Purinergic Signals Regulate Daily S-Phase Cell Activity in the Ciliary Marginal Zone of the Zebrafish Retina

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Abstract Regeneration and growth that occur in the adult teleost retina have been helpful in identifying molecular and cellular mechanisms underlying cell proliferation and differentiation. Here, it is reported that S-phase cell number, in the ciliary marginal zone (CMZ) of the adult zebrafish retina, exhibits daynight variations with a mid-light phase peak. Oscillations persist for 24 h in constant darkness (DD), suggesting control by a circadian component. However, variations in the S-phase nuclei number were rapidly dampened and not present during and after a second day in DD. An ADPBS treatment significantly enhanced S-phase activity at night to mid-light levels, as assessed by in vivo BrdU incorporation in a 2-h interval. Moreover, daylight increase in S-phase cell number was completely abolished when extracellular nucleotide levels or their extracellular hydrolysis by ectonucleoside triphosphate diphosphohydrolases (NTPDases) were significantly disrupted or when a selective antagonist of purinergic P2Y1 receptors was intraocularly injected before BrdU exposure. Extracellular nucleotides and NTPDase action were also important for maintaining nocturnal low levels of S-phase activity in the CMZ. Finally, we showed that mRNAs of NTPDases 1, 2 (3 isoforms), and 3 as well as of P2Y1 receptor are present in the neural retina of zebrafish. NTPDase mRNA expression exhibited a 2-fold increment in light versus dark conditions as assessed by quantitative RT-PCR, whereas P2Y1 receptor mRNA levels did not show significant day-night variations. This study demonstrates a key role for nucleotides, principally ADP as a paracrine signal, as well as for NTPDases, the plasma membrane-bound enzymes that control extracellular nucleotide concentration, for inducing S-phase cell entry in the CMZ—normally associated with retinal growth—throughout the light-dark cycle.

Key words zebrafish retina, ciliary marginal zone, extracellular nucleotides, S-phase cell rhythm

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Eyes in teleost fish grow throughout life, and the mature retina matches this growth by stretching and the continuous genesis of new cells (Johns and Fernald, 1981; Stenkamp, 2007). Cell proliferation and differentiation occur in a neuroepithelial zone at the interface between neural retina and ciliary epithelium, called the ciliary marginal zone (CMZ) (Lyall, 1957; Reh and Fischer, 2001). Within mature retina, rod photoreceptors differentiate from precursor cells within the outer nuclear layer (ONL) (Johns and Fernald, 1981). In embryonic development, retinal cells clonally originate from a common cell precursor. However, whether a unique cell precursor exists in the CMZ and the nature of the extrinsic and intrinsic signals that regulate retinogenesis are subjects of intense investigation. Light, for a photosensitive tissue such as the retina, might regulate normal growth and cell proliferation like in other zebrafish larval tissues (Dekens et al., 2003). In fact, daily variations of rod precursor proliferation during the light-dark (LD) cycle has been described in a teleost fish (Chiu et al., 1995). Several studies in the vertebrate retina have revealed that a circadian clock drives many retinal functions (Cahill and Besharse, 1995).

Extracellular nucleotides/nucleosides (eNCT/S) in the retina are important cues for neuron-glia communication (Newman, 2006). ATP is released by exocytosis in the nervous system (Burnstock, 2006) and via gap junction hemi-channels in the retina (Pearson et al., 2005). Di- and triphosphate nucleotides such as ATP, ADP, and UTP are extracellularly dephosphorylated by ectonucleoside triphosphate diphosphohydrolases (ENTPDases). The coordinated action of ENTPDases and an ecto-5'-nucleotidase dephosphorylates extracellular ATP to adenosine (Hunsucker et al., 2005). eNCT/S act via specific purinergic or P receptors in the plasma membrane (Burnstock, 1978; 2007). Studies on retinal development in the chick embryo suggest a role for eNCT/S in regulating progenitor cell proliferation (Pearson et al., 2005; França et al., 2007). Interestingly, NTPDase2 and ADP participate in eye and retina morphogenesis in Xenopus laevis (Massé et al., 2007).

Therefore, we investigated the role of extracellular ATP and its metabolites in regulating S-phase cell activity in the CMZ of the zebrafish retina, which we have found follows day-night oscillations during the LD cycle that persist for 24 h in constant darkness (DD). The number of S-phase nuclei was highest at the mid-light phase and mid-subjective day. S-phase cell number was regulated by eNCT in both light and

darkness. mRNAs of NTPDases 1, 2, and 3 are expressed in the zebrafish retina, and their levels showed significant day-night variations.

MATERIALS AND METHODS

Materials

Materials included paraformaldehyde; PBS (137 mM NaCl, 2.7 mM KCl, 5.7 mM phosphate, pH:7.4); Tween-20; MS-222; sucrose; apyrase (EC-3.6.1.5); hexokinase (EC-2.7.1.1); 5-bromo-2'-deoxyuridine (BrdU); ADPβS; 8-sulfophenyl theophylline (8-SPT) (Sigma, St Louis, MO); MRS2211, MRS2179, NF110, NF157, and ARL67156 trisodium salt (Tocris, Bristol, UK); mouse monoclonal anti-BrdU antibody (Roche, Mannheim, Germany); goat anti-mouse ALEXA 488 antibody, random primers, dNTPs, MMLV, DTT, RNAseOUT, and sense and antisense primers (Invitrogen, Carlsbad, CA); Total RNA Extraction Kit (RBC Biosciences, Taiwan); GoTaq-DNA polymerase, dNTPs-mix, DNasa-I (Promega, Madison, WI); realtime PCR mix (Biodynamics, Buenos Aires, Argentina); and ultra pure agarose (GenBiotech, Buenos Aires, Argentina).

