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Molecular Cloning of a Novel Membrane Glycoprotein, Pal, Specifically Expressed in Photoreceptor Cells of the Retina and Containing Leucine-Rich Repeat

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We have isolated a novel retina-specific gene in a screen for genes of which expression is not apparent neonatally in rat retina but is abundant postnatally on day 14 (P14). This gene, named Pal, encodes a putative type I transmembrane protein containing five leucine-rich repeats (LRRs), a single C2-type Ig-like domain, and a single fibronectin type III domain and is considered to be a new member of the LRR and Ig superfamily. No expression of Pal was found in rat retina at P1, but it was detected at P7 and markedly increased with subsequent development. These expression patterns of Pal appeared to be correlated with the development of the photoreceptor outer segments, because in the adult rat retina it was specifically localized in these segments. Ultrastructurally, Pal immunoreactivity was distributed diffusely on the disk membrane in the lamellar regions. On the basis of its structural features and localization pattern, Pal may act as a receptor for a certain trophic factor or for an adhesion molecule participating in morphogenesis. The human homolog of Pal was mapped to chromosome 10q23.2–23.3 using fluorescence in situ hybridization.

Key words: retina; Pal; leucine-rich repeat; immunoglobulin superfamily; fibronectin III; photoreceptor outer segment

The development of the neural retina goes through steps of neuronal specification and differentiation. In the rodent retina, the majority of ganglion cells, cones, and horizontal cells develop before birth, whereas most rods develop after birth (for review, see Reh, 1992; Cepko et al., 1996). Around postnatal day 7 (P7), the outer nuclear layer and the inner nuclear layer are separated by the outer plexiform layer, and the rod outer segments in which phototransduction is carried out begin to appear. At approximately the same time, the levels of expression of photoreceptor-specific genes such as rhodopsin (Hicks and Barnstable, 1987), cGMP-gated ion channel (Ahmad et al., 1990), and arrestin (Ni et al., 1992) increase rapidly. On P9 a small a-wave can first be detected physiologically by electroretinogram (ERG), and 2 d later a positive b-wave also appears (Grun, 1982). Just before the ERG can be recorded, avoidance behavior is noted as the first reaction to light (Grun, 1982). Thus, significant changes occur in the retina leading to both morphological and functional maturation in the first 10 d after birth, and the genes that begin to be expressed in the retina at this stage could play crucial roles in the specialized functions of the retina.

Isolations of retina-specific genes have been attempted over the last decade. Such genes include rhodopsin (Nathans and Hogness, 1984), transducin (Lochrie et al., 1985), and cGMP-gated channel (Cook et al., 1987; Korschen et al., 1995), which are involved in phototransduction, and peripherin/rds (Connell and Molday, 1991). These genes of which expression is not apparent neonatally in rat retina but is abundant postnatally on day 14 (P14) when development of the retina is morphologically and functionally complete. We isolated several cDNA fragments using the differential display technique. One of these was a novel gene encoding a membrane glycoprotein containing a leucine-rich repeat (LRR), a C2-type Ig-like domain, and a fibronectin (FN) type III domain, and the transcript of which was specifically expressed in the retina. This novel retina-specific protein was considered to be a new member of the LRR and Ig superfamily and was named Pal (photoreceptor-associated LRR superfamily). We report here the structural properties and the expression patterns of this gene in the retina, plus we discuss its potential function.

MATERIALS AND METHODS

Differential display. Total RNA was isolated from P1 and P14 rat retina, and the differential display was performed as described previously (Imaizumi et al., 1994). Briefly, total RNA (3 μg) was converted to cDNA with Moloney murine leukemia virus reverse transcriptase (Life Technologies, Rockville, MD). Subsequently, each pool of cDNA was amplified by PCR with 200 different arbitrary primers. After separation by 5% PAGE, cDNA bands that were amplified abundantly only from the cDNA derived from the P14 retina were recovered from the gel, ream-
plified with the corresponding primer, and cloned into the pGEM-T vector (Promega, Madison, WI).

dNA library screening. A rat retina cDNA library constructed in the Uni-ZAP XR vector (Stratagene, La Jolla, CA) was screened using the cDNA fragment obtained from the differential display as a probe by the standard methods. To obtain the sequence of human homolog of Pal, a human retina cDNA library (Clontech, Palo Alto, CA) was screened using the rat cDNA probe containing a full-length Pal coding region.

