



## Vertebrate Bistable Pigment Parapinopsin: Implications for Emergence of Visual Signaling and Neofunctionalization of Non-visual Pigment

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Opsins are light-sensor proteins, each absorbing a specific wavelength of light. This, in turn, drives a specific G protein-mediated phototransduction cascade, leading to a photoreceptor cell response. Recent genome projects have revealed an unexpectedly large number of opsin genes for vision and non-visual photoreception in various animals. However, the significance of this multiplicity of opsins remains largely unknown, except for that of cone visual opsins, which are diversified with respect to spectral sensitivity to achieve vertebrate color vision. Here, an implication of multiplicity is discussed, with focus on an ultraviolet-sensitive non-visual opsin-parapinopsin-that underlies pineal wavelength discrimination in lower vertebrates. Parapinopsins are phylogenetically close to vertebrate visual opsins, which have bleaching properties, but interestingly, parapinopsins are bleach-resistant bistable pigments, which photo-convert to stable photoproducts that revert to their original dark state by subsequent light absorption, similar to invertebrate rhodopsins. The unique characteristics of parapinopsin as an evolutionary intermediate between bistable and bleaching pigments provide insight into the evolutionary transition between signaling molecules that interact with two types of opsin-based pigments. Furthermore, the parapinopsin gene was duplicated in the teleost lineage and the spectral sensitivities of the duplicated parapinopsins were different from each other. On the basis of these results, together with the histochemical findings of parapinopsin, a plausible link between the diversification of a non-visual opsin parapinopsin and diverse pineal functions, wavelength discrimination, and melatonin secretion, implying why multiple opsins exist in animals, is proposed.

Keywords: bistable pigment, pineal photoreception, signal transduction, G protein, arrestin, wavelength discrimination, melatonin secretion, gene duplication

## INTRODUCTION

Many animals utilize light information for vision and non-visual functions such as circadian photoentrainment. In animal photoreception, opsin-based pigments, which are composed of a protein moiety opsin and chromophore retinal, serve as light sensors to capture light and initiate G protein-mediated phototransduction cascades in photoreceptor cells. Thousands of opsin genes

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have been identified in various animals and are phylogenetically and functionally classified into eight groups that diversified during early eumetazoan evolution (Terakita, 2005; Peirson et al., 2009; Yau and Hardie, 2009; Lamb, 2013; Koyanagi and Terakita, 2014). In addition, recent genome projects have revealed an unexpectedly large number of opsin genes in various animals, indicating that most animals have multiple opsins (Davies et al., 2015). In visual systems, the biological significance of this diversification is clear; multiple types of visual opsins with different spectral sensitivities are responsible for color vision and other visual functions (Nathans et al., 1986; Okano et al., 1992; Nagata et al., 2012; Terakita and Nagata, 2014; see also Figure 3A). However, the significance of the diversification of non-visual opsins remains unclear, despite the importance of understanding the molecular mechanism and evolution of various non-visual photoreceptions. General characteristics and diversities of the opsin family have been previously reviewed (Terakita, 2005; Peirson et al., 2009; Yau and Hardie, 2009; Lamb, 2013; Koyanagi and Terakita, 2014; Davies et al., 2015); therefore, here we addressed this issue based on the molecular properties of opsins by focusing on a pineal photopigment, parapinopsin.