Animals

Adult zebrafish (*Danio rerio*) of about 3 cm and 1.5 g were maintained at 28.5 °C in a 14:10 h LD cycle (lights on 0700 h and off 2100 h) and fed twice a day. A group of zebrafish, used for night intervals, was raised in an inverted phase (I) 14:10 h LD cycle (lights on 2000h and off 1000 h) in a separate room. Other groups of zebrafish were transferred to DD. Animals were euthanized by immersion in MS-222 anesthetic solution (0.02% wt/vol) and then decapitated and enucleated, at 4 different intervals defined as Zeitgeber time (ZT2, ZT9, ZT16, and ZT21; ZT0 indicates lights on and ZT14 lights off) during the LD cycle or first, second, and fourth days in DD. A group of zebrafish from the (I) LD cycle were euthanized at ZT9 (i.e., mid-light for this group). The Committee on Animal Research at University of Buenos Aires approved protocols for animal use and care.

Intravitreal Injections

5-Bromo-2'-deoxyuridine administration. BrdU, a thymidine analogue, incorporates into DNA during the

S-phase of the cell cycle. Anesthetized zebrafish were intraocularly injected with 0.5 µL of a sterile saline (0.65%) solution containing 20 µg/µL of BrdU, 2 h before euthanasia. BrdU was injected in control or previously treated eyes after 20 h of each treatment. Zebrafish in darkness were injected and euthanized in dim far red light.

eNCT hydrolyzing enzyme treatment. Apyrase dephosphorylates diphosphate and triphosphate nucleotides whereas hexokinase hydrolyzes ATP to phosphorylated D-hexose plus ADP. We injected 0.5 µL of 25 U/ mL apyrase, hexokinase, or heat-inactivated enzymes (boiled for 5 min) once. Zebrafish were euthanized 22 h later to prepare eyecups. Apyrase or hexokinase remains active within the vitreous at least for 20 h (Battista et al., 2009).

Ecto-ATPase activity inhibitor treatment. ARL67156, an ATP analogue, selectively inhibits ecto-ATPase activity without activating purinergic receptors (Robson et al., 2006). We injected 100 µM ARL67156 (Westfall et al., 1997; Newman, 2003) via the intravitreal route in $0.5 \mu L$ of saline solution (pIC₅₀ = 4.62 and 5.1 in human blood and rat vas deferens, respectively). Zebrafish were euthanized 22 h later.

eNCT/S agonist and antagonist treatment. We injected $5 \mu M$ MRS2211, $1 \mu M$ MRS2179, $10 \mu M$ 8-SPT, or $5 \mu M$ NF157 plus 5 µM NF110, and 22 h later zebrafish were euthanized (after 2 h of BrdU exposure). Additionally, a group of zebrafish were injected with 25 μM ADPβS, a slightly hydrolysable ADP analogue. Control eyes from different zebrafish were injected with sterile saline.

MRS2211. P2Y13 receptor antagonist (IC₅₀ = 1.07 μ M) displays greater than 20-fold selectivity over P2Y1. Agonist relative specificity: ADP>>ATP. MRS2179: P2Y1 receptor antagonist ($IC_{50} = 0.330 \mu M$; $K_i = 100 nM$); selective over P2X1/X3/X2/X4 and P2Y2/Y4/Y6 receptors. Agonist relative specificity: ADPβS> ADP>>ATP (Burnstock, 2007). NF110: high-affinity P2X3 receptor antagonist (IC₅₀ = 0.527 μ M; K_i = 36/82/4144 nM for ATP-activated P2X3/X1/X2, respectively); shows no activity on P2Y1/Y2/Y11 receptors (IC₅₀ >10 μ M). NF157: Potent antagonist of P2Y11 receptors (IC₅₀ = 0.463 μ M); selective for ATPactivated P2Y11 and P2X1 over P2Y1/Y2, P2X2/ X3/X4/X7 receptors. Agonist relative specificity: ATPγS>ATP (Burnstock, 2007). 8-sulfophenyl theophylline (8-SPT): adenosine P1 receptors antagonist $(K_i = 1.2 \mu M)$.

Immunocytochemistry

To prepare eyecups, the cornea, lens, and vitreous were removed. Eyecups were fixed with 4% paraformaldehyde for 1 h at room temperature (RT); incubated in 5%, 10% (30 min), and 20% sucrose (overnight); and embedded in tissue freezing medium. We collected 16-µm cryosections onto gelatin-subbed slides, which were kept at -20 °C. Retinal sections were incubated in 2 M HCl for 15 min at 37 °C and thoroughly washed with PBST (PBS plus 0.1% Tween-20). Then, sections were incubated 1 h at RT in 5% normal goat serum (NGS) and overnight with an antibody against BrdU (1:200 in 3% NGS) at 4 °C. Sections were washed and incubated in darkness for 2 h at RT, with a fluorescent antibody (1:400 in 3% NGS). Omission of antibodies was performed as negative controls.

