

The *Xenopus* ortholog of the nuclear hormone receptor *Nr2e3* is primarily expressed in developing photoreceptors

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ABSTRACT *Nr2e3* is a nuclear hormone receptor that is involved in rod photoreceptor differentiation. The *Nr2e3* gene was previously identified in humans, mice, zebrafish and chicken. In all species, *Nr2e3* expression is restricted to the retina and is believed to have a role in rod photoreceptor specification and maintenance. Here we report the identification and characterization of the *Xenopus Nr2e3*. We found that *Nr2e3* is primarily expressed in developing rod photoreceptors. In contrast to other species, *Nr2e3* is also expressed in the notochord and pineal gland during *Xenopus laevis* development.

KEY WORDS: *Nr2e3*, nuclear hormone receptor, photoreceptor, retina, pineal gland, notochord, *Xenopus*

Nr2e3 belongs to the superfamily of nuclear hormone receptors that function as ligand-dependent transcription factors (Escriva *et al.*, 2000, Kobayashi *et al.*, 1999). Nuclear receptors all share three canonical domains: an N-terminal variable domain, a zinc finger DNA binding domain (DBD) and a ligand binding domain (LBD) (Escriva *et al.*, 2000, Zhao *et al.*, 1998). Despite their common structure, nuclear receptors have distinct functions in different tissues (Escriva *et al.*, 2000, Giguere, 1999, Kobayashi *et al.*, 1999).

Many of the nuclear receptors identified to date, including *Nr2e3*, lack a known ligand and are thus named orphan nuclear receptors (Giguere, 1999). The superfamily of orphan nuclear receptors includes *COUP TF I (Nr2f1)*, *COUP TF II (Nr2f2)* and *Tlx (Nr2e1)*, a homolog of the *Drosophila* terminal/gap gene *tailless* (Chen *et al.*, 2005, Kobayashi *et al.*, 1999). *Nr2e3* was initially identified for its sequence similarity to *Tlx (Nr2e1)* (Kobayashi *et al.*, 1999).

Nr2e3 expression was found to be restricted to the retina and observed only in the outer nuclear layer (ONL) where the nuclei of photoreceptors are located (Chen *et al.*, 2005, Kobayashi *et al.*, 1999). It was initially observed in the human retina that *Nr2e3* is specifically expressed in rod photoreceptors (Bumsted O'Brien *et al.*, 2004) and later the same pattern of expression was observed in the rodent retina (Cheng *et al.*, 2004).

In the embryonic mouse retina, *Nr2e3* expression is observed early in rod precursors between E18.0-18.5, before the expression of any rod differentiation markers (Chen *et al.*, 2005, Cheng *et al.*, 2004). Similarly, in the developing zebrafish

retina *Nr2e3* expression becomes restricted to rod photoreceptors by 4 days post-fertilization (dpf), although transient expression of *Nr2e3* was detected in both cone and rod photoreceptors at 2 dpf (Chen *et al.*, 2005). Functionally, *Nr2e3* has been established to be a transcriptional activator of rod photoreceptor genes (Cheng *et al.*, 2004, Peng *et al.*, 2005). *Nr2e3* heterodimerizes with *Nr1d1* and in conjunction with *Crx* and *Nrl*, regulates the activity of the *Rhodopsin* promoter (Cheng *et al.*, 2004).

Enhanced S-cone syndrome (ESCS) is an autosomal recessive hereditary disease in which the S-cones (blue) are hypersensitive (Jacobson *et al.*, 1991, Marmor *et al.*, 1990). The hypersensitivity of S-cones in ESCS is due to a greater number of S-cones than the more normally abundant L or M-cones (Hood *et al.*, 1995). In humans, different mutations in *NR2E3* have been correlated with ESCS (Haider *et al.*, 2000, Hayashi *et al.*, 2005, Milam *et al.*, 2002, Nakamura *et al.*, 2002, Nakamura *et al.*, 2004, Sharon *et al.*, 2003, Wright *et al.*, 2004), although mutations in *NRL* have also been correlated with ESCS (Wright *et al.*, 2004). In many of the cases where the *NR2E3* gene is mutated, an increased number of S-cones, retinal dysplasia and degeneration was observed (Haider *et al.*, 2000). Mice harboring the naturally occurring *rd7* mutation also exhibit retinal dysplasia and degeneration (Akhmedov *et al.*, 2000, Haider *et al.*, 2001), reminiscent of the phenotype of the mutations observed in humans. Later studies showed that *rd7* mice harbor a deletion encompassing exons 4 and 5 of the *Nr2e3* message (Akhmedov *et al.*, 2000, Haider *et al.*, 2001)

