



Published in final edited form as:

Adv Exp Med Biol. 2018 ; 1074: 345–350. doi:10.1007/978-3-319-75402-4_42.

The Retinal Circadian Clock and Photoreceptors Viability

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Abstract

Circadian rhythms are present in most leaving organisms and these rhythms are not just a consequence of the day/night fluctuation, but rather they are generated by endogenous biological clocks with a periodicity of about 24 hrs. In mammals, the master pacemaker of circadian rhythms is localized in the suprachiasmatic nuclei of the hypothalamus (SCN). The SCN controls circadian rhythms in peripheral organs. The retina also contains circadian clocks which regulate many aspects of retinal physiology, independently of the SCN. Emerging experimental evidence indicates that the retinal circadian clocks also affect ocular health and a few studies have now demonstrated that disruption of retinal clocks may contribute to the development of retinal diseases. Our study indicates that in mice lacking the clock gene *Bmal1* photoreceptor viability during aging is significantly reduced. *Bmal1* knock-out mice at 8-9 months of age have 20-30% less nuclei in the outer nuclear layer. No differences were observed in the other retinal layers. Our study suggests that the retinal circadian clock is an important modulator of photoreceptor health.

XX.1 Introduction

Circadian rhythms have been observed in animals, plants, fungi and even cyanobacteria. In mammals, including humans, the master pacemaker controlling 24-hour rhythms is localized in the suprachiasmatic nuclei of the hypothalamus (SCN). The SCN is responsible for orchestrating circadian clocks in peripheral organs to regulate physiological functions such as behavior, sleep, body temperature, blood pressure and hormone release (Herzog and Tosini 2001). Accumulating evidence indicates that dysfunction of the circadian rhythms due to genetic mutations or environmental factors (i.e., jet-lag or shift work) may contribute to the development of many serious diseases, including cancer and type-2 diabetes (Evans and Davidson 2013).

The retinal circadian clock was the first extra-SCN circadian oscillator to be discovered in mammals (Tosini and Menaker 1996). The molecular clockwork mechanism of the retinal

clock is similar to what has been reported for the SCN (Tosini et al. 2008), albeit it appears that the retinal clock is less robust (Ruan et al. 2012; Jaeger et al. 2015). Several studies have also established that many aspects of retinal physiology and function are under the control of retinal circadian clocks (see McMahon et al. 2014 for a review) and new experimental evidence suggests that other ocular structures (e.g., cornea, retinal pigment epithelium) also possess circadian clocks that control important physiological functions (Yoo et al. 2005; Baba et al. 2010; Baba et al. 2015, Buhr et al. 2015). Interestingly, as seen in the SCN, it appears that the neural retina communicates the photic information to the other ocular structures via humoral signals (e.g., melatonin and dopamine, Ruan et al. 2008, Baba et al. 2015) since most of these ocular structures are not capable of direct light transduction (Baba et al. 2010).

XX.2 The Retinal Circadian Clock and Ocular health

Similar to molecular circadian clock in SCN, the retinal clock also consists of auto-regulatory transcriptional/translational negative feedback loops involving several clock genes and their protein products which generate approximately 24 hours cycle. The primary core loop involves two basic helix-loop-helix-PAS domain transcription factors, BMAL1 and CLOCK, which heterodimerize and bind to E-box elements in promoter region to enhance transcription of *Period 1* and *2*, and *Cryptochrome 1* and *2*. The protein products, PERIOD and CRYPTOCHROME together then inhibit their own transcription by blocking CLOCK/BMAL1-mediated transactivation (see Tosini et al. 2008, Figure 1 for a schematic illustration). The second feedback loop involves the negative and positive transactivation of five other genes, *Rev-erb α , β* and *Ror α , β , c* via REV-ERB/ROR response element (RRE) promoter elements in promoter regions. REV-ERB as a negative element inhibits *Clock* and *Bmal1* transcription whereas ROR as a positive element promotes *Clock* and *Bmal1* transcription. The transcriptions of *Rev-erbs* and *Rors* are regulated via E-box elements in their promoter regions. REV-ERBs and RORs compete for binding to RRE in the *Bmal1* promoter regions to regulate rhythmic expressions of *Bmal1*. These intertwining oscillation signals also regulate transcription of other clock controlled genes via E-box or RRE elements, and the products of these genes serve as circadian clock outputs (Takahashi et al. 2008). In the mouse, clock genes are rhythmically expressed in the different retinal layers (Hiragaki et al. 2014). In the photoreceptor layer only the cones appear to express all the circadian clock proteins (Lui et al. 2012).