## PARAPINOPSIN, A BISTABLE ULTRAVIOLET-SENSITIVE PIGMENT FOR THE PINEAL WAVELENGTH DISCRIMINATION IN LOWER VERTEBRATES

Lower vertebrates can discriminate wavelengths of light with pineals and related organs independently of image-forming color vision in the eyes (Dodt and Meissl, 1982). This so-called pineal "color discrimination" is achieved through antagonism primarily between ultraviolet (UV) and visible light reception, which cause inhibitory and excitatory effects, respectively, on neuronal firing of a specialized type of ganglion cell. Detection of the ratio of UV to visible light in environmental light has been observed in pineal organs of lamprey (Morita and Dodt, 1973) and some teleosts (Morita, 1966; Falcon and Meissl, 1981), as well as in pineal-related organs of frogs (frontal organ; Dodt and Heerd, 1962; Korf et al., 1981) and reptiles (parietal eye; Jenison and Nolte, 1980). We demonstrated that parapinopsin, which was first identified in catfish pineal and parapineal organs (Blackshaw and Snyder, 1997), is the UV-sensitive pigment underlying pineal UV reception in lamprey, through spectroscopic, immunohistochemical, and electrophysiological analyses (Koyanagi et al., 2004). Parapinopsins were also identified from rainbow trout and clawed frog, in which the pineal and related organs were reported to exhibit antagonistic chromatic responses to UV and visible light (Morita, 1966; Korf et al., 1981). In addition, parapinopsin expression was recently reported in the pineal organ of a closely related species of rainbow trout (Nakane et al., 2013). Thus, parapinopsin is considered to be a common molecular basis for UV reception in pineal wavelength discrimination. This idea is supported by the observation that parapinopsin is expressed in photoreceptor cells of the iguana parietal eye, where the ratio of UV to visible light is detected (Wada et al., 2012).

Pineal UV-sensitive opsins, parapinopsins are phylogenetically classified into the same group as vertebrate visual opsins such as rhodopsins and cone visual opsins. It is well known that vertebrate visual opsins are bleaching pigments; the photoproduct releases its retinal chromophore over time and bleaches. Interestingly, although parapinopsins are closely related to vertebrate visual opsins, parapinopsins are bistable pigments, which is common for invertebrate visual opsins; the photoproduct is stable and reverts to the original dark state by subsequent light absorption (Koyanagi et al., 2004). Spectroscopic analysis of the photoproduct properties of varied opsins revealed that the bistable property is common in animal opsins and the bleaching property was acquired during the course of vertebrate visual opsin evolution (Terakita et al., 2004). Accordingly, parapinopsins could be an evolutionary intermediate between bistable pigment and bleaching pigment, providing an opportunity to study the detailed aspects of the evolution of vertebrate visual signaling (Figure 1A).

# TRANSITION OF SIGNALING MOLECULES THAT INTERACT WITH OPSINS

In general, opsin-based pigments activate G proteins upon light absorption to initiate phototransduction cascades and the G protein-mediated signalings are terminated by binding of arrestins. Therefore, it is of interest to compare signaling molecules that couple to visual and non-visual opsins to understand the evolution of visual signaling. In vertebrate visual cells, light-absorbed visual opsin-based pigment interacts with a G protein and an arrestin that are specialized for light sensing, called visual G protein transducin (Gt) and visual arrestin, which starts and shuts off G protein-mediated visual signaling, respectively (Lamb, 2013; Figure 1A). However, a Gt-like visual G protein or visual arrestin has not yet been identified in the genomes of invertebrates, including ascidians (Nakagawa et al., 2002; Yoshida et al., 2002), which are among the invertebrates most closely related to vertebrates. These facts suggest that opsin-Gt and opsin-visual arrestin interactions were established during the course of vertebrate visual opsin evolution but questions remain regarding the detailed timing of the establishment.

In the molecular phylogenetic tree of the opsin family, one group (**Figure 1A**) is formed by vertebrate visual opsins composed of rhodopsins (RH) and four types of cone opsins, namely: middle-wavelength-sensitive (MWS) opsins, short-wavelength-sensitive 2 (SWS2) opsins, and long-wavelength-sensitive (LWS) opsins, along with several non-visual opsins, pinopsins (P), VAL-opsins (VAL), parapinopsins (PP), and parietopsins (PT). The group is generally referred to as the Gt-coupled opsin group because not only visual opsins (Lamb et al., 2016) but also pinopsins (Nakamura et al., 1999; Matsushita et al., 2000; Su et al., 2006) couple with Gt, and frog VAL-opsin has been reported to activate Gt *in vitro* (Sato et al., 2011). In contrast, parietopsin has been reported to couple with Go type G protein



