
5 Pharmacological Properties and Biological 6 Functions of the GPR17 Receptor, 7 a Potential Target for Neuro-Regenerative 8 Medicine 9 10

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14

15 Abstract

16 In 2006, cells heterologously expressing the “orphan” receptor GPR17
17 were shown to acquire responses to both uracil nucleotides and cysteinyl-
18 leukotrienes, two families of signaling molecules accumulating in brain or
19 heart as a result of hypoxic/traumatic injuries. In subsequent years, evi-
20 dence of GPR17 key role in oligodendrogenesis and myelination has
21 highlighted it as a “model receptor” for new therapies in demyelinating
22 and neurodegenerative diseases. The apparently contrasting evidence in
23 the literature about the role of GPR17 in promoting or inhibiting
24 myelination can be due to its transient expression in the intermediate
25 stages of differentiation, exerting a pro-differentiating function in early
26 oligodendrocyte precursor cells (OPCs), and an inhibitory role in late
27 stage maturing cells. Meanwhile, several papers extended the initial data
28 on GPR17 pharmacology, highlighting a “promiscuous” behavior of this
29 receptor; indeed, GPR17 is able to respond to other emergency signals like
30 oxysterols or the pro-inflammatory cytokine SDF-1, underlying GPR17
31 ability to adapt its responses to changes of the surrounding extracellular
32 milieu, including damage conditions. Here, we analyze the available
33 literature on GPR17, in an attempt to summarize its emerging biological
34 roles and pharmacological properties.

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Keywords

Differentiation • GPCR • Multiple sclerosis • Myelination •
Oligodendrocyte precursor cells

Abbreviations

39	CNS	central nervous system
40	cysLT	cysteinyl-leukotrienes
41	EAE	experimental autoimmune encephalomyelitis
42		
43	ERK1/2	extracellular signal-regulated kinases 1 and 2
44		
46	FACS-	frontal affinity chromatography-
47	MS	mass spectrometry
48	GPCRs	G-protein coupled receptors
49	HM	homology modeling
50	Lys	lysolecithin
51	MCAo	middle cerebral artery occlusion
52	MS	multiple sclerosis
54	NC-	Nomenclature Committee of the
55	IUPHAR	International Union of
56		Pharmacology
57	OLs	oligodendrocytes
58	OPCs	oligodendrocyte precursor cells
59	MBP	myelin basic protein

1 Introduction: The History of GPR17

62 In 2006, a paper was published where it was
63 demonstrated that cells heterologously
64 expressing the “orphan” receptor GPR17 (i.e., a
65 molecularly identified, 339 amino acid-long
66 G_i-protein-coupled receptor that still lacked a
67 defined ligand) acquired responses to both uracil
68 nucleotides (such as UDP, UDP-glucose,
69 UDP-galactose) and cysteinyl-leukotrienes
70 (cysLTs, like LTC₄ and LTD₄) (Ciana et al.
71 2006), two chemically unrelated families of sig-
72 naling molecules that are known to massively
73 accumulate in organs like the brain or the heart
74 as a result of hypoxic/traumatic injuries. Uracil
75 nucleotides and cysLTs were already known to
76 exert multiple biological effects via the

activation of separate G-protein-coupled 77
receptors (GPCRs): the eight recognized P2Y 78
receptor subtypes (the P2Y_{1,2,4,6,11,12,13,14} 79
receptors, (Abbracchio et al. 2006) and the two 80
CysLT1 and CysLT2 receptors. Interestingly, the 81
GPR17 sequence had been originally described 82
as the result of a cloning strategy based on the 83
use of RT-PCR degenerate oligonucleotide 84
primers designed on the sequences of the P2Y₁ 85
and P2Y₂ receptors, with the final aim of 86
identifying new members of this receptor family 87
(Blasius et al. 1998). GPR17 was later found to 88
be at an intermediate structural and phylogenetic 89
position between already known P2Y and CysLT 90
receptors, and GPR99, recently proposed as the 91
third CysLT receptor (also known as 92
2-oxoglutarate receptor 1, OXGR1) (Kanaoka 93
et al. 2013) (Fig. 1), in the so called “purine 94
receptor cluster” of class A GPCRs (Fredriksson 95
et al. 2003). GPR17 also emerged as represented 96
the closest receptor to a common ancestor that, 97
during evolution, could have generated both P2Y 98
and CysLT receptors (Ciana et al. 2006; 99
Parravicini et al. 2008; Parravicini et al. 2010). 100
To further highlight GPR17 structural similarity 101
to the other members of the P2Y family, a partial 102
sequence of the rat receptor was initially 103
identified from rat striatum by employing oligo- 104
nucleotide primers specifically designed on the 105
sequence of human P2Y₁₁ (Lecca and 106
Abbracchio 2008). Of note, a human GPR17 107
long splice variant encoding a receptor with a 108
28-amino acid longer NH₂ terminal (for a total 109
of 367 amino acids instead of 339) had been also 110
identified in very early studies aimed at discov- 111
ering new members of the chemokine receptor 112
family (in this respect, see also Sect. 2.1) 113
(Blasius et al. 1998). Genomic analysis revealed 114
a three-exon structure of the hGPR17 gene, with 115
two putative open reading frames. While the 116
“short isoform” derives from splicing of the 117

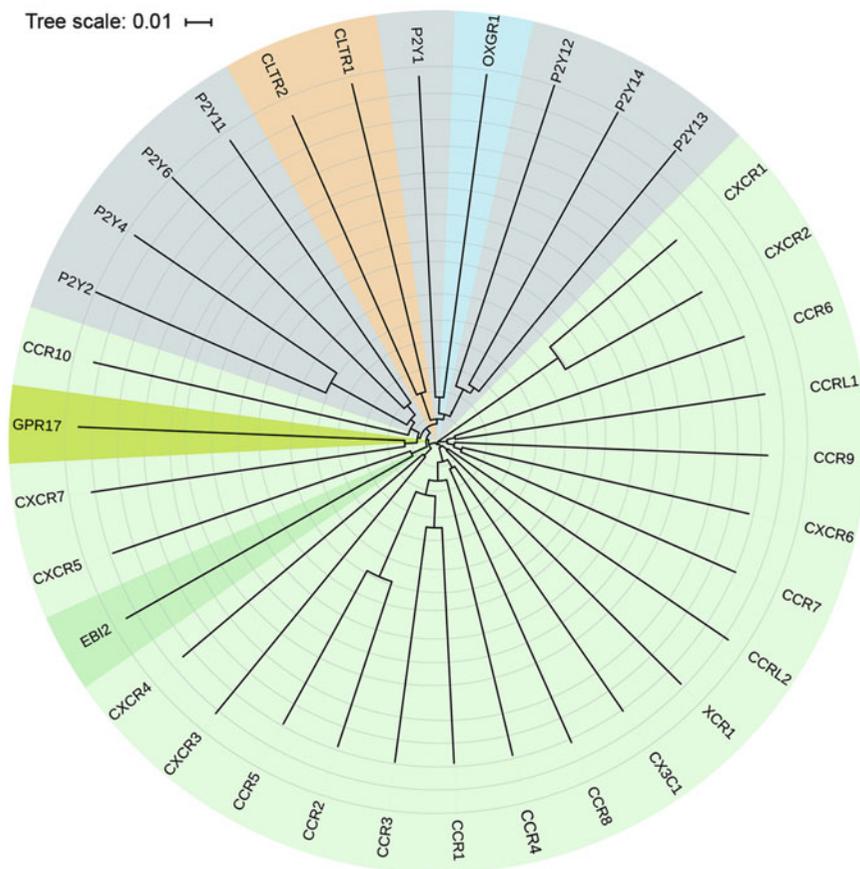


Fig. 1 Phylogenetic tree illustrating the relationship of GPR17 to selected structurally related class-A GPCRs. The evolutionary relationship analysis is based on a multiple sequence alignment performed on homologous GPCR sequences using TM-Coffee, a module of the T-Coffee package optimized for transmembrane proteins (Chang et al. 2012). Receptors belonging to the same

family are clustered according to the following color code: *grey* for purinergic receptors (P2Y), *orange* for cysteinyl-leukotriene receptors (CysLT), *light green* for chemokine receptors (CXCRn, CCRn, XCRn), *emerald green* for Epstein-Barr virus-induced G-protein coupled receptor 2 (EBI2), *sky blue* for 2-Oxoglutarate receptor 1 (OXGR1/GPR99), *apple green* for GPR17

118 second exon, the “long one” contains all three
 119 exons of the hGPR17 gene, leading to a transcript
 120 which is 1104 bp in length (Blasius et al. 1998;
 121 Pugliese et al. 2009). Interestingly, quantitative
 122 gene expression studies revealed that GPR17
 123 short isoform is expressed more abundantly in
 124 the brain than the long one (a tenfold increase),
 125 whereas the opposite was observed in heart and
 126 kidney. Pharmacological profile of the long iso-
 127 form also showed that some differences exist
 128 between the two GPR17 receptor isoforms
 129 (Pugliese et al. 2009; Benned-Jensen and
 130 Rosenkilde 2010).

