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6	P2X7 receptor: an emerging target in CNS diseases		
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24 **Key words:** P2X7 receptor, ATP, neurodegenerative diseases, psychiatric

25 disorders

26 Abstract

27

28 The ATP-sensitive homomeric P2X7 receptor (P2X7R) has received particular 29 attention as a potential drug target because of its widespread involvement in 30 inflammatory diseases as a key regulatory element of the inflammasome 31 complex. However, it has only recently become evident that P2X7Rs also play a pivotal role in central nervous system (CNS) pathology. There is an explosion of 32 33 data indicating that genetic deletion and pharmacological blockade of P2X7Rs 34 alter responsiveness in animal models of neurological disorders, such as stroke, 35 neurotrauma, epilepsy, neuropathic pain, multiple sclerosis, amyotrophic 36 lateralsclerosis, Alzheimer's disease, Parkinson's disease, and Huntington's 37 disease. Moreover, recent studies suggest that P2X7Rs regulate the 38 pathophysiology of psychiatric disorders, including mood disorders, implicating 39 P2X7Rs as drug targets in a variety of CNS pathology.

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It has been known for almost a decade that P2X7Rs convey important physiopathological functions in the CNS [1]. The aim and scope of the present review are to summarize the latest developments in the description of these functions, to redirect interest to those fields, where there are still significant gaps in our present understanding and to promote further development of those therapeutic areas, in which P2X7R is the most promising as a potential drug target.

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## 51 The structure and molecular physiology of P2X7Rs

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53 P2X7Rs are ATP-gated, non-selective cation channels belonging to the family of 54 ionotropic P2X receptors. P2X7Rs function in homo-trimeric form and most mammalian P2X7R subunits comprise 595 amino acids [2]. The common 55 56 structural motifs of P2X7Rs are the two transmembrane domains (TM1, TM2), a 57 large, glycosylated, cysteine-rich extracellular loop, a short intracellular N-58 terminal domain, and an intracellular C-terminal domain, which is longer than that 59 of other P2X receptor subunits. Within the family of P2X receptors, so far only the crystal structure of zebrafish (zf)P2X4.1R has been solved in the closed [3] and 60 ATP-binding, open state [4]; nevertheless, its considerable homology with 61 62 mammalian P2X7Rs allowed for the structural modelling of the latter [2]. The molecular architecture of an individual P2X7R subunit is akin to a leaping 63 64 dolphin, with the extracellular loop forming the body, and the TM domains 65 forming the tail. When co-assembled as a trimeric unit, P2X7R has a chalice-like

66 structure, overarching the channel pore (Figure 1A). There are three ATP binding sites localized at the interface of the three subunits; occupancy of at least two of 67 68 the three sites is necessary for the activation of the receptors [5]. The adenine base and the  $\beta$ - and  $\gamma$ -phosphate groups of ATP form hydrogen bonds with the 69 70 respective amino acid residues of the ATP binding pocket, as suggested for the 71 zfP2X4.1R. However, because a residue corresponding to Leu217, which 72 interacts with the ribose moiety, is missing in the mammalian P2X7R, the affinity 73 of ATP to P2X7Rs is more than a hundredfold lower than to other P2XR-74 subtypes [2]. On the other hand, non-conserved residues surrounding the ATP 75 binding site might confer differences in agonist sensitivity between mammalian 76 P2XR species, (i.e. rat P2X7Rs display substantially higher sensitivity to ATP and 77 BzATP than their human and mouse counterparts [6]). A distinctive feature of the mouse P2X7R is that it can be activated by extracellular nicotinamide adenine 78 79 dinucleotide (NAD<sup>+</sup>) by ADP-ribosylation with the ADP-ribosyltransferase 2 80 ectoenzyme [7]. In contrast, less is known about the binding site of antagonists, 81 although potent and selective antagonists of P2X7Rs are now widely available. 82 Earlier data indicated that P2X7R subunits are able to form heterotrimers with 83 P2X4Rs [8], but more recent studies did not confirm this (e.g. [9]).

There are several splice variants of mammalian P2X7Rs, all of which are widely expressed in the nervous system. Hence, a naturally occurring truncated isoform of the human P2X7R (P2X7B) has been found in the CNS [10]; a Cterminally truncated variant of mouse P2X7R has also been identified, which partly retains its functionality, when expressed in tissues of the *P2rx7* gene

deficient mice [11]. Another mouse isoform is the P2X7(k) variant, which in
contrast to P2X7(a), is sensitive to ADP-ribosylation [12, 13].

The gene encoding the human P2X7R (*P2RX7*) is also well known to exhibit a number of non-synonymous single nucleotide polymorphisms (NS-SNPs), which results in a change in amino acid sequence and the expression of different human P2X7 variants, further increasing the structural diversity of P2X7Rs. The functional consequence of several individual NS-SNPs has been determined in native and recombinant systems and their association with various human CNS disease states has been extensively investigated in genetic linkage studies [14].