Counting of BrdU-Positive Nuclei

Eyecups were cut in the sagittal plane (nasal to temporal). Each slide contained sections from different portions of the eyecup. We identified BrdU-positive nuclei focusing through the section in both CMZ under microscopic observation. We counted all BrdUpositive nuclei in 10 to 12 nonadjacent sections from each eye. Nuclei in the nonlaminated retina at both ends were considered CMZ nuclei. Differences among groups were analyzed using 1-way ANOVA, followed by Tukey-Kramer multiple comparison test or Student t test.

Microscopy

Counting was performed with an epifluorescence microscope (Olympus BX50, Japan; objective 40X-NA: 0.75). Confocal images were captured with an FV1000 Fluoview spectral microscope (Olympus; objective SAPO-60X oil-NA: 1.35). Two-dimensional confocal and differential interference contrast (DIC) images were obtained from Z-stacks by using Fluoview software.

Reverse Transcription (RT) and Polymerase Chain Reaction (PCR)

Groups of 5 zebrafish were euthanized at ZT7–9 (mid-light phase) in bright light and ZT14-16 (1:30 h of darkness) in dim far red light. Ten neural retinas were homogenized in a buffer of a RNA purification kit and considered a sample. Three to 4 sample replicates

Table 1. Specific primers for each target gene and reference gene (β-actin).

Gene	Sense	Antisense
NTPDase1	TCACAGACACAGCGAAACCC	GAATGTGCAACGGAGGTAAGC
NTPDase2 mq	GAGCAGCAGCACGTAGCC	GACCTCAGCCGACTCTTTGG
NTPDase2 mg	CTTACGCCTCTTTGGACACG	AGTCAGATGGATAGCAGGGATG
NTPDase2 mv	TGATGGAGTCTTTCAGCCCAAC	AGCAGCGTCCTCTAAATGAGC
NTPDase3	GGCTGGATCACTGTCAACTACC	CCCAAATCCAATGACCCAACTG
P2Y1 receptor	TATGGACAATGCTCCGCTTAGG	CGCTCTCATGTTCAGGTTCTTC
β-actin	TCCCAAAGCCAACAGAGAGAAG	GTCACACCATCACCAGAGTCC

Primers were selected by using exon sequences from zebrafish genome (Ensembl database). Selected primer pairs did not hybridize with other NTPDase enzyme or isoform (mq, mg, mv).

were examined for each interval. RNA was quantified and DNAse-I treated (30 min, 25 °C). cDNA was reverse transcribed from RNA with random primers and quantified with a NanoDrop 3300 spectrophotometer (Thermo Fisher Scientific, Waltham, MA). Either 1 or 2 µg of cDNA was used for real-time quantitative PCR (qPCR) (RotorGene 6000; Quiagen, Valencia, CA). Specific primers were selected using Beacon Designer Software (Premier Biosoft International, Palo Alto, CA) from zebrafish genome reported in Ensembl database (Table 1). Products of standard or qPCR were checked by electrophoresis in 2% agarose gels. No-RT controls, in which the MMLV-reverse transcriptase was omitted, were amplified by qPCR with every primer pair. No-RT samples showed background fluorescence levels. No-RT or cDNA-containing samples were run in triplicates. β-Actin (internal reference gene) expression showed no significant variations among intervals and was amplified in parallel with each gene of interest (target genes). Slopes from qPCR kinetic curves obtained as the logarithm of relative fluorescence after *n* cycles (Rn) plotted against the number of cycles (n) were calculated for each sample for β -actin and target genes. Equation 1 describes the linear range in the exponential phase of the kinetic curve:

$$\lg Rn = \lg (1 + E) n + \lg Ro \tag{1}$$

where Ro is the initial fluorescence amount and E is the PCR efficiency. Within the linear range, we defined a trendline of 4 to 5 data points, with the highest possible slope and correlation coefficient, and calculated the efficiency for each reaction ($E = 10^{slope} - 1$). Averaged amplification efficiencies for each sample were 1 ± 0.02 (mean \pm SD) for reference or interest genes (Schefe et al., 2006). Samples with efficiencies out of this interval were considered outliers and not included in data analysis.

Quantitative PCR Data Analysis

Analysis was performed throughout the $2^{-\Delta \Delta CT}$ method as described by Livak and Schmittgen (2001). An arbitrary fluorescence threshold of 0.06 was used to obtain C_{T} values from each curve. We chose cDNA samples from mid-light intervals (ZT7–9)

as the interest sample and beginning of darkness (ZT14–16) as the calibrator sample. Both C_T values were previously normalized with β-actin. We obtained a ratio that indicates fold change of mRNA levels of each target gene at ZT7-9 over ZT14-16. For the calibrator sample, 2-AACT value equals 1. Relative values from each pool of 10 retinas are reported. Standard deviations (SD) from triplicates were propagated by using standard methods and confidence intervals were calculated as: 2^{-(ΔΔCT ± propagated SD)} (Livak and Schmittgen, 2001).