In 2006, there were already papers reporting
 functional interactions between “classical” P2Y
 and CysLT receptors. For example, under some
 conditions, the CysLT1 receptor antagonist
 montelukast effectively antagonized the
 responses evoked by purinergic P2Y receptors
 Conversely, the P2Y₁₂ receptor had been reported
 to be also activated by LTE₄ (Paruchuri et al. 2009),
 suggesting the existence of some kind of ligand/
 receptor promiscuity between the P2Y and
 CysLT receptor families. On this basis, the iden-
 tification of GPR17 as the first dual member of
 the “purine receptor cluster” able to respond to

145 both purinergic and cysLT ligands (Ciana et al.
 146 2006) represented the demonstration of a further
 147 level of interaction between these two chemi-
 148 cally unrelated, but functionally interconnected,
 149 systems. Later studies extended the response pro-
 150 file of this receptor to other classes of endoge-
 151 nous “emergency” molecules connected to
 152 oxidative stress, neuroinflammation and
 153 neurodegeneration, i.e., oxysterols and chemo-
 154 kine stromal derived factor-1 (SDF-1)
 155 (Parravicini et al. 2016), further highlighting the
 156 promiscuous behaviour of GPR17. Of note, a
 157 phylogenetic analysis among structurally related
 158 class-A GPCRs (Parravicini et al. 2008, 2010,
 159 2016; Sensi et al. 2014), suggest that, besides
 160 P2Y and CysLT receptors, GPR17 holds a tight
 161 evolutionary relationship also with chemokine
 162 receptors and Epstein-Barr virus-induced
 163 G-protein coupled receptor 2 (EBI2) (Fig. 1).
 164 The possibility that GPR17 can be activated by
 165 diverse family of ligands underlines the rele-
 166 vance of a new transversal signaling mechanism
 167 that synchronizes all these emergency molecules
 168 and their receptors under specific neurodegener-
 169 ative conditions. Such a high promiscuity in
 170 receptor behaviour is often found in receptors
 171 involved in immunological responses and may,
 172 at least in part, depend on GPR17 ability to form
 173 dimers with other related receptors, thus widen-
 174 ing the array of pharmacological responses (see
 175 also Sect. 2.2.1).

176 In subsequent years, new data have revealed a
 177 key role for GPR17 in oligodendrogenesis and
 178 myelination (Lecca et al. 2008; Chen et al. 2009).
 179 However, while some authors have provided evi-
 180 dence for a stimulatory role of GPR17 in the
 181 specification and maturation of oligodendrocyte
 182 precursor cells (OPCs), some others have pro-
 183 posed an inhibitory role. Here, we aim at
 184 analyzing all the available literature on GPR17
 185 in an attempt to provide an overview of the
 186 different biological and pharmacological data
 187 emerged from all these papers.

2 GPR17 Characterization 188

2.1 Receptor Structure, Amino Acid Homology with Phylogenetically Related GPCRs and Binding Sites 189-191

GPR17 displays the typical 7-transmembrane (TM) domain topology of GPCRs, with an amino acid identity with the known P2Y and CysLT receptors between 21 and 48% (Abbracchio et al. 2006; Lecca and Abbracchio 2008). All these receptors show partial or complete conservation of a H-X-X-R/K amino acid motif in TM6 (and also of a K-E-X-X-L motif in TM7, in the case of P2Y₁₂, P2Y₁₃, P2Y₁₄) that are important for ligand recognition and have been proposed to represent specific molecular signatures for these receptors (Lecca and Abbracchio 2008) (see also below). Homology Modelling (HM) studies combined with other *in silico* tools have been performed to raise hypothesis on the molecular interaction between GPR17 and its putative endogenous ligands (Parravicini et al. 2008, 2010; Calleri et al. 2010), as well as to identify new potential ligands (Eberini et al. 2011) (see also Sect. 2.2.2). In these studies, *in silico* receptor modeling was performed using different templates, according to the progressive availability of new high-resolution GPCR structures. Starting from bovine rhodopsin, that, since 2000, has represented for many years the only atomistic scaffold for the structural investigation of GPCRs, the recent explosion in the resolution of GPCR crystal structures has given access to detailed structural information previously unavailable, allowing the construction of more and more accurate GPR17 models (Fig. 2).

For example, in 2010, crystallization of human CXCR4 (Wu et al. 2010) provided a significant improvement in the accuracy of GPR17 modelling because this structure enabled to reliably describe the extracellular regions of the receptor, especially extracellular loop 2 (ECL2) and the disulphide bridge linking the N-terminal to ECL3, known to be crucial in ligand molecular recognition (Wheatley et al. 2012), but for which none of the earlier templates was suitable. More



Fig. 2 Three-dimensional homology model of the human GPR17. Topological domains are represented as ribbon and coloured according to their secondary structure: *magenta* for alpha-helices; *yellow* for beta-sheets; *white* for loops and grey for turns

233 recently, modelling of GPR17 has been further
 234 improved thanks to the atomic resolution of the
 235 structures of two members of the P2Y receptor
 236 family: the human P2Y₁₂ (Zhang et al. 2014) and
 237 P2Y₁ (Zhang et al. 2015).

238 Globally, all the *in silico* results on the short
 239 isoform of GPR17 suggest that its nucleotide
 240 binding pocket is similar to that described for
 241 the other P2Y receptors (including the TM6
 242 HXXR/K motif designated to accommodate the
 243 phosphate moieties of nucleotide ligands
 244 (Parravicini et al. 2008; Jiang et al. 1997), and
 245 that this site is also shared by other small
 246 molecules identified as GPR17 ligands, including
 247 oxysterols and new synthetic compounds (Sensi
 248 et al. 2014; Eberini et al. 2011). According to
 249 these studies, also the nucleotide-derivative
 250 antagonist cangrelor binds to the same binding
 251 pocket, behaving as a competitive antagonist for
 252 orthosteric ligands.

253 In both P2Y and CysLT1 and CysLT2
 254 receptors, ligand binding is critically dependent
 255 on the basic Arginine residue belonging to the

conserved TM6 motif (Parravicini et al. 2010; 256
 Temporini et al. 2009). Computational studies 257
 suggested that this also holds true for Arg255 of 258
 GPR17 (Parravicini et al. 2008). To assess the 259
 actual role of this residue in receptor binding, this 260
 basic amino acid was mutated to isoleucine, and 261
 an *in silico* mutant GPR17 receptor (R255I) was 262
 generated. Using steered molecular dynamics 263
 simulations (SMD), forced unbinding of the 264
 endogenous ligand UDP from both wild type 265
 (WT) and R255I receptor models of GPR17 266
 was modeled *in silico*. The energy required to 267
 unbind UDP from the nucleotide binding pocket 268
 of GPR17 was higher for the WT than for the 269
 mutated R255I receptor, and the exit of the 270
 ligand from its intracellular cavities occurred 271
 earlier in the R255I model compared to the WT 272
 receptor. Generation and expression of the 273
 mutated receptor in 1321N1 cells confirmed 274
 also *in vitro* that the mutation was not silent 275
 (Calleri et al. 2010). 276

Besides the orthosteric binding site, *in silico* 277
 studies suggested that GPR17 also possesses an 278
 “accessory” binding site in a region formed by 279
 extracellular loops ECL2, ECL3 and the N- 280
 terminal, which also faces the extracellular space. 281
 This external accessory binding site could guide 282
 small agonist ligands to the deeper principal 283
 binding site in a multistep mechanism of activa- 284
 tion. Thanks to further *in silico* investigations, 285
 that showed the possibility of GPR17 to be 286
 stimulated also by a large peptide ligand such 287
 as SDF-1, the extracellular recognition site has 288
 been extensively characterized and GPR17 rec- 289
 ognition mechanism has been compared to those 290
 of some peptide receptors (Parravicini et al. 291
 2016), in which a two-step model of receptor 292
 activation, passing through both an extracellular 293
 and a TM binding site, has been proposed 294
 (Rajagopalan and Rajarathnam 2004). 295

296 Due to the intrinsic inaccuracy of the standard
 297 template-based HM techniques in predicting
 298 conformations of highly flexible and unaligned
 299 loop sequences in absence of adequate templates,
 300 no modeling studies are yet available for the long
 301 isoform of GPR17. Nevertheless, we can specu-
 302 late that the N-terminal may influence the bind-
 303 ing affinity of nucleotide agonists via a different

304 conformation of the external accessory binding
305 site, resulting in a slightly different pharmaco-
306 logical profile of the long isoform with respect to
307 the short one (Pugliese et al. 2009).

308 **2.2 Pharmacology and Signaling** 309 **Pathways**

310 **2.2.1 Putative Endogenous Ligands** 313 **and Transduction Systems**

314 In the initial studies, only GPR17 short isoform
315 has been characterized, and both the human, rat
316 (Ciana et al. 2006) and the previously unidenti-
317 fied mouse GPR17 receptors (Lecca and Ceruti
318 2008) were shown to respond to UDP,
319 UDP-glucose, UDP-galactose and LTC₄ and
320 LTD₄, with comparable profiles that were highly
321 conserved across species (the chemical structures
322 of UDP-glucose and LTD₄ are reported in
323 Fig. 3). Interestingly, the concentration ranges
324 at which uracil nucleotides and cysLTs activated
325 GPR17 (i.e., μ M and nM ranges, respectively)
326 were fully consistent with those necessary for
327 these endogenous ligands to activate their
328 already known cognate P2Y and CysLT
329 receptors (Abbracchio et al. 2006; Brink et al.
330 2003). Very similar agonist responses were
331 detected in a number of different cell lines
332 (1321N1, CHO, COS-7, HEK-293 cells). The
333 1321N1 cells was the most appropriate cells to
334 test GPR17 responses, since they are one of the
335 few cell lines that do not endogenously express
336 any functional purinergic or CysLT receptors.
337 Responses were highly specific, since no
338 response was ever found in cells transfected
339 with the empty vector. The antagonist response
340 profile of GPR17 was also rather peculiar. Acti-
341 vation by uracil nucleotides was reversed by
342 some typical purinergic antagonists like the
343 P2Y₁ antagonist MRS2179 or the P2Y₁₂ antago-
344 nist cangrelor (Fig. 3). Conversely, responses to
345 cysLTs were inhibited by typical CysLT receptor
346 antagonists like the already marketed drug
347 montelukast (Fig. 3) and pranlukast (Ciana
348 et al. 2006).