98 The activation of P2X7Rs results in the opening of the channel pore, allowing the passage of small cations (Na<sup>+</sup>, Ca<sup>2+</sup>, and K<sup>+</sup>). In addition, a hallmark feature 99 100 of the P2X7R is the opening of a non-selective pore in response to repeated or 101 prolonged activation, allowing the permeation of large molecular weight organic 102 cations up to 600-800 Da. The pore forming property of P2X7Rs can be studied 103 by the uptake of high molecular weight cations, such as NMDG<sup>+</sup>, or dyes, such 104 as Yo-Pro-1 or ethidium bromide; nevertheless, its molecular mechanism has 105 remained a highly debated issue, with two alternative, but non excluding 106 possibilities, both having substantial experimental support (Figure 1 B, C). The first potential mechanism is the progressive dilation of the P2X7R-gated channel 107 108 itself. A conformational change of the receptor-protein could be the structural 109 basis for channel dilation, as previously confirmed for other P2XRs (P2X2, P2X4) by electrophysiological methods [15]. In agreement with the pore dilation theory, 110 111 the carboxyl terminal domain [16] and the TM2 region of the P2X7R protein are

essential for pore formation [17]. Moreover, recent studies revealed that the open channel conformation of the P2X7R can allow the passage of negatively charged fluorescent dyes with molecular diameters of up to 1.4 nm [18], and occupation of one or two agonist binding sites favors transition to the desensitized state, whereas occupation of the third binding site favors the transition to the sensitized/dilated state [19].

118 The alternative mechanism involves the recruitment of an additional pore-119 forming protein, most likely the pannexin-1 hemichannel (Panx1). Evidence 120 derived from studies using genetic knockdown of Panx1 indicate that this protein 121 is indispensable for the pore formation (e.g. [20]) and can be selectively affected pharmacologically by colhicine [21]. However, other data conflict with the 122 123 involvement of Panx1 in the formation of the membrane pore (e.g. [22]). 124 Therefore, it appears that although recruitment of pannexin hemichannels is a 125 downstream signaling event closely linked to P2X7R activation, it is not an 126 absolute requirement [23]. A potential dissolution of conflicting results is that 127 different P2X7R splice variants display distinct pore forming properties [12, 23].

The opening of the large pore might eventually result in membrane blebbing and cell death; however, this is not an obligatory consequence of P2X7R activation. Pore formation might gain significance in the pathological sensitization underlying chronic pain as highlighted by a recent study [24]. This paper reported that mutations of the gene encoding the P2X7R, which result in hypofunctional pore formation, affect chronic pain sensitivity in both mice and humans. Moreover treatment with a peptide corresponding to the P2X7R C-terminal domain, which

blocks pore formation, but not cation channel activity, selectively reduced allodynia only in mice with the pore-forming P2rx7 allele. These findings illustrate that the pore formation associated with P2X7R, by itself could be a potential target of personalized therapy to combat chronic pain disorders.

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## 140 Tissue and cell type specific distribution of P2X7Rs

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142 P2X7Rs are expressed by many cell types, including cells of hematopoietic origin 143 (lymphocytes, monocyte-macrophages, microglia) and intrinsic cells of the 144 nervous system (neurons, astrocytes, oligodendrocytes, Schwann cells). P2X7R binding sites have been explored in autoradiographic studies using the 145 146 radioligand [<sup>3</sup>H]-A-804598, and a dense P2X7R binding was found throughout the 147 brain and spinal cord [25], including hypothalamic nuclei, thalamic nuclei, 148 hippocampus, spinal trigeminal nucleus and tract, cortical regions, cerebellum 149 and caudate putamen [25]. Nevertheless, the cell-type specific localization of the 150 P2X7Rs in the CNS has been the subject of a long-standing debate, which has not reached general consensus even after a decade: immunohistochemical 151 152 findings are inhomogeneous and contradict findings obtained by physiological 153 and neurochemical methods. Whereas early studies found a prominent 154 expression of P2X7R immunoreactivity (IR) on excitatory nerve terminals [26], 155 and later studies confirmed these findings throughout the CNS [27, 28]; other 156 groups questioned these findings, revealing P2X7R-immunoreactivity in brain 157 sections obtained from P2X7R deficient animals [29]. Subsequently however,

158 functional splice variants of rodent P2X7R [11, 12] were identified which are likely 159 to be responsible for P2X7-pseudo-immunoreactivities, found in the brain of P2X7R<sup>-/-</sup> mice. These variants represent either gain- or loss-of function P2X7Rs, 160 161 and may explain the high variability of responses induced by P2X7R stimulation. 162 Other studies reported an activity-dependent expression pattern of P2X7Rs, 163 induced or upregulated following an insult such as a seizure [30], ischemia [31], sleep deprivation [32], undernourishment [33], or morphine tolerance [34]. A 164 165 recent study utilizing single particle tracking photoactivated localization 166 microscopy (sptPALM) revealed that Dendra2 tagged P2X7Rs transfected to 167 hippocampal neurons formed two dynamic populations within the extrasynaptic membrane of proximal dendrites: one was composed of rapidly diffusing 168 169 receptors and another stabilized within nanoclusters, both being rarely 170 appositioned to synaptic sites [35].

