Y_{13}$ receptors, or if coexpression of the receptors changes the pharmacology. Importantly, co-

expression of both P2Y receptors in ooctyes resulted in an intermediate profile where, for example, 2MeSADP-activated current is greater than the current in P2Y₁₃ expressing oocytes but less than that of P2Y₁₂ expressing oocytes. In addition, the current inactivation during wash out of 2MeSADP in ooctyes expressing both receptors also shows an intermediate profile. Also, the ADP-induced current is not enhanced when both receptors are expressed together, also suggesting that there is minimal interaction (**Fig. 6b**).

P2Y₁₂ is known to be sensitive to inactivation by compounds that modify cysteine residues, such as the active metabolite of the anti-thrombotic drug clopidogrel²⁰. Evidence suggests that P2Y₁₃ is not able to bind the active metabolite of clopidogrel¹⁴, however the sensitivity of this receptor to thiol reagents is unknown. We applied PCMBS to P2Y₁₂ and P2Y₁₃ expressing oocytes (**Fig. 6c**), a mercury compound thought to modify cysteine residues in a manner similar to clopidogrel. For these experiments, the mu-opioid receptor (muOR) was co-injected to show that inactivation of P2Y receptors does not result in the inability of the oocyte to be activated by another G_I-coupled receptor. P2Y₁₂ expressing oocytes showed complete inhibition of ADP-induced current by 1 mM PCMBS, whereas P2Y₁₃ expressing oocytes showed only 30% ADP-induced current inhibition. Since these two receptors share a very similar nucleotide sequence yet display such different sensitivities to thiol reagents, they provide a unique setting to pinpoint the required structural components needed by clopidogrel to inactivate the P2Y₁₂ receptor in platelets.

Although its expression profile has not been well defined, $P2Y_{13}$ mRNA transcript was found to be absent from platelets²¹ and thus does not contribute to nucleotide-induced thrombosis. However, $P2Y_{13}$ mRNA was reported to be expressed in brain²¹ and therefore we were interested to see whether it is expressed in microglia. To address this question, we isolated primary microglia from wild-type neonatal mouse brain, isolated mRNA, and performed PCR with $P2Y_{13}$ specific primers (**Fig. 7a**). Indeed, $P2Y_{13}$ transcripts were apparent in these cells. *In situ* hybridization of adult brain with a $P2Y_{13}$ specific probe (**Fig. 7b**) confirmed low level expression by microglia and showed that no other CNS cell type expressed $P2Y_{13}$ transcripts. Interestingly, it seems that microglia do not express $P2Y_{13}$ protein *in vivo*, since we failed to detect receptor protein using a $P2Y_{13}$ specific antibody (**Fig. 7c**). These results suggest that $P2Y_{13}$ most likely does not contribute significantly to microglial biology *in vivo*.

Due to the close proximity of the P2Y₁₂ and P2Y₁₃ receptors, we were interested to see if perhaps the P2Y₁₃ locus was disturbed in our P2Y₁₂–knockout mice. In addition, we were curious to know whether P2Y₁₃ transcripts showed the same down-regulation after microglial activation as P2Y₁₂. Therefore, we took brains injected with either PBS (vehicle) or LPS to induce inflammation (see above) and performed a northern blot with a P2Y₁₃ specific probe. Surprisingly, we found that P2Y₁₃ transcripts were down-regulated after LPS injection in a similar manner to P2Y₁₂, and knockout animals expressed less transcripts as well as a second, higher molecular weight species. Therefore, we surmise that the $P2Y_{13}$ locus, although protein is never transcribed *in vivo*, is under the same transcriptional regulatory elements as $P2Y_{12}$. In addition, it is possible that targeted deletion of $P2Y_{12}$ in the knockout animals resulted in an alteration of a $P2Y_{13}$ regulatory element. In any case, it is unlikely that this receptor plays a major role in the biology of these cells due to the fact that receptor protein is apparently never made.

Discussion

A major problem in the field of microglial biology is the inability to detect whether a cell located in close proximity to an injury site is a local CNS microglia or a macrophage that has infiltrated from the periphery. The activated microglia and peripheral macrophage display very similar cellular markers, making it nearly impossible to distinguish between them. Therefore, it is difficult to ascertain the physiological functions of microglia at various stages of activation. P2Y₁₂, however, displays a unique expression profile in that it is expressed only by CNS resident microglia and absent from macrophages. In addition, it becomes downregulated upon microglial activation, making this a very good marker to visualize microglia in the highly ramified, resting state. Also, it may be the best known marker to distinguish activated from non-activated microglia, since most other markers are not completely turned on or off but rather are only up- or down-

regulated. The physiological significance of $P2Y_{12}$ receptor down-regulation is unknown, however it is likely that this receptor functions in the microglial resting state or at a very early time in the activation process.

The discovery of new GPCRs that are closely related to $P2Y_{12}$ and that lie in close proximity on chromosome 3 has elucidate a new class of G_i-coupled purinergic receptors. $P2Y_{13}$ is the most closely related and shares many pharmacological properties. Two important differences in the pharmacological profiles of these two receptors is 1) $P2Y_{13}$ is not activated as strongly by 2MeSADP as $P2Y_{12}$ and 2) $P2Y_{13}$ is not as sensitive to thiol reagents as $P2Y_{12}$, such as the commonly prescribed anti-thrombotic drug clopidogrel. Uncovering the structural requirements for clopidogrel's actions on preventing thrombosis would be very interesting, and these two receptors provide a unique capacity in which to study this mechanism.

Methods

peptide. This antibody showed immunoreactivity for mouse, rat, and human P2Y₁₂ as assessed by immunofluorescence and western blotting of transfected HEK293T cells expressing each of these receptors. Brain and spleen tissue from adult (3 months) mice was fixed with 4% paraformaldehyde and sectioned (30 µm thick) on a freezing cryotome, blocked for 30 min in 10% normal goat serum/0.1% triton in PBS and stained with P2Y₁₂, CD11b (eBiosciences), or GFAP (Pharmingen) primary antibodies overnight at 4° C, washed and visualized with goat anti-rabbit Alexa 546 or 594, goat anti-rat Alexa 594, or goat anti-mouse Alexa 633 secondary antibodies (Molecular Probes), respectively. Images were taken on a Zeiss Pascal or LSM 510 Meta confocal microscope using 40x oil- or 20x multi-immersion objective, or a Nikon wide-field fluorescent microscope using a 20x lens.