RESULTS

S-Phase Nuclei Number in the CMZ Shows Day-Night Variations That Persist for 24 h in DD

We counted BrdU-positive nuclei (S-phase nuclei) within the following 2-h intervals: ZT0–2 and ZT7–9 in light, ZT14–16 and ZT19–21 in darkness (Figure 1A). Despite the scant number of BrdU-positive nuclei per CMZ/section, this parameter showed daily variations with a peak at ZT7-9, that is, after 7 to 9 h of light $(4.78 \pm 0.98 \text{ nuclei}, n = 8 \text{ zebrafish})$. After 2 h of darkness (ZT14-16), S-phase cell number decreased to a minimum (1.49 \pm 0.30 nuclei, n = 5 zebrafish), which was significantly different from BrdU-positive nuclei after 7 h of darkness (ZT19–21: 2.80 ± 0.56 nuclei, n =8 zebrafish). At dawn, ZT0-2, BrdU-positive nuclei were (2.34 \pm 0.78, n = 7 zebrafish) not different from night values. The ratio of S-phase nuclei at the midlight peak to the trough at night was 3.2 to 1. Upon transfer to DD, S-phase cell number showed significant variations among intervals for 24 h (ZT0–2: 2.72 ± 0.44; ZT7-9, 4.83 ± 0.43 ; ZT14-16, 1.05 ± 0.07 ; ZT19-21, 1.80 ± 0.32 ; n = 4 zebrafish per interval; Figure 1B) with a maximum to minimum ratio of 4.6 to 1. Subjective

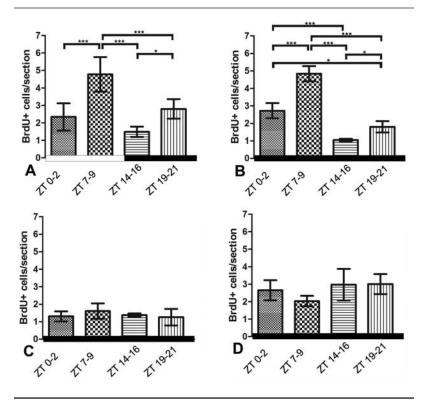


Figure 1. Number of BrdU-positive nuclei plotted as a function of time of day (ZT) in a 14:10 h LD cycle (A) or during the first (B), second (C), or fourth (D) day in DD. Zebrafish were handled either in light or under far red illumination. Values are BrdU-positive cell number per ciliary marginal zone (CMZ) per retinal section and are mean \pm SD (n = 4–8 zebrafish per ZT interval; ZT0 is lights on and ZT14 lights off). Brackets indicate significant differences between intervals. ***p < 0.001; *p < 0.05, Tukey-Kramer multiple comparison test after 1-way ANOVA.

day values were significantly higher than values at subjective night. Confocal images with a representative S-phase nuclei number at ZT7-9 and ZT14-16 in LD and 1 day in DD are shown in Supplementary Figure S1A-D and E-H, respectively. In contrast, BrdU-positive nuclei number during the second day in DD (ZT0-2: 1.30 ± 0.29 ; ZT7-9: 1.60 ± 0.43 ; ZT14-16: 1.37 \pm 0.09; ZT19–21: 1.26 \pm 0.47, n = 4 zebrafish per interval; Figure 1C) or the fourth day in DD $(ZT0-2: 2.65 \pm 0.57; ZT7-9: 2.03 \pm 0.30; ZT14-16: 2.97 \pm$ 0.90; ZT19–21: 3.00 ± 0.57 , n = 6 zebrafish per interval; Figure 1D) did not show significant variations among intervals. During subjective days 2 and 4, BrdUpositive nuclei failed to increase to the LD mid-light peak (Figure 1A) versus DD through all ZT intervals (Figure 1C and D; p < 0.001, Tukey-Kramer after ANOVA). Moreover, during the second day in DD, values were not statistically different from the night minimum in LD (Figure 1A vs. 1C). In contrast, BrdU-positive nuclei number during day 4 in DD

was significantly higher at ZT14-16 and ZT19-21 (Figure 1D) than at ZT14-16 in LD (Figure 1A) (p < 0.05, Tukey-Kramer after ANOVA). Data from day 4 in DD intervals (except for ZT7-9) were significantly higher than those observed during the second day in DD (p < 0.05, *p* < 0.01; Figure 1C vs. 1D).

Supplementary Figure S2 depicts S-phase nuclei number in the CMZ of 2 groups of zebrafish injected with BrdU at ZT7 and euthanized at ZT9 (mid-light for both groups of fish). One of the groups was kept in a different room in a 12-h shifted phase relative to the other. BrdU-positive nuclei number in both groups did not show significant differences (ZT7–9: 4.78 ± 0.98 , n = 8; ZT7–9(I): $4.72 \pm 0.15 \ n = 3 \text{ zebrafish}$).

Effect of eNCT Hydrolyzing **Enzymes and an EctoATPase** Selective Blocker on S-Phase Cell **Activity During the LD Cycle**

Figure 2 shows the effect of apyrase or hexokinase activity on S-phase nuclei number at different intervals. Apyrase treatment significantly decreased BrdUpositive nuclei number at the CMZ at both

night and daylight as compared to heat-inactivated apyrase-treated retinas (ZT7–9: 3.99 \pm 0.26, n=3; ZT14–16: 1.69 \pm 0.30, n = 3) and saline controls (ZT7– 9: 4.20 ± 0.26 , n = 3; ZT14–16: 1.67 ± 0.10 , n = 3). Control data were not significantly different (ZT7–9: p = 0.3788 and ZT14–16: p = 0.9864), and they were pooled in Figures 2A and B. At ZT7-9, S-phase cell number in apyrase-treated retinas diminished below control night levels (apyrase ZT7–9: 0.61 ± 0.20 , n = 4[Figure 2A] vs. control ZT14–16: 1.68 \pm 0.20, n = 6[Figure 2B] and ZT19–21: 2.19 \pm 0.34, n = 3 [Figure 2C]; p < 0.001). The low number of BrdU-positive nuclei at night was further decreased to either 1 or zero in the majority of the quantified sections (ZT14-16: apyrase 0.60 ± 0.15 , n = 4 vs. control 1.68 ± 0.20 , n = 6 [Figure 2B]; ZT19–21: apyrase 0.39 \pm 0.12, n = 4vs. control 2.19 \pm 0.34, n = 3 [Figure 2C]). In contrast, intravitreal activity of hexokinase showed no significant effect on S-phase nuclei number either in light or dark conditions as compared to the heat-inactivated