In contrast to immunohistochemistry, the available evidence on functional 171 172 P2X7Rs on different cell types of the CNS is convincing. Functional studies, verifying P2X7Rs on neurons, astrocytes and microglia are presented in Table 1. 173 The most parsimonious explanation for the contradictory findings is that the 174 175 expression of P2X7Rs dynamically changes in response to experimental variables such as age or different levels of stressful stimuli prior to sample 176 177 collection (freshly prepared vs. fixed sections). Moreover, under in vivo conditions 178 even mild stimuli, such as saline injection, may cause a dramatic change in the expression level of P2X7Rs. 179

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# 181 **Physiopathology of P2X7 receptors**

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183 P2X7R function can be studied with a selection of pharmacological and genetic tools (Box 1). The activation of P2X7Rs is followed by Ca<sup>2+</sup> influx and a variety of 184 cellular responses depending on the cell type investigated (Figure 2). Outside the 185 186 nervous system, the most prominent role of P2X7R is in the regulation of cytokine response to inflammatory challenge. In fact, P2X7R is a key regulatory 187 element of the inflammasome molecular complex, providing the external stimulus 188 189 necessary for the posttranslational modification and subsequent release of the 190 pro-inflammatory cytokine IL-1β. The role of P2X7Rs has been confirmed in the 191 regulation of central cytokine response after LPS priming [36]. This effect could 192 be involved in physiological and pathological actions controlled by P2X7Rs, such 193 as memory formation [37]; sleep [32], fever [38], hyperalgesia [39] and 194 depression [40, 41].

195 However, a major caveat in our understanding of the physiopathology of 196 P2X7R function is how the endogenous activation of P2X7Rs is achieved, given 197 the low affinity of the endogenous agonist ATP. ATP is present in the synaptic 198 vesicles and is co-released as a co-transmitter with various other transmitters in 199 the autonomic nervous system under physiological conditions [42]. This holds 200 also true to a certain extent for central synapses and the increase in extracellular 201 ATP in response to normal neuronal activity might transiently reach the high 202 micromolar concentration required for the activation of P2X7R, at least in the 203 synaptic cleft. However, a more widespread activation of P2X7Rs is expected

204 under pathological conditions, when tissue damage, trauma or other pathological 205 signals provide an ATP-rich extracellular milieu, which might lead to the 206 activation of extrasynaptic and extraneuronal P2X7Rs. In addition, the possibility 207 of constitutive activity without the presence of the endogenous agonist cannot be 208 excluded either and should be further investigated. In the CNS, the best 209 characterized consequence of P2X7R activation is the release of 210 neurotransmitters, in particular of glutamate to the extracellular space [43]. This 211 effect could be evoked both from synaptosomes [44] and from astrocytes [45]. In 212 nerve terminals and cell lines expressing recombinant P2X7Rs, the P2X7R 213 mediated glutamate release appears to be both exocytotic and non-exocytotic, 214 [46, 47]. P2X7R mediated excitatory amino acid efflux can be detected in acutely 215 prepared brain slices by neurochemical (e.g. [48, 49]) and electrophysiological 216 techniques [50]. In rat hippocampal (hilar neurons; [51] CA1 neurons [52]), and 217 midbrain slices (locus coeruleus; [50]), stimulation of P2X7Rs by BzATP elicited 218 an increase of the frequency but not amplitude of spontaneous excitatory 219 postsynaptic currents (sEPSCs) and miniature (m)EPSCs. Occasionally [49, 50] 220 the P2X7R-mediated glutamate release was sensitive to blockade by fluorocitric 221 acid, a glia-selective metabolic poison, and to antagonists of glutamate receptors. 222 These findings imply that glutamate release induced by P2X7R stimulation from 223 neurons could also be indirect, mediated by glutamate release from astrocytes, 224 acting subsequently on glutamatergic nerve terminals.

To add further complexity to neuron-glia and glia-neuron P2X7R signaling,
 P2X7R stimulation elicits or reinforces the release of ATP, thereby providing an

227 auto-stimulatory loop. This effect was observed in retinal ganglion cells [53] 228 hippocampal brain slices [49] and cultured spinal cord astrocytes [54]. The 229 mechanism of P2X7R-driven ATP release could be exocytotic, as observed by 230 total internal reflection microscopy in neuroblastoma cells [55], whereas in other 231 studies it appears to involve connexin and/or pannexin hemichannels [49, 54].

A further interesting function of P2X7Rs is to regulate differentiation and cellfate during development. P2X7Rs are expressed by both embryonic [56] and adult neural progenitor cells (NPCs) in the subventricular zone of the lateral ventricle [57]. Whereas stimulation of P2X7Rs induces neuronal differentiation in embryonic NPCs [56], other studies indicated that P2X7Rs stimulate gliogenesis [58]. In contrast, the activation of P2X7Rs on adult, cultured NPCs decrease cell proliferation and induce necrotic/apoptotic cell death [57].

Of note, a very recent study showed that P2X7Rs regulate ion channel density and protein composition/function of the axon initial segment, a key structural element of neuronal excitability and in consequence action potential initiation in cultured hippocampal neurons and brain slices [59].

It has been known for a long time that P2X7R activation might lead to cell death through pore formation as it has been described for peripheral immune cells. However, a more recently emerging view is that P2X7Rs also convey trophic function against cell-death promoting physiological or pathological stimuli: for example the microglial "suicide" P2X7R promotes cell cycle progression and proliferation [60, 61], and this receptor might act as a scavenger for the removal of apoptotic cells in the absence of its ATP ligand [62, 63].