Ex vivo hippocampal slice preparation and imaging. Hippocampal tissue slices (400 µm thick) were prepared from P4–7 mice and stained as described previously^{22,23} using anti-P2Y₁₂ primary antibody (1:500), and goat anti-rabbit Alexa 594 (Molecular Probes) secondary antibody (1:1000). Slices were stained with anti-P2Y₁₂ antisera at 0 h, 8 h, or 24 h after cutting, and imaged on a Zeiss LSM 510 Meta, Leica TCS NT, or Leica SP2 AOBS confocal microscope with a 20x multi immersion or air lens. (*n* = 3 slices per time-point, ~ 40 cells per slice at the CA3 region.) 15 *z*-steps spaced 2 µm apart were collected per image (30 µm total depth) and a maximum projection created. P2Y₁₂ images were overlayed with GFP images to analyze expression level. To determine
expression level, cell regions were drawn by outlining the cell in the GFP channel, overlaying that region on the $P2Y_{12}$ image, and calculating the mean gray level. Only cells with the entirety of their projections and cell body residing within the slice were quantified. Cellular morphology was determined by counting the number of ramifications projecting immediately from the cell body (primary projections.)

LPS injections and analysis of P2Y₁₂ expression. Mice were anesthetized with ketamine/zylazine and placed in a stereotaxic apparatus. A small incision was made in the scalp and a small hole drilled through the skull over the area of injection. The needle was positioned at 1 mm anterior, 1.5 mm lateral, and 3 mm ventral to bregma and 1 µl of 5 mg/ml LPS (Sigma) or PBS (control) was bilaterally injected into the striatum. The wound was closed and mice allowed to recover for 4 d prior to tissue collection. For immuno- and in situ hybridization histochemistry, PFA-fixed (20-30 µm thick) brain sections were incubated with anti-P2Y₁₂ antibody, or a digoxygenin-labeled cRNA probe containing the entire mouse P2Y₁₂ coding region (Sacl-Nsil fragment). For in situ hybridization experiments, GFP expression was visualized with an anti-GFP antibody (Molecular Probes). For northern blot and RT-PCR analyses, mice were perfused with PBS and RNA was extracted from a coronal brain slab that included a region 2 mm posterior and anterior to the injection site. Poly A+ RNA was purified with a micro-FastTrack kit (InVitrogen) and analyzed as previously described²⁴ using a ³²P-labeled BbvCI cDNA fragment containing the P2Y₁₂

coding region. Real-time RT-PCR was performed on first-strand cDNA generated from DNAse-treated total RNA using the following primer pairs for P2Y₁₂: 5'-CATTGCTGTACACCGTCCTG-3' and 5'-GGCTCCCAGTTTA GCATCAC-3'. Quantification of amplified products was carried out as previously described²⁴.

P2Y₁₃ expression analysis. Quantitative RT-PCR, *in situ* hybridization, northern blot and immunohistology were performed as described above. The following primers pairs were used for PCR analysis:

P2Y ₁₂ :	5'-CCTCAGCCAATACCACCTTCTCCCC-3'
	5'-CGCTTGGTTCGCCACCTTCTTGTCCCTT-3'
P2Y ₁₃ :	5'-GGGACACTCGGATGACACAGCTGC-3'
	5'-GCCAGAAAGAGAGTTGCTTCTTTAGCAATAAACAGC-3'
CX₃CR1:	5'-TTCACGTTCGGTCTGGTGGG-3'

5'-GGTTCCTAGTGGAGCTAGGG-3'

Anti-P2Y₁₃ receptor polyclonal antibody was generated by immunizing rabbit with a synthetic peptide corresponding to the mouse P2Y₁₃ C-terminus (NH₂-Cys-Thr-Ala-Gly-Ser-Ser-Glu-Asp-His-His-Ser-Ser-Gln-Thr-Asp-Asn-Ile-Thr-Leu-Ala-OH Anaspec, Inc). Antibody was affinity purified using a Sulfolink coupling gel (Pierce) to immobilize the antigenic peptide. **Pharmacology of P2Y receptors by two-electrode voltage clamp electrophysiology.** *Xenopus laevis* oocytes were injected with mP2Y₁₂ (0.1 ng), mP2Y₁₃ (0.1 ng), Kir 3.1 (2 ng), Kir 3.4 (2 ng), or muOR (0.1 ng) mRNA as described². Kir 3.1 and 3.4 were included in all experiments to allow for detection of GPCR-induced flow of K⁺ ions across the membrane. The muOR was added as a positive control for inhibition assays. 0.5 ng P2Y₁₂ and 0.5 ng P2Y₁₃ was used for experiments in which the receptors were coexpressed Three to seven days after injection, electrophysiology recordings were performed using Geneclamp 500 amplifier (Axon Instruments). Cells were clamped –80 mV membrane potential while the recording chamber was perfused with solution containing KCI (70 mM), NaCI (20 mM), MgCl₂ (3 mM), HEPES (5 mM), ph 7.4, at room temperature. Ligands (ATP, ADP, ADP-ribose, AP4A, 2MeSADP, damgo, PCMBS, Sigma) were added to the recording solution as indicated.

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Figure Legends

Figure 1. P2Y₁₂ immunoreactivity is localized to the surface of CNS microglia. (a) P2Y₁₂ antibody (red) was used to visualize receptors in the cortex of adult wild-type (*P2ry12^{+/+}*) or P2Y₁₂-deficient (*P2ry12^{-/-}*) mice. P2Y₁₂ immunoreactivity co-localized with GFP fluorescence (green), which was expressed under control of the CX₃CR1 promotor to specifically label microglia in these mice. P2Y₁₂ staining was excluded from astrocytes, which were identified by GFAP (blue) expression. No P2Y₁₂ immunoreactivity was observed in sections from *P2ry12^{-/-}* mice. Scale bar = 20 µm. (b) Staining of spleen and brain sections with anti-P2Y₁₂ (red) and anti-CD11b (green) antisera demonstrates absence of P2Y₁₂ expression in peripheral macrophages. For spleen, scale bar = 25 µm; for brain scale bar = 20 µm. (c) Higher magnification images of brain microglia reveal P2Y₁₂ (red) localization to the plasma membrane, whereas soluble GFP (green) is spread throughout the cell body and cytoplasm. Scale bar = 5 µm.