Emerging evidence suggests that retinal circadian clocks and their output signals contribute to retinal disease and pathology, as well as normal retinal function. For example, diabetic retinopathy is associated with reduced clock gene expression in the retina (Busik et al. 2009), circadian disruption recapitulates diabetic retinopathy in mice (Bhatwadekar et al. 2013), and removal of *Period2* induces dysfunction in the retinal microvasculature (Jadhav et al. 2016).

Trophic signaling by the retinal clock and its outputs seem to play a role in the regulation of eye growth and refractive errors (reviewed in: Stone et al. 2013). A recent study has also reported that mice lacking *Period1* and *Period2* show significant alteration in the distribution

of cone photoreceptors (Ait-Hmyed et al. 2013) and mice lacking *Rev-erb α* show a significant alteration in photoreceptor response to light (Ait-Hmyed et al. 2016).

Finally, it is worthwhile mentioning that the retinal clock influences the susceptibility of photoreceptors to light induced damage (Organisciak et al. 2000) and recent genomic studies have also implicated the clock genes *Rev-erbα* and *Rora* in retinal functioning (Mollema et al. 2011) and age-related macular degeneration (Jun et al. 2011).

XX.3 Bmal1 and retinal cell viability

As previously mentioned, *Bmal1* gene (also known as *Arntl*) is a key component of the mammalian circadian clock. *Bmal1* knock-out mice (*Bmal1*^{-/-}) do not show any circadian rhythmicity (Bunger et al. 2000) and develop several pathologies (Kondratov et al. 2006). *Bmal1*^{-/-} mice show premature aging and their lifespan is significantly reduced (about 9 months) (Kondratov et al. 2006). In the mouse retina, Bmal1 is expressed in many cell types, (Ruan et al. 2008), but within the photoreceptor layer BMAL1 was only detected in the cones (Liu et al. 2012). Storch et al. (2007) reported that many genes (more than a thousand) show a daily rhythm in mouse retina, but a large fraction of these genes are no longer rhythmically expressed or have reduced amplitude in *Bmal1*^{-/-} mice. In *Bmal1*^{-/-} mice, the day/night (circadian) changes in the amplitude of the photopic b-wave are no longer present (Storch et al. 2007). The same result has been also obtained from the mice lacking *Bmal1* only in the retina (*Chx10-Cre-Arntl*^{loxP/loxP} mice; Storch et al. 2007), thus indicating that retinal *Bmal1* is required for the circadian rhythm in visual processing. Interestingly, the photoreceptors of these mice (2-3 months) appear to be normal and unaffected by the lack of *Bmal1* (Storch et al. 2007). Additional studies have reported that in mice lacking the *Bmal1* gene there is a significant increase in the rate of cataract development and corneal inflammation during aging (Kondratov et al. 2006).

Previous studies have shown that the effects of circadian disruption become evident during the aging process (Baba et al. 2009; Musiek et al. 2013). Hence we decided to investigate whether removal of the *Bmal1* gene affects retinal cell viability during aging. Eyes from *Bmal1*^{-/-} and control mice at two different ages (3 months and 7-8 months) were obtained and then the morphometric analysis of the retina was performed according to a well-established method in our laboratory (see Baba et al. 2009 for details). As expected, and previously reported by Storch et al. (2007), young *Bmal1*^{-/-} did not show any significant variation in the number of cells in the outer nuclear layer or in any other retinal layers (Figure 1) whereas in older *Bmal1*^{-/-} mice we observed a significant reduction in the number of photoreceptor cell nuclei (about 20-30%) with respect to control. No changes were detected in the other retinal layers (Figure 1). Previous studies have shown that *Bmal1* can interact with a large number of genes (more than 1000) and therefore the phenotypes observed in *Bmal1* KOs may or may not be the consequence of a dysfunctional circadian clock (Rey et al. 2011). Thus we decided to investigate the retinal morphometry in another mouse model in which the circadian clock has been disabled. A previous investigation reported that *Clock/Npas2* KO mice do not have a functional circadian clock (DeBruyne et al. 2007, Musiek et al. 2013). Eyes from young (3months) and old (9 months) *Clock/Npas2* KO mice were obtained from Dr. David Weaver's laboratory (University of Massachusetts