in the lizard parietal eye (Su et al., 2006). In addition, it is known that mosquito Opn3 and pufferfish TMT opsin in the Opn3 group, which is the closest group to the Gt-coupled opsin group, activate Gi and Go type G proteins but not Gt in vitro (Koyanagi et al., 2013; Figure 1A). These observations, together with the phylogeny that the parietopsin subgroup underwent relatively earlier branching in the Gt-coupled opsin group, suggest that the ancestral opsin in this group was not coupled to Gt type G protein (Gt-"uncoupled" opsin; Figure 1A). We recently investigated whether parapinopsins couple with Gt by focusing on pufferfish, zebrafish, and lamprey pineal organs. Biochemical analyses demonstrated that the lamprey and teleost parapinopsins activated Gt in vitro in a light-dependent manner, similar to vertebrate visual opsins (Kawano-Yamashita et al., 2015). In the teleosts, which possess rod and cone types of Gt, Gt1, and Gt2 (Lamb et al., 2016), parapinopsin was colocalized with Gt2. In contrast, in the lamprey, which does not possess Gt2 gene (Muradov et al., 2008), parapinopsin was colocalized with Gt1 type transducin, GtS, suggesting that lamprey parapinopsin

couples with GtS instead with Gt2 (Kawano-Yamashita et al., 2015). Although there is a small difference in the Gt species, the results showed that parapinopsins couple with Gt in lamprey and teleost pineal organs. Taken together, opsin-Gt coupling could have been established before the split between parapinopsins and other opsins, including visual opsins (Figure 1A). Thus, the emergence of opsin-Gt coupling predates that of the bleaching pigment, suggesting no apparent correlation between the two events and rather, biological significance of the coupling of bistable pigment to Gt. A possible explanation for the significance is that Gt-mediated signal transduction is required for lightdependent hyperpolarization of photoreceptor cells because parapinopsin-expressing photoreceptor cells in the lamprey pineal organ exhibited hyperpolarization responses (Koyanagi et al., 2004) similar to the bleaching pigment-containing visual and non-visual cells in vertebrates. Interestingly, Gt activation by parapinopsin is provoked and terminated by UV and subsequent orange light irradiation, respectively, due to the bistable nature of parapinopsin (Kawano-Yamashita et al., 2015). A possible contribution of this behavior in pineal wavelength discrimination via Gt-mediated hyperpolarization of photoreceptor cells may provide clues to understanding why a bistable pigment acquired the ability to couple with Gt.

Investigation of arrestin as a negative regulator of parapinopsin-Gt interaction also provided functional and evolutionary clues. In the lamprey pineal organ, the fact that cells expressing parapinopsin and rhodopsin are localized in the dorsal and ventral regions of the pineal organ, respectively (Koyanagi et al., 2004), enabled a comparison of the arrestins that bind to parapinopsin and rhodopsin in the same organ. Surprisingly, ß-arrestin, which is known to bind to activated G protein-coupled receptors (GPCRs) other than opsin-based pigments (Lohse et al., 1990), was localized to parapinopsinexpressing cells (Kawano-Yamashita et al., 2011). The result contrasts the localization of visual arrestin in rhodopsinexpressing cells. In addition, the light-dependent translocation of ß-arrestin to cell membranes in parapinopsin-expressing cultured cells and to the outer segments of pineal parapinopsinexpressing photoreceptor cells was observed, suggesting that ß-arrestin binds to parapinopsin to arrest parapinopsin signaling (Kawano-Yamashita et al., 2011; Figure 1A). Interestingly, under light illumination, the granules containing parapinopsin and ß-arrestin were observed in parapinopsin-expressing cell bodies (Kawano-Yamashita et al., 2011). Because it is well known that ß-arrestin mediates clathrin-mediated GPCR internalization (Lohse et al., 1990; Ferguson et al., 1996; Goodman et al., 1996), the results suggest that the granules were generated in a lightdependent manner by ß-arrestin-mediated internalization of parapinopsins from the outer segments. The ß-arrestin-mediated internalization of parapinopsin enables to eliminate the stable photoproduct and restoring cell conditions to the original dark state. In contrast, vertebrate visual signaling does not require the elimination of photoproducts via internalization because vertebrate visual pigments convert to unstable photoproducts that spontaneously decay by chromophore dissociation. Taken together with the evolutionary scenario that the bleaching pigment evolved from a bistable pigment, vertebrate visual arrestin may have evolved from a "ß-like" arrestin by losing its clathrin-binding domain and function as an internalization mediator after emergence of the bleaching pigment.