Figure 2. Microglia from P2Y₁₂-deficient mice have normal morphology and prevalence within the CNS. (a) Microglia were quantified by counting GFP-positive cell bodies in three regions of the brain (striatum, cortex, and hippocampus), retina, and spinal cord in cryostat sections from wild-type (gray) or P2Y₁₂-deficient (black) Cx3cr1 ^{+/GFP} mice (n = 2 mice/genotype; 10-30 sections per animal). The number of microglia per 100 μ m² area is shown (mean ± s.e.m.). (b) The area occupied by processes originating from individual microglia

was quantified from whole mount retinas (n = 1 mouse/genotype; 12 sections per animal; mean \pm s.e.m.).

Figure 3. Loss of P2Y₁₂ expression accompanies microglial transform from highly ramified to amoeboid state. (a) Average P2Y₁₂ immunoreactivity per microglial cell was measured in hippocampal slices from P2ry12+/+Cx3cr1+/GFP neonatal mice at 0 h, 8 h, and 24 h after preparation. (b) Average number of primary projections per microglial cell was determined at indicated times following hippocampal slice preparation (n = 50-80 cells per time point; values represent mean \pm s.e.m.). (c) P2Y₁₂ expression is linearly correlated with morphological state (number of primary projections) of microglia at 8 h time-point $(R^2 = 0.93)$. Data derive from 3 separate experiments and were normalized to background fluorescence of the slice. (d) Analysis of GFP fluorescence (green) and immunoreactivity (red) in P2Y₁₂ hippocampal slices from $P2ry12^{+/+}Cx3cr1^{+/GFP}$ neonatal mice shows complete loss of $P2Y_{12}$ expression by 24 h. Scale bar = 50 μ m. White boxes highlight representative microglia, morphology of which is depicted in panels (e-h). Scale bar = $15 \mu m$.

Figure 4. Activated microglia lack $P2Y_{12}$ expression *in vivo*. (a) Striatal sections from vehicle (PBS)- or lipopolysaccharide (LPS, 5 µg)-injected $P2ry12^{+/+}Cx3cr1^{+/GFP}$ mice were examined for $P2Y_{12}$ immunoreactivity (red). Note the absence of $P2Y_{12}$ staining 4 days after LPS injection. Similar results were obtained in 5 independent experiments. Scale bar = 20 µm. (b) *In situ*

hybridization with P2Y₁₂ antisense probe showed a substantial loss of P2Y₁₂ mRNA expression following LPS injection. Anti-GFP antibody was used to visualize microglia (green). Scale bar = 20 μ m. (c) Northern blot analysis showed a decrease in P2Y₁₂ mRNA expression in tissue taken from a region surrounding the LPS injection site relative to PBS-injected control. Probe specificity was verified by lack of signal from *P2ry12^{-/-}* tissue. Cyclophilin transcripts were analyzed to verify equivalent sample loading.

Figure 5. Relationship of P2Y₁₂ to other purinergic receptors. (**a**) A phylogenetic tree of mammalian P2Y receptors reveals two distinct classes. Class 1 includes G_q -coupled receptors P2Y₁, P2Y₂, P2Y₄, P2Y₆, and G_s -coupled receptor P2Y₁₁. Class 2 includes the purported G_i -coupled receptors P2Y₁₂, P2Y₁₃, P2Y₁₄, GPR 87, and GPR 171. (**b**) A diagram of *Mus musculus* chromosome 3q24-25 shows the close proximity of P2Y₁₂ and the newly discovered G_i -coupled receptors.

Figure 6. Pharmacological profiles of the P2Y₁₂ and P2Y₁₃ receptors. (a) *Xenopus laevis* oocytes expressing the P2Y₁₂ or P2Y₁₃ receptor (with Kir 3.1 and Kir 3.4 to allow for GPCR-activated ion flow across the membrane) were exposed to Ringer's solution containing the nucleotides indicated, and the resulting change in current was measured by two-electrode voltage clamp electrophysiology. Note the differences in current amplitude after 2MeSADP, ADP-ribose, and AP4A. Scale bar represents 100 s, 200 nA. Oocytes expressings Kirs without GPCRs showed no nucleotide-evoked currents (data

not shown). All oocyte experiments were performed in the presence of Kir 3.1 and Kir 3.4. (b) Co-expression of P2Y₁₂ and P2Y₁₃ receptors results in an intermediate pharmacological profile, suggesting that there is little or no physiological interaction between the receptors. Scale bar represents 100 s, 100 nA. (c) $P2Y_{12}$ is more sensitive to inhibition by the thiol reagent PCMBS than P2Y₁₃, supporting the observation that the latter receptor is non-responsive to the anti-thrombotic, cysteine modifying drug clopidogrel. P2Y₁₂ expressing oocytes showed an inhibition of ADP-induced current after a brief application of PCMBS. However, PCMBS did not inhibit damgo-induced activation of the muOR receptor, another G_i-coupled receptor that was co-injected as a positive control. Interestingly, P2Y₁₃ expressing oocytes showed a greatly reduced inhibition of ADP-induced current by PCMBS whereas the muOR activation remained unchanged in these cells. As expected form pervious results, so expression of P2Y₁₂ and P2Y₁₃ receptors showed an intermediate phenotype, and the absence of either puringeric receptor resulted in a complete loss of ADP-induced current (negative control). Scale bars represent 50 s or 100 s, 50 nA or 200 nA as indicated.

Figure 7. Microglial expression of P2Y₁₃. (a) RT-PCR products were produced with primer pairs designed to specifically amplify P2Y₁₂ or P2Y₁₃ coding regions (or CX₃CR1 as a positive control) using total RNA from microglia cultured from wild-type (+/+) or P2Y₁₂-deficient mice. Amplification of RNA-derived reverse transcripts was confirmed by lack of signal in the absence of reverse

