

Identification of components of a phosphoinositide signaling pathway in retinal rod outer segments

(phospholipase C/G protein/visual transduction)

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ABSTRACT Phototransduction in retinal rods involves a G protein-coupled signaling cascade that leads to cGMP hydrolysis and the closure of cGMP-gated cation channels that are open in darkness, producing a membrane hyperpolarization as the light response. For many years there have also been reports of the presence of a phosphoinositide pathway in the rod outer segment, though its functions and the molecular identities of its components are still unclear. Using immunocytochemistry with antibodies against various phosphoinositide-specific phospholipase C (PLC) isozymes ($\beta 1-4$, $\gamma 1-2$, and $\delta 1-2$), we have found PLC $\beta 4$ -like immunoreactivity in rod outer segments. Similar experiments with antibodies against the α -subunits of the G_q family of G proteins, which are known to activate PLC $\beta 4$, have also demonstrated $G_{\alpha 11}$ -like immunoreactivity in this location. Immunoblots of total proteins from whole retina or partially purified rod outer segments with anti-PLC $\beta 4$ and anti- $G_{\alpha 11}$ antibodies gave, respectively, a single protein band of the expected molecular mass, suggesting specific labelings. The retinal locations of the two proteins were also supported by *in situ* hybridization experiments on mouse retina with probes specific for the corresponding mouse genes. These two proteins, or immunologically identical isoforms, therefore likely mediate the phosphoinositide signaling pathway in the rod outer segment. At present, $G_{\alpha 11}$ or a $G_{\alpha 11}$ -like protein represents the only G protein besides transducin (which mediates phototransduction) identified so far in the rod outer segment. Although absent in the outer segment layer, other PLC isoforms as well as $G_{\alpha q}$ (another G_q family member), are present elsewhere in the retina.

Visual transduction in the retina takes place in the outer segments of rod and cone photoreceptors and is known to involve a cGMP signaling cascade (see ref. 1 for a recent review). In this process, light isomerizes the visual pigment into an active form, which, via the G protein transducin, stimulates a cGMP-phosphodiesterase to lead to cGMP hydrolysis. In darkness, cytoplasmic cGMP binds to and opens cGMP-activated cation channels on the plasma membrane of the outer segment. These open channels sustain an inward dark current to keep the cell partially depolarized and maintain a steady release of glutamate from the synaptic terminal of the photoreceptor. In the light, the hydrolysis of cGMP leads to the closure of these channels, producing a membrane hyperpolarization as the light response and reducing the glutamate release from the cell.

Over the years, however, there have also been reports of a phosphoinositide signaling pathway in the rod outer segment. In this pathway, the membrane phospholipid phosphatidylinositol-

4,5-bisphosphate is hydrolyzed by phospholipase C (PLC) enzymes to release the second messengers inositol-1,4,5-trisphosphate and diacylglycerol, with the former leading to intracellular Ca^{2+} release and the latter activating the enzyme protein kinase C (PKC). In rod outer segment preparations, phosphoinositide-specific PLC activity has been observed (2–7) and reported to be activated by light (2–5). This enzyme in the rod outer segment has also been characterized biochemically (6–8) and immunologically (8). Finally, PKC activity in the same location has been reported (9–12), with putative substrates including rhodopsin (9, 13–15), the α -subunit of transducin (16), the inhibitory γ -subunit of the cGMP phosphodiesterase (17–19), guanylate cyclase (20), and arrestin (21), all of which are involved in rod phototransduction. Despite these findings, the identities of the components of the phosphoinositide pathway have remained unknown. To address this question, we have undertaken an immunocytochemical approach, using specific antibodies directed against different PLC isoforms. As presently known, these enzymes can be divided according to their amino acid sequences and functional properties into three families, designated β , γ , and δ (see refs. 22–24 for recent reviews). The PLC β family is coupled to seven-transmembrane-helix receptors through the α -subunits of the pertussis toxin-insensitive G_q family of G proteins and/or through G protein $\beta\gamma$ -subunits. The PLC γ family, on the other hand, is activated by growth factor receptors via tyrosine phosphorylation. The activation mechanism for the PLC δ family remains unknown. In this study, we have found that, even though almost all of the PLC isozymes examined are in the retina, only PLC $\beta 4$ is present in the rod outer segment. This has led us also to examine several α -subunits of the G_q family of G proteins for colocalization in the rod outer segment. Only $G_{\alpha 11}$ -like immunoreactivity was found in this location. These findings have previously been reported in abstract form (25).