DNA sequencing and analysis. Sequencing was performed by the dideoxy chain-termination method, using the Taq dye primer cycle sequence kit (Perkin-Elmer, Norwalk, CT) and the 373A DNA sequencer (Applied Biosystems, Foster City, CA). The final sequence was confirmed from both strands. Homology searches were performed using the FASTA and BLAST programs in any available DNA or protein data bases. Possible protein motifs in the conceptual amino acids sequence were determined using the MOTIFS program with the PROSITE dictionary.

Northern blot analysis. Total RNA was extracted from developing (P1, P7, and P14) and adult (P56) rat retina and P56 rat testes. Ten micrograms of each total RNA were fractionated by electrophoresis through a 1.0% agarose/formaldehyde gel and transferred onto Immobilon-N membrane (Millipore, Bedford, MA). The membrane was hybridized with a labeled cDNA probe. After washing, the membrane was dried and autoradiographed.

In situ hybridization. Frozen sections (12 μm thick) of the rat retina (P1, P7, P14, and P56) were made and thaw-mounted onto poly-l-lysine-coated slides. Digoxigenin-labeled cRNA probes (antisense and sense) were prepared by in vitro transcription using T7 and T3 RNA polymerase from partial Pal cDNA subcloned into pBlueScript SK+ (Stratagene) in the presence of digoxigenin-labeled d-UTP (Boehringer Mannheim, Germany). Hybridization and posthybridization procedures were performed as described previously (Imaizumi et al., 1994).

Generation of anti-Pal polyclonal antibodies. To obtain polyclonal antibody against Pal, a peptide with the sequence SREPSEHQETQMGRF (amino acids 419-434) (see Fig. 1) was synthesized and conjugated to keyhole limpet hemocyanin. Two rabbits were immunized by conventional methods, and antisera was obtained. The antibody was purified by affinity chromatography using a ProtOn kit (Multiple Peptide Systems, San Diego, CA) column with immobilized Pal-derived synthetic peptide.

Preparation of membrane fractions from rat retina and immunoblotting. Frozen rat retinas were homogenized in PBS and centrifuged at 18,000 × g for 10 min. The pellets were solubilized and incubated in 10 mM Tris-acetate, pH 8.0, 1 mM EDTA, and 0.5% Nonidet P-40 for 90 min.

Preparation of membrane fractions from rat retina and immunoblotting. Frozen rat retinas were homogenized in PBS and centrifuged at 18,000 × g for 10 min. The pellets were solubilized and incubated in 10 mM Tris-acetate, pH 8.0, 1 mM EDTA, and 0.5% Nonidet P-40 for 90 min. After washing, the sections were incubated for 2 hr with FITC-labeled goat anti-rabbit IgG (Jackson ImmunoResearch Laboratories, West Grove, PA) or Alexa 546-labeled goat anti-rabbit IgG (Molecular Probes, Eugene, OR) for immunofluorescent labeling.

For immunohistochemistry of retina, the eye cups made from P0, P7, P14, and P56 rat were fixed with Zamboni solution by overnight immersion at 4°C, soaked in 30% sucrose/PBS for 4 hr, and then frozen. Retinas were sliced into 12 μm sections and adhered to poly-l-lysine-coated slides. Sections were washed in 0.02 M PBS and incubated overnight at 4°C using a 1:100 dilution of polyclonal antibody against Pal and in 0.1 μM PBS containing 3% BSA and 0.3% Triton X-100. After washing in 0.02 M PBS, the sections were incubated for 2 hr with anti-Pal polyclonal antibody, 500× for anti-Pal monoclonal antibody (Sigma), and 500× for anti-GRP78 monoclonal antibody (StressGen Biotechnologies, Victoria, British Columbia, Canada). After washing in 0.02 M PBS, the sections were incubated for 2 hr with FITC-labeled goat anti-rabbit IgG (Jackson ImmunoResearch Laboratories, West Grove, PA) or Alexa 546-labeled goat anti-rabbit IgG (Molecular Probes, Eugene, OR) for immunofluorescent labeling.

RESULTS

Isolation and sequence analysis of Pal cDNA

After screening by differential display, 12 cDNA fragments were found to be markedly amplified in the P14 rat retina. Some of these were known photoreceptor-specific genes, including rhodopsin, G-protein α subunit, and interphotoreceptor matrix proteoglycan (IMP)-1. In addition, three novel genes were identified. In the present study, one of these novel genes, which we named Pal, was selected for further analysis. The size of the initial cDNA fragment of Pal subcloned into the pGEM-T vector was 293 bp. This clone was used to screen a rat
The largest cDNA insert obtained was 4.1 kb, which was consistent with the size of the Pal transcripts detected by Northern blot analysis (4.3 kb; see Fig. 2).