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## 251 **P2X7R** as a potential target in neurological diseases

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ATP is released in large quantities following any kind of cell injury, and the ensuing stimulation of the low affinity P2X7R results in necrosis/apoptosis or proliferation as the two opposing end-points of neuroinflammation. P2X7R antagonists are potential therapeutics of traumatic brain injury, stroke, epilepsy, neuropathic pain, and neurodegenerative illnesses, because in these cases secondary cell damaging conditions accompany the primary pathological condition.

Middle cerebral artery occlusion, the most widely used animal model of 260 261 cerebral ischemia, results in cell death in the core of the affected neuronal tissue, while around it, in the so called penumbra, the cellular damage is reversible. Both 262 263 infarct size and neurological deficits were reduced by P2X7R antagonists [64, 264 65]. In combination with the sequential up-regulation of P2X7R-IR in microglia 265 and then in astrocytes and neurons, this receptor-type was considered to be a primary target of the considerable amounts of ATP released. Similar results were 266 reported for subarachnoid hemorrhage [66], traumatic brain [67, 68] or spinal 267 cord injury [69] and ischemic retina degeneration [70]. However, a later study 268 269 failed to reconfirm the protective action of P2X7R in spinal cord injury [71]. 270 Reperfusion after transient global cerebral ischemia exacerbates the 271 consequences of oxygen/glucose deprivation (OGD) due to microglial and 272 astroglial activation [72]. The ensuing neuroinflammatory reaction is also

alleviated by P2X7R antagonists [73, 74]. BBG partially reversed the OGDinduced anoxic depolarization and cell damage in cultured oligodendrocyte cells
[75]. Accordingly, left common carotid artery occlusion decreased P2X7Rimmunoreactivity at oligodendrocyte precursor cells in cerebral cortex, subcortical
white matter and hippocampus [76].

Status epilepticus (SE)-like seizures, modelled in rodents by pilocarpine or 278 279 kainate, up-regulate P2X7R-immunoreactivity in microglial cells [77] astrocytes 280 and neurons [78]; quantification by western-blotting confirmed these results [79, 281 80]. Utilizing the intra-amygdala application of kainate as an epileptic stimulus 282 [79, 80], it was shown that (1) Bz-ATP facilitated and prolonged the EEG activity 283 caused by seizures, and (2) P2X7R antagonists had a neuroprotective effect 284 after epilepsy due to suppression of IL- $\beta$  production and microglial response. 285 More recent findings suggest that the effect of P2X7Rs during SE depends on 286 the nature of the chemical stimulus used. A-438079 decreased pilocarpin-287 induced seizure susceptibility in mice by interrupting a direct facilitatory 288 interaction between P2X7- and muscarinic receptors [81] or blockade of the 289 release of the protective TNF- $\alpha$  [82]. P2X7R activation also influenced leukocyte 290 infiltration [83] and reactive astrogliosis following SE [84].

The involvement of P2X7Rs in different models of inflammatory and neuropathic pain and the potential therapeutic effect of P2X7R antagonists are well documented [85]. Down regulation of P2X7Rs with siRNA or BBG prevented the induction of spinal long-term potentiation *in vitro* and at the same time alleviated mechanical allodynia in naive rats *in vivo* [39]. Central sensitization of

296 nociceptive neurons could be produced by intrathecal superfusion of Bz-ATP and 297 was depressed by P2X7R antagonists [86]. Additional studies extended these 298 findings to mechanisms participating in the development of neuropathic or 299 orofacial pain [87-89], bone cancer pain [90] and migraine [91]. Recent studies 300 highlighted the association between human P2X7R variants with chronic pain 301 sensitivity [24].

302 Multiple sclerosis (MS) is a chronic degenerative disease of the CNS that is 303 characterized by focal lesions with inflammation, infiltration of immune cells, 304 demyelination, oligodendroglial death and axonal damage [92]. A putative 305 association of the P2X7R gene with this illness was indicated by the most 306 frequent expression of the gain-of-function T allele of rs17525809 polymorphism 307 of the receptor, which yields an Ala-76 to Val change in its extracellular domain 308 [93]. The overexpression of P2X7Rs was detected in experimental autoimmune 309 encephalomyelitis (EAE), an animal model of SM [94], whereas the amelioration 310 of EAE was found in P2X7R deficient animals [95, 96], but see [97]. Further, 311 pannexin-1 knockout mice with restricted ability to mediate pore development/dye 312 uptake after P2X7R stimulation, also displayed a delayed onset of clinical signs 313 of EAE and decreased mortality when compared with wild-type mice [98].

Amyotrophic lateral sclerosis (ALS) is characterized by the progressive degeneration of motor neurons in the spinal cord, brainstem and motor cortex, leading to respiratory failure and death of the affected patients within a few years of diagnosis [99]. Microglia and astrocytes are major contributors to motor neuron dysfunction in ALS through the maintenance of a chronic inflammatory response.