METHODS

Antibodies. The anti-PLC $\beta 1$, $\beta 2$, $\beta 3$, $\beta 4$, $\gamma 2$, and $\delta 2$ antibodies are rabbit polyclonal antisera, and the anti-PLC $\gamma 1$ and $\delta 1$ antibodies are mouse monoclonal antibodies. Two anti-PLC $\beta 4$ antibodies were used and gave identical results in the immunocytochemistry and immunoblotting experiments described here. Of the two, the first (Ab1) is against purified bovine retinal PLC $\beta 4$ (26), and the second (Ab2) is a mixture of two antibodies against, respectively, synthetic peptides (AVFDRYEEESFV and VKLE-AEMDRRATV) based on the N and C termini of the cloned rat PLC $\beta 4$ protein (27). In immunoblots, both Ab1 and Ab2 (as an example, see Fig. 1 for Ab2) clearly recognized the purified PLC $\beta 4$ protein from bovine retina as well as HPLC fractions from rat uterus that showed high PLC activity. All other anti-

Abbreviation: PLC, phospholipase C.

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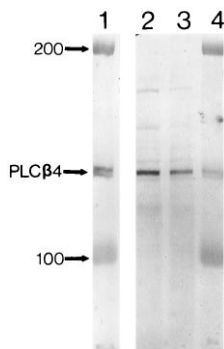


FIG. 1. Immunoblotting of purified PLC β 4 from bovine retina (lanes 1 and 4) and HPLC fractions from rat uterus that show high phospholipase C activity (lanes 2 and 3), using the Ab2 antibody. The numbers indicate protein markers of molecular masses 100 and 200 kDa. See refs. 26 and 27 for the methods of collecting the HPLC fractions and assaying phospholipase C activity.

bodies are against the respective bovine proteins. The characterization of these antibodies has been described elsewhere (26–29). Three rabbit polyclonal antibodies against several α -subunits of the G_q family of G proteins were used, kindly provided by John H. Exton (Vanderbilt University, Nashville, TN). These antipeptide antibodies, designated E973, E976, and 24, recognize $G_{\alpha q}$, $G_{\alpha 11}$, and $G_{\alpha 14}$, respectively; the specificities of two of these antibodies (E973, E976) have been described (30, 31). Finally, the antibody PMc1D1, kindly provided by Robert S. Molday (University of British Columbia, Vancouver) and directed against the rod cGMP-gated channel (32), was used to check the degree of purity of partially purified rod outer segment preparations (see *Results*).

Immunocytochemistry. All immunocytochemistry was performed on frozen retinal sections prefixed with 4% paraformaldehyde (33). The described results are for bovine retina, but similar results were obtained from rat, mouse, and rabbit retinas. For immunoperoxidase staining, the sections were first incubated with 5% normal goat serum (Vector Laboratories) in PBS for 1 hr at room temperature to reduce background staining. The sections were then incubated overnight at 4°C with the primary antibody (1:10,000 dilution for anti-PLC β 1, 1:2000 for anti-

PLC β 2, 1:3000 for anti-PLC β 3, 1:2000 for anti-PLC β 4, 1:500 for anti-PLC γ 1, 1:5000 for anti-PLC γ 2, 1:400 for anti-PLC δ 1, and 1:2000 for anti-PLC δ 2, $G_{\alpha q}$, and $G_{\alpha 11}$; no specific staining with anti- $G_{\alpha 14}$ was found). This was followed by two washes in PBS, each for 30 min. Triton X-100 (0.3%) was added to all incubation and wash buffers. Depending on the primary antibody, the sections were then incubated with a biotin-conjugated goat anti-rabbit or anti-mouse secondary antibody (Vector Laboratories; 1:200 dilution) for 2 hr at room temperature, then washed twice in PBS for 30 min each, followed by a 1-hr incubation with an avidin-biotin-peroxidase complex (Vector Laboratories; 1:100 dilution) in PBS. After two more washes for 30 min each, the stain was developed with a substrate solution of 20 ml PBS, 0.1 ml 3% H_2O_2 , and 10 mg diaminobenzidine. The staining reaction was terminated by washing with PBS, and the sections were covered with 50% glycerol in PBS. For each antibody, the specificity of staining was also confirmed by preadsorbing the corresponding antigen to the antibody, which resulted in no staining.

Immunoblotting. Immunoblotting experiments on total proteins from whole bovine retina or partially purified bovine rod outer segments were carried out with the antibodies against PLC β 4, $G_{\alpha 11}$, $G_{\alpha q}$, and PLC γ 2, as well as PMc1D1. For total proteins, a bovine retina was homogenized in 5% SDS, 1 mM benzamide, 0.5 mM phenylmethylsulfonyl fluoride, and 1 mM EDTA in Tris-buffered saline (TBS). Insoluble material was removed by centrifugation at $13,000 \times g$ for 10 min. Proteins were assayed using Pierce reagent. SDS/PAGE was performed on 1.5-mm thick, 5–16% polyacrylamide gels, and the separated proteins were transferred to nitrocellulose membranes (34). The blots were blocked with 5% nonfat dry milk in TBS for 2 hr before an overnight incubation at 4°C with the primary antibodies diluted in 3% BSA in TBS (1:2000 for all three antibodies). Afterwards, they were washed three times for 10 min each in 5% nonfat dry milk in TBS, then incubated with a horseradish peroxidase-linked, goat anti-rabbit or anti-mouse secondary antibody for 1 hr at room temperature. After three washes of 10 min