349 GPR17 responses were demonstrated by using
350 [³⁵S]GTP γ S binding, a typical functional assay

for agonists acting at G_i coupled receptors 351
(Kotani et al. 2001; Marteau et al. 2003; 352
Fumagalli et al. 2004). Under some 353
circumstances, activation of GPR17 could also 354
increase intracellular calcium levels via a phos- 355
pholipase C mediated pathway; however, this 356
effect occurred only in about 30% of 1321N1 357
transfected cells, suggesting preferential cou- 358
pling to the adenylyl cyclase pathway (Ciana 359
et al. 2006). In subsequent studies, GPR17 pecu- 360
liar profile was confirmed in many other distinct 361
assays independently performed in different 362
laboratories. Concentration-dependent inhibition 363
of forskolin-stimulated adenylyl cyclase was also 364
shown in oligodendrocyte precursor cells 365
(OPCs), the cell type natively expressing 366
GPR17 at highest levels (Fumagalli et al. 367
2011a) (Table 1). Inhibition of cAMP was fully 368
counteracted by the same antagonists utilized in 369
the [³⁵S]GTP γ S binding. In 2009, another paper 370
appeared where, in GPR17 expressing 1321N1 371
cells, enhancement of an outward rectifying K⁺ 372
current was shown upon addition of either uracil 373
nucleotides or cysLTs (Pugliese et al. 2009). 374
These effects were blocked by MRS2179, 375
cangrelor and montelukast. A few years later, 376
these same authors showed that similar delayed 377
rectifier K⁺ currents were stimulated in a concen- 378
tration dependent manner by GPR17 ligands in a 379
subpopulation of OPCs and 380
pre-oligodendrocytes, but not in terminally 381
mature cells, fully in line with the transient 382
expression of GPR17 during OPC specification 383
(in this respect, see also Sect. 3.1.1) (Coppi et al. 384
2013). This effect was blocked by MRS2179 and 385
cangrelor and sensitive to the K⁺ channel blocker 386
tetraethyl-ammonium. Importantly, the latter 387
also inhibited oligodendrocyte maturation, to 388
support previous literature data on the impor- 389
tance of these currents in OPC differentiation. 390

Fewer studies are available on hGPR17 long 391
isoform. In the electrophysiological study 392
already mentioned above, no significant 393
differences between the short and long isoforms 394
were detected (Pugliese et al. 2009). In 2010, 395
Bened-Jensen and Rosenkilde independently 396
confirmed the ability of heterologously 397
expressed GPR17 to respond to uracil 398

t.1 **Table 1** GPR17 signaling in native systems and relevant pharmacology

t.2	Tested ligand	Type of cells	Signaling	EC ₅₀ /IC ₅₀ values	Reference
t.3	UDP-glucose	Rat primary OPCs	Inhibition of cAMP production	IC ₅₀ : 424.7 ± 125 nM	Fumagalli et al. (2011a)
t.4		Rat primary OPCs	Outward K ⁺ currents	EC ₅₀ : 4.6 μM	Coppi et al. (2013)
t.5		Rat primary OPCs	Association to GRK5	N.A.	Daniele et al. (2014)
t.6		Rat primary OPCs	β-arrestin dependent ERK1/2 activation	N.A.	Daniele et al. (2014)
t.7		Oli-neu cells	Clathrin-mediated endocytosis	N.A.	Fratangeli et al. (2013)
t.8		PC12 cells	ERK1/2 and p38 phosphorylation	N.A.	Daniele et al. (2010)
t.9	UDP LTD ₄	Rat primary OPCs	Inhibition of cAMP production	IC ₅₀ : 1.29 ± 0.07 μM	(Fumagalli et al. 2011a)
t.10		Rat primary OPCs	Inhibition of cAMP production	IC ₅₀ : 2.85 ± 0.89 nM	Fumagalli et al. (2011a)
t.11		Rat primary OPCs	Association to GRK2	N.A.	Daniele et al. (2014)
t.12		Rat primary OPCs	CREB activation	N.A.	Daniele et al. (2014)
t.13		PC12 cells	ERK1/2 and p38 phosphorylation	N.A.	Daniele et al. (2010)
t.14		Oli-neu cells	Clathrin-mediated endocytosis	N.A.	Fratangeli et al. (2013)
t.15	LTE ₄	Rat primary OPCs	Inhibition of cAMP production	IC ₅₀ : 51.8 ± 6.6 pM	Fumagalli et al. (2011a)
t.16	MDL29,951	Rat primary OPCs	Inhibition of cAMP production	N.A.	Hennen et al. (2013)
t.17		Rat primary OPCs	Ca ²⁺ _i increase	N.A.	Hennen et al. (2013)

t.18 N.A.: Not available

407 2008) and in frontal affinity chromatography-
 408 mass spectrometry (FAC-MS) studies (Calleri
 409 et al. 2010; Temporini et al. 2009). Much lower
 410 potencies to uracil nucleotides were observed for
 411 the long receptor isoform, with a 50–170 fold
 412 increase in EC₅₀ (Bened-Jensen and Rosenkilde
 413 2010). Moreover, no responses to cysLTs were
 414 detected either on the short or long isoform, nor
 415 were cysLTs able to induce GPR17 removal
 416 from the membrane and internalization. This is
 417 in contrast with subsequent studies on cells
 418 natively expressing the receptor (Fratangeli
 419 et al. 2013) (see below). This may depend on
 420 differences in the conformation/ability of the
 421 recombinant receptor to respond to agonists com-
 422 pared to the native one, as well as on the fact that,
 423 in heterologously expressing systems, constitu-
 424 tive activity of transfected receptors may

significantly alter ligand behavior (Kenakin 425
 2001; Im 2013) (see also Conclusions). In this 426
 respect, Benned-Jensen and Rosenkilde indeed 427
 reported a notable constitutive activation of 428
 recombinant GPR17 resulting in potent inhibi- 429
 tion of forskolin stimulated adenylyl cyclase in 430
 the absence of any endogenous ligand (Bened- 431
 Jensen and Rosenkilde 2010). 432

At the same time, another paper suggested 433
 GPR17 as a negative regulator of the CysLT1 434
 receptor (Maekawa et al. 2009). This effect was 435
 proposed to depend on the formation of a 436
 CysLT1-GPR17 heteromer, as suggested by 437
 co-immunoprecipitation studies in CHO cells. 438
 The interaction between GPR17 and CysLT1 439
 was further confirmed in primary human mono- 440
 cyte cells and in a rodent knock out GPR17 441
 model, thus extending to GPR17 the previously 442

443 reported interaction and promiscuity between
444 different members of the “purine receptor cluster”.
445 These data indicate that, besides working on
446 its own, GPR17 may also modify the function of
447 other related receptors by the formation of
448 heteromers.

449 In 2010, the first paper describing the
450 characteristics of GPR17 in a *native* system (rat
451 pheochromocytoma PC12 cells) was published
452 (Daniele et al. 2010) (Table 1). GPR17 was not
453 expressed in undifferentiated PC12 cells but was
454 specifically induced by a 10-day NGF treatment,
455 suggesting a role in the control of neuronal differentiation.
456 Both UDP-glucose and LTD₄
457 induced a significant pro-survival effect on
458 PC12 cells. By *in vitro* silencing experiments
459 with small interfering RNAs and by using receptor
460 antagonists, these effects were confirmed to
461 be mediated by the selective activation of
462 GPR17. In differentiated PC12 cells,
463 UDP-glucose and LTD₄ caused a significant
464 increase in extracellular signal-regulated kinases
465 1 and 2 (ERK1/2) phosphorylation. ERK activation
466 induced by the two agonists occurred with
467 different kinetics: LTD₄ induced a transient ERK
468 activation that returned to basal value within
469 120 min. In contrast, ERK phosphorylation
470 induced by UDP-glucose was maintained over
471 basal values for 120 min and the activation kinetics
472 appeared to be biphasic with two peaks, one
473 at 15 and the other one at 120 min. In addition,
474 incubation of cells with the purinergic antagonist
475 cangrelor completely counteracted UDP-glucose
476 effects at all tested incubation times (Daniele
477 et al. 2010). These data confirmed the responses
478 to uracil nucleotides and cysLTs already seen on
479 the recombinant receptors, and suggested, for the
480 first time, that endogenous GPR17 ligands can
481 couple to distinct G proteins and intracellular
482 pathways, a finding that was later confirmed by
483 other studies (Hennen et al. 2013; Daniele et al.
484 2014). The signaling pathways of native GPR17
485 are summarized in Table 1.

486 In 2011, in another study that independently
487 confirmed the purinergic component of GPR17,
488 Buccioni and coworkers (Buccioni et al. 2011)
489 exploited an innovative and non-radioactive

functional cAMP assay to monitor GPR17 activation
(and the effects of various ligands) through changes
in intracellular cAMP concentrations by using a mutant
form of Photinus pyralis luciferase into which a cAMP-
binding protein moiety had been inserted (GloSensor
cAMP reagent). In HEK293 cells stably transfected
with the GloSensor reagent, transient expression of
hGPR17 resulted in the appearance of highly specific
concentration-dependent responses to both UDP,
UDP-glucose and UDP-galactose and to a series of
UDP and ATP derivatives that behaved as either
agonist or antagonists, with EC₅₀ values that were
very similar to those obtained in parallel on [³⁵S]
GTPγS binding. In this system, cysLTs were not
tested, due to the high constitutive expression of
traditional CysLT receptors in the HEK293 cells
(Ciana et al. 2006; Buccioni et al. 2011).