Medical School) and the retinal morphometry was investigated using the same method mentioned for *Bmal1*^{-/-} mice retinas. As shown in Figure 1, *Clock/Npas2* KO mice showed an almost identical phenotype as *Bmal1*^{-/-} animals. The fact that almost identical results were obtained in *Bmal1* and *Clock/Npas2* KOs indicates that the reduced photoreceptor viability observed in *Bmal1*^{-/-} is likely due lack of a functional circadian clock in these cells and not to a possible pleiotropic effect of *Bmal1*. Our preliminary data indicate that dysfunctions of circadian clock genes may affect the photoreceptor cell viability during aging.

XX.4 Conclusions

The circadian clock is responsible for a wide variety of physiological functions and a large number of studies have now demonstrated that retinal circadian clocks regulate many functions in the eye. Experimental evidence also suggests that dysfunction of the retinal clocks may promote ocular diseases. In addition, our preliminary data indicate that genetic removal of clock genes affects photoreceptor viability during aging. Further studies are required to fully understand the role of circadian clock and its associated gene product in the regulation of retinal function and health.

Acknowledgments

Research in the author's laboratories is supported by NIH grants GM116760 to KB, EY018640 to CR, R01EY004864 and P30EY006360 to PMI, EY022216, EY026291 to GT, and an unrestricted departmental grant from Research to Prevent Blindness (Emory Department of Ophthalmology).

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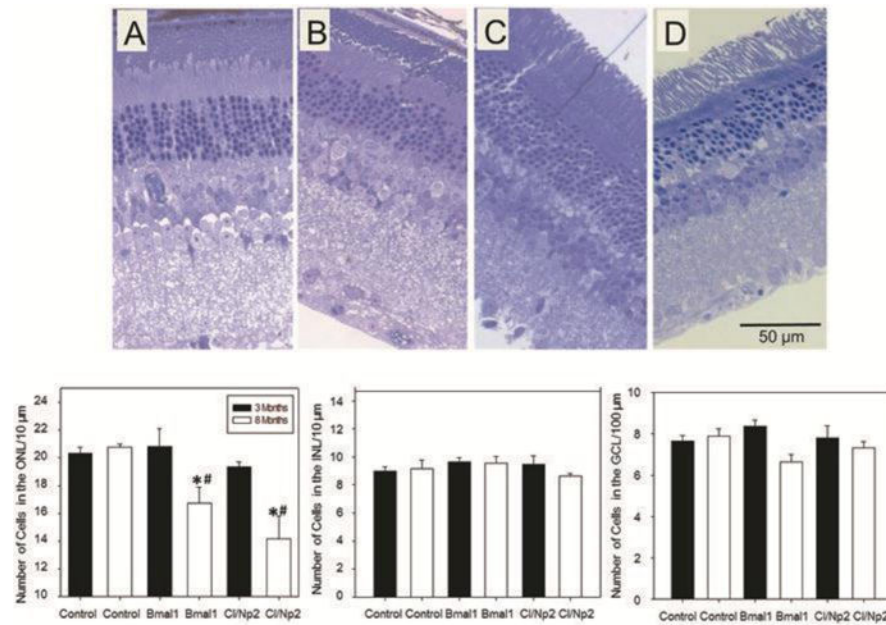


Fig. 1. Morphometric analysis of retinae obtained from young (3 months old, black bars) and old (8 months old, white bars) of control, *Bmal1*^{-/-} and *Clock*^{-/-}/*Npas2*^{-/-} (*Cl/Np2*) mice. The number of cells in the ONL of 8 months old *Bmal1* (B) and *Clock/Npas2* KOs (D) were significantly lower than the number of cells in 3 months old mice (A, C) of the same genotype (One-way ANOVA following *post hoc* test * $p < 0.05$) and age matched control group (# $p < 0.05$). No differences in the number of cells were observed in the INL and/or GCL ($n = 3-4$). The number of cells in the ONL of young *Bmal1* and *Clock/Npas2* KO mice was not different from the number of cells in control mice of the same genotype (C57BL/6). The microphotographs in A, B, C and D represent a typical example of a section obtained from 3 months old *Bmal1* KO (A), 8 months old *Bmal1* KO (B), 3 months old *Clock/Npas2* KO (C) and 8 months old *Clock/Npas2* KO (D).