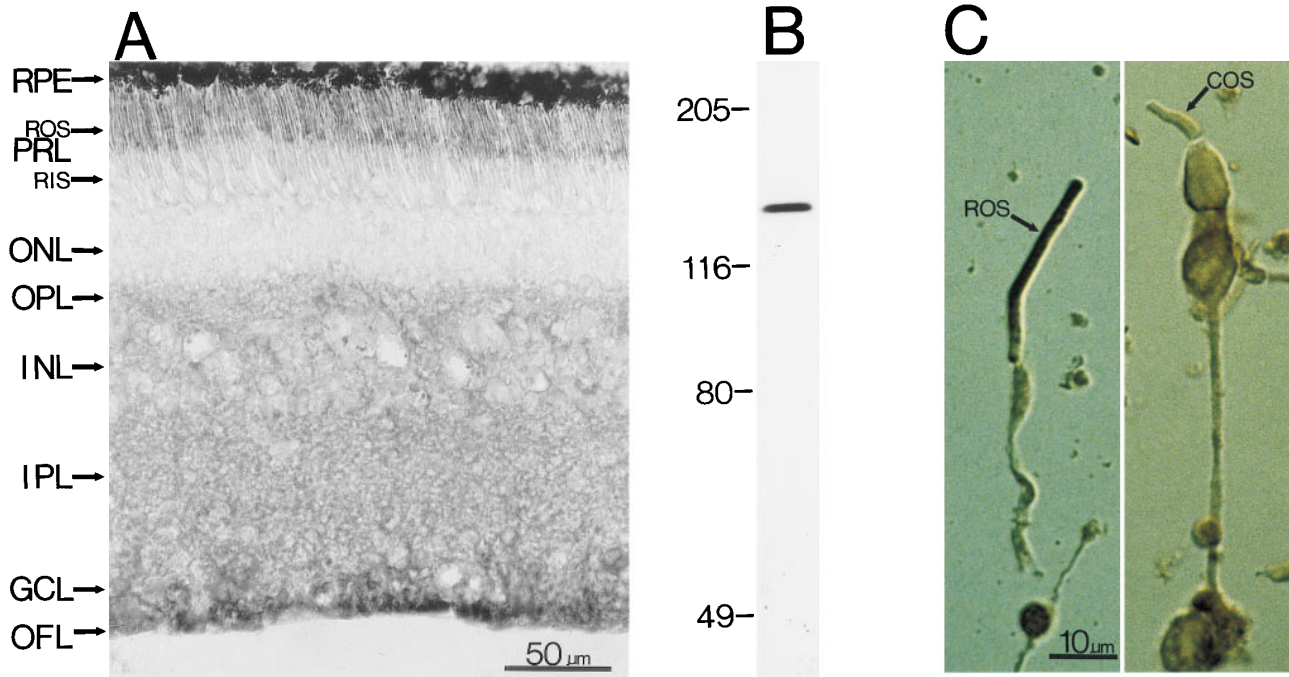


FIG. 2. (A) Immunostaining of a cross-section of the bovine retina with an antibody against PLC β 4. (Nomarski differential interference contrast optics; 8- μ m frozen section.) The anatomical layers are as follows: RPE, retinal pigment epithelium; PRL, photoreceptor layer containing predominantly the outer segments (ROS) and inner segments (RIS) of rods, though cones are also present in far fewer numbers; ONL, outer nuclear layer containing the cell bodies of the photoreceptors; OPL, outer plexiform layer; INL, inner nuclear layer; IPL, inner plexiform layer; GCL, ganglion cell layer; OFL, optic fiber layer. (B) Immunoblotting of total bovine retinal proteins with the anti-PLC β 4 antibody. Sizes of molecular mass standards ($\times 10^{-3}$) are shown adjacent to the lane. (C) Dissociated rod (Left) and cone (Right) cells stained with the same antibody. Only the rod outer segment (ROS) shows staining, but apparently not the cone outer segment (COS).