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A

xtNr2e3	1	-----MSSPTGSLVST--EDSQSVPS---APGPAPN---PMLLCRVCGDSSSGKH
chickenNr2e3	1	-----MAASPAGSVVSAGLEDSPSGLSP---APGKALS---PVLCCRVCGDSSSGKH
mouseNr2e3	1	-----MSSPTVAASTMPVSVAAASKKESPRGWGLGEDPTGVGSLQCRVCGDSSSGKH
humanNr2e3	1	METRPALMSSPTVA--AAPAAGAASR--KESPRGWGLGEDPTGVGSLQCRVCGDSSSGKH
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xtNr2e3	45	GIYACNGCSGFFKRSVRRLLIYRCQAGTGLCPVDKAHRNQCAQLRKKCLQAGMNDAVQ
chickenNr2e3	48	GIYACNGCSGFFKRSVRRLLIYRCQAGTGLCPVDKAHRNQCAQLRKKCLQAGMNDAVQ
mouseNr2e3	53	GIYACNGCSGFFKRSVRRLLIYRCQAGTGLCPVDKAHRNQCAQLRKKCLQAGMNDAVQ
humanNr2e3	60	GIYACNGCSGFFKRSVRRLLIYRCQAGTGLCPVDKAHRNQCAQLRKKCLQAGMNDAVQ
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xtNr2e3	105	NERQPRSTAQVRLDSIDLDTSRSEHLATRDPPPS-CFOGN-NLRTPIPSAISGTLSPF
chickenNr2e3	108	NERQPRSTAQVRLDSIDLDTSRSEHLATRDPPPS-CFOGN-NLRTPIPSAISGTLSPF
mouseNr2e3	113	NERQPRSTAQVRLDSIDLDTSRSEHLATRDPPPS-CFOGN-NLRTPIPSAISGTLSPF
humanNr2e3	120	NERQPRSTAQVRLDSIDLDTSRSEHLATRDPPPS-CFOGN-NLRTPIPSAISGTLSPF
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xtNr2e3	163	HNHRFMASLMTAETCAKLEPEDADENIDVTSNDPEPFTDFQISGFPTSSPEGVYETSAR
chickenNr2e3	168	HNHRFMASLMTAETCAKLEPEDADENIDVTSNDPEPFTDFQISGFPTSSPEGVYETSAR
mouseNr2e3	164	HNHRFMASLMTAETCAKLEPEDADENIDVTSNDPEPFTDFQISGFPTSSPEGVYETSAR
humanNr2e3	171	HNHRFMASLMTAETCAKLEPEDADENIDVTSNDPEPFTDFQISGFPTSSPEGVYETSAR
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xtNr2e3	223	LLFMVAVKAKNLPVFSNLPRDQVILLEEAWSELFLLCATQWSMPLDSCPLLSPDLSSQ
chickenNr2e3	228	LLFMVAVKAKNLPVFSNLPRDQVILLEEAWSELFLLCATQWSMPLDSCPLLSPDLSSQ
mouseNr2e3	214	LLFMVAVKAKNLPVFSNLPRDQVILLEEAWSELFLLCATQWSMPLDSCPLLSPDLSSQ
humanNr2e3	227	LLFMVAVKAKNLPVFSNLPRDQVILLEEAWSELFLLCATQWSMPLDSCPLLSPDLSSQ
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xtNr2e3	283	--VHGKSVSSITIDVRLQETISRFRKSLNVDPTFACMKAVLFLFKPETRGLKDPQIENLQ
chickenNr2e3	286	--VHGKSVSSITIDVRLQETISRFRKSLNVDPTFACMKAVLFLFKPETRGLKDPQIENLQ
mouseNr2e3	274	--VHGKSVSSITIDVRLQETISRFRKSLNVDPTFACMKAVLFLFKPETRGLKDPQIENLQ
humanNr2e3	287	--VHGKSVSSITIDVRLQETISRFRKSLNVDPTFACMKAVLFLFKPETRGLKDPQIENLQ
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xtNr2e3	341	DQSQVMLQSHSRAHPSQPVRFKGLLLPSLRFITAEIRIELLFFRKTIGNTPMEKLLCD
chickenNr2e3	343	DQSQVMLQSHSRAHPSQPVRFKGLLLPSLRFITAEIRIELLFFRKTIGNTPMEKLLCD
mouseNr2e3	332	DQSQVMLQSHSRAHPSQPVRFKGLLLPSLRFITAEIRIELLFFRKTIGNTPMEKLLCD
humanNr2e3	347	DQSQVMLQSHSRAHPSQPVRFKGLLLPSLRFITAEIRIELLFFRKTIGNTPMEKLLCD
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xtNr2e3	401	MFKN
chickenNr2e3	403	MFKN
mouseNr2e3	392	MFKN
humanNr2e3	407	MFKN