319 Transgenic mice expressing a mutant protein Cu<sup>+</sup>/Zn<sup>+</sup> superoxide dismutase 320 SOD1-G93A, which directly enhances the activity of the main reactive oxygen 321 species producing enzyme in microglia (NADPH oxidase 2: NOX2) is used widely 322 as a model of ALS [100]. P2X7R activation by BzATP induced the death of motor 323 neurons in mixed astrocytic/neuronal cultures prepared from wild-type mice [101]. Further, apyrase, an enzyme degrading ATP or BzATP, decreased neuronal 324 325 death observed in cultures prepared from SOD-G93A spinal cord. Bz-ATP also 326 increased the activity of NOX2, leading to motor neuron damage, an effect which 327 did not occur in primary microglia cultures of SOD-G93A mice lacking P2X7Rs [102]. 328

329 A neuropathological hallmark of Alzheimer's disease (AD) is the appearance of 330 plaques consisting of extracellular  $\beta$ -amyloid peptide (A $\beta$ ) surrounded by reactive microglial cells [103]. A $\beta$  triggered increases in intracellular Ca<sup>2+</sup>, ATP release, 331 IL-1ß secretion and plasma membrane permeabilization in microglia from wild-332 type but not P2X7R<sup>-/-</sup> mice [104]. These findings and the neuroprotective effects 333 334 of BBG against intrahippocampally injected A $\beta$  suggest that A $\beta$  activates a 335 purinergic autocrine/paracrine stimulatory loop of which the P2X7R is an 336 obligatory component. In fact, in vivo inhibition of the P2X7R in mice transgenic 337 for mutant human APP indicated a significant decrease of the number of 338 hippocampal amyloid plaques [105].

339 Parkinson's disease (PD) is a motor disease affecting the striatal 340 dopaminergic system, by damaging dopaminergic neurons in the substantia 341 nigra. In the disease model induced by unilateral intrastriatal injection of 6-

342 hydroxydopamine, BBG and A-438079 prevented the ensuing synaptotoxicity, 343 gliosis and neurotoxicity [106]. In another study, A-438079 prevented the 344 depletion of striatal dopamine stores by 6-hydroxydopamine treatment, but this 345 was not associated with a reduction of dopaminergic cell loss [107]. Similarly, the 346 effects of P2X7R antagonists appeared to depend on the neurotoxin used, 347 because in MPTP- or rotenone-induced Parkinson models, the genetic deletion of 348 the P2X7R did not increase survival rates of mice compared to wild-type 349 counterparts [108].

Huntington's disease (HD) is an autosomal dominant neurodegenerative disorder caused by a triplet repeat expansion coding for a polyglutamine sequence in the N-terminal region of the huntingtin protein. A higher P2X7R level was documented by western-blot analysis in the striatum of transgenic mice models of this disease [109]. In addition, P2X7R antagonists prevented neuronal apoptosis and attenuated body weight loss and motor-coordination deficits.

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#### 357 **P2X7R** as a potential target in psychiatric disorders

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Mood disorders arise from complex interactions between genetic, developmental and environmental factors [110, 111]. Linkage studies suggested that variations of the chromosome 12q24.31 containing candidate genes for P2X7R, P2X4R and calmodulin-dependent protein kinase b (CaMKKb) may be associated with major depressive, bipolar and anxiety disorders. It has repeatedly been reported that the NS-SNP rs2230912 coding for the P2X7R-Glu460Arg is associated with

365 major depressive disorder [112, 113]. Further, relevant SNP mutations identified 366 by linkage studies were introduced into the human recombinant P2X7R and were expressed in human embryonic kidney cells [114]. The measurement of their 367 368 functional properties by the patch-clamp technique indicated that some of them, 369 including Glu460Arg, exhibited a strong impairment of the current response to 370 ATP, while other mutants demonstrated significant increases in sensitivity. In 371 contrast, other studies failed to confirm the allelic or genotypic association of 372 rs2230912 or other SNPs of P2X7R with mood disorders [115, 116]. The reasons 373 for this discrepancy are presently unknown. Eventually, variations in the P2X7R 374 gene were described to be associated with cognitive manic symptoms in bipolar 375 disorders [117], but not in schizophrenia [118].

376 Production of TNF- $\alpha$  and IL-6 is initiated by the activation of Toll-like receptors 377 (TLRs) by e.g. bacterial lipopolysaccharide. The formation of IL-1 $\beta$  also requires 378 TLR4 induction of gene transcription but requires an additional step, the 379 processing of pro-IL-1 $\beta$  to the mature form of IL-1 $\beta$ , which is then released via NLRP3 referred to as the "inflammasome" [110, 119]. P2X7Rs are indispensable 380 381 activators of NLRP3. Inflammatory cytokines have been suggested to play key 382 roles in the development of depressive behavior. Their levels are elevated in 383 depressed patients [110, 120] and rodents exposed to stressful stimuli [111]. 384 These cytokines are potent activators of the hypothalamic-pituitary-adrenal axis 385 through which the secretion of hypothalamic corticotropin releasing hormone 386 (CRH), pituitary adrenocorticotropic hormone (ACTH) and corticosterone are 387 stimulated. In this respect it is interesting to note that P2X7R stimulation also

directly leads to increased ACTH secretion from the terminals of hypothalamicmagnocellular neurons [121].