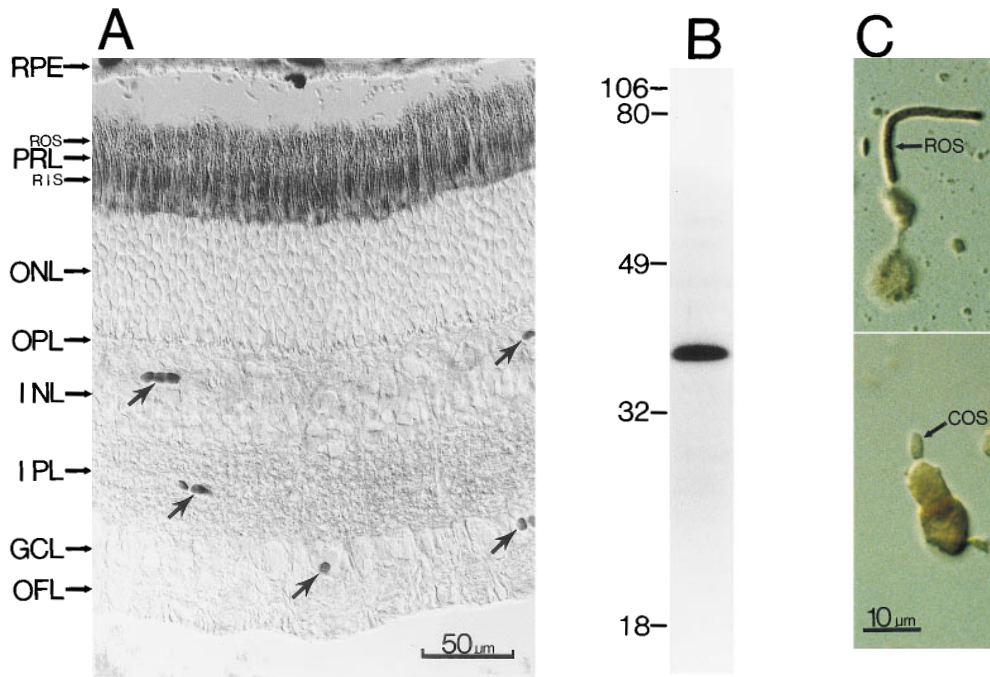


FIG. 3. (A) Immunostaining of a cross-section of the bovine retina with an antibody against $G_{\alpha 11}$. (Nomarski optics; 8- μ M frozen section.) The anatomical layers are as in Fig. 2. The arrows indicate blood vessels. (B) Immunoblotting of total bovine retinal proteins with the anti- $G_{\alpha 11}$ antibody. Sizes of molecular mass standards ($\times 10^{-3}$) are shown adjacent to the lane. (C) Dissociated rod (Top) and cone (Bottom) cells stained with the same antibody. Again, only the rod outer segment (ROS), but apparently not the cone outer segment (COS), shows staining.

each in TBS, the blots were developed using 4-chloro-1-naphthol as the substrate. Proteins from partially purified rod outer segments were treated in a similar way. The method for purifying rod outer segments has been described elsewhere (35).

In situ Hybridization. Frozen sections of paraformaldehyde-fixed mouse retina were prepared as described above, but with standard procedures to eliminate RNase activity. The sections were rinsed in 2 \times standard saline citrate (SSC; 1 \times SSC = 0.15 M sodium chloride/0.015 M sodium citrate, pH 7) and incubated

in 2–10 mg/ml proteinase K in Tris-EDTA for 15–20 min at 37°C. The slides were then washed in 2 \times SSC and treated with 0.1 M triethanolamine/0.25% acetic anhydride. The sections were then covered with a sense or antisense RNA probe diluted to 1 mg/ml in hybridization buffer, coverslipped, and sealed with nail polish. These sealed slides were incubated overnight at 50°C. The coverslips were then removed and the slides washed in 2 \times SSC. After the sections were incubated in 20 mg/ml RNase A in RNase buffer for 30 min, the slides were washed successively in 2 \times SSC, 1 \times SSC, and 0.1 \times SSC at 50°C. After washing at room temperature in PBS/0.1% Tween 20, the sections were blocked in 5% normal goat serum. An anti-digoxigenin antibody conjugated with alkaline phosphatase (Boehringer Mannheim, 1:500 diluted in PBS) was then added to the slides and incubated overnight. The slides were developed for 2–4 hr with color reaction and sealed with coverslips.

To prepare RNA probes for *in situ* hybridization, PCRs were performed using mouse brain cDNA as template to obtain fragments corresponding to published sequences for nucleotides 441–763 of mouse $G_{\alpha 11}$ and 313–622 of rat PLC $\beta 4$. The PCR products were then subcloned into a TA cloning vector (pCR II; Invitrogen). The resulting plasmids showed 100% and 97% identities, respectively, to published mouse $G_{\alpha 11}$ and rat PLC $\beta 4$ sequences. Digoxigenin-labeled RNA probes, sense or antisense, were made with a commercial kit (Boehringer Mannheim).

Dissociated Cells. The isolated bovine retina was incubated at room temperature in DMEM (GIBCO) supplemented with 10 units/ml papain (Worthington), 1.2 mM EDTA, and 5.5 mM cysteine. After 45 min of incubation, the retina was washed with DMEM containing bovine serum albumin (0.1 mg/ml). Dissociation of the treated retina into individual cells was then effected by gentle trituration with a wide-bore transfer pipette. Aliquots of freshly dissociated cells were placed in a test tube and fixed for 2 hr with 4% paraformaldehyde in phosphate buffer at 4°C. The fixed cells were pipetted onto poly-D-lysine-coated slides and left to settle for 2 hr. The subsequent immunostaining procedures were identical to those described for retinal sections.