transcriptase (RT). We also performed quantitative RT-PCR to ascertain whether P2Y₁₃ mRNA expression is markedly altered in P2Y₁₂-deficient mice, reflecting either a compensatory increase or a decrease due to the close proximity (~5 kilobases) of the P2Y₁₃ gene to the targeted locus. We found a slight reduction (~ 2-fold) in P2Y₁₃ signal amplified from brain tissue of P2Y₁₂deficient mice compared to wild-type littermates. (b) In situ hybridization of brain sections from P2ry12^{+/+}Cx3cr1^{+/GFP} mice using a P2Y₁₃-specific anti-sense probe revealed weak, but detectable signals in GFP-positive microglia. (c) Transiently transfected HEK293T expressing cells mouse P2Y₁₃ cDNA show immunoreactivity with anti-P2Y₁₃ antibody, whereas vector (pcDNA3)-transfected controls do not. Brain sections from $P2ry12^{+/+}Cx3cr1^{+/GFP}$ mice lacked detectable immunoreactivity when stained with anti-P2Y₁₃ antibody. GFP fluorescence shows location of microglia. Scale bars = 50 µm. (d) Northern blot analysis of brains from wild-type or P2Y₁₂-knockout mice injected with LPS to induce microglial activation or PBS (vehicle) using a P2Y₁₃-specific probe. Upon microglial inflammation, P2Y₁₃ transcript level decreases in a manner similar to P2Y₁₂. Examination of P2Y₁₂-knockout animals reveals that P2Y₁₃ transcript levels are diminished even in the healthy, non-activated state, and a second, higher molecular weight species is apparent. A cyclophilin-specific probe was used as a loading control.

















a <u>Phylogenetic tree of mammalian P2Y receptors</u>



b

Chromosome 3q24-25









Brain



d

P2Y13

cyclophilin

WT PBS WT LPS PBS LPS

CHAPTER 3

NUCLEOTIDE-EVOKED CHEMOTAXIS OF MICROGLIA

REQUIRES P2Y₁₂ RECEPTORS

NUCLEOTIDE-EVOKED CHEMOTAXIS OF MICROGLIA REQUIRES P2Y₁₂ RECEPTORS

In light of the proposed roles for purinergic receptors in regulating microglial motility, we asked whether P2Y12-deficient mice show deficits in cellular responses to exogenously applied nucleotides. Based on our observation that the P2Y₁₂ receptor is preferentially expressed in the "resting" state, we used experimental paradigms that allow us to examine cellular responses immediately following nucleotide stimulation. First, we used an in vitro culture system to address the potential involvement of P2Y₁₂ in ATP/ADP-mediated actin-based lamellipodial extension. When microglia are isolated from neonatal rodent brain and subjected to serum starvation for several hours, they revert to a presumptive "resting" or "inactivated" state based on morphological and immunological criteria¹. Under these conditions, the vast majority ($\sim 87\%$) of microglia from wild-type mice displayed robust membrane ruffling upon application of ADP or ATP (50 µM), consistent with previous observations². In contrast, cells from P2rv12^{-/-} mice showed no response to ADP or ATP above that elicited by UTP (50 μ M) or vehicle alone (~ 7%). Application of macrophage colony stimulating factor (M-CSF, 200 ng/ml) produced equivalent responses in wild-type and $P2Y_{12}$ -deficient microglia (~ 74%), demonstrating that the absence of $P2Y_{12}$ does not generally disrupt signalling mechanisms required for actin polymerization (Fig. 1a, b). Interestingly, pre-incubation with LPS (0.1 mg/ml)

significantly diminished the nucleotide-evoked membrane ruffling of wild-type microglia *in vitro* (Fig. 2).

To examine directed motility in response to a localized nucleotide source, we placed purified microglia in a Dunn chemotaxis chamber³ and observed their behavior in a gradient of ADP or ATP (0 to 50 μ M) using phase contrast time-lapse microscopy. Microglia from wild-type mice showed a clear and robust polarization or chemotaxis toward the nucleotide source within the 30 min observation time, whereas P2Y₁₂-deficient microglia showed no evidence of membrane ruffling, polarization, or directed movement (**Fig. 1c, d**).

We next asked whether similar phenotypes would be observed when microglia are examined in a more native neuronal environment. To do this, we used the hippocampal slice system described above to monitor microglial chemotactic behavior by time-lapse microscopy following bath application of nucleotides (1 mM ADP). As the nucleotide diffuses into the slice, a gradient is formed which induces microglial chemotaxis towards the periphery of the slice. In brain slices from neonatal (P4–7) wild-type mice, microglia demonstrated robust process extension and cellular migration towards the periphery of the slice that initiated within minutes of nucleotide exposure and was sustained for over 6 h. In contrast, slices prepared from neonatal P2Y₁₂-deficient mice show no directed polarization, process extension, or migration in response to the nucleotide stimulus, consistent with our *in vitro* chemotaxis data (**Fig. 3**). In fact, microglia

from mutant mice retract their processes and travel towards the injured neuronal cell body layer of the CA3, consistent with the actions of microglia in wild-type slices in the absence of nucleotides⁴.

Finally, we asked whether similar deficits could be observed in the intact living brain of adult mice. We therefore placed a microelectrode containing ATP (20 mM) into the neocortex of $P2ry12^{+/+}Cx3cr1^{+/GFP}$ or $P2ry12^{-/-}Cx3cr1^{+/GFP}$ adult mice and followed changes in microglial morphology using two-photon time-lapse microscopy. As previously described⁵, microglia in adult wild-type animals showed very active extension of cellular processes toward the nucleotide source. In striking contrast, $P2Y_{12}$ -deficient mice showed greatly reduced responses during an equivalent 40 min post-injection period, demonstrating that the $P2Y_{12}$ receptor is absolutely required for ATP-mediated process extension of microglia *in vivo* (**Fig. 4**).

Discussion

Taken together, our studies show that the $P2Y_{12}$ receptor is essential for the ability of microglia to respond to extracellular nucleotides by process extension or whole cell movement. Our use of several experimental paradigms revealed that this is true whether microglia are exposed to extracellular nucleotides in the context of a homogeneous culture, or in their native neuronal environment.

Moreover, our data suggest that $P2Y_{12}$ is important in this process at all stages of postnatal development on through adulthood.

Methods

Primary microglia cell culture, membrane ruffling, and chemotaxis assays. Primary microglia were isolated from P1–3 mice using a modified version of the mixed glial culture technique as previously described⁶. Briefly, brains were dissected, homogenized by passing through an 18 G needle, and cultured in DME-H21 with 10% heat-inactivated FBS and penicillin/streptomycin. After 12–14 d, cultures were gently shaken by hand for 15 min and microglia collected as floaters, resulting in > 99% purity as assessed by GFP expression. Isolated microglia were spotted onto glass chamber slides (Fisher) for membrane ruffling assays or fibronectin-coated coverslips (Becton Dickinson) for chemotaxis assays, washed, and incubated for 4 h in serum-free DME-H21.