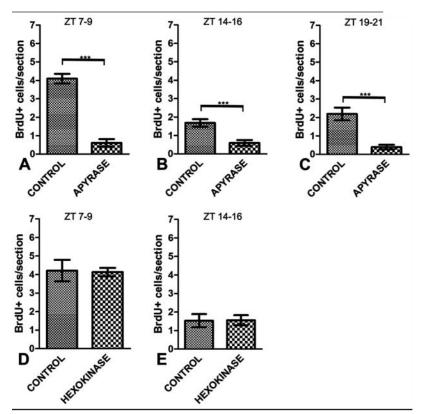


Figure 2. Apyrase or hexokinase activity effect on BrdU-positive nuclei number analyzed at daily intervals during a 14:10-h LD cycle. Eyes underwent intravitreal injection of saline, apyrase, or heat-inactivated apyrase (A-C) or hexokinase or heat-inactivated hexokinase (D, E). Twenty hours later, they were injected with a solution containing BrdU (20 μg/μL). Zebrafish were euthanized 2 h later at the end of the indicated ZT interval (ZT0 is lights on and ZT14 lights off). Values represent the number of BrdU-positive cells per ciliary marginal zone (CMZ)/retinal section and are mean ± SD. Controls: ZT7-9 and ZT14-16 saline plus heat-inactivated apyrase (n = 6 zebrafish per ZT); ZT19–21 heat-inactivated apprase (n = 3 zebrafish); ZT7-9 and ZT14-16 heat-inactivated hexokinase (n = 4-5 zebrafish). Apyrase- or hexokinase-treated retinas: n = 4 zebrafish per ZT interval. Brackets indicate significant differences between treatments. ***p < 0.001 by Student t test.

hexokinase-treated retinas (ZT7–9: hexokinase 4.13 \pm 0.22 and control 4.21 \pm 0.58, n = 4; ZT14–16: hexokinase 1.56 ± 0.27 , n = 4 and control 1.53 ± 0.36 , n = 5) (Figure 2D, E). Furthermore, when ARL67156 known to inhibit NTPDase activity at the concentration tested—was injected within the vitreous (Figure 3 A-C), it significantly decreased BrdU-positive nuclei number at mid-light to levels below those normally observed at night (ARL67156: ZT7–9, 0.91 \pm 0.22, n =4 [Figure 3A] vs. control: ZT14–16, 1.68 \pm 0.20, n = 6[Figure 3B] and ZT19–21, 2.12 \pm 0.28, n = 5 [Figure 3C]; p < 0.001). ARL67156 treatment also decreased BrdU-positive nuclei number at night (ARL67156: ZT14–16, 0.29 \pm 0.05, n = 4 [Figure 3B]; ZT19–21, 0.32 \pm 0.11, n = 4 [Figure 3C]). Supplementary Figure S1 shows confocal images with a representative number of S-phase cells in the CMZ at ZT7-9 in absence (I, J) or presence (K, L) of active apyrase and without (M, N) or with ARL67156 treatment (O, P).

Specific eNCT Regulate S-Phase Cell Activity in the CMZ During the LD Cycle

Figure 3A-C depicts data showing that MRS2179—a selective antagonist of ADP-activated P2Y1 metabotropic receptors—significantly reduced BrdUpositive nuclei number at mid-light, to levels below control values recorded at night (MRS2179: ZT7-9, 0.76 ± 0.34 , n =4 [Figure 3A] vs. control: ZT14–16, 1.68 \pm 0.20, n = 6 [Figure 3B] and ZT19–21, 2.12 ± 0.28 , n = 5 [Fig 3C]; p < 0.001). MRS2179, like apyrase or ARL67156 treatments, significantly decreased BrdU-positive nuclei counts at night to less than a dividing nucleus in a 2-h window per retinal section (MRS2179: ZT14-16, 0.51 \pm 0.17 [Figure 3B] and ZT19–21, 0.60 \pm 0.39 [Figure 3C]; n = 4 zebrafish per group). In contrast, P2 receptor antagonists such as MRS2211, which blocks ADP action on P2Y13 receptors, showed no significant effect (MRS2211: ZT7-9, 3.95 ± 0.21 [Figure 3D]; ZT14–16, 1.97 ± 0.10 [Figure 3E], n = 4 zebrafish per group). Antagonists for ionotropic P2X1/2/3 and metabotropic P2Y11 receptors also failed in demonstrating