RESULTS

PLC $\beta 4$ - and $G_{\alpha 11}$ -like Immunoreactivities in Rod Outer Segments. Among all of the PLC isozymes we have examined

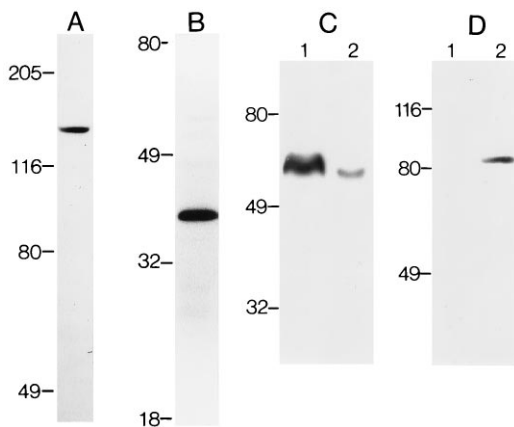


FIG. 4. Immunoblotting of total proteins from partially purified bovine rod outer segments. (A) With the anti-PLC $\beta 4$ antibody. (B) With the anti- $G_{\alpha 11}$ antibody. (C) Experiment examining the degree of purity of the partially purified rod outer segment preparation. Lane 1, partially purified rod outer segments; lane 2, total retinal proteins. Both are stained with the PMc1D1 antibody against the rod cGMP-gated channel, known to be present predominantly in the rod outer segment (38). Densitometric measurements indicated that the intensity of staining was approximately 10:1 between lanes 1 and 2. (D) Experiment and display similar to C, but stained with an antibody against PLC $\delta 2$, which is predominantly in Müller cells rather than rod outer segments (see Results). Sizes of molecular mass standards ($\times 10^{-3}$) are shown adjacent to the lanes.

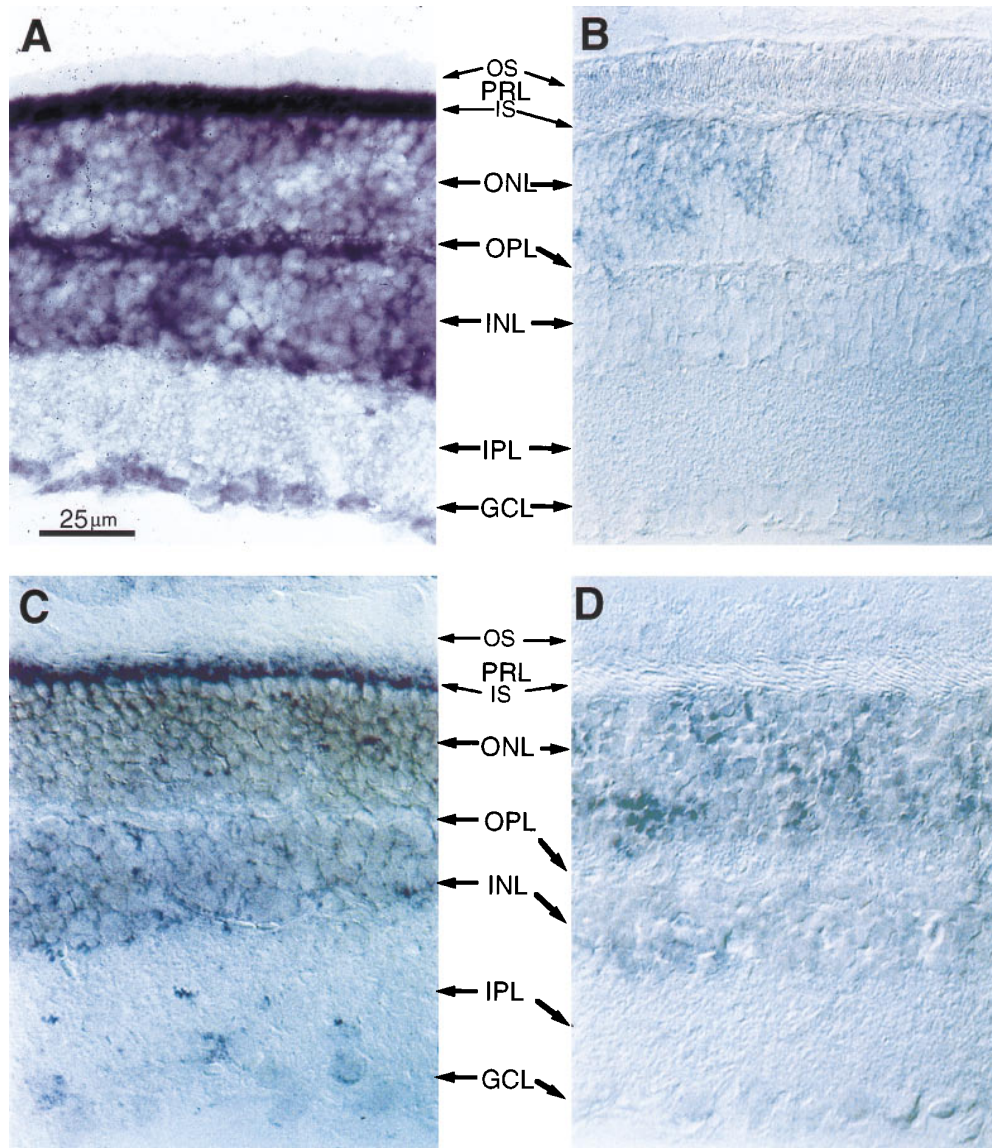


FIG. 5. *In situ* hybridization of mouse retinal sections with digoxigenin-labeled riboprobes. The anatomical layers are as in Fig. 2. (A) Bright-field optics; with antisense riboprobe for PLC β 4. (B–D) Nomarski optics; 8- μ M frozen sections. (B) With sense riboprobe for PLC β 4. (C) With antisense riboprobe for G α 11. (D) With sense riboprobe for G α 11.