B

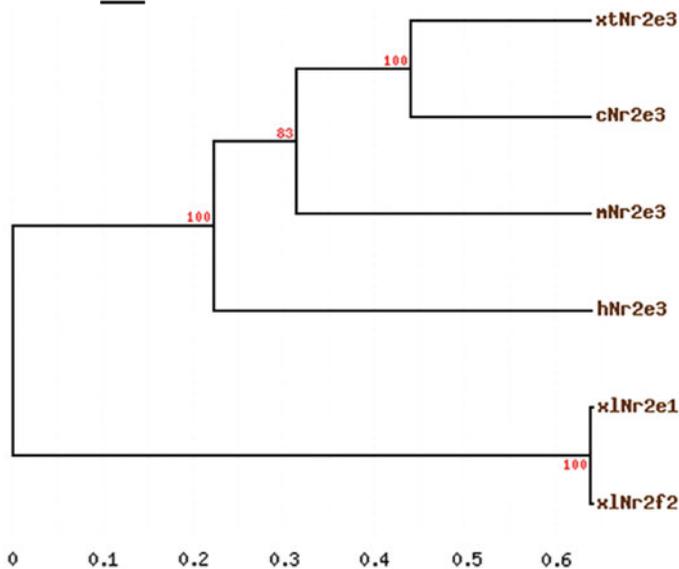


Fig. 1. Amino acid sequence comparison of Nr2e3 between different species. (A) The sequence of the *Xenopus tropicalis* Nr2e3 (xtNr2e3) gene product (Accession number: DQ906161) is compared to that of the chicken, mouse and human. Single underline denotes the conserved ligand binding domain (LBD) and the double underline denotes the conserved DNA binding domain (DBD). **(B)** Phylogenetic analysis of xtNr2e3 and the related orphan nuclear receptors Nr2e1 (Tlx) and Nr2f2 (COUP TF II). Phylogenetic analysis was performed from ClustalW alignment using the Phylip tree format in the Tree Top program (http://www.genebee.msu.edu/services/phtree_reduced.html). Species abbreviations: xt, *Xenopus tropicalis*; c, chicken; m, mouse; h, human; xl, *Xenopus laevis*.

and causes excess cone cell proliferation (Akhmedov *et al.*, 2000, Haider *et al.*, 2001).

Here we report for the first time the identification and characterization of the *Xenopus tropicalis* Nr2e3 ortholog. We also examine the expression pattern of Nr2e3 during *Xenopus laevis* development.

Results and Discussion

The predicted *X. tropicalis* Nr2e3 (xtNr2e3) gene product is most similar (76%) to the published chicken Nr2e3 protein (Figure 1A), but also has a high degree of sequence conservation with the mouse (68%) and human (67%) Nr2e3 proteins (Figure 1A). *xtNr2e3* encodes both a canonical DBD and LBD that are found throughout the orphan nuclear receptor superfamily of proteins (Figure 1A). In a phylogenetic analysis, xtNr2e3 is clustered with Nr2e3 from other species and it is excluded from the *X. laevis* Nr2e1 (Tlx) and Nr2f2 branches (Figure 1B). These findings suggest that our isolated clone encodes the frog ortholog of Nr2e3, since it is most closely related to the other vertebrate Nr2e3 gene products.