2.2.2 GPR17 Non-conventional Ligands

In the last years, the increasing number of class-A
GPCR solved structures allowed the scientific
community to recognize some common features
that are crucial for their operability (Levit et al.
2014); however, these studies also revealed an
unexpected heterogeneity and complexity in
GPCR recognition, challenging the classical
pharmacology paradigms of the ‘monogamous’
interaction between a specific class of natural
ligands and a single GPCR (Haupt et al. 2013).
In line with the growing promiscuity of GPCRs,
ligand dependent transactivation has been
demonstrated for GPR17, already known as a
“dual” receptor: similarly to EB12 (Hannedouche
et al. 2011; Liu et al. 2011a), and the CXC
chemokine receptor 2 (CXCR2) (Raccosta et al.
2013). Specifically, it was shown that, GPR17
could act as a molecular target for oxysterols,
oxidized derivative of cholesterol that, in the
CNS, are involved in activities not strictly
associated with cholesterol metabolism. Of
note, these activities are particularly relevant
for neurodegenerative disorders, including
demyelinating (Raccosta et al. 2013; Garenc

537 et al. 2010). More in detail, three selected
538 oxysterols (27-Hydroxycholesterol, 7
539 α -Hydroxycholesterol and 22R-Hydroxycho-
540 lesterol) were tested in 1321N1 cells stably
541 expressing GPR17, showing that all the tested
542 compounds were able to stimulate GTP γ S bind-
543 ing, in a concentration-dependent manner, with
544 EC₅₀ values of 4.99 ± 0.78 nM, 0.70 ± 0.09 nM
545 and 0.21 ± 0.03 nM, for 27-Hydroxycholesterol,
546 7α -Hydroxycholesterol and 22R-Hydroxycho-
547 lesterol, respectively.

548 Stimulus of cell membranes with different
549 oxysterol concentrations after treatment with
550 the purinergic ligand UDP-glucose showed a
551 left-shift of the concentration-response curves
552 or an enhancement of their maximal [³⁵S]
553 GTP γ S binding stimulation, suggesting that
554 these ligands may cooperate under
555 neuroinflammatory conditions.

556 In parallel, the effect of different concentra-
557 tion of the GPR17 receptor antagonist cangrelor
558 on oxysterol-stimulated [³⁵S]GTP γ S binding was
559 evaluated, demonstrating that cangrelor can
560 counteract GPR17 activation by oxysterols
561 through a competitive mechanism, with IC₅₀
562 values in a sub-nM range. These results are also
563 in agreement with *in silico* data suggesting a
564 common orthosteric molecular recognition
565 mechanism for oxysterols and other small
566 GPR17 ligands, despite different local
567 arrangements in the TM binding site (Sensi
568 et al. 2014).

569 Among other non-conventional ligands, fur-
570 ther evidence showed that SDF-1, historically
571 known as the endogenous ligand for CXCR4
572 and CXCR7 receptors, is able to transactivate
573 GPR17 *in vitro*, specifically increasing the [³⁵S]
574 GTP γ S binding to membrane of GPR17-
575 expressing cells, with affinity constant values of
576 0.14 ± 0.03 nM. The effect of SDF-1 in
577 modulating GPR17 responses *in vitro* was further
578 assessed in primary OPC cultures natively
579 expressing GPR17. In this model, treatment
580 with physiological concentrations of SDF-1 sig-
581 nificantly increased the number of cells
582 expressing the Myelin Basic Protein MBP com-
583 pared to control, thus accelerating OPC differ-
584 entiation towards a mature phenotype. The specific

involvement of GPR17 in these effects was 585
unequivocally demonstrated by further 586
experiments showing that, in presence of the 587
GPR17 antagonist cangrelor, SDF-1 induced no 588
increases of either [³⁵S]GTP γ S binding to cell 589
membranes, or MBP-expression in OPC cultures. 590
Moreover, the mechanism by which GPR17 and 591
SDF-1 can directly interact to each other has 592
been predicted and extensively characterized *in* 593
silico through molecular modeling (Parravicini 594
et al. 2016). 595

596 These results are in line with literature data,
597 that propose a role of SDF-1 in orchestrating
598 OPC differentiation and maturation also via
599 CXCR4/CXCR7-axis (Li et al. 2012; Patel et al.
2010; Carbajal et al. 2011). 600

601 Interestingly, not only GPR17, CXCR4 and
602 CXCR7, but also others chemokine receptors,
603 like CXCR2, have demonstrated roles in
604 regulating OPCs. As previously mentioned (see
605 Sect. 2.1), besides sharing the same ligands,
606 GPR17 and chemokine receptors are phylogen-
607 etically related to each other, and all participate
608 to CNS reparative responses. This raises the
609 hypothesis that, under neurodegenerative demy-
610 elinating conditions, oxysterols and other
611 pro-inflammatory ligands, such as SDF-1, act as
612 non-conventional molecules with a transversal
613 regulatory role, representing a conserved,
614 “unspecific” signaling mechanism, by which
615 emergency molecules synchronize multiple
616 receptors involved in inflammatory/immune
617 responses.

2.2.3 New GPR17 Synthetic Ligands 628

619
620 In 2009 and 2010, two papers reported the devel-
621 opment of a new FAC-MS binding method for
622 the analysis of GPCRs (Calleri et al. 2010;
623 Temporini et al. 2009). In this assay, UDP was
624 found to bind to GPR17 with a Kd value of
625 1612.0 ± 708 nM that was very similar to the
626 Kd value (1140.0 nM) obtained by Ciana et al.
627 and Lecca et al. in the [³⁵S]GTP γ S-binding
628 (Ciana et al. 2006; Lecca and Ceruti 2008).
629 This paper also unveiled a number of previously
630 unreported GPR17 ligands, some of which were
631 able to increase [³⁵S]GTP γ S binding, with
632

633 potency values in the μM and sub-nM range. For
634 example, the ATP analogue 2-Phenylethynyladenosine-5'-monophosphate Compound N. 4
635 behaved as a very potent agonist with an EC_{50}
636 value of 36 pM. In contrast, other ligands (e.g.: N
637 ⁶-Benzoyl-2'-deoxyadenosine 3',5'-Bis phosphate, referred by the authors as Compound
638 N. 12) did not induce any increase in [³⁵S]
639 GTP γ S binding, but counteracted stimulation
640 induced by UDP-glucose with an antagonist profile and an affinity constant in the nM range
641 comparable to that reported for its analogue derivative MRS2179. Both the newly identified
642 agonists and antagonists displayed similar
643 behavior in the FAC-MS binding assay (Calleri
644 et al. 2010). A comparison between these data
645 and [³⁵S]GTP γ S binding results have been also
646 reported in a recent review article on GPR17
647 (Marucci et al. 2016).

652 In the same year, an advanced *in silico* HM
653 procedure combined with high-efficiency virtual
654 screening of more than 120,000 compounds from
655 the Asinex Platinum Collection (<http://www.asinex.com/>), a lead-like structural library, on
656 the modeled receptor led to the selection of
657 5 chemically diverse molecules (the ASINEX
658 compounds, see Fig. 3 for the chemical structure
659 of one representative compound,
660 2-[[5-(2-Methoxyphenyl)-4-(4-methoxyphenyl)-
661 4H-1,2,4-triazol-3-yl]thio]-N-phenylpro-
662 panamide, also referred as ASN 1), that were
663 completely unrelated to already known ligands.
664 These compounds were tested *in vitro* in the [³⁵S]
665 GTP γ S binding assay, revealing a sub-nM
666 potency for GPR17 (Eberini et al. 2011) (see
667 also below). None of these compounds could
668 have been expected 'a priori' to act on GPR17,
669 and all of them behaved as much more potent
670 ligands than GPR17 endogenous activators
671 (Eberini et al. 2011). Finally, in 2013,
672 MDL29,951 was reported as an additional small
673 molecule agonist at GPR17 (Hennen et al. 2013)
674 (Fig. 3). In a variety of different heterologous
675 expression systems, MDL29,951-stimulated
676 GPR17 engaged the entire set of intracellular
677 adaptor proteins for GPCRs: G proteins of the
678 G α i, G α s, and G α q subfamily, as well as
679 β -arrestins. This was visualized as alterations in

681 the concentrations of cyclic adenosine 681
682 monophosphate and inositol phosphate, 682
683 increased Ca^{2+} flux, phosphorylation of ERK1/
684 2, as well as multifaceted cell activation 684
685 recorded with label-free dynamic mass redistribution and impedance biosensors. pEC_{50} values
686 for MDL29,951 at GPR17 ranged between 5 and
687 8.80, depending upon the transfected cell type
688 and the used read out. MDL29,951-stimulated
689 GPR17 effects were counteracted in a
690 concentration-dependent manner by pranlukast
691 and, to a lesser extent, by montelukast. This is
692 fully in line with the activities of these
693 antagonists on recombinant GPR17 in previous
694 studies, in which pranlukast was significantly
695 more potent than montelukast in antagonizing
696 LTD₄-stimulation of GPR17 (Ciana et al.
697 2006). In OPCs, MDL29,951 rapidly mobilized
698 intracellular Ca^{2+} in a concentration-dependent
699 manner and engaged both G α i and G α q, but not
700 G α s signaling pathways, further suggesting
701 differences in GPR17 responses between
702 transfected and native systems (see also
703 Conclusions). This is at variance from previous
704 studies reporting G α i coupling and decreases of
705 intracellular cAMP as a primary transduction
706 pathway of GPR17 in OPCs (Daniele et al.
707 2014; Fumagalli et al. 2011b). However, it has
708 to be emphasized that, despite being selective for
709 GPR17 inside the "purine receptor cluster"
710 (Hennen et al. 2013), MDL29,951 also significantly
711 interacts with the glycinergic site of the
712 glutamate NMDA receptor (Salituro et al. 1992).
713 This may be at the basis of the ability of
714 MDL29,951 to activate multiple signaling
715 pathways in both transfected cells and in OPCs,
716 and of the data reported for this compound on
717 myelination (see also Sect. 3.1.1). 718

2.2.4 Agonist-Induced Desensitization and Internalization 729 720

723 In 2011, the first complete agonist-induced
724 GPR17 desensitization/resensitization study was
725 published (Daniele et al. 2011). By using [³⁵S]
726 GTP γ S binding and cAMP measurements in
727 1321N1 cells expressing hGPR17, both
728 UDP-glucose and LTD₄ were shown to induce a
729 time- and concentration-dependent loss of 729

730 GPR17 response (homologous desensitization).
731 GPR17 homologous desensitization was
732 accompanied by internalization of receptors
733 inside cells, as assessed by biotin labeling of
734 cell surface receptors. Desensitization occurred
735 in a time-dependent manner, with similar kinet-
736 ics for both agonists. Upon agonist removal,
737 receptor resensitization occurred with the typical
738 kinetics of GPCRs. Finally, activation of GPR17
739 by UDP-glucose induced a partial heterologous
740 desensitization of LTD₄-mediated responses (but
741 not *vice versa*), suggesting that nucleotides have
742 a hierarchy in producing desensitizing signals.