(PLC β 1–4, γ 1–2, and δ 1–2), only PLC β 4-like immunoreactivity is present in the photoreceptor outer segment layer. Fig. 2A shows the staining in a bovine retinal section. Examination of dissociated photoreceptors indicates that only rod outer segments, but apparently not cone outer segments, are stained (Fig. 2C). PLC β 4-like staining is also present in the outer plexiform layer and the inner retina. The retinal staining seems specific, because, besides preadsorption experiments as negative controls (see *Methods*), immunoblotting of total proteins from bovine retina gave a single band of molecular mass (\approx 130 kDa) appropriate for PLC β 4 (Fig. 2B).

Because PLC β 4 has been shown to be activated by the α -subunits of the G $_q$ family of G proteins (36, 37), we have examined the distributions of some of these α -subunits (G α_{aq} , G α_{11} , and G α_{14}) in the retina as well. Among these, only G α_{11} -like immunoreactivity was found in rod outer segments. Fig. 3A shows this staining in bovine retina. It is present mostly in the outer and inner segment layers (stained structures in the inner retina marked by arrows are blood vessels); again, dissociated-cell studies indicate that only rod outer segments, but apparently not cone outer segments, show staining (Fig. 3C). Immunoblotting of total proteins from bovine retinas with the anti-G α_{11} antibody

likewise gave a single protein band of molecular mass (\approx 42 kDa) appropriate for this G protein, suggesting specific labeling (Fig. 3B).

Two other kinds of experiments were carried out to confirm the presence of PLC β 4 and G α_{11} in rod outer segments. First, we performed immunoblotting of proteins from partially purified rod outer segment preparations (Fig. 4) with the anti-PLC β 4 and anti-G α_{11} antibodies. In each case, a single protein band of appropriate molecular mass was observed (Fig. 4A and B). Labelings with an antibody against the rod cGMP-gated channel (which is present almost exclusively in the rod outer segment) and the antibody against PLC δ 2 (which is absent in the rod outer segment; see below) indicated that the partially purified rod outer segment preparations we used were indeed concentrated with rod outer segment proteins (Fig. 4C and D). Second, *in situ* hybridizations were performed on mouse retinal sections using antisense cRNA probes specific for mouse PLC β 4 and G α_{11} mRNA (see *Methods*). The mouse was chosen because nucleotide sequence information was available in the database for G α_{11} in this species and for PLC β 4 in the closely related rat. For PLC β 4, the hybridization signal is present in the inner segment layer (where most photoreceptor mRNA is situated) and the outer nuclear