390 The interrelationship between inflammatory cytokines, P2X7Rs and mood 391 related behavior has been intensively studied in animal models. The genetic 392 deletion of P2X7Rs resulted in antidepressive-like behavior in the forced swim 393 and tail suspension tests and alleviated amphetamine induced hyperactivity [40, 394 41]. Although P2X7Rs are present at peripheral/central immunocytes, glial cells 395 and neurons, it was shown that macrophages and microglia are not responsible 396 for the detected changes in mood measured by tail suspension test and amphetamine-induced hyperlocomotion in P2X7R<sup>-/-</sup> mice [41]. On a larger scale, 397 398 several potential mechanisms were identified for the antidepressant phenotype of P2X7R<sup>-/-</sup> mice, such as the absence of P2X7R-mediated glutamate release, 399 400 elevated basal brain-derived neurotrophic factor (BDNF) production, enhanced 401 neurogenesis and increased serotonin bioavailability in the hippocampus [48]. It 402 has also been observed that P2X7Rs are downregulated in the hippocampus in response to chronic stress [122] and P2X7R<sup>-/-</sup> mice exhibited impaired adaptive 403 coping responses to repeated stress [123], which enlighten the potential role of 404 405 P2X7Rs as a protective adaptive mechanism in the process leading to mood 406 disorders.

The above data illustrate that P2X7R seems to be activated in a number of different pathological conditions raising the possibility that the receptor is one common avenue of cellular stress signaling pathways (Figure 2). However, one should keep in mind that the pathophysiology of CNS diseases is very complex

411 involving a multiplicity of mediators and signaling pathways and the P2X7R is 412 only one among the multiple signaling pathways activated. Moreover, the 413 significance of this avenue is probably not uniform in all CNS pathologies and 414 could be more prominent in certain disease conditions (e.g. chronic pain, status 415 epilepticus) than in other ones (e.g. Parkinson's disease), depending on the 416 expression of P2X7Rs in the brain area afflicted. Finally, important physiological 417 functions mediated by P2X7Rs should not be neglected. For instance, taking into 418 account that the purportedly necrotic/apoptotic P2X7Rs also convey trophic and 419 adaptive changes, their role might vary or even reverse during the course of the 420 same disease, because neuroinflammation, regulated by P2X7Rs has also a 421 double-faced role. In fact, inflammation initially is a protective reaction and 422 becomes detrimental only, when it progresses to an excessive or chronic phase. 423 These aspects serve as explanations to conflicting results with P2X7R inhibition 424 on the disease outcome (e.g. [95-97]) and should also be addressed when 425 P2X7R is considered as a potential human drug-target.

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# 428 Current development of P2X7R ligands

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Although end-products of the pioneering developments of P2X7R antagonists,
such as CE-224,535 [124] and AZD 9056 [125] have not proved efficacious in
Phase II trials in rheumatoid arthritis patients, clinical studies revealed an
acceptable safety and tolerability profile of such antagonists as a whole [124-

434 126], opening up the possibility of developing P2X7R-targeting compounds in
435 new areas, such as CNS disorders.

436 In recent years, a number of different classes of small molecular weight, drug-437 like P2X7R ligands have been developed (Table 2), and P2X7Rs have been 438 gualified as the most "druggable" target within the P2X receptor family [85, 127]. More recently, the development of centrally penetrating potent P2X7R 439 440 antagonists has also been reported (Table 2). In addition, systematic search 441 through compound libraries resulted in the further discovery of novel P2X7R 442 antagonists and allosteric modulators utilizable either for basic research or drug 443 development. Analyses of natural compounds have also resulted in several 444 valuable P2X7R ligands (Table 2).

445

# 446 **Concluding remarks**

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448 In conclusion, P2X7R mediated pathways appears to be a common avenue of 449 many CNS disorders of different aetiology and P2X7R antagonists are potential 450 drugs to treat them. Their immense advantage may lie in the absence or low 451 density of P2X7Rs in healthy tissue and therefore in the limited systemic side 452 effects of these compounds. However, major caveats in our understanding of the 453 physiopathological functions of central P2X7Rs should be further elucidated (Box 454 2). Though the majority of known antagonists fail to pass the blood-brain barrier, 455 BBG and some new and high affinity P2X7R antagonists readily enter the CNS 456 [128]. Further, recently identified negative allosteric modulators of P2X7Rs (e.g.

457 certain phenothiazine-type antipsychotic drugs), already registered for human458 use [129], may become important therapeutic tools.

459 The future development of new P2X7R antagonists has to take into 460 consideration that P2X7R isoforms may exhibit large variability between different 461 species in their agonist/antagonist sensitivities. Therefore, the classic search for 462 new pharmacologically active compounds based on the use of laboratory 463 animals, may lead to spurious negative or positive results. A further complicating 464 factor is the presence of numerous splice variants and SNPs widely distributed in 465 the animal and human organism; their sensitivities to pharmacological blockade 466 is often different from that of the wild-type receptor. Hence, the development of new and therapeutically valuable P2X7R antagonists is a tedious task but the 467 468 reward may be enormous.