Sequence analysis showed that this cDNA was a novel gene and had an open reading frame of 1869 bp encoding a putative protein of 623 amino acid residues (Fig. 1A). Its hydropathy profile revealed that Pal had two hydrophobic segments. The amino terminus segment was considered to serve as a signal sequence, and another hydrophobic region (amino acids 522–547) represented a transmembrane domain. Comparative protein database analysis showed that it has three distinct motifs in the putative extracellular region: LRR, Ig C2-like domain, and FN III-like domain (Fig. 1B).

Figure 1. Primary structural properties of Pal. A. The deduced amino acid sequences of rat and human Pal. Numbering of amino acid residues is indicated in the right column. The putative signal peptide is underlined, and the transmembrane region is boxed. Identical amino acids are indicated by colons, and conservative changes are shown as periods. The LRRs, Ig C2, and FN III domains are indicated with arrows underneath the included sequences. B. Schematic diagrams showing the structural features of Pal, ISLR, Trk, and NCAM.

Toward the amino end, Pal contained five contiguous LRRs (amino acids 64–192) characterized by the specific spacing of leucine and asparagine residues. The last repeat contained nine more amino acid residues than the common 24-residue repeat. These repeats were flanked by a cysteine-rich amino-flanking consensus region (amino acids 24–63) and a carboxy-flanking region (amino acids 193–253). One C2-type Ig-like domain (amino acids 254–346) was identified following the carboxy-flanking region. It had two cysteine residues for a putative disulfide bond at the canonical position, which is characteristic of the C2 subcategory. There was also one FN III domain (amino acids 410–521) following the Ig-like motif. In the intramembrane region, however, no recognizable structural features such as kinase domains were detected.

This overall structure containing these three motifs is considered to represent a new class of putative transmembrane protein, although combinations of two of the three motifs in transmem-
brane proteins have been identified as Trk (Schneider and Schweiger, 1991) or NCAM (Small et al., 1987) (Fig. 1B). A homology search showed the closest relationship to ISLR, which is a recently cloned putative membrane protein in the human retina containing both the LRR and Ig-like motif (Nagasawa et al., 1997).

The human homolog of Pal cDNA also contained an open reading frame of 623 amino acids. Comparison of the human and rat deduced protein sequences showed that they were highly homologous, with an overall homology of 78.8%. The functional domains found in the rat Pal sequence were also contained in the human sequence.

The sequences of rat and human Pal cDNAs have been deposited with the DNA data bank of Japan/European Molecular Biology Laboratory/GenBank nucleotide sequence databases under accession numbers AB028461 and AB031547, respectively.

Expression of Pal mRNA in the retina

To confirm Pal expression in the developing retina, RNA from P1, P7, P14, and P56 rat retina was analyzed by Northern blotting using rat Pal cDNA as a probe (Fig. 2A). No signals were found at P1. A single transcript of 4.3 kb was observed at P7, with the signal showing marked increases with subsequent development. In agreement with the findings of Northern blot analysis, in situ hybridization of developing rat eye sections demonstrated the same expression pattern (Fig. 2B); i.e., no positive signals were detected at P1 but the expression was found at P7 in the presumptive photoreceptor layer. Increased signal intensity was seen in the outer nuclear layer and the inner segment at P14 and P56. These observations indicated that Pal mRNA was specifically expressed in the photoreceptor cells.

To examine the tissue specificity of Pal expression, various adult rat tissue RNAs were analyzed by Northern blotting. Among the 10 different tissues analyzed, expression of Pal mRNA was observed only in the retina (Fig. 2C), indicating that Pal is a retina-specific gene.

Cellular and subcellular distribution of Pal in the retina

To examine the localization of Pal protein in rat retina, we generated a polyclonal antibody against the Pal peptide sequence (419 – 434). We performed immunoprecipitation and immunoblotting assays of membrane fractions of rat retina. Anti-Pal antibody reacted with a single band of 75 kDa in the membrane fractions of the retina (Fig. 3A) and did not cross-react with other molecules containing the FN III domain. In contrast, immunoblotting of brain membrane fractions showed no Pal immunoreactivities. Thus, we confirmed that Pal was specifically expressed in the retina and that this antibody could successfully detect the endogenous Pal protein.