During *X. laevis* development, the expression of Nr2e3 is first observed at stage 30/31 in the center of the eye (Figure 2A). In sections taken from the wholemount *in situ* embryos, Nr2e3 expression can be confirmed to be present in the central retina (Figure 2H). This pattern of Nr2e3 expression is consistent with the development of the retina in a central-to-peripheral manner (Rapaport and Stone, 1984, Stone *et al.*, 1985), with photoreceptor cell birth following the same pattern of development (Chang and Harris, 1998, Rapaport *et al.*, 2001). At stage 33/34, Nr2e3 expression is no longer observed in the central retina, but is expressed strongly in the developing peripheral retina (Figure 2B). By stage 35/36, Nr2e3 expression is observed only in the neural retina and is clearly excluded from the lens (Figure 2C arrow). This expression pattern differs from that of Tlx (Nr2e1), which is expressed in the retina, forebrain and midbrain between stages 30/31 to 35/36 (Figure 2 D-F) (Holleman *et al.*, 1998). Nr2e3 expression was also observed in the pineal gland (Figure 2 A-C arrowheads). The pineal gland contains photoreceptor-like cells (Vigh *et al.*, 2002) and the expression of Nr2e3 in this organ is similar to the expression of Rx in the pineal gland at this stage (Figure 2G arrow) (Casarosa *et al.*, 1997, Mathers *et al.*, 1997). In addition, Nr2e3 expression was also detected in the

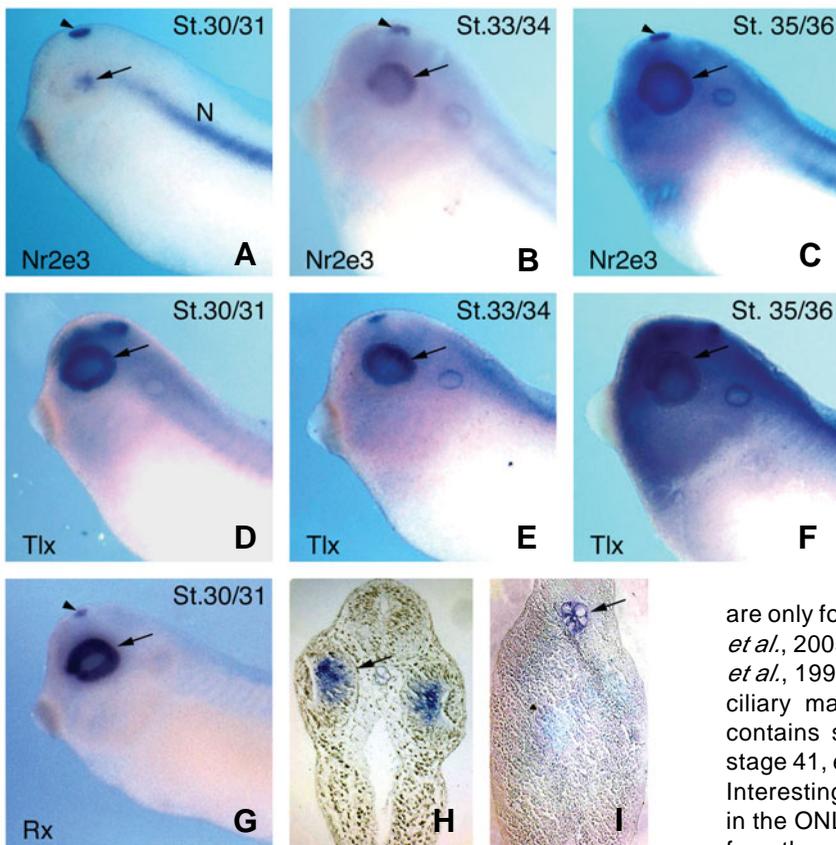


Fig. 2. Expression of *Nr2e3* in *X. laevis* embryos during development. (A-H) Lateral views of embryos subjected to wholemount in situ hybridization using antisense riboprobes specific for *Nr2e3* (A-C), *Tlx* (D-F), or *Rx* (G). Arrow denotes retinal expression and the arrowhead denotes pineal gland expression. (H, I) Transverse sections of an embryo at stage 30/31 after wholemount in situ hybridization. (H) Transverse section through the head showing retinal expression (arrow). (I) Transverse section around the gut area showing strong notochord expression (arrow). In (A-G), riboprobes and developmental stages are indicated in the lower left and upper right corner of each panel, respectively.