### BIOLOGICAL SIGNIFICANCE OF NON-VISUAL OPSIN MULTIPLICITY

In the case of visual opsins, the diversification with respect to spectral sensitivity is generally involved in the evolution of color vision (e.g., the evolution of trichromatic vision in the primate lineage, **Figure 3A**) but biological significance of the multiplicity of opsins underlying non-visual photoreception remains unclear. The parapinopsin-involving findings also allow further discussion. We recently found another parapinopsin gene, *PP2*, in the zebrafish and pufferfish genomes (Koyanagi et al., 2015), in addition to orthologs of the catfish and rainbow trout parapinopsins, *PP1*s, which have been previously reported (Blackshaw and Snyder, 1997; Koyanagi et al., 2004). A comprehensive survey of parapinopsin genes in vertebrate genome databases revealed that the spotted gar possess only one parapinopsin gene (**Figure 1B**). Since the spotted gar diverged before the teleost-specific whole-genome duplication (Hoegg et al., 2004; Crow et al., 2006), this suggests that the gene duplication giving rise to *PP1* and *PP2* occurred via teleost-specific whole-genome duplication (Taylor et al., 2003; Kuraku and Meyer, 2009). Phylogenetic analyses of the Gt-coupled opsin group, including parapinopsins, and synteny analysis of genomic regions around parapinopsin gene(s) in teleosts, including the spotted gar, supported this hypothesis (Koyanagi et al., 2015).

The molecular properties of two teleost parapinopsins also allowed us to speculate the biological significance of diversification of the parapinopsin gene. Zebrafish and pufferfish PP1s and spotted gar parapinopsin had absorption maxima at 360-370 nm in the UV region (Koyanagi et al., 2015), indicating that they serve as UV-sensitive pigments, similar to lamprey parapinopsin (Koyanagi et al., 2004; Figure 2A). In contrast, zebrafish, pufferfish, and rainbow trout PP2s exhibited absorption maxima in the visible light region, with peaks at  $\sim$ 480, 460, and 460 nm, respectively, indicating that teleost PP2 is a blue light-sensitive pigment (Koyanagi et al., 2015; Figure 2B). Regardless of the absorption spectra in the dark state, all of these teleost parapinopsins were bistable pigments, like lamprey parapinopsin and unlike vertebrate visual pigments. These findings demonstrate the diversification of vertebrate bistable pigment with respect to spectral sensitivity (UV and blue sensitivity). In addition, PP1 and PP2 exhibit mutually exclusive expression in the pineal organs of zebrafish, pufferfish, and rainbow trout (Koyanagi et al., 2015). The expression profiles indicate that PP1- and PP2-expressing cells capture light independently, which enables PP2 to be involved in pineal functions apart from wavelength discrimination, which is supported by UV-sensitive parapinopsins. PP1- and PP2expressing cells were characterized with the transgenic zebrafish in which either PP1- or PP2-expressing cells were labeled (Koyanagi et al., 2015). In the transgenic zebrafish, PP1expressing cells basically possessed neuronal processes, which is consistent with their involvement in wavelength discrimination (Figure 2C). However, PP2-expressing cells rarely possessed neuronal processes, suggesting that PP2 is involved in nonneural responses rather than in neural responses, including wavelength discrimination (Figure 2D). Interestingly, further histochemical investigations revealed that PP2-expressing cells contain serotonin and aanat2, a key enzyme involved in melatonin synthesis from serotonin, whereas PP1-expressing cells did not contain either molecule (Figures 2E,F). These findings suggest that blue-sensitive PP2 is involved in lightregulation of melatonin secretion (Koyanagi et al., 2015).