The distribution of Pal protein in the retina was examined immunohistochemically. No immunoreactive cells were observed in retina at P1, but the presumptive photoreceptor outer segments showed weak Pal immunoreactivity at P7 (Fig. 3B). At P14 and P56, anti-Pal antibody intensely labeled the outer segments of the photoreceptor cells, showing that Pal protein is localized and functions in the outer segments of the photoreceptor cells. Pre-
immune sera did not stain photoreceptor cells in this region (Fig. 3C). In addition, no immunoreactivity was seen when the antibody was preabsorbed with an excess amount of antigen (data not shown).

To demonstrate ultrastructural localization of Pal in photoreceptor cells, the immunogold labeling method for electron microscopy was used. LR White resin-embedded rod photoreceptor cells were labeled with the polyclonal antibody and immunogold particles. Labeling was found in the outer segment (OS, left) but not in the inner segment (IS, middle). The particles were specifically localized on the disk membranes (right). Scale bar, 200 nm.

**Molecular topology of Pal in mammalian cells**

Pal contains a signal sequence at its N-terminal region. The molecular topology of Pal on the disk membrane was presumed to be that the N-terminal region was oriented toward the disk lumen. To confirm this, we examined the orientation of Pal on the membrane in HeLa cells that were transiently transfected with the expression plasmids for Pal tagged with the FLAG epitope.

Cells expressing Pal showed no morphological changes. Pal immunoreactivities were not observed on the plasma membrane under either permeabilized or nonpermeabilized conditions, suggesting that Pal is not translocated to the plasma membrane (Fig. 5). In contrast, when the cells were permeabilized by Triton X-100, both anti-Pal antibody, which recognizes the N-terminal side of Pal, and anti-flag antibody, which recognizes the C-terminal side of Pal, detected Pal molecules at the perinuclear regions (Fig. 5). The double staining of Pal and GRP78, ER markers, showed that these proteins were colocalized in the ER. These data indicate that Pal is predominantly localized on the ER. We also examined Pal immunoreactivities under conditions that just permeabilized the plasma membrane, using saponin as the detergent (Fig. 5) (Radhakrishna and Donaldson, 1997). Anti-flag antibody detected the Pal molecules. In contrast, anti-Pal antibody failed to detect the Pal molecules, indicating that the N-terminal portion of Pal is oriented toward the lumen of the ER.
and the C-terminal portion is oriented toward cytoplasm. From the membrane architecture of the outer segment, it was considered that the molecular topology of the transmembrane proteins were identical at the ER and the disk of photoreceptor cells. Taken together, this suggested that the N-terminal portion of the Pal molecule could be oriented toward the interior of the disks.

**Biochemical characteristics of Pal protein**

Pal is considered to be capable of binding to itself or to other adhesion molecules because it contains a C2-type Ig-like domain and an FN domain. We examined the homodimerization activity of Pal by transient transfection.

The expression plasmids for Pal tagged with the FLAG epitope sequence were transfected into 293T cells. Western blotting with anti-FLAG antibody showed both a major 75 kDa band and a minor 150 kDa band (Fig. 6A). A Pal expression plasmid tagged with the HA epitope (Pal-HA) was also constructed. 293T cells were transiently transfected with the indicated plasmids. Transfection was performed with equal amounts of plasmid DNA using empty plasmid as a control. To confirm this, expression plasmids were transfected into HEK293T cells (right). Western blotting with anti-FLAG antibody showed both a major 75 kDa band and a minor 150 kDa band (arrowheads). Asterisk indicates nonspecific signals.

**Chromosomal mapping**

The initial fluorescence in situ hybridization (FISH) experiment resulted in specific labeling of the long arm of chromosome 10. A second experiment was conducted in which a probe, which had previously been mapped to 10q25, was cohybridized with Pal. Measurements of specifically labeled chromosome 10 demonstrated that Pal was located at a position that was 49% of the distance from the centromere to the telomere of chromosome arm 10q, an area that corresponds to 10q23.2–23.3 (Fig. 7). A total of 80 metaphase cells were analyzed, with 70 exhibiting specific labeling.
DISCUSSION