743 The pattern of GPR17 desensitization and
744 internalization was fully confirmed and further
745 expanded in differentiated oligodendroglial
746 Oli-neu cells that natively express GPR17
747 (Fratangeli et al. 2013) (Table 1). Agonist-
748 induced internalization, intracellular trafficking
749 and membrane recycling of GPR17 were
750 analyzed by biochemical and immunofluores-
751 cence assays using an *ad hoc*-developed new
752 antibody against the extracellular N-terminal of
753 GPR17. Both UDP-glucose and LTD₄ increased
754 GPR17 internalization, although with different
755 efficiency. At early time points, internalized
756 GPR17 co-localized with transferrin receptor,
757 whereas at later times it partially co-localized
758 with the lysosomal marker Lamp1, suggesting
759 that a portion of GPR17 is targeted to lysosomes
760 upon ligand binding. Internalization of GPR17
761 occurred via clathrin-dependent endocytosis
762 (Fratangeli et al. 2013). Analysis of receptor
763 recycling and degradation demonstrated that a
764 significant fraction of GPR17 is recycled to the
765 cell surface. These results provided the first data
766 on the agonist-induced trafficking of native
767 GPR17 in oligodendroglial cells and may have
768 implications in fine-tuning cell responses to
769 demyelinating and inflammatory conditions
770 when these ligands accumulate at lesion sites
771 (see also Sect. 3.1.2). More recently, GPR17
772 downregulation by uracil nucleotides and cysLTs
773 was confirmed in primary cultured OPCs, and the
774 role of the GRK/ β -arrestin machinery in receptor
775 desensitization and intracellular signaling was
776 also extensively investigated (Daniele et al.
777 2014). It was shown that, following OPCs

778 treatment with the two classes of purinergic and 778
779 cysLT ligands, different GRK isoforms were 779
780 recruited. Specifically, cysLT-mediated GPR17 780
781 desensitization mainly involved GRK2 via a G 781
782 protein-dependent mechanism (Daniele et al. 782
783 2014). This kinase promoted transient binding 783
784 of the receptor to β -arrestins, rapid ERK phos- 784
785 phorylation and sustained nuclear CREB activa- 785
786 tion. Furthermore, GRK2, whose expression 786
787 paralleled that of the receptor during the differ- 787
788 entiation process, was required for cysLT- 788
789 mediated OPCs maturation (see also Sect. 3.2.). 789
790 On the other hand, purinergic ligands exclusively 790
791 recruited GRK5 via a G protein- 791
792 independent/ β -arrestin-dependent mechanism. 792
793 This kinase induced a stable association between 793
794 the receptor and β -arrestin, followed by slower 794
795 and sustained ERK stimulation and marginal 795
796 CREB activation (Daniele et al. 2014). These 796
797 results show that, through activation of GPR17 797
798 and recruitment of specific GRK isoforms, 798
799 purinergic and cysLT ligands engage distinct 799
800 intracellular pathways. 800

801 Recently GPR17 desensitization (and its rela- 801
802 tionship to terminal OPC maturation) has been 802
803 linked to activation of mTOR (the “mammalian 803
804 target of rapamycin”), which has long been 804
805 known to be involved in myelination. During 805
806 OPC differentiation, mTOR regulates 806
807 GRK-mediated desensitization of GPR17 by pro- 807
808 moting the nuclear translocation of the ubiquitin 808
809 ligase MDM2, which had been previously only 809
810 involved in cancer via regulation of p53 activity 810
811 and now emerges as a new interesting actor in 811
812 oligodendrogenesis (Fumagalli et al. 2015). Spe- 812
813 cifically, treatment of OPCs with either the 813
814 mTOR inhibitor rapamycin, or with nutlin-3, a 814
815 small molecule inhibitor of Mdm2-p53 815
816 interactions, was shown to keep MDM2 in the 816
817 cytosol, where it could bind to GRK2 and sustain 817
818 its degradation, thus impairing the physiological 818
819 desensitization of GPR17 (Fumagalli et al. 819
820 2015). Important, prevention of GPR17 desensi- 820
821 tization was also associated to a defect of OPC 821
822 maturation, confirming that aberrantly elevated 822
823 GPR17 levels in late stage OPCs blocks cells at 823
824 immature stages (Fumagalli et al. 2015). 824

In another study, GPR17 plasma membrane recycling and stability was shown to be also modulated by SNX27, a recently identified protein of the endosome-associated retromer complex, whose functions in oligodendrocytes had never been studied. It was found that, after endocytosis, GPR17 is either sorted into lysosomes for degradation or recycled to the plasma membrane. Balance between degradation and recycling was important for modulation of receptor levels at the cell surface, and thus for the silencing or maintenance of GPR17-signaling pathways, that, in turn, affect OPC differentiation (see also Sect. 3.2). The endocytic trafficking of GPR17 was mediated by interaction of SNX27 with a type I PDZ-binding motif located at the C-terminus of the receptor. Of note, SNX27 knock-down reduced GPR17 plasma membrane recycling in differentiating oligodendrocytes while accelerating terminal cell maturation. Interestingly, trisomy-linked down-regulation of SNX27 in the brain of Ts65Dn mice, a model of Down syndrome, correlated with a dysfunction in GPR17⁺ cells and an increase in mature oligodendrocytes, which, however, failed in reaching full maturation, eventually leading to hypomyelination (Meraviglia et al. 2016). Thus, disruption of SNX27/GPR17 interactions leading to alterations of GPR17 membrane trafficking might contribute to pathological oligodendrocyte differentiation and myelination defects present in Down syndrome (Meraviglia et al. 2016).

3 Role of GPR17 in Central Nervous System Pathophysiology

3.1 GPR17 Specific Roles in Oligodendroglial Functions and Myelination

3.1.1 Physiological Roles

In the healthy intact brain, GPR17 expression is predominantly in oligodendrocyte (OL) cells. The very first demonstration that, in the adult

brain, GPR17 is highly expressed by a sub-population of endogenous quiescent parenchymal OPCs dates back to 2008 (Lecca et al. 2008) and has sparked a lot of interest on GPR17 role in CNS myelination. Specifically, GPR17 was shown to be present in ramified early neural cell precursors dispersed throughout brain's gray and white matter that also positively stained for typical early OPC markers. Since then, increasing evidence has progressively accumulated to show a pivotal role of GPR17 in OPC maturation, with different and apparently paradoxical effects during different phases of the maturation process (Chen et al. 2009; Fumagalli et al. 2011a) (see also below).

In vitro studies on purified rat postnatal OPC cultures showed that GPR17 expression coincides with a specific temporal window of the OL differentiation process. It covers two distinct phases: a first phase, during which early differentiation markers like NG2, A2B5, PDGF receptor-alpha and the immature PLP isoform DM-20 are still present (early stage 2 OPCs in Fig. 4), and a subsequent phase characterized by more ramified, still immature pre-oligodendrocytes (stages 3 and 4 in Fig. 4), where NG2 has been downregulated and more advanced markers like O4, O1 and the proteolipid myelin protein PLP are present (Fumagalli et al. 2011a). Based on these data, GPR17 is currently utilized by other independent scientists to specifically label pre-immature OLs at these two transition stages (Mitew et al. 2013; Nakatani et al. 2013; Crociara et al. 2013; Ferrara et al. 2016).

Of note, GPR17 expression progressively increases during the transition of OPCs to pre-OLs (when it is maximally expressed in cellular processes), but is then gradually silenced and never found in fully morphologically mature OLs (Fumagalli et al. 2011a) (see Fig. 4). Accordingly, *in vivo*, GPR17 is present in a subset of NG2/Olig2-positive OPCs expressing the first myelin proteins, but not in more mature cells expressing myelin basic protein (MBP). Also during rodent brain development, GPR17 expression in OPCs precedes myelin production. Interestingly, GPR17 immunoreactivity appears first