The studies demonstrated that teleost PP1 is a UV-sensitive pigment and its pineal expression is conserved in three kinds of teleosts (Koyanagi et al., 2015), including rainbow trout, which has been shown to have pineal wavelength discrimination ability (Morita, 1966). Accordingly, these observations strongly suggest that PP1 underlies UV reception for pineal wavelength discrimination in teleosts. A series of findings including the involvement of parapinopsins in UV reception in the lamprey



FIGURE 2 | Functional difference between PP1 and PP2. The absorption spectra of zebrafish PP1 (A) and PP2 (B). Neuronal processes (arrowheads) were observed in many GFP-labeled PP1-expressing cells (C), but rarely in RFP-labeled PP2-expressing cells (D) in the zebrafish pineal organ. Distribution of serotonin-containing cells compared to that of PP1-expressing cells (E) and PP2-expressing cells (F) in the pineal organ of adult zebrafish. PP2-expressing cells contain serotonin, a precursor of melatonin, whereas PP1-expressing cells do not. The scale bars represent 20 µm (C,D) and 50 µm (E,F). All data are derived from Koyanagi et al. (2015).



pineal organ and iguana parietal eye (Koyanagi et al., 2004; Wada et al., 2012), suggests that UV-sensitive parapinopsin is an elemental molecule for wavelength discrimination involving UVreception in the pineal and related organs of non-mammalian vertebrates. In contrast, PP2 is suggested to underlie a different pineal function: light-regulated melatonin secretion. Because parapinopsin of spotted gar, which diverged before teleostspecific whole-genome duplication (Hoegg et al., 2004; Crow et al., 2006), is a UV-sensitive pigment (Koyanagi et al., 2015) as well as other vertebrate parapinopsins (Koyanagi et al., 2004; Wada et al., 2012), blue-sensitive parapinopsin likely evolved from a UV-sensitive parapinopsin (Koyanagi et al., 2015; **Figure 3B**). Accordingly, the findings provide a clear example of the functional differentiation among duplicated non-visual opsins; UV-sensitive PP1 could have retained the original function, and blue-sensitive PP2 acquired a novel functional role via a neofunctionalization event (**Figure 3B**). The diversification of non-visual opsins with respect to spectral sensitivity in the teleost lineage shows good contrast to that of visual opsins, which is involved in the evolution of color vision (**Figure 3**).

## CONCLUSION

A series of findings on the bistable pigment parapinopsin has improved our understanding of the evolution of both visual and non-visual opsins, and partly revealed biological significance of the multiplicity of opsins in animals. We emphasize here that

parapinopsin is the first identified bistable pigment in vertebrates (Kovanagi et al., 2004). After its discovery, many bistable pigments, such as Opn4 or melanopsin, circadian photopigment in mammals, Opn3, Opn5, and peropsin have been identified in various vertebrates and invertebrates (Koyanagi et al., 2005, 2013; Mure et al., 2007; Terakita et al., 2008; Nagata et al., 2010; Yamashita et al., 2010), suggesting that the bistable property is common in animal opsins (Terakita et al., 2004). Mutational analyses of bistable pigments, including parapinopsin, revealed the displacement of the counterion, important amino acid residue for visible light absorption of opsin-based pigment, from Glu181 to Glu113 (bovine rhodopsin numbering system) during the evolution of vertebrate visual pigments (Terakita et al., 2000, 2004). Comparison of characteristics between a bleaching pigment (bovine rhodopsin) with Glu113 counterion and a bistable pigment (parapinopsin) with Glu181 counterion revealed the correlation between the G protein activation ability and the magnitude of light-induced conformational change. The results suggested that counterion displacement resulted in higher G protein activation ability, generated by larger light-induced conformational change of the pigment (Tsukamoto et al., 2009). In addition, counterion displacement could contribute to the evolution of red-sensitive pigment because displacement from

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Glu181 to Glu113 allows acquisition of His181, an essential amino acid residue for red-sensitivity of the pigment through chloride ion binding (Terakita et al., 2000, 2004). Despite the wide distribution of the bistable nature in animal opsins and a better understanding of the molecular properties of the bistable pigment, its biological significance remains unclear. Studies of the contributions of bistability *in vivo*, such as that of parapinopsin in pineal wavelength discrimination, may provide insight into the physiological importance of the bistable property.

## **AUTHOR CONTRIBUTIONS**

All authors made substantial and intellectual contributions to this work. MK and AT wrote the paper. All authors read and approved the final manuscript.

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**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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