We have isolated a novel retina-specific gene, Pal. Primary structure analysis indicated that the Pal protein contains unique functional domains such as LRRs, Ig C2, and FN III. LRR is a repetitive motif made up of several copies of amino acid residues with the conserved sequence LxxLxxLxxNlxLxxLxxxxFx. This motif is thought to be involved in highly specific protein–protein interactions or cell adhesion (for review, see Kobe and Deisenhofer, 1994). The Ig C2 and FN III domains have also been implicated in adhesion or binding to other proteins (for review, see Williams and Barclay, 1988; Holness and Simmonds, 1994). These motifs can be adapted to specific functions by variation and repetition and by combination with other motifs. Members containing both LRR and the Ig C2 domain have only been cloned from a few genes such as Trk receptors [trkA (Martin-Zanca et al., 1989); trkB (Klein et al., 1989); trkC (Lam-balle et al., 1991)], LIG-1 (Suzuki et al., 1996), ISLR (Nagasawa et al., 1997), and Drosophila ekc1/2 (Musacchio and Perrimon, 1996). Most of these proteins play significant roles in differentiation and development of tissues such as the CNS. However, other than in Trk receptors, the mechanisms by which these genes are involved in such phenomena have not been determined. Trk receptors are transmembrane glycoproteins containing LRR, Ig C2, and a cytosolic tyrosine kinase domain (Schneider and Schweiger, 1991). Neurotrophins selectively bind to Trk receptors and transduce signaling through phosphorylation by their tyrosine kinase domain (Kaplan and Stephens, 1994). The LRR and Ig-like domain of Trk receptors have been indicated as being important for specific ligand binding (Pérez et al., 1995; Urfet et al., 1995; Windisch et al., 1995). It has also been shown that Trk receptors function as homodimers (Jing et al., 1992, Schlessinger and Ullrich, 1992). Although Pal lacks the tyrosine kinase domain, its domain organization is similar to that of Trks. In addition, Pal molecules expressed in mammalian cells revealed dimerization similar to Trks. Therefore, it is possible that Pal may act as a receptor for a certain trophic factor.

The cells expressing Pal were photoreceptor cells in the retina. In the development of the retina, Pal expression begins at P7 and increases with developmental stage; i.e., the expression of Pal corresponds to the fact that photoreceptor cells respond to light and perform phototransduction. Thus, it is likely that Pal might take part in phototransduction, or be indirectly associated with the phototransduction system through morphogenesis or maintenance of photoreceptors.

Ultrastructurally, the protein encoded by the Pal gene was distributed in the lamellar region of the disk in photoreceptor cells, a region in which rhodopsin captures a photon and initiates the phototransduction. The N-terminal region of Pal containing LRRs, Ig C2, and FN III is presumed to be oriented toward the intradiskal space from the results of overexpression of Pal in HeLa cells. Several possible roles are suggested from its structure and localization. First, Pal may bind to some molecules in the intradiskal space and transduce the signals to cytoplasm like a receptor for trophic factors. Second, Pal may interact with other disk membrane proteins such as rhodopsin at its intradiskal domain. Rhodopsin is distributed diffusely and abundantly on the lamellar region of disks. The contact to disk membrane proteins may contribute to maintain the membrane structure of disks. Another possibility is that Pal might play a role as an adhesion molecule. NCAM is representative of such an adhesion molecule containing both Ig C2 and FN III domains. NCAM also shows homophilic interaction and is considered to stimulate neuronal outgrowth and differentiation by activating some signaling pathways (Walsh and Doherty, 1997). Therefore, Pal also may mediate signals through homophilic binding. However, we cannot exclude other functions of Pal on photoreceptor disks. Further studies are required to determine the precise roles of Pal on the disk membrane. To clarify the roles of this molecule, the generation of Pal-deficient mice is currently being undertaken.

Mutations in approximately 20 genes have been found to cause forms of nonsyndromic RP (for review, see van Soest et al., 1999). A recent update of all genes involved in retinal diseases can be found through RetNet (http://www.sph.uth.tmc.edu/RetNet/). Because almost half of the identified RP genes encode photoreceptor-specific proteins, the human homolog of Pal is a potential candidate disease gene for inherited retinal degenerations. The Pal gene was mapped to chromosome 10q23.2–23.3 by FISH. Although no RP genes have been mapped to this chromosomal location, it is believed that other genes may be involved in RP. In addition, genes involved in Usher syndrome types 1f and 1d are known to be localized to 10 and 10q, respectively (Wayne et al., 1996, 1997). Usher syndrome is an autosomal recessive disorder, and affected individuals develop sensorineural hearing deficiencies and progressive RP. Because we did not examine the expression of Pal in auditory systems, it is not known whether mutations in Pal cause damage to these systems. Further investigations are needed to elucidate whether mutations in the Pal gene are linked to Usher syndrome or other diseases.

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