Membrane ruffling assays were performed as described (Honda, et. al., 2001) by replacing DME-H21 with fresh DME-H21 containing no stimulus (negative control) or ADP, ATP, or UTP (50 μ M), 2-MeSADP (0.5 μ M, Sigma), or M-CSF (200 ng/ml, BD Biosciences). Microglia were incubated for 5 min at 37° C, fixed with 4% PFA, and stained with rhodamine- or FITC-conjugated phalloidin (Molecular Probes). To determine the effect of LPS pre-incubation on

nucleotide-evoked membrane ruffling, primary microglia were prepared as described and incubated in serum-free media with or without LPS (0.1 mg/ml) for 6 h immediately before a brief (5 min) ATP application (50 μ M). Cells were then fixed with 4% PFA and stained with FITC-conjugated phalloidin.

Chemotaxis assays were performed using the Dunn chamber³ to form a nucleotide gradient from 0 to 50 μ M ADP or ATP. Distance and direction of movement by the cell's leading edge was monitored over a 30 min period by phase contrast time-lapse microscopy; the average distance migrated in the absence of a stimulus (< 0.8 μ m) was subtracted to obtain final values. Images were processed and analyzed using the National Institutes of Health ImageJ software.

Ex vivo hippocampal slice preparation and imaging. To examine the microglial response to applied nucleotides, acute hippocampal slices were prepared as previously described and incubated in imaging media alone or with 1 mM ADP. Microglia were visualized by GFP fluorescence or by staining with fluorescently tagged isolectin-IB4, and imaged as described⁴. For each image in the time-series, 15 *z*-steps spaced 2 μ m apart were collected per image (30 μ m total depth) and a maximum projection created for each. Images were taken at 5 min intervals for up to 8 h on Leica TCS NT, or Leica SP2 AOBS confocal microscope with a 20x lens. Image processing and analysis was performed using NIH Image J software.

In vivo imaging of microglia. GFP expressing microglia were imaged by twophoton time-lapse microscopy as described^{5,7}. Briefly, ATP (20 mM) was diluted in artificial CSF for intracranial injections. A *z*-stack (~ 100 μ m depth, 2 μ m *z*steps) was acquired every 4 min for 40 min. A maximum projection of the 15 *z*steps centered on the laser ablation or injection site was constructed for each time point (30 μ m total depth). Microglial response to ATP injection was determined as described⁵. Changes in GFP fluorescence were monitored as processes entered a circular zone of 70 μ m in diameter centered over the injection needle and normalized to an outer region 136 μ m in diameter.

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Figure Legends

Figure 1. Nucleotide-evoked membrane ruffling and chemotaxis are abolished in cultured microglia lacking P2Y₁₂ receptors. (a) Primary microglia from wild-type $(P2ry12^{+/+})$ or P2Y₁₂-deficient $(P2ry12^{-/-})$ mouse brains were exposed to vehicle (DMEH21), ADP (50 µM) or macrophage-colony stimulating factor (M-CSF, 200 na/ml). Membrane ruffling was visualized within 5 min by staining with rhodamine-phalloidin (red) to detect actin polymerization (white arrowheads). Scale bar = $20 \, \mu m$. (b) Fraction of microalia undergoing membrane ruffling in the presence of vehicle (DMEH21), UTP (50 µM), ADP (50 µM), ATP (50 µM), 2MeSADP (5 μ M), and M-CSF (200 ng/ml). Asterisks indicate P < 0.001; $n \ge 9$ trials per genotype for each experimental condition, $n \ge 150$ cells per trial; Student's *t*-test. Values represent mean \pm s.e.m. (c) Microglia from wild-type and P2Y₁₂ mutant mice were placed in a gradient of ATP (50 to 0 µM) using a Dunn chemotaxis chamber and visualized by phase contrast time-lapse microscopy. Microglia from wild-type mice showed robust chemotaxis toward the nucleotide source (white arrowheads indicate leading edge of migrating cell at 5 and 30 min time points), whereas P2Y12-deficient microglia showed a dramatically decreased response. Scale bar = 20 µm. (d) Scatter plot indicates distance (µm) travelled by the leading edge of wild-type (gray) or mutant (black) microglia in a gradient of ADP or ATP after 30 min. Values are plotted in the x, y directions relative to location of the nucleotide source.

Figure 2. Pre-treatment with LPS diminishes ATP-induced microglial ruffling. Microglia were cultured from brains of wild-type mice and exposed to vehicle (media) or LPS (0.1 mg/ml) for 6 h in serum-free media followed by a 4 h starvation period. Cells were subsequently exposed to ATP (50 μ M) for 5 min and membrane ruffling assessed by staining with FITC-conjugated phalloidin (see Methods). Representative images (a) and quantification (b) illustrates significant decrease in nucleotide responsiveness following LPS exposure. Scale bar = 20 μ m. (*n* = 4 trials per experimental condition, 200-400 cells per trial, **P* < 0.01)

Figure 3. P2Y₁₂-deficient microglia do not respond to exogenous nucleotides in brain slices. (a) Microglia in hippocampal slices from neonatal (P4–7) GFPexpressing mice were examined by time-lapse confocal imaging. $P2ry12^{*/*}$ microglia displayed robust process extension (red arrowheads) and whole-cell locomotion (yellow arrowheads) towards the periphery of the slice (right side of each frame) at various time points following addition of ADP (1 mM) to the bath. Slices from $P2ry12^{-/-}$ mice showed no such response over 6 h; green arrowheads show microglial locomotion away from the nucleotide source and towards the CA3 pyramidal cell body layer in $P2ry12^{-/-}$ slice. (b) Microglial branch length (µm) in wild-type and P2Y₁₂-deficient slices was measured at each time-point indicated in the presence of 1 mM ADP (n = 5 branches/cell). Positive length change indicates extension, whereas negative length change indicates retraction. Note that all wild-type branches extend, whereas all P2Y₁₂-deficient

branches retract within the time indicated. (c) Microglial locomotion in wild-type and P2Y₁₂-deficient slices in the presence (with ADP) or absence of nucleotides. Distance travelled (μ m) indicates net cell body displacement towards (positive values) or away from (negative values) the nucleotide source. In the absence of exogenous nucleotides, wild-type or P2Y₁₂-deficient microglia migrate toward the pyramidal region, presumably in response to neuronal injury within this cell body layer. Twenty cells residing in the *stratum pyramidale* and *oriens* of the CA3 area of the hippocampus were examined over a 30 min period.