an ATP-induced effect on S-phase cell activity $(NF110/157: ZT7-9, 3.97 \pm 0.07 [Figure 3E]; ZT14-16,$ 1.73 ± 0.13 [Figure 3D]; n = 4 zebrafish per group). Likewise, 8-SPT an antagonist for adenosine-activated P1 receptors did not alter BrdU-positive nuclei number at any interval examined (8-SPT: ZT7-9, 3.87 \pm 0.64 [Figure 3E]; ZT14–16, 1.94 \pm 0.32 [Figure 3D]; n = 4 zebrafish per group). For MRS2211, NF110/157 and 8-SPT control values were ZT7-9, 4.18 \pm 0.55; ZT14–16, 1.68 \pm 0.25, n = 4 zebrafish per group. Consistent with this, 1 intravitreal injection of ADPβS, an agonist for P2Y1 receptors, caused a significant increase of the number of BrdU-positive nuclei in the CMZ at night (control: ZT14–16, 1.63 \pm 0.22, n = 4 [Figure 3F]). S-phase nuclei number in the presence of ADPβS at ZT14–16 was statistically

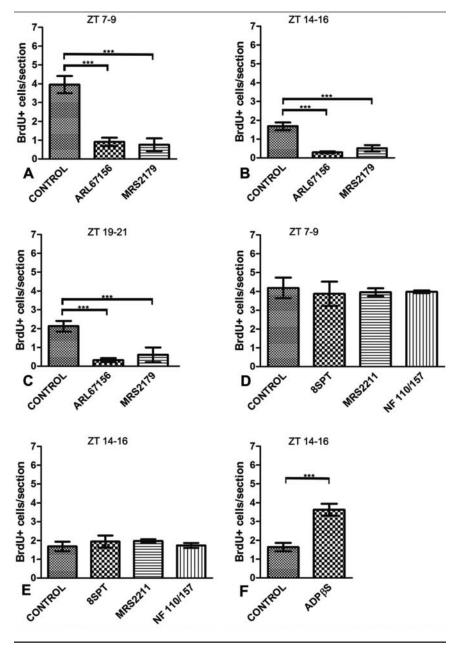


Figure 3. P receptor agonist and antagonists and an ectoATPase activity inhibitor effect on BrdU-positive nuclei number at different intervals during a 14:10-h LD cycle. Eyes underwent intravitreal injection with saline, ARL67156, or MRS2179 (A-C); saline, 8-SPT, MRS2211, or NF110/157 (D, E); or saline or ADPβS (F). Twenty hours later, they were injected with a solution containing BrdU (20 µg/µL). After 2 h, zebrafish were euthanized at the end of the indicated ZT interval (ZT0 is lights on and ZT14 lights off). Values represent BrdU-positive cell number per ciliary marginal zone (CMZ)/retinal section and are mean \pm SD. (A) ZT7–9: n = 4–7 zebrafish. (B) ZT14–16: n = 4–6 zebrafish. (C) ZT19–21: n = 4–5 zebrafish. (D) ZT7-9 and (E) ZT14-16: n = 4 zebrafish per group. (F) ZT14-16: n = 4-5 zebrafish. Brackets indicate significant differences between treatments. ***p < 0.001, as assessed by Tukey-Kramer multiple comparison test after 1-way ANOVA or Student t test.

indistinguishable from mid-light values in absence of exogenous nucleotides (ADPβS: ZT14–16, 3.62 ± 0.32, n = 5 [Figure 3F] vs. control: ZT7–9, 3.95 \pm 0.45

n = 7 [Figure 3A]; p = 0.1996). Supplemen-tary Figure S1 depicts 2-dimensional confocal images with a representative S-phase nuclei number in the CMZ at ZT14-16 in absence (Q, R) or presence (S, T) of ADPβS.

NTPDase mRNA Levels Show **Day-Night Variations** in the Zebrafish Retina

RT-PCR experiments demonstrated mRNA expression of NTPDases 1, 2, and 3 as well as of ADP-activated P2Y1 metabotropic receptor in the zebrafish neural retina (Figure 4C). One band was observed with each primer pair, and amplicon lengths matched expected molecular weights. Furthermore, mq, mv, and mg isoforms for NTPDase2 previously described in other tissues were identified. To reliably detect mg isoform, we used 2 μg of cDNA, whereas 1 μg was sufficient to identify mv and mq isoforms or NTPDases 1 and 3. NTPDase mRNA levels during the LD cycle showed a higher expression in mid-light phase (ZT7-9) than at the beginning of night, after 1:30 h of darkness (ZT14–16) (Figure 4A, B). Expression level of NTPDase mRNAs between intervals was on average about 2- to 3-fold. In contrast, P2Y1 receptor mRNA levels did not show apparent day-night variations.

DISCUSSION

Here we report daily variations in S-phase progenitor cell activity in the CMZ of the adult zebrafish retina, with highest values at the

mid-light phase. The observed day-night oscillations persist for 24 h in free-running conditions. S-phase activity in the CMZ is strongly regulated by