469

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903

904	Boxes
905	
906	Box 1. Tools to study P2X7 receptors
907	
908	The continuously evolving interest in this receptor resulted in the generation
909	of various tools to study its function. P2X7Rs could be identified based on
910	the following distinctive pharmacological features:
911	• The affinity of the endogenous agonist ATP is low, in the high micromolar-
912	millimolar range.
913	• BzATP is a more potent agonist than ATP itself. It has been frequently
914	used mistakenly as a selective agonist of P2X7R. This is, however, not
915	valid, because BzATP also binds to other P2X receptors with high affinity.
916	• The effect of ATP and BzATP are potentiated by a low Ca <sup>2+</sup> /no Mg <sup>2+</sup> -
917	containing external medium.
918	• There are several potent antagonists available, such as A-438079, A-
919	740003, the negative allosteric modulator AZ-10606120 and Brilliant blue
920	G (BBG); among them BBG is selective in concentrations below 1 $\mu M.$
921	This antagonist is also a useful tool in <i>in vivo</i> experiments. The
922	penetration of BBG through the blood-brain barrier has already been
923	determined and using doses not higher than 50 mg/kg, the resultant brain
924	concentration remains below 1 $\mu$ M [105]. It should be noted, however,
925	that many P2X7R antagonists, including BBG also inhibit Panx1
926	channels. Therefore, BBG alone is inadequate to prove the involvement

927 of P2X7Rs [130]. In this respect, a valuable compound could be Brilliant
928 blue FCF, which inhibits Panx1, but not P2X7R [131].

- Novel radioligands, i.e. [<sup>3</sup>H]A-804598 are also available to characterize
   the affinity of newly developed compounds to rodent P2X7Rs [25].
- 931 In addition to pharmacological approaches,
- genetic knock-down by siRNA has been increasingly used to silence
   P2X7Rs in the past years in both *in vitro* and *in vivo* studies (e.g. [34, 39]).
- Mouse lines, genetically deficient in P2X7Rs, initially generated by the companies Glaxo (LacZ gene and neomycin cassette insertion into exon 1; [132]) and Pfizer (Neo insertion in exon 13, close to the carboxyl terminal; [133]), have also been widely used. However, none of these mouse lines could be regarded as fully deficient in P2X7Rs, as individual splice variants evaded inactivation [11, 12].
- For studies of P2X7R function in morphologically identified neurons,
   astrocytes or microglia, the GFP-P2X7 reporter mouse seems to be a
   crucial tool [134].

943

## 945 **Box 2. Outstanding Questions**

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947 Despite the large interest in P2X7Rs and the correspondingly high number of 948 publications dealing with this receptor, many questions still remain unresolved.

• The C-terminus of the P2X7R has been implicated in regulating receptor function including signaling pathway activation, cellular localization, proteinprotein interactions, and post-translational modification [135]. It would be important to learn the three-dimensional structure of the P2X7R C-terminal tail, which is yet to be determined [4].

Although repetitive or long-lasting stimulation of P2X7Rs by ATP allows the
 passage of 600-800 Da organic molecules through the cell membrane, the
 mechanism of pore opening is still a matter of debate. There are good
 arguments favouring an accessory protein, with Panx1-hemichannels
 probably involved in this effect, but the cationic channel-dilation theory is also
 an attractive alternative.

Original work based on co-immunoprecipitation with epitope tagged subunits
 demonstrated that overexpressed recombinant P2X1-6 subunits could form
 hetero-oligomeric complexes, while P2X7 was able to form only homomeric
 receptor channels [136]. However, it remains to be established whether true
 functional P2X4/7 heteromers are formed in native systems, which might have
 great significance for CNS immune functions e.g. in microglia.

A lot of controversy has arisen on the issue of whether P2X7Rs are located
 exclusively at microglia and astroglia in the CNS or also at neurons (see the

968 discussion on "Tissue and cell type specific distribution of P2X7Rs"). The 969 solution of this enigma might be that under normal conditions P2X7Rs are 970 dormant but after various types of damaging conditions (mechanical trauma, 971 ischemia, inflammation, etc.) they become unmasked, mostly at central 972 immunocytes but probably also at neurons. Already the tissue damage 973 afflicted to cells during the culturing procedure or the preparation of brain 974 slices may be sufficient to induce the expression of previously absent 975 P2X7Rs.

Although endogenous activation of P2X7Rs under disease conditions has
 repeatedly been proven, its exact mechanism is not fully understood, given
 the low affinity of ATP. The possibility of constitutive activity of this receptor as
 well as its potential endogenous ligands other than ATP should be explored.

Whereas available gene deficient mouse models are not fully deficient in
 P2X7Rs, more advanced mouse models, such as cell-type specific and/or
 inducible knockouts, optogenetic constructs, as well as humanized mouse
 models reproducing human gene polymorphisms in rodents are yet to be
 generated for probing P2X7R function.

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# 987 Tables

- 988 Table 1. Examples from recent studies verifying functional P2X7Rs on different
- 989 cell types of the rodent central nervous system.