Figure 4. Loss of P2Y₁₂ receptors abrogates response of microglia to ATP injection in the living brain. (a) Microglia from wild-type ($P2ry12^{+/+}$) or P2Y₁₂-deficient ($P2ry12^{-/-}$) GFP-expressing mice were visualized *in vivo* using two-photon time-lapse microscopy. Exuberant process extension towards a point source of ATP (20 mM from injection needle, red) was observed in the cortex of wild-type, but not mutant mice. Scale bar = 20 µm. (b) Quantification of ATP-evoked process extension in wild-type (grey) and P2Y₁₂-deficient (black) brains over a 40 min time course (n = 4 mice per genotype). *P < 0.05, **P < 0.01, ***P < 0.001; Student's *t*-test. Values represent mean ± s.e.m.







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CHAPTER 4

$\mathsf{P2Y}_{12}$ MEDIATES THE EARLY STAGES OF MICROGLIAL

MORPHOLOGICAL RESPONSE TO TISSUE INJURY

P2Y₁₂ MEDIATES THE EARLY STAGES OF MICROGLIAL MORPHOLOGICAL RESPONSE TO TISSUE INJURY

Role of P2Y₁₂ in microglial response to cortical laser-ablation

Tissue damage and loss of cellular integrity can lead to the release of nucleotides and other intracellular factors¹, one or more of which may serve to activate nearby microalia. In light of the robust phenotype that we have observed in response to exogenous nucleotides, we asked whether P2Y₁₂ is also required for the ability of microglia to rapidly detect and respond to tissue injury^{2,3}. We therefore utilized the in vivo imaging technique previously described to visualize the response of GFP expressing microglia to localized phototoxicity induced by the two-photon laser. Similar to what we observed with ATP injection, microglia from wild-type mice showed immediate and robust responses characterized by the extension of branches toward the site of damage, whereas P2ry12-- mice showed dramatically reduced chemotactic responses over an equivalent 40 min period. Interestingly, when microglia from mutant mice were examined at 2 h post-injury, the degree of process extension into the damaged area approached that observed with wild-type controls at the 32 min time-point (Fig. 1a, b), demonstrating that lack of P2Y₁₂ receptors significantly delays, but does not abolish the ability of microglia to respond to local tissue damage.

Recent imaging studies have shown that even in the absence of an injury stimulus, microglia exhibit appreciable, but non-directed process extension, possibly representing homeostatic surveillance of their environment^{2,3}. It has been proposed that this response is mediated by a purinergic receptor because it can be abrogated by apyrase or a high concentration of extracellular nucleotides². To ask whether this baseline motility requires functional P2Y₁₂ receptors, we analyzed absolute length change of randomly selected processes from wild-type or P2Y₁₂-deficient microglia. No significant differences were observed between genotypes (**Fig. 1c**), suggesting that P2Y₁₂ is involved primarily in mediating responses to cellular injury.

Physiological role of P2Y₁₂ in microglial response to injury in vivo

Here we have shown that microglia from P2Y₁₂-knockout mice do not chemotax or project process extensions towards sites of ATP *in vitro*, *ex vivo*, or *in vivo*, and therefore these mice are a valuable tool to assess the contribution of nucleotides to microglial activation after injury. We have also shown that microglia from P2Y₁₂-deficient mice are still able, albeit slowly, to morphologically respond to laser-induced damage in the living brain. However, the contribution of nucleotides to the pathological events that occur following other CNS diseases or injuries has not been characterized. To this aim, we performed various animal models that induce microglial activation in P2Y₁₂-deficient mice to determine the physiological significance of this signaling pathway on severity of disease.
When the sciatic nerve of the mouse is lesioned, the primary afferent terminals that project into the spinal cord will undergo degenerative atrophy⁴, and microglia in lamina I of the dorsal horn will become activated⁵. As previously mentioned, P2Y₁₂ is expressed in platelets where it functions in coagulation after vascular injury, and as expected P2Y₁₂-deficient mice display a slight bleeding disorder. This is important to keep in mind while performing animal models because increased bleeding by knockout mice may exacerbate the injury. In this regard, the sciatic nerve injury is ideal in that it allows for the surgery to be performed on peripheral tissue while the affects of the injury can be observed centrally. This surgery can be adapted to induce mechanical allodynia, a model of human peripheral neuropathies such as sciatica, an recent evidence suggests that microglia play a central role in this pain process⁵. In this spared nerve model, the sural and perineal branches of the sciatic nerve are cut just below the knee at the point of branching, while the tibial branch is spared. Microglial activation occurs following both the whole-sciatic nerve and spared nerve model, however the former provides a more robust response.

To determine microglial response to injury, we performed the whole sciatic nerve lesion model on wild-type and $P2Y_{12}$ -deficient mice. Since pervious results suggested that $P2Y_{12}$ plays a role at early stages of the microglial response to injury, we decided to look at early time-points of activation. At 1 d post-surgery, microglia began to accumulate around the affected site, and gradually continued to increase over the next few days. At all time-points acquired, there was no

significant difference in the increase of microglial cell density on the ipsilateral side between wild-type and P2Y₁₂-deficient animals (**Fig. 2**). To determine whether P2Y₁₂-deficient mice display altered mechanical allodynia, we performed the spared nerve model and used a behavioral test^{4,5} to look for changes in pain hypersensitivity. At 24 h to 2 weeks post-surgery, there was no significant difference in mechanical allodynia observed between wild-type and P2Y₁₂-deficient animals (data not shown), suggesting that this receptor does not play a major role in pain.

We have also looked at $P2Y_{12}$ -deficient animals in other models such as stroke⁶, glioma⁷, stab wound⁸, herpes virus infection in the retina⁹, and experimental autoimmune encephalomyelitis¹⁰ (an animal model of multiple sclerosis.) In all of the models, we were unable to determine a physiological consequence of receptor loss. However, these animal models offer much lower time-resolution than *in vivo* imaging, so it is possible that a difference in microglial migration towards an injury site simply went unnoticed.

Discussion

Although other molecules such as fractalkine, macrophage colony stimulating factor (m-CSF) or cannabinoids have been implicated in microglial migration¹¹⁻¹³, the specific molecules that mediate morphological responses to injury *in vivo*

remain elusive. Recent studies have suggested that ATP is released at a CNS injury site¹⁴ and may be involved in microglial reponse². Since P2Y₁₂-deficient mice show no microglial morphological response to applied ATP in any of the experimental paradigms we tested, we may use these mice to determine the contribution of ATP signaling in injury models.