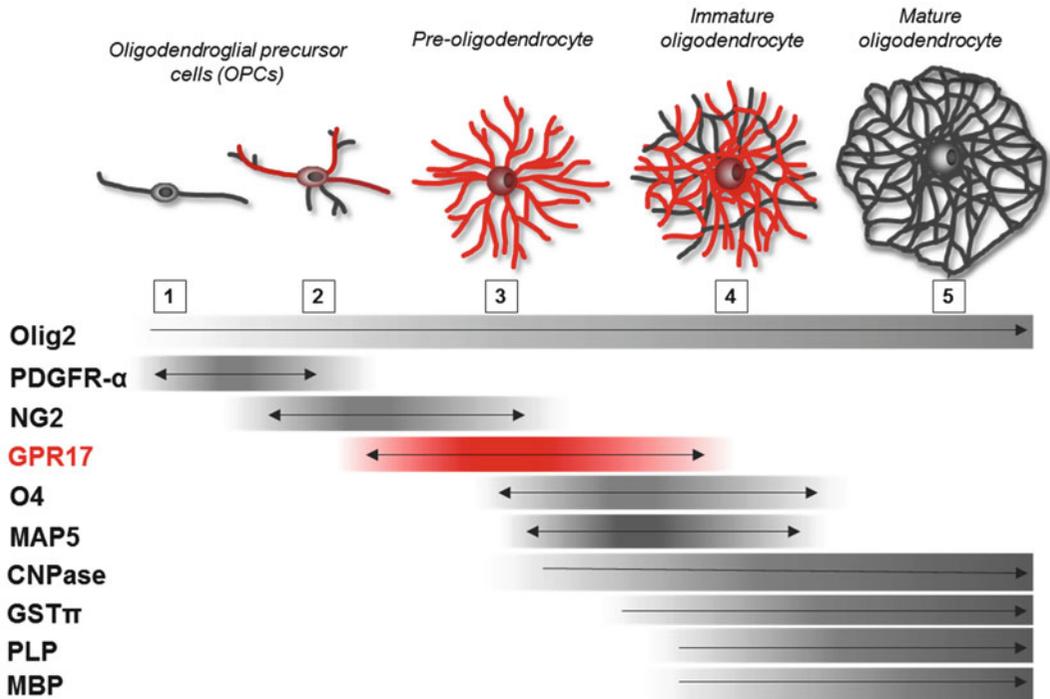


Fig. 4 Transient GPR17 expression during oligodendroglial differentiation. The expression pattern of GPR17 (in red) during oligodendroglial differentiation is shown in parallel to other known oligodendroglial markers (other colours). Progressive differentiation stages are indicated with numbers from 1 to 5. From a functional

point of view, GPR17 exerts opposing stage-specific roles: a positive role for differentiation in early OPCs and a negative function for OL maturation in late OPCs. In late OPCs, gradual silencing of GPR17 is needed to allow OPCs to complete their maturation (see text for more details)

918 in the cell body, partially coinciding with
919 markers of the Golgi apparatus, and then gradu-
920 ally extends to cellular processes (Boda et al.
921 2011). Early after birth, the expression of the
922 receptor is low, but progressively expands to
923 cover the 80% of OPCs at the end of the third
924 week of life. Afterwards, GPR17 is down-
925 regulated while myelination proceeds (Boda
926 et al. 2011).

927 The transient nature of GPR17 expression in
928 OPCs suggests that the receptor may display
929 stage-specific roles during OL development.
930 Intriguingly, as already reported for the
931 Wnt/ β -catenin pathway (Fancy et al. 2009; Ye
932 et al. 2009) and more recently proposed for the
933 transcription factor Olig2 (Mei et al. 2013),
934 GPR17 exhibits opposing functions on OL dif-
935 ferentiation in relation to its expression stage. In
936 cultured cortical postnatal rat OPCs, early recep-
937 tor obliteration with small interfering RNAs

profoundly affected their ability to generate 938
939 mature OLs, suggesting that cells are retained at
940 a less differentiated stage (Fumagalli et al.
941 2011a) (Fig. 4). Although the molecular
942 mechanisms at the basis of these events have
943 not been yet investigated, these data highlight a
944 pivotal role of GPR17 in the initial phases of the
945 differentiation process. They support the hypoth-
946 esis that, at these stages, GPR17 may be impor-
947 tant to keep cells at an immature state which
948 may, in turn, be necessary to prepare them for
949 myelination (Fumagalli et al. 2011a). In contrast,
950 cultured cortical progenitors from GPR17 knock-
951 out E15.5 mouse embryos differentiated earlier
952 toward mature OLs compared to control cells
953 (Chen et al. 2009). The reasons for these
954 discrepancies remain unknown, although it may
955 be hypothesized that compensatory mechanisms
956 are activated as a result of early embryonic
957 GPR17 knock out. Of course, the generation of

958 conditional transgenic mice in which deletion of
959 GPR17 in OPCs could be induced under con-
960 trolled conditions at specific ages will help
961 clarifying this issue.

962 Lecca and coworkers also clearly showed that
963 GPR17 is no longer present in morphologically
964 mature MBP-positive cells (Lecca et al. 2008),
965 raising for the first time the possibility that loss of
966 GPR17 at advanced differentiation stages is a
967 prerequisite to allow cells to complete terminal
968 maturation. Subsequent *in vivo* data showed that
969 myelinogenesis is indeed defective in transgenic
970 mice overexpressing GPR17 under the promoter
971 of 2',3'-Cyclic-nucleotide 3'-phosphodiesterase
972 (CNase), a relatively advanced OL marker
973 (Chen et al. 2009). These animals exhibited
974 motor disabilities, tremors and precocious death
975 within the second week of life. The forced and
976 un-timely expression of GPR17 at a maturation
977 stage (i.e., in CNase⁺ cells), at which GPR17 is
978 normally already downregulated, might have cre-
979 ated conflicting signals leading to defective ter-
980 minal maturation. Thus, interference with the
981 stage-restricted expression of GPR17 resulting
982 in un-programmed receptor expression in late
983 OPCs completely alters the differentiation pro-
984 gram of these cells. This hypothesis is fully in
985 line with the demonstration that OPCs
986 incorporating a vector for GPR17 over-
987 expression maintained an immature morphologi-
988 cal phenotype and never expressed the mature
989 marker CNase, and with data showing that,
990 under conditions where terminal OPC maturation
991 is impaired, such as demyelinating diseases (see
992 Sect. 3.1.2) or treatment with the mTOR inhibi-
993 tor rapamycin, that reduces OPC maturation,
994 GPR17 is markedly up-regulated (Fumagalli
995 et al. 2015; Tyler et al. 2011).

996 Both intrinsic and extrinsic mechanisms could
997 contribute to GPR17 stage-specific functions
998 during oligodendrocyte differentiation. GPR17
999 can be extrinsically regulated by physiological
1000 ligands accumulating in the extracellular milieu:
1001 activation of early OPCs (stages 2 and 3 in Fig. 4)
1002 with GPR17 endogenous putative ligands (i.e.,
1003 UDP-glucose or LTD₄) indeed promoted conver-
1004 sion to more mature cells expressing myelin

1005 markers (Lecca et al. 2008; Fumagalli et al. 1005
1006 2011a; Ceruti et al. 2011). Consistent with these 1006
1007 data, GPR17 antagonists like cangrelor (Fig. 3) 1007
1008 delayed the ability to generate mature cells 1008
1009 (Lecca et al. 2008; Fumagalli et al. 2011a), 1009
1010 suggesting that GPR17 endogenous ligands are 1010
1011 basally released in culture and are responsible for 1011
1012 the observed spontaneous OPC *in vitro* matura- 1012
1013 tion. In another independent study, while not 1013
1014 modifying the potential of adult multipotent neu- 1014
1015 ral stem cells, montelukast, which also acts as a 1015
1016 GPR17 antagonist (Ciana et al. 2006; Benned- 1016
1017 Jensen and Rosenkilde 2010; Lecca et al. 2008), 1017
1018 markedly increased their proliferation rate, 1018
1019 suggesting that GPR17 antagonism induces 1019
1020 retention of cells at a more undifferentiated 1020
1021 stage (Huber et al. 2011). 1021

1022 As already mentioned, besides cAMP inhibi- 1022
1023 tion, GPR17 has been also shown to specifically 1023
1024 mediate activation of delayed rectifier K⁺ 1024
1025 currents (Table 1) in a sub-population of OPCs 1025
1026 and O4⁺ pre-OLs, but not in mature OLs. This 1026
1027 effect was shown to contribute to the terminal 1027
1028 maturation of OPCs and to their migratory 1028
1029 abilities. 1029

1030 In contrast with the above studies, 1030
1031 MDL29,951, the new putative GPR17 agonist 1031
1032 mentioned above, was reported to inhibit, rather 1032
1033 than stimulate, OL maturation (Hennen et al. 1033
1034 2013). However, it is worth to note that, due to 1034
1035 the *transient* expression of GPR17 in culture, the 1035
1036 timing of OPC manipulation and treatment is 1036
1037 crucial for obtaining comparable results. On the 1037
1038 other hand, as already mentioned, MDL29,951 is 1038
1039 not a selective ligand for this receptor, and inde- 1039
1040 pendent effects could be due to its antagonistic 1040
1041 activity at the glycinergic site of the glutamate 1041
1042 NMDA receptor (Salituro et al. 1992), which has 1042
1043 been indeed reported to promote OPC differenti- 1043
1044 ation (Li et al. 2013). 1044

1045 Globally, these findings suggest that GPR17 1045
1046 exerts opposing stage-specific roles: a positive 1046
1047 role for differentiation in early OPCs and a nega- 1047
1048 tive function for OL maturation in late OPCs. 1048
1049 They also suggest that, in late OPCs, physiologi- 1049
1050 cal GPR17 silencing is needed to allow cells to 1050
1051 complete their maturation program. The latter 1051

1052 may occur via either GPR17 desensitization/
 1053 internalization by endogenous agonists or by
 1054 GPR17-mediated engagement of intracellular
 1055 pathways culminating in nuclear events, or
 1056 both. Blockade of GPR17 mRNA translation
 1057 into the receptor protein a specific microRNA
 1058 has been also recently reported to contribute to
 1059 GPR17 regulation during OPC maturation
 1060 (Lecca et al. 2016).