hybridization on retinal sections of embryos at stages 38, 41 and 45 (Figure 3). At stage 38, when the retina is not yet obviously laminated, *Nr2e3* expression is very robust in the photoreceptor layer (Figure 3A), consistent with the previous observation that *Nr2e3* transcripts are only found in photoreceptors (Akhmedov *et al.*, 2000, Chen *et al.*, 2005, Haider *et al.*, 2000, Haider *et al.*, 2001, Kobayashi *et al.*, 1999). However, *Nr2e3* expression is excluded from the ciliary marginal zone (CMZ), the region of the retina that contains stem cells and retinal progenitors (Figure 3A). By stage 41, expression of *Nr2e3* is apparent throughout the ONL. Interestingly, by stage 45, *Nr2e3* expression was only observed in the ONL just adjacent to the CMZ (Figure 3C), which differs from the expression of *Nr2e3* in the rodent and zebrafish retina where *Nr2e3* is expressed throughout the ONL once photoreceptors have differentiated and matured (Chen *et al.*, 2005, Cheng *et al.*, 2004). In contrast, *Tlx* expression is restricted to the CMZ, where *Nr2e3* is absent (Figure 3D-F). *Tlx* expression decreases as development progresses (Figure 3D-F).

notochord of the developing embryos (Figure 2 A-C, I-arrow). Expression of *Nr2e3* in pineal gland and notochord has not been reported in other species.

To further analyze the expression of *Nr2e3* in the developing retina at later developmental stages, we performed *in situ*

Double *in situ* hybridization showed overlapping expression of *Nr2e3* and *Rhodopsin* in the ONL at stage 45 (Figure 4A, B arrow). *Nr2e3* expression is observed in the photoreceptor cell body, since *Nr2e3* transcripts are restricted to photoreceptors (Akhmedov *et al.*, 2000, Chen *et al.*, 2005, Haider *et al.*, 2001, Kobayashi *et al.*, 1999). Interestingly, co-expression of *Rhodopsin* and *Nr2e3* is observed in rods at the periphery of the neural retina, but excluding the CMZ (Figure 4B). *Nr2e3* is expressed in rod precursors before differentiation and prior to expression

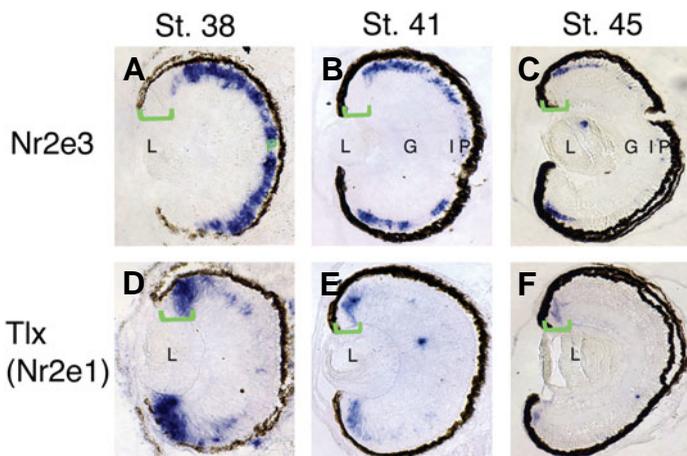


Fig. 3. (Left) *Nr2e3* and *Tlx* (*Nr2e1*) expression in retinal sections. In situ hybridization using paraffin sections of embryos at various stages during *X. laevis* development probed for *Nr2e3* (A-C) and *Tlx* (*Nr2e1*) (D-F) expression. *Nr2e3* is expressed in the neural retina, but not in the CMZ (A-C). Retinal expression of *Tlx* is restricted to the CMZ by stage 45 (arrows in (D-F)). Retinal layers and the lens are labeled with a letter abbreviation: L, lens; G, ganglion cell layer; I, inner nuclear layer; P, photoreceptor layer.