Our *in vivo* imaging studies have shown that the P2Y₁₂ receptor is required for microglia to rapidly respond to laser-induced cortical damage, and other signaling pathways may mediate the later stages of this response. Whole animal injury models have not yet elucidated a physiological role for this receptor in microglial accumulation around an injury site or mechanical allodynia after spared nerve injury. It is possible that the time-resolution of these models is not sensitive enough to pick up a difference in microglial migration to the site of injury, if in fact microglia from P2Y₁₂-knockout mice only take \sim 4 fold longer to arrive at the affected site. However, these mice may be a useful tool to help determine whether the immediate microglial response to injury is critical to neuronal survival.

Methods

In vivo imaging of microglia. GFP expressing microglia were imaged by twophoton time-lapse microscopy as described². A *z*-stack (~ 100 μ m depth, 2 μ m

z-steps) was acquired every 4 min for 40 min. A maximum projection of the 15 *z*steps centered on the laser ablation was constructed for each time point (30 μ m total depth). Changes in GFP fluorescence were monitored as processes entered a circular zone of 35 μ m in diameter centered on the ablation and normalized to an outer region of 97 μ m in diameter.

Sciatic nerve lesion. The sciatic nerve of wild-type or P2Y₁₂-deficient mice were lesioned using a modified protocol as described⁴. In brief, adult mice were anesthetized with isoflurane and placed under a dissecting microscope. The eyes were kept lubricated and a heating lamp was used to maintain body temperature. An incision was made through the skin and muscle of the right hind limb directly over the hollow area behind the knee. The sciatic nerve was located at the point in which it branches to form the perineal, tibial, and sural nerves. A suture was tied around the entire sciatic nerve above the branch point, and a 1 mm section of the nerve was removed below the suture. The muscle was then sutured closed and skin stapled. The mouse was sacrificed at 1, 2 or 3 d possurgery for microglial analysis. The animal was perfused with 4% PFA, the spinal cord was removed and sectioned (20 µm) at lumbar region 4, 5, and 6, and microglia were visualized by GFP fluorescence. The total number of microglia in a region of gray matter (within 100 µm from the tip of the dorsal horn) was counted (NIH ImageJ software), and the increase in cells on the ipsilateral versus contralateral side was plotted over time.

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Figure Legends

Figure 1. P2Y₁₂ receptors mediate the initial response of microglia to focal laser ablation in the living brain. (a) Tissue ablation by the two-photon laser (white zone in center) results in microglial process extension towards the site of injury. Microglia from mutant mice showed greatly reduced responses over a 40 min observation period. Scale bar = 20 µm. (b) Quantification of process extension toward the site of laser ablation (n = 4 mice per genotype). P2Y₁₂-deficient mice were further analyzed at 2 h post-injury (n = 3). *P < 0.05, **P < 0.01, ***P <0.001; Student's *t*-test. Values represent mean ± s.e.m. (c) Microglia from P2Y₁₂-deficient animals display normal baseline motility in the living brain. The average processes length change (µm) over a 10 min time scale is displayed on the y-axis (n = 3 animals per genotype, 8-16 processes per animal.)

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Figure 2. $P2Y_{12}$ -deficient mice display normal microglial response to sciatic nerve lesion. (a) The whole sciatic nerve of adult wild-type (WT) or $P2Y_{12}$ knockout (KO) mice was lesioned, and the resulting microglial accumulation in lamina I of the dorsal horn of the spinal cord was imaged by GFP fluorescence. Ipsilateral side is left, contralateral side is right; dorsal horns are down, ventral horns are up in each image. Each section is nicked on the ventral horn of the contralateral side. Number of days post-surgery (1, 2, or 3 d) is indicated. Note the increase in microglial cell density on the ipsilateral side. Scale bar represents 100 μ m. (b) Quantification of increase in microglial accumulation on the

ipsilateral side over contralateral side after 1, 2, or 3 d post-injury. For 2 d, P < 0.05. For 1 and 3 d, n = 1 animal, 5-6 sections each; for 2 d, n = 6 animals, 5-6 sections each. Wild-type (WT) is represented by blue diamond, P2Y₁₂-knockout (KO) Is represented by purple square.

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CHAPTER 5

CONCLUSIONS AND FUTURE DIRECTIONS

CONCLUSIONS AND FUTURE DIRECTIONS

Conclusions

Recent evidence suggests that ATP functions as an important signalling molecule in the microglial response to laser induced damage in the CNS¹. We now provide genetic proof that nucleotides report tissue injury by directly activating P2Y₁₂, a G_i-coupled purinergic receptor on microglia. In this study, three experimental paradigms were used to examine the effects of extracellularly applied nucleotides on microglial behavior, ranging from isolated microglia to intact neural systems. In the most reduced of these, namely primary microalial culture, our analysis showed that applied nucleotides can stimulate migration and elicit morphological changes without contributions from other CNS cell types, such as astroglia or neurons. In the more native environment of the hippocampal slice, where traumatic injury to neural tissue is produced during sample preparation, robust P2Y12-dependent polarization and migratory responses to exogenously applied ATP were observed as well, even in the context of endogenously produced injury signals. The most intact of these systems, in vivo imaging of the cortex, clearly showed that P2Y₁₂ mediates microglial responses to local nucleotide injection in the adult brain. Taken together, these observations suggest that the P2Y₁₂ receptor play an important role in modulating microglial response to extracellular nucleotides.

It has recently been reported that inhibitors of connexin channels on astrocytes abrogate microglial response to cortical laser ablation¹, suggesting that astrocytes are activated by ATP and signal to microglia via a different second messenger, or that astrocytes amplify ATP release by positive feedback. In either case, it was unclear whether ATP released from the injury site activates microglia directly, or by promoting the release of other non-nucleotide factors. Our studies suggest that nucleotides have profound effects on microglial polarity and movement through direct activation of P2Y₁₂ receptors on these cells, and nucleotides do not initiate the release of other non-nucleotide chemotactic signals. Recent studies have also shown that "resting" microglia are actually quite dynamic and that this activity is dependent on ATP signalling^{1,2}. Our results indicate that the P2Y₁₂ receptor is not required for this baseline activity, which may therefore be mediated by another purinergic receptor subtype on these cells.