1063 3.1.2 Dysregulation in Demyelinating 1062 Neurodegenerative Diseases

1065 The demonstration that levels of endogenous
 1066 nucleotides and cysLTs are massively increased
 1067 upon CNS trauma and ischemia and their
 1068 hypothesized roles as danger signals after injury
 1069 (Davalos et al. 2005; Haynes et al. 2006) has
 1070 raised the hypothesis that GPR17 may act as a
 1071 crucial mediator of reactivity to acute injury.
 1072 While physiologically GPR17 is mostly an
 1073 oligodendroglial receptor, after acute injury,
 1074 GPR17 is sequentially induced in dying neurons
 1075 inside and at the borders of the ischemic/trau-
 1076 matic lesion, in infiltrating microglia/
 1077 macrophages and in activated parenchymal
 1078 OPCs in the lesion's surrounding areas, with
 1079 similar expression patterns in different models
 1080 of pathology. In more detail, in both rats and
 1081 mice, 24 h after permanent middle cerebral artery
 1082 occlusion (MCAo), GPR17 is up-regulated in
 1083 neurons damaged by the ischemic insult inside
 1084 the ischemic core (Ciana et al. 2006; Lecca et al.
 1085 2008). When the penumbra area is well visible
 1086 and most of the neurons in the core are dead,
 1087 GPR17 appears on highly activated microglia
 1088 and blood-borne macrophages at the borders of
 1089 the lesion (Lecca et al. 2008). This has been
 1090 independently confirmed to also occur in a tran-
 1091 sient MCAo rodent model, where the number of
 1092 GPR17 expressing cells was significantly
 1093 upregulated in two distinct phases, 24 h and
 1094 7 days after reperfusion, consistent with an
 1095 early acute neuronal injury followed by a late
 1096 microgliosis (Zhao et al. 2012). It is known that
 1097 OPCs are extremely sensitive to the pathophysi-
 1098 ological state of the brain, and that they react to
 1099 many different types of experimentally induced
 1100 insults. Starting from 72 h after the insult, in the

regions surrounding the ischemic area and in the 1101
 ipsilateral corpus striatum of MCAo mice, a 1102
 higher number of GPR17-expressing OPCs was 1103
 indeed found compared to contralateral hemi- 1104
 sphere (Lecca et al. 2008), suggesting an 1105
 increased proliferation rate in response to 1106
 demyelination. 1107

Dysregulated expression of GPR17 has been 1108
 described also after traumatic injury, in both 1109
 brain (Boda et al. 2011) and spinal cord (Ceruti 1110
 et al. 2009). In stab wound, a model of cortical 1111
 trauma, early after lesion, the density of GPR17- 1112
 expressing OPCs in gray matter was reduced 1113
 compared to contralateral cortex, consistent 1114
 with a global oligodendroglial loss. At later 1115
 times, GPR17⁺ cells increased significantly in 1116
 number around the lesion in both gray and 1117
 white matter, likely due to the expansion of the 1118
 NG2 cell pool, which, in turn, reflects an attempt 1119
 to replace dead OPCs. This reactivity lasted up to 1120
 7 days and then declined over time, going back to 1121
 basal levels at 14 days after lesion. This pattern 1122
 has been confirmed in human samples from 1123
 patients with traumatic brain injury (Franke 1124
 et al. 2013). In both neurosurgical and autopsy 1125
 specimens, GPR17 expression was evident inside 1126
 the contused core and progressively declined 1127
 distally according to a spatio-temporal gradient. 1128
 Inside and around the core, GPR17 labeled dying 1129
 neurons, reactive astrocytes, and activated 1130
 microglia/macrophages. In peri-contused paren- 1131
 chyma, GPR17 was found on OPCs, some of 1132
 which had proliferated, indicating 1133
 re-myelination attempts. In agreement with the 1134
 above data, in a double transgenic model of 1135
 Alzheimer's disease (the APPS1 mouse) a 1136
 high number of GPR17-positive cells 1137
 accumulated close to amyloid plaques in gray 1138
 matter, revealing receptor up-regulation as a fea- 1139
 ture of oligodendroglial reactivity also in this 1140
 pathological condition (Boda et al. 2011). 1141

Similar GPR17 changes have been reported 1142
 also in typically de-myelinating diseases such as 1143
 in models of multiple sclerosis (MS). In this 1144
 disease, remyelination occurs after the initial 1145
 myelin damage, but it fails after multiple demye- 1146
 lination episodes, which eventually leads to axo- 1147
 nal degeneration and progressive disability 1148

1149 (Franklin and Ffrench-Constant 2008). Interest-
1150 ingly, synthesis of cysLTs is increased in MS
1151 plaques and in the spinal cord of mice subjected
1152 to experimental autoimmune encephalomyelitis
1153 (EAE), an immune-mediated model of demyelin-
1154 ation (Whitney et al. 2001). Of note,
1155 montelukast, an antagonist at both CysLT1 and
1156 GPR17, attenuated CNS infiltration of inflamma-
1157 tory cells and the clinical symptoms of EAE
1158 (Wang et al. 2011). However, the exact contribu-
1159 tion of GPR17 to these effects has not been
1160 investigated in detail. Overexpression of the
1161 GPR17 transcript has been observed in both
1162 EAE mice and in a cohort of human MS tissues
1163 (Chen et al. 2009). GPR17 expression was sig-
1164 nificantly increased in MS plaques as compared
1165 with white matter from non-neurological donor
1166 samples and normal-appearing white matter from
1167 MS donors. In a similar way, acute damage to
1168 myelin induced by lysolecithin (Lys) injection in
1169 corpus callosum induced a strong overexpression
1170 of GPR17 at the lesion site 10 days after injury
1171 (Boda et al. 2011). Thus, independently of the
1172 original cause, GPR17 is abnormally
1173 up-regulated in MS and some models of neuro-
1174 degenerative conditions characterized by myelin
1175 disruption (Fumagalli et al. 2016).

1176 On this basis, it could be hypothesized that,
1177 after damage, GPR17 is initially induced to pro-
1178 mote the growth and differentiation of OPCs;
1179 however, at later stages, due to lack of appropri-
1180 ate environmental stimuli, presence of inflamma-
1181 tory signals and/or intrinsic factors,
1182 physiological GPR17 downregulation is
1183 impeded, thus freezing cells at a stand-by stage,
1184 where they are neither proliferating nor
1185 differentiating. When this happens, interventions
1186 targeting GPR17 may help bypassing this check-
1187 point and facilitate terminal maturation. Since
1188 GPR17 is a *membrane* receptor that, at variance
1189 from other intrinsic regulators of oligoden-
1190 drogenesis, can be easily targeted and
1191 manipulated with pharmacological agents, it is
1192 envisaged that agents counteracting GPR17 aber-
1193 rant expression under these conditions could
1194 induce OPCs to resume myelination and promote
1195 neurorepair. To support this hypothesis, in
1196 MCAo animals, administration of GPR17

antagonists such as cangrelor or montelukast 1197
(Ciana et al. 2006; Lecca et al. 2008), or 1198
GPR17 silencing due to *in vivo* delivery of spe- 1199
cific antisense oligonucleotides (Ciana et al. 1200
2006; Lecca et al. 2008) or small interfering 1201
RNAs (Zhao et al. 2012) resulted in a significant 1202
reduction in brain's ischemic volume. Use of 1203
GPR17 anti-sense oligonucleotides also reduced 1204
damage and improved functional recovery in a 1205
model of spinal cord injury, in line with the 1206
hypothesis that GPR17 is aberrantly 1207
overexpressed as a consequence of damage 1208
(Ceruti et al. 2009). 1209

In contrast to what observed in MCAo, in a rat 1210
neonatal model of ischemic periventricular 1211
leukomalacia (PVL), a common cerebral white 1212
matter injury, the GPR17 agonist UDP-glucose 1213
(and not an antagonist) significantly contributed 1214
to myelin sheaths recovery and improved motor 1215
functions, learning and coordination in PVL pups 1216
(Mao et al. 2012). The reason for this discrep- 1217
ancy may reside in the different outcome of the 1218
ischemic insult in neonatal brain compared to 1219
adults. It could be hypothesized that, in neonatal 1220
pups, existing OPCs, which are very sensitive to 1221
ischemic death, are immediately killed by the 1222
ischemic insult, with no obvious GPR17 1223
upregulation; conversely, being these cells 1224
generated at distinct waves during the first 1225
weeks of life at much higher rates compared to 1226
adulthood, a GPR17 agonist (instead of an antag- 1227
onist) would allow to properly activate newborn 1228
OPCs, thus favouring the formation of myelin 1229
sheaths and neurological recovery. 1230

Several of the still obscure aspects of GPR17 1231
pathophysiology have been linked to the diffi- 1232
culty of establishing a causal relationship 1233
between GPR17 expression and myelination 1234
in vivo. Since GPR17 is no longer expressed in 1235
mature myelinating OLs (Lecca et al. 2008; 1236
Fumagalli et al. 2015), it was impossible to dem- 1237
onstrate that cells that have expressed GPR17 in 1238
their earlier life can indeed myelinate. Only 1239
recently, the generation of the first 1240
GPR17^{iCreER}^{T2}-GFP reporter mouse line for 1241
fate mapping studies has allowed to follow the 1242
final destiny of GPR17⁺ cells during both physi- 1243
ological differentiation and in disease, thanks to 1244

1245 the inducible expression of the green fluores-
 1246 cence protein (GFP). In these mice, upon tamox-
 1247 ifen induced recombination, OPCs expressing
 1248 GPR17 at that very specific moment, become
 1249 green and can be traced as such for the entire
 1250 animal's life. Use of these mice has allowed to
 1251 show that, in normal brain, GFP⁺ cells differenti-
 1252 ate very slowly (needing about 3 months to reach
 1253 maturity), but after acute insults, they rapidly
 1254 reacted to damage with proliferation and migra-
 1255 tion toward the injured site, thus representing a
 1256 'reserve pool' of adult quiescent progenitors
 1257 maintained for repair purposes (Vigano et al.
 1258 2016). A full characterization of the long-term
 1259 events occurring in the brain of ischemic MCAo
 1260 GPR17iCreER^{T2}-GFP mice has shown that,
 1261 despite massive recruitment of GFP⁺ green
 1262 OPCs at the ischemic site, only a few percentage
 1263 of these cells become mature myelinating OLs,
 1264 likely due to local unfavourable inflammatory
 1265 milieu (Vigano et al. 2016; Bonfanti et al. 2017).
 1266 More recently, it has been demonstrated that
 1267 GPR17 over-activation inhibited oligodendro-
 1268 cyte survival by reducing intracellular cAMP
 1269 levels and inducing expression of the
 1270 pro-apoptotic gene *Xaf1*. GPR17 overactivation
 1271 also negatively regulated protein kinase A sig-
 1272 naling pathway and expression of the transcrip-
 1273 tion factor c-Fos. In line with these data, in the
 1274 lysolecithin-mediated demyelination injury
 1275 model, the pharmacological inhibition of
 1276 GPR17 with pranlukast increased oligodendro-
 1277 cyte survival and promoted immature oligoden-
 1278 drocyte differentiation through the upregulation
 1279 of Epa1, the exchange factor directly activated
 1280 by cAMP (Ou et al. 2016). These data are fully
 1281 consistent with our results in other injury models
 1282 characterized by demyelination and abnormal
 1283 GPR17 upregulation (summarized in Fumagalli
 1284 et al. 2016), suggesting that under these
 1285 conditions GPR17 inhibition has potential for
 1286 treatment of demyelinating diseases (Ou et al.
 1287 2016).