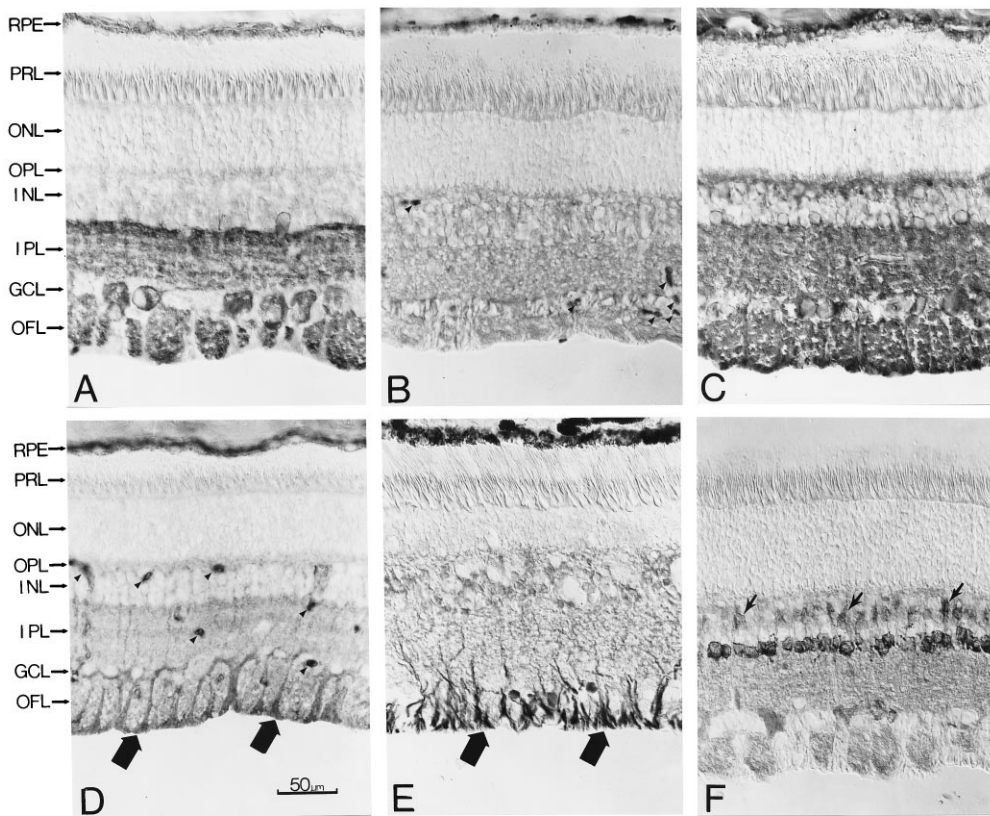


FIG. 6. Immunostaining of bovine retinal cross-sections for other PLC isozymes and also $G_{\alpha q}$. (Nomarski optics; 8- μ M frozen sections.) The anatomical layers are as in Fig. 2. (A) PLC β 1. (B) PLC β 3. (C) PLC γ 1. (D) PLC δ 1 (thick arrows indicate stained endfeet of Müller glial cells; arrowheads indicate blood vessels). (E) PLC δ 2 (arrows indicate stained endfeet of Müller glial cells). (F) $G_{\alpha q}$ (arrows indicate stained bipolar cells).

layer (which contains the cell bodies of photoreceptors), as well as cell bodies in the inner nuclear layer and the ganglion cell layer (Fig. 5A). The intense signal in the photoreceptors is thus consistent with the immunocytochemical results. The hybridization in the inner nuclear layer and the ganglion cell layer likewise agrees with the immunostaining in the inner retina. For $G_{\alpha 11}$, the hybridization signal is confined mostly to the inner segment layer (Fig. 5C) and is also in agreement with the immunocytochemistry. Control experiments with sense cRNA probes did not give any staining (Fig. 5B and D).

Other PLC Isozymes and G_q Family Members in the Retina.

Although PLC β 4 is the only isoform present in the outer segment layer, we have found other isoforms elsewhere in the retina. For PLC β 1, the immunoreactivity is confined mostly to the inner plexiform layer, some amacrine and ganglion cells, and the optic fiber layer (Fig. 6A). No PLC β 2 immunostaining was observed, which is in agreement with previous immunoblotting results (28). Finally, PLC β 3 staining has a weak, diffuse appearance in the bovine retina, spanning the outer plexiform layer and more proximal layers (Fig. 6B). As with PLC β 4, PLC β 1 and β 3 are activated by the α -subunits of the G_q family of G proteins (22–24). It is thus relevant that their locations in the inner retina are roughly coincident with that of $G_{\alpha q}$, another G_q family member present in the retina (see below). PLC β 3, and to a much lesser degree PLC β 1, can also be activated by G protein $\beta\gamma$ -subunits (23, 24), but in this study we have not pursued this point further in the retina.

For the PLC γ isoforms examined, only PLC γ 1 staining was detected. The staining is even more intense and widespread than that of PLC β 4, but is absent in rod outer segments (Fig. 6C). In the brain, this enzyme is also widespread (39, 40). Experiments with dissociated cells have confirmed staining in different retinal cell types, including photoreceptors, though the photoreceptor staining is confined to synaptic terminals, which are situated in the outer plexiform layer (data not shown). The PLC γ isoforms are known to be activated by growth-factor receptors through tyrosine phosphorylation (22–24). Such receptors are widespread in the retina (e.g., see ref. 41).

The antibodies against PLC δ 1 and PLC δ 2 both stained Müller glial cells exclusively, in particular their endfeet regions (Fig. 6D and E, large arrows; in Fig. 6D, small arrowheads indicate blood vessels). In the brain, these enzyme isoforms are also present predominantly in glial cells (42, 43). Their specific associations with glial cells may imply some unique functions.

As for other members of the G_q family, we have found $G_{\alpha q}$ but not $G_{\alpha 14}$ immunoreactivity in the retina. The $G_{\alpha q}$ immunostaining is not in photoreceptors but is primarily in the inner nuclear layer (Fig. 6F). The stained cell bodies in the latter location appear to be predominantly amacrine cells, though some bipolar cells (arrows) and some ganglion cells also show staining. Despite their high homology to each other and colocalizations in most tissues (44–46), recent work has suggested that $G_{\alpha 11}$ and $G_{\alpha q}$ can have differential functions, by, for example, coupling to different receptors (47). Our finding that the two proteins have differential localizations in the retina supports this point.

Overall, the combined distribution of all of the PLC isozymes we have examined partially match that of the IP $_3$ receptor we have found previously, which is predominantly in the two plexiform layers and Müller glial cells (33). One prominent difference is the absence of IP $_3$ receptor staining in the photoreceptor outer segment layer (33, 48). Minor differences also exist in the inner retina. It is possible that the antibody used in our previous study does not recognize all forms of the IP $_3$ receptor. Another possibility is that there is a functional bias toward the PKC branch instead of the IP $_3$ branch of the phosphoinositide pathway in certain cell locations, such as the case in the rod outer segment.