Recent studies have shown that laser induced damage in the cortex of living mice evokes a microglial response that is blocked by broad-spectrum inhibitors of purinergic signalling, apyrase, or a saturating concentration of ATP¹. Thus, it has been proposed that microglial response to injury involves nucleotide signalling through one or more puringeric receptors. Our results support this hypothesis and demonstrate that the P2Y₁₂ receptor plays an important role in this response. While P2Y₁₂-deficient mice display a markedly diminished response to focal injury over a 40 min time-period, this deficit becomes less apparent at later time-

points, suggesting that other signalling mechanisms compensate for loss of $P2Y_{12}$ *in vivo*. Other purinergic receptor subtypes may account for such compensation, but signalling by non-nucleotide factors (such as M-CSF, cannabinoids, or fractalkine) seems more likely since $P2Y_{12}$ -deficient mice exhibit a complete loss of nucleotide-evoked migration both *in vitro* and *in vivo*. Indeed, this is reminiscent of what is seen in the cardiovascular system, where ADP plays an important role in regulating initial morphological changes in platelets as they transform from an inactive to active state³⁻⁵. In humans or mice, loss of P2Y₁₂ function results in prolonged bleeding times, but platelets can eventually clot since other factors (such as thromboxane A₂) also contribute to the activation process⁴. Despite such functional redundancy, drugs that inhibit platelet P2Y₁₂ receptors (such as clopidogrel) have significant cardio-protective effects by decreasing the recurrence of stroke and heart attack, illustrating the importance of this receptor to the overall response to tissue injury.

Microglia express several metabotropic and ionotropic purinergic receptor subtypes⁶, suggesting that nucleotides may play various roles in modulating inflammatory responses to a range of CNS injuries or diseases. For example, pharmacological studies have implicated P2X receptors on microglia in the development of neuropathic pain⁷ or the release of neuro-modulating agents, such as TNF- α and IL-1 $\beta^{8,9}$. In addition, recent studies have identified a second ADP-activated G_i-coupled receptor, P2Y₁₃, with similar pharmacological properties¹⁰ that lies in close proximity to the P2Y₁₂ locus. Interestingly, P2Y₁₃

was less sensitive to cysteine modifying agents (such as clopidogrel) than P2Y₁₂ in electrophysiology assays. As such, analysis of these receptors may uncover the specific structural requirements necessary for clopidogrel's anti-thrombotic actions. While we can detect low levels of P2Y₁₃ transcript in microglia, we were unable to observe protein expression *in vivo* using a P2Y₁₃-specific antibody. These results, together with our functional studies, suggest that P2Y₁₂ is a primary site through which nucleotides mediate the rapid microglial responses that we describe here, although a contribution by P2Y₁₃ cannot be formally excluded.

Our findings support the involvement of P2Y₁₂ receptors at early stages of the microglial response to neural injury, perhaps resembling their role in platelet activation. In the latter case, the physiological consequences of P2Y₁₂ receptor inactivation (genetic or pharmacologic) can be identified through quantitative measurements of blood coagulation rates throughout the course of the clotting process. In the case of microglia, however, long-term physiological consequences of CNS injury and disease are generally assessed over a period of days or weeks, and thus the identification of P2Y₁₂-dependent behavioral or anatomical phenotypes may be difficult to define if, indeed, this receptor plays an early, but redundant role in microglial response to injury. Whether and how a delay of microglial response (as observed in $P2ry12^{-t-}$ mice) would affect neuronal survival near an injury site has yet to be determined. However, the clinical benefit of P2Y₁₂ antagonists in preventing heart attack and stroke

suggests that modulation of this receptor may also have physiological benefit in the context of neural injury and disease.

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Future Directions

While our studies have provided evidence that P2Y₁₂ receptors function in microglial activation, the physiological consequence of receptor loss is still unknown. In the future, we would like to look more closely at animal models to determine if P2Y₁₂-knockout mice display altered cytokine release, clearance of cellular debris, neuronal survival or regeneration, pain sensitivity, or infiltration of other immune cells. Our studies have focused primarily on microglial migration towards the injury site, and not on other aspects of the microglial inflammatory response.

Here we have shown that $P2Y_{12}$ expression is dynamic and down-regulated upon microglial activation, the physiological significance of which we have not yet determined. For example, do microglia decrease receptor expression so that cells that have reached their destination become desensitized to nucleotide signals and do not leave the affected site? In addition, it would be interesting to determine the cellular factors that influence $P2Y_{12}$ expression in the CNS. To address this question, we would like to take activated mircoglia that do not express $P2Y_{12}$ and expose them to the CNS environment by placing them on top of a hippocampal slice or in the presence of homogenized brain. If $P2Y_{12}$ expression is regained, we may then fractionate the material until individual

factors that are sufficient to induce expression are found. In addition, we may take non-derived bone marrow cells (that contain microglial and macrophage precursors) and expose them to CNS factors to determine if a specific molecule(s) is sufficient to induce expression. It may also be interesting to characterize the P2Y₁₂ promoter region and look for known transcription factor binding sites or regulatory elements that may modulate protein levels.

The most significant problem facing the field of microglial biology today is the inability to tell these cells apart from other monocyte-derived cells. Microglia and macrophages share strikingly similar molecular profiles, and consequently there are no good markers to distinguish between these two myeloid cell types. P2Y₁₂ is the first marker that can easily differentiate between these cells since protein expression is very robust in microglia of the healthy brain but completely absent from peripheral macrophages. In this regard, we may use the P2Y₁₂ promoter to drive expression of various genes of interest specifically in microglia of the CNS. For example, we may insert of the lacZ gene into the P2Y₁₂ open reading frame and make a mouse knock-in construct for targeted expression *in vivo*. Of course, platelets will also express this reporter gene, but it is very easy to differentiate between microglia and this megakaryocytic cell type.

In this same manner, we may be able to make microglial specific knockout mice of loxP-flanked genes by expressing Cre recombinase under the P2Y₁₂ promoter. Previous studies have used a similar approach in which Cre recombinase is

driven off the CD11b promoter, a molecular marker shared by both microglia and macrophages¹¹, but these studies are unable to differentiate between the affects of gene deletion in these related cell types. Thus, the P2Y₁₂ locus provides the solution to a major problem in studying the microglial inflammatory response to injury and diseases of the CNS.

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