Cell type/Brain area,	Technique	Refs		
preparation				
Neurons				
Cerebral cortex, purified	neurochemistry, Ca <sup>2+</sup>	[44]		
synaptosomes	fluorimetry			
Midbrain, synaptic terminals	Ca <sup>2+</sup> microfluorimetry	[137]		
Neurohypophysis, nerve	patch clamp	[138]		
terminals	electrophysiology			
Caudal brainstem, nerve	neurochemistry	[139]		
terminals				
Hippocampus, isolated hilar	patch clamp	[51]		
neurons	electrophysiology			
Retina, isolated ganglion cells	patch clamp	[53]		
	electrophysiology			
Suprachiasmatic nucleus,	Ca <sup>2+</sup> imaging	[140]		
isolated neurons				
Embryonic spinal cord, cultured	neurochemistry	[141]		
neurons				
Cortex, cultured neurons	neurochemistry	[142]		
Astrocytes				

Cortex, in situ	patch clamp	[143]		
	electrophysiology			
Cortex, cultured	patch clamp	[144]		
	electrophysiology			
Cerebellum, cultured	neurochemistry	[145]		
Human, cultured	Ca <sup>2+</sup> fluorimetry	[146]		
Bergmann glia				
Cerebellum, in situ	patch clamp	[147]		
	electrophysiology, Ca <sup>2+</sup>			
	imaging			
Satellite glia				
Immature dorsal root ganglion,	electrophysiology	[148]		
isolated				
Microglia				
Cortex, in situ	patch clamp	[149]		
	electrophysiology			
N9 microglia, cultured	neurochemistry	[150]		

- 992 Table 2. Non-comprehensive list of different classes of P2X7 receptor
- 993 antagonists and allosteric modulators. For more information see [151]
- 994

Class/Compound	Function	Refs		
Novel, small molecule				
(1H-pyrazol-4-yl)	antagonist	[152, 153]		
acetamides				
benzamides	antagonist	[154, 155]		
tetrasubstituted-	antagonist	[156]		
imidazoles				
2-oxo-N-(phenymethyl)-	antagonist	[157]		
4-				
imidazolinecarboxamides				
Novel, small molecule, CNS active				
JNJ-47965567	antagonist	[128]		
polycyclic carboranes	antagonist	[158]		
Identified by screening compound libraries				
clemastine	Positive allosteric	[159]		
	modulator			
perazine-type	Negative allosteric	[129]		
antipsychotic drugs	modulator			
ivermectine	Negative allosteric	[160]		
	modulator			

Natural compounds	Natural compounds			
teniposide	antagonist	[161]		

997 Figure Legends

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999 Figure 1. The simplified schematic structure of the P2X7R in open state (A) and 1000 during pore formation (B and C). The P2X7R functions as a homo-trimer, forming 1001 a chalice-like structure, while the individual P2X7R subunit is akin to a leaping 1002 dolphin. The agonist binding sites are located at the subunit interfaces and the 1003 occupation of two out of three binding sites is necessary for opening of the 1004 channel. In addition to ATP, which is the presumed endogenous agonist, the mouse P2X7R receptor could also be activated by NAD<sup>+</sup> through ADP-1005 1006 ribosylation. The activation of the receptor-ion channel leads to the inward flux of 1007 cationic current. Prolonged and /or repeated activation of P2X7R and occupation 1008 of the third agonist binding site renders the membrane permeable for high molecular weight organic cations and dyes such as NMDG<sup>+</sup> and Yo-Pro-1<sup>+</sup> (B 1009 1010 and C). B. One potential mechanism of the pore formation is the dilation of the 1011 P2X7R-mediated channel pore itself. C. Alternatively, but not exclusively, 1012 additional pore forming proteins, such as pannexin (Panx1) might be recruited, 1013 which seem to be indispensable for pore formation under certain circumstances.

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Figure 2. Common disease mechanism by P2X7R mediated pathways in CNS disorders of different etiology. P2X7 receptors are expressed on nerve terminals, astrocytes and microglia and they are upregulated upon various disease conditions. Stress signals, such as hypoxia/ischemia (metabolic limitations),

1020 mechanical injury, and bacterial or chemical toxins elicit the endogenous 1021 activation of P2X7R and leads to a self-amplifying ATP release and to further activation of P2X7 receptors on neighbouring cells. Following the influx of Ca<sup>2+</sup> 1022 1023 through the receptor ion channel complex, P2X7 receptor activation (a) releases 1024 glutamate from nerve terminals and astrocytes by both exocytotic and non-1025 exocytotic mechanisms, which may give rise excitotoxicity; (b) leads to the 1026 posttranslational processing of pro-IL-1 $\beta$  to the leaderless, mature IL- $\beta$  and to its 1027 further release by the NLRP3 inflammasome and that of other cytokines, which contribute to neuroinflammation; (c) enhance ROS production and thereby 1028 1029 aggravate protein misfolding and neuronal damage; (d) leads directly or 1030 indirectly to cell death and the following reactive astrogliosis (e) directly or 1031 indirectly downregulates the production of BDNF and the following 1032 neuroplasticity. These key mechanisms could be manifested and contribute to 1033 disease pathology in Alzheimer's disease (AD), Parkinson's disease (PD), 1034 Huntington's disease (HD), status epilepticus (SE), amyotrophic lateral sclerosis 1035 (ALS), multiple sclerosis (MS), stroke, pain and mood disorders in different forms 1036 and proportion, depending on the etiology. GLU, glutamate, ROS, reactive 1037 oxygen species.

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1043 Figure 1