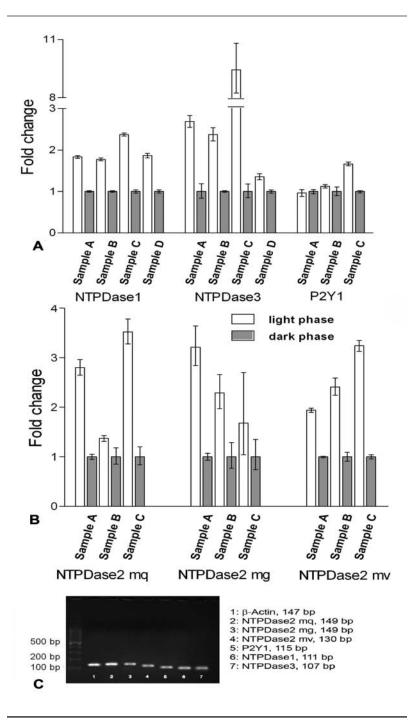


Figure 4. NTPDases and P2Y1 receptor expression in the zebrafish retina identified by real time quantitative RT-PCR (qPCR). RNA was purified from 3-4 pools (samples A-D) of 10 retinas each obtained at the mid-light phase (ZT7-9) or beginning of the dark phase (ZT14-16) of a 14:10-h LD cycle (ZT0 is lights on and ZT14 lights off). Total RNA was treated with DNAse-I, retro-transcribed to cDNA, and amplified by qPCR. One microgram of cDNA, except for NTPDase2 mg isoform (2 μ g), was used as template for qPCR reactions, with primers for β -actin (reference) or specific primers for each target gene. Fold change is the media from triplicate values (for each reaction) normalized to an internal reference and a calibrator sample (ZT14–16) that was calculated by the 2^{-ΔΔCT} method. Propagated errors were estimated from triplicate SD by using standard methods. Error ranges (brackets) were calculated as 2-(AACT ± propagated SD). Amplified cDNA products were separated by electrophoresis in a 2% agarose gel (C). NTPDase2 isoforms: mq, mg, mv.

eNCT levels and hence by the activity of plasma membrane-bound NTPDases. NTPDase mRNA expression is differentially regulated during the LD cycle.

S-Phase Cell Activity in the CMZ Shows Day-Night Oscillations in LD That Persist for One Day in Constant Darkness

It has been reported that the proliferative activity of rod precursors in the ONL of a teleost fish retina (Haplochromis burtoni) exhibits circadian driven variations with a peak at night (Chiu et al., 1995). Here we showed day-night variations of S-phase nuclei number during a regular LD cycle that are at least 3 times higher at daylight than in darkness. A further control was made because, as aforementioned, zebrafish for light intervals were maintained in a separate room with an inverted (I) LD phase from zebrafish for night intervals. Results indicated that BrdU-positive nuclei number peak is resynchronized by light to a 12-h shifted LD cycle (as shown in Supplemental Figure S2), and the observed differences between groups at daylight or night intervals are not due to any uncontrolled environmental variable.

Rod precursors in the ONL (Johns and Fernald, 1981) or proliferative cells in the INL (Julian et al., 1998; Battista et al., 2009) were virtually unlabeled under our experimental conditions (2 h of BrdU exposure); therefore, we could not assess day-night variations in these cell populations. S-phase activity in the CMZ, which involves a different neurogenic cell population, seems to be coincident with a higher behavioral activity (Cahill et al., 1998) and visual sensitivity at daylight (Li and Dowling, 1998). Mitotic division in vertebrates occurs several hours after S phase—in our experiments, at the end of the light phase or beginning of the dark phase. Therefore, BrdU in our experimental conditions labeled S-phase nuclei that did not reach the mitotic phase (Matsuo et al., 2003). Oscillations of S-phase cell

activity persisted 24 h in DD, suggesting that they are driven at least in part by a circadian pacemaker (Cahill, 2002; Dekens et al., 2003). Nonetheless, this putative rhythm rapidly dampened and disappeared from the second day on in DD. BrdU-positive nuclei number throughout day 2 in DD was low. In contrast, during day 4 in DD, S-phase cell activity showed a global nonoscillatory increase, suggesting a lightindependent cell cycle regulation to sustain cell proliferation and hence retinal growth. Light does not regulate cell cycle entry instantaneously, because S-phase cell activity showed similar levels the first 2 h of light (ZT0-2) or in darkness (ZT19-21). Together, these results might indicate that S-phase nuclei entry in the CMZ is driven by a weak circadian oscillator that maintains DD cyclic activity for 1 day. Results also suggest that S-phase cell activity is partially lightdriven, because LD cycle absence causes a delayed suppression of cyclic oscillations on day 2. A combination of a clock-driven and a delayed light-driven extracellular diffusible signal and/or retinal cell activity, probably including photoreceptor or another kind of photosensitive cells (Cheng et al., 2009), might be responsible for increasing daylight S-phase activity in the CMZ.

Apyrase and ARL67156 Exert a Similar Effect on Daily Variations of S-Phase Cell Activity

Apyrase hydrolyses triphosphate and diphosphate nucleotides and hence terminates their action on P2 receptors. The action of apyrase by producing AMP indirectly favors adenosine formation. However, the lack of effect of 8-SPT suggests that endogenous extracellular adenosine does not regulate S-phase nuclei number in the CMZ. Therefore, the inhibitory effect of apyrase on S-phase activity in the CMZ, in either the light or dark phase, might result from hindering nucleotide action on P2 receptors. In contrast, hexokinase treatment, which hydrolyzes extracellular ATP, did not affect S-phase nuclei number. This might indicate that ATP is not essential for controlling S-phase activity in the CMZ. Another possible candidate is ADP because it binds to specific P receptors—unlike AMP—and apyrase quickly hydrolyzes it. Hexokinase activity may produce ADP; however, no effect of hexokinase was observed in either light or darkness. ADP produced by hexokinase activity at daytime could be redundant with extracellular ADP produced by other enzymatic reactions. Our data suggest that

at night, there is less extracellular ADP; nonetheless, hexokinase was ineffective in producing enough ADP to exert a significant effect on S-phase cell activity. This might be due to a very low extracellular ATP concentration at night. Alternatively, ADP produced within the vitreous by hexokinase could be rapidly hydrolyzed by endogenous ectoATPase activity (Battista et al, 2009).