DISCUSSION

The existence of a phosphoinositide signaling pathway in the outer segments of vertebrate photoreceptor cells has been a topic of interest for many years, though the molecular identities of its components remained unknown. Our results suggest that PLC β 4 and $G_{\alpha 11}$, or immunologically identical isoforms, may be the constituents of this pathway. The amino acid sequences of the cloned bovine and rat PLC β 4 (49, 50) suggest that it is the mammalian homolog of a *Drosophila* PLC enzyme present pre-

dominantly in the eyes (51). A defect in this *Drosophila* enzyme results in a mutant called *norpA*, the photoreceptors of which fail to give any electrical response to light (52, 53); thus, this protein, called PLC-*norpA*, appears to be crucial for invertebrate phototransduction. The phototransduction pathways in *Drosophila* or other invertebrate rhabdomeric photoreceptors, nonetheless, are still unclear. By contrast, the central role of a cGMP cascade in phototransduction is well established in vertebrate rods (see *Introduction*), though Ca^{2+} also exerts an important negative-feedback control on this pathway. Thus, in rods, the phosphoinositide pathway can, in principle, interact with the cGMP cascade through IP_3 production and regulation of intracellular Ca^{2+} . So far, however, the evidence for such a mechanism is largely lacking. For example, the observed light-triggered production of IP_3 in rod outer segments is very transient (4, 5); in fact, there is no uniform agreement as to whether light actually influences phosphoinositide turnover (6, 7). So far, there also is no evidence for the presence of an IP_3 receptor in the rod outer segment, as pointed out in *Results*. Finally, the present understanding of rod phototransduction, which is quite detailed, does not need to invoke any IP_3 -mediated Ca^{2+} mobilization (1). Taken together, these points suggest that the IP_3 branch of the phosphoinositide pathway may be insignificant in the rod outer segment, at least with respect to its canonical mode of function (i.e., via an IP_3 receptor). The evidence for a function of the diacylglycerol-PKC branch of the pathway, on the other hand, is more substantial (9–21). In particular, its proposed role in light adaptation through phosphorylation of both bleached and unbleached rhodopsin (13–15) is intriguing, though confirmation from physiological experiments has yet to be made. There is also the possibility that the phosphoinositide pathway is involved in functions other than phototransduction in the outer segment, and triggered by a chemical signal rather than light.

A discrepancy between our study and a previous study by Ferreira and Pak (54) should be mentioned here. Using antibodies different from ours against PLC β_4 , these investigators have localized the enzyme to cones and not rods. The reason for this discrepancy is presently unclear, especially because earlier *in situ* hybridization experiments by the same group (49) have identified substantial message in the outer nuclear layer of the rod-dominant bovine retina, a result seemingly consistent with our present finding.

The colocalizations of PLC β_4 - and $G_{\alpha 11}$ -like immunoreactivities in the rod outer segment agree with the biochemical finding that PLC β_4 is specifically activated by α -subunits of the G_q family but not by rod transducin (28, 36, 37). Incidentally, $G_{\alpha 11}$ (or a very homologous isoform) is at present the only G protein besides transducin identified in the rod outer segment. An obvious question is whether $G_{\alpha 11}$ bears any functional relation to transducin in this location, and which $G_{\beta\gamma}$ subunits $G_{\alpha 11}$ is coupled to. Rod transducin ($G_{\alpha t1}$) is known to be coupled to $G_{\beta 1}$ and $G_{\gamma 1}$, the only G protein β - and γ -subunits so far identified in rod outer segments (55). Thus, it will be interesting to know whether the same or different β - and γ -subunits are involved in the function of $G_{\alpha 11}$. Another issue has to do with the apparent absence of PLC β_4 (at least according to our results) and $G_{\alpha 11}$ stainings in cone outer segments. Given its presence in rods, the phosphoinositide pathway is most likely present in cones as well, thus raising questions about the identities of the corresponding PLC enzyme and G_{α} protein in cones. One G protein β -subunit ($G_{\beta 3}$), but two γ -subunits ($G_{\gamma 2}$ and $G_{\gamma 8}$), have recently been identified in cone outer segments (55–57). Biochemical analysis suggests that, functionally, $G_{\gamma 8}$ is likely the cone homolog of rod $G_{\gamma 1}$ and coupled to cone transducin ($G_{\alpha t2}$) (57). Perhaps, then, $G_{\gamma 2}$ is coupled to the yet-to-be-identified G_q member in cone outer segments. Two α -subunits of the G_q family not examined in our study are $G_{\alpha 15}$ and $G_{\alpha 16}$, but these two proteins are thought to be predominantly in hematopoietic tissues (58, 59). Possibly, a novel

G_q family member exists. The same applies to the PLC isoform in cone outer segments. It is interesting that distinct molecular species seem to exist in rods and cones for many proteins of known function.

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