3.2 GPR17 in Brain Rejuvenation

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A recent report has investigated the roles of
 GPR17 in age-associated cognitive decline
 (Marschallinger et al. 2015). Authors have first
 shown that oral administration of montelukast
 (an antagonist of both CysLTR1 and GPR17,
 see above), for 6 weeks to moderately old rats
 (20 months old) resulted in structural and func-
 tional rejuvenation of aged brains, as
 demonstrated by restoration of blood brain bar-
 rier integrity, reduced microglia activation in the
 brain, increased levels of hippocampal
 neurogenesis and significantly improved learning
 and memory tasks. Important, montelukast had
 no effects on the behaviour and cognitive
 abilities of young animals, suggesting that its
 actions specifically target an aging associated
 defect (see also below). Regression and correla-
 tion analyses showed that montelukast-induced
 learning improvement in the old animals was
 independent of the changes in microglia mor-
 phology but rather depended on the rate of
 neurogenesis measured as increased number of
 proliferating neuroblasts in hippocampal dentate
 gyrus. Interestingly, authors also provided immu-
 nohistochemical evidence for the presence of
 GPR17, but not CysLTR1, in a subset of
 doublecortin (DCX)⁺ newborn neurons in hippo-
 campal dentate gyrus, suggesting a role in the
 proliferation and specification of these cells.
 Studies on neurospheres obtained from mice
 lacking FOXO1, a GPR17 regulating transcrip-
 tion factor, and from GPR17^{-/-} mice indeed
 confirmed montelukast effects be due to action
 on GPR17/DCX⁺ neuroblasts in hippocampal
 dentate gyrus, leading to increased neurogenesis.
 Globally these data suggest that, under normal
 conditions, GPR17 exerts a negative control on
 the proliferation of neural progenitors in the hip-
 pocampus; in aged animals, due to the overall
 decrease of neurogenesis, GPR17 inhibition of
 proliferation becomes detrimental and
 contributes to memory impairment. Under such
 pathological conditions, montelukast can restore
 neurogenesis by alleviating GPR17 inhibitory
 effect (Marschallinger et al. 2015).

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1334 **3.3 GPR17 in Gliomas**

1335 OL markers such as Olig2, PDGFR α and NG2
 1336 are often expressed in glioma cells. Little is
 1337 known about the origin of these tumors, but it is
 1338 possible that they arise from dysregulated OPCs
 1339 (Liu et al. 2011b). Considering that OPCs are the
 1340 only proliferating population in the adult brain,
 1341 defects in differentiation mechanisms favouring
 1342 cell proliferation could be a primary cause of
 1343 gliomas. A complementary strategy for tumor
 1344 treatment is to promote pathways for maintaining
 1345 quiescence and/or driving terminal differentia-
 1346 tion of the tumoral progenitors. In this respect,
 1347 a recent microarray analysis of mouse and human
 1348 gliomas aimed at unveiling new candidates pro-
 1349 moting differentiation or quiescence has
 1350 highlighted GPR17 as a new potential target
 1351 (Dougherty et al. 2012). In glioma cells, treat-
 1352 ment with UDP, UDP-glucose or LTD₄ indeed
 1353 reduced the formation of glioma spheres
 1354 suggesting that GPR17 stimulation can represent
 1355 a good strategy to drive the differentiation of
 1356 highly proliferative uncommitted tumor cells to
 1357 the oligodendroglial fate, negatively affecting
 1358 both tumor cell proliferation and self-renewal
 1359 (Dougherty et al. 2012). These data are in line
 1360 with the fact that most of the OPCs expressing
 1361 GPR17 in brain are quiescent (Lecca et al. 2008),
 1362 and support the pro-differentiative effects of its
 1363 putative endogenous ligands (see also Sects. 2.1
 1364 and 3.2).

1365 **4 Conclusions**

1366 GPR17 has emerged as a new GPCR of great
 1367 interest for drug development. It is almost exclu-
 1368 sive localization to OPCs, the myelin forming
 1369 cells and the only (slowly) proliferating cell pop-
 1370 ulation in the intact brain, has highlighted
 1371 GPR17 as a novel pharmacological target for
 1372 demyelinating diseases. At variance from other
 1373 myelinating genes, GPR17 is a membrane recep-
 1374 tor, thus amenable for pharmacological modula-
 1375 tion, which has attracted a lot of interest for the
 1376 development of new therapeutic approaches to

MS and other neurodegenerative diseases 1377
 characterized by myelin disruption. The recent 1378
 demonstration that GPR17 is also expressed by a 1379
 subset of hippocampal neural progenitors 1380
 involved in cognitive functions does not detract 1381
 from the potential interest of GPR17 ligands in 1382
 neurodegenerative diseases, since, as 1383
 demonstrated by the montelukast study 1384
 (Marschallinger et al. 2015), these ligands may 1385
 be active only when specific pathological GPR17 1386
 changes are present. 1387

The recent studies on GPR17 revealed its 1388
 transient expression in OPCs and a more com- 1389
 plex role than expected: a pro-differentiating role 1390
 in early OPCs and a negative function on matu- 1391
 ration in late stage OPCs. Thus, the apparently 1392
 contrasting *in vitro* data obtained with different 1393
 GPR17 stimulatory agents (Lecca et al. 2008; 1394
 Fumagalli et al. 2011a; Hennen et al. 2013) 1395
 may depend on the specific differentiation stage 1396
 at which these compounds have been added to 1397
 cultured OPCs. It may well be that the function 1398
 of GPR17 is different in the intact and diseased 1399
 brain, based on the availability of its endogenous 1400
 ligands. If uracil nucleotides, cysLTs, oxysterols 1401
 and chemokines like SDF-1 are indeed among 1402
 the signaling molecules able to activate GPR17 1403
in vivo (see also below), we envisage that their 1404
 role would be more likely unveiled under patho- 1405
 logical conditions, where these ligands massively 1406
 accumulate at lesion sites inside the CNS. 1407

Experiments in a wide variety of rodent 1408
 models of neurodegeneration have shown that, 1409
 independently of the nature of the insult (ische- 1410
 mic, traumatic or toxic) and of the presence of 1411
 any concomitant neuronal pathology, demyelin- 1412
 ating conditions invariably led to GPR17 1413
 upregulation. We believe that this dysregulation 1414
 reflects an initial attempt to repair the lesion by 1415
 stimulating OPCs differentiation via GPR17, but 1416
 that this attempt is later invalidated by the inabil- 1417
 ity of maturing cells to downregulate/internalize 1418
 the receptor, which, in turn, leads a differentia- 1419
 tion blockade. On this basis, it is envisaged that 1420
GPR17 antagonists would be useful in MS and 1421
 neurodegenerative diseases. By counteracting 1422
 GPR17 aberrant dysfunction, antagonists would 1423
 help OPCs to complete their maturation, thus 1424

1425 re-establishing endogenous remyelination, as
1426 recently also confirmed (Ou et al. 2016).

1427 Due to the still ambiguous state of the phar-
1428 macology for this receptor, the Nomenclature
1429 Committee of the International Union of Phar-
1430 macology (NC-IUPHAR) has not yet officially
1431 de-orphanized this GPCR (Davenport et al.
1432 2013). However, as also emphasized by
1433 NC-IUPHAR, much of the work in this area has
1434 been based on recombinant expression systems
1435 using different host cells and transfection
1436 methodologies compared to data derived from
1437 native cells. In recombinant “artificial” cell
1438 systems, activity tests are highly dependent on
1439 the experimental conditions utilized and subject
1440 to several artifacts, especially in the case of
1441 receptors’ constitutive activation, a typical fea-
1442 ture of several GPCRs including GPR17
1443 (Bened-Jensen and Rosenkilde 2010; Maekawa
1444 et al. 2009; Qi et al. 2013; Eggerickx et al. 1995;
1445 Uhlenbrock et al. 2002; Rosenkilde et al. 2006;
1446 Qin et al. 2011; Im 2004) that can profoundly
1447 alter ligand behavior (Kenakin 2001; Davenport
1448 et al. 2013).

1449 In terms of drug development, neither uracil
1450 nucleotides nor CysLTs are suitable to this pur-
1451 pose, because neither ligand class is competent to
1452 discriminate between the functions of purinergic
1453 receptors, CysLT receptors, and GPR17 *in vivo*,
1454 where multiple receptors are often co-expressed.
1455 Nevertheless, the already available *in vivo* rodent
1456 data reporting positive neuro-reparative effects
1457 induced by commercially available montelukast
1458 or pranlukast (Yu et al. 2005a, b), which are
1459 potent (although non selective) GPR17
1460 antagonists, foster the search for further GPR17
1461 ligands (Eberini et al. 2011; Hennen et al. 2013)
1462 and may represent an important advancement for
1463 patients with neurodegenerative diseases.

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1469 interest.

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Author Queries

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