Like the apyrase effect, ARL67156 significantly diminished S-phase nuclei number in the CMZ by inhibiting extracellular hydrolysis of nucleotides. Therefore, these results also suggest that ATP presence by itself in the extracellular milieu is not sufficient to regulate S-phase entry. Presence of ecto ATPase activity mainly driven by NTPDases is crucial for hydrolysis of eNCT and key for S-phase induction in the CMZ at daylight and night.

P2Y1 Receptors Mediate Extracellular Nucleotide Effect on S-Phase Cell Activity in the CMZ

A treatment with ADP β S (25 μ M) led to significant increases in BrdU-positive cell number at nighttime. An elevated level of this ADP analogue for 20 h was a signal sufficient to increase S-phase cell number at night to mid-light values. There is common agreement that P2Y1 receptors have a greater selectivity for ADP than ATP (Burnstock, 2007). In concordance with ADPβS results, a selective antagonist for P2Y1 receptors MRS2179 significantly inhibited S-phase cell number mid-light increase. In contrast, antagonists selective for other ADP-activated (P2Y13) or ATP-activated receptors (P2X1/2/3 and P2Y11) were ineffective. These results suggest that the enhanced daylight S-phase cell number is principally mediated by ADP action on P2Y1 metabotropic receptors. Further evidence supporting this finding indicated that ADP is a key signal for injury-induced cell proliferation in the zebrafish retina (Battista et al., 2009). Moreover, we have also demonstrated—for the first time—that P2Y1 receptor mRNA is expressed in the zebrafish retina.

RNA Messengers for Different Plasma Membrane-Bound NTPDases Are Expressed in the Zebrafish Retina and Exhibit Daily Variations

We have previously found by immunocytochemistry that NTPDases 1 and 2 are differentially localized in mature layers as well as in the CMZ of the zebrafish retina (Ricatti et al., 2009). We show here that NTPDase1, NTPDase3, and 3 isoforms of NTPDase2 mRNAs previously described in the brain (Rico et al., 2006; Appelbaum et al., 2007) are also expressed in the zebrafish neural retina. NTPDases are the main regulating system of eNCT concentration (Zimmermann, 2006). NTPDase1 hydrolyzes triphosphate and diphosphate nucleotides with an activity ratio of 1 to 1. NTPDase2, in contrast, hydrolyzes principally ATP accumulating extracellular ADP, whereas NTPDase3 exhibits an intermediate activity (Kukulski et al., 2005). In this report, we described day-night variations of mRNA levels for all NTPDases examined, except for the mg isoform, which only showed a significant change in 2 of 3 samples and seems to be the less abundantly expressed because, as mentioned before, we had to use twice the amount of cDNA to detect it. NTPDase gene expression at mid-light doubles in average the expression observed after a short time in darkness. Presumably, NTPDase activity might be higher at daylight than at night consistent with a tighter regulation of extracellular nucleotide levels.

Different Levels of eNCT as Well as NTPDase Activity Might Be Critical to Drive CMZ S-Phase Cell Activity During the LD Cycle

A recent report has described that larval zebrafish electroretinographic activity is virtually null at night and is under circadian control (Emran et al., 2010). The authors suggest that metabolic activity is significantly diminished at night to conserve energy. In the adult zebrafish, a decreased retinal activity at night has also been reported (Li and Dowling, 1998). Presumably, less activity and lowest metabolism might imply a reduced content of extracellular ATP (which depends on intracellular ATP concentration). This fact, in addition to a lower NTPDase activity, might contribute to low levels of extracellular ADP at night. Alternatively, although we did not detect day-night variations of P2Y1 receptor mRNA levels, a decreased response to eNCT at night cannot be ruled out. Therefore, the assumed highest levels of eNCT as well as an increased NTPDase mRNA expression, which might be under circadian control, could induce S-phase cell entry in the CMZ (Martins and Pearson, 2008) at daylight or mid-subjective day for 24 h in DD. ARL67156, apyrase, hexokinase, and P receptor antagonist treatments indicated that the BrdU-positive nuclei increase chiefly depends on activation of P2Y1 receptors by ADP as well as NTPDase extracellular activity. The low number of S-phase nuclei in the CMZ at night is also regulated by extracellular ADP—probably at much lower concentrations—because exogenous ADPβS was able to increase S-phase activity to daylight levels. When eNCT were scavenged or NTPDase activity inhibited, S-phase entry in the CMZ decreased to almost null values. It is tempting to suggest that the very few remaining proliferating cells might be stem cells, whose cell cycle regulation is independent of eNCT/S.

We have demonstrated that NTPDase mRNA expression levels show day-night variations. These might be driven by a weak endogenous oscillator and/ or the photic stimulus to regulate eNCT day-night concentration and hence precursor cell S-phase entry in the CMZ. It remains to be shown, however, if light, a circadian mechanism, or both are indeed capable of regulating eNCT concentration or ATP release by retinal cells.

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NOTE

Supplementary online material for this article is available on the Journal of Biological Rhythms website: http://jbr.sagepub.com/supplemental.

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