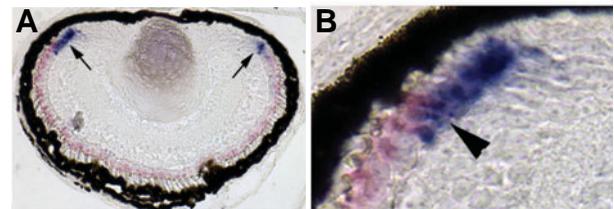


Fig. 4. (Right) Co-expression of *Nr2e3* and *Rhodopsin* in rods. (A) Double in situ hybridization for *Nr2e3* (purple) and *Rhodopsin* (red) in a section of an eye from a stage 45 embryo. Arrows denote the overlap between *Nr2e3* and *Rhodopsin* expression. (B) Higher magnification (optical and digital) of the picture in (A) showing the region of overlap between *Nr2e3* and *Rhodopsin* expression (arrowhead).

of *Rhodopsin*, which is first detected at stage 33/34 in *Xenopus laevis* (Chang and Harris, 1998).

Materials and Methods

Isolation and cloning of the *X. tropicalis* Nr2e3 ortholog

A BLAT search of the *Xenopus tropicalis* genome in the DOE Joint Genome Institute (JGI) website (<http://www.jgi.doe.gov/>) yielded a genomic locus for the *Nr2e3* gene (Scaffold 103: 742787-749884). By sequence comparison of the intron and exon arrangements of the chicken *Nr2e3*, primers (5'- caatgagttctccccacaggatc; 3'- gattgattgtctgaacatatcacaag) were designed against the first exon and last exon predicted by the BLAT search. Total mRNA was purified from heads isolated from stage 38 *Xenopus tropicalis* tadpoles (NASCO Sciences; Fort Atkinson, WI, USA) using TRIZOL (Invitrogen). RT-PCR was performed from the total RNA using the SuperScript One Step RT PCR Kit (Invitrogen). The amplified product was subcloned (TOPO TA Kit, Invitrogen), sequenced and confirmed to be full length *Nr2e3* (Submitted to PubMed; accession number: DQ906161). *Nr2e3* sequences were aligned using the ClustalW program from the European Bioinformatics Institute (EBI) website (<http://www.ebi.ac.uk/clustalw/>).

In situ hybridization

The *Nr2e3* probe used for wholemount *in situ* hybridization was generated from a *Xenopus laevis* expressed sequence tag (EST) (Accession number: CD328661) that was identified by its degree of sequence homology to the chicken Nr2e3 gene product in a BLAST search. To generate the riboprobe the Nr2e3est/pCMV-Sport6-ccdb plasmid was cut with using NotI and EcoRI restriction enzymes (New England Biolabs) to release the predicted coding sequence from the EST. The excised coding region from the Nr2e3est/pCMV-Sport6-ccdb plasmid was then cloned into the NotI and EcoRI sites of pBlueScriptII KS to generate the plasmid Nr2e3cds/pBlueScript. Nr2e3cds/pBlueScript was then linearized with EcoRI (New England Biolabs) and the antisense riboprobe made using T7 RNA polymerase plus (Ambion). Antisense riboprobes for *Rhodopsin*, *Tlx*, *Rx* were generated as described previously (Holleman *et al.*, 1998, Mathers *et al.*, 1997, Pan *et al.*, 2006, Zuber *et al.*, 2003).

Embryos were generated by *in vitro* fertilization (Sive *et al.*, 2000), staged and fixed as described previously (Nieuwkoop and Faber, 1994, Sive *et al.*, 2000). Wholemount *in situ* was performed using whole embryos as previously described using digoxigenin-labeled antisense riboprobes (Sive *et al.*, 2000, Turner and Weintraub, 1994). Briefly, hybridization and post-hybridization washes were performed at 65°C and after color reaction in BM-Purple (Roche) the embryos were post-fixed in Modified Bouin's fixative (picric acid was omitted), dehydrated and bleached (Seufert *et al.*, 2005, Shimamura *et al.*, 1994). For section *in situ* hybridization, paraffin sections (8 µm) of retinas were processed using digoxigenin-labeled antisense riboprobes as previously described (Shimamura *et al.*, 1994, Viczian *et al.*, 2003). Double section *in situ* hybridization was performed using digoxigenin-labeled *Nr2e3* and fluorescein-labeled *Rhodopsin* antisense riboprobes (2 µg/ml each). After hybridization, the retinal sections were incubated with anti-digoxigenin-AP antibody overnight (Roche), followed by the color reaction with BM Purple (Roche). After color development, sections were subjected to post-hybridization washes (beginning after RNase treatment) and incubated with anti-fluorescein-AP antibody (Roche) overnight. Rhodopsin expression was visualized using Vector Red alkaline phosphatase substrate (Vector Laboratories).

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