

Muscarinic Receptor Activation Promotes the Membrane Association of Tubulin for the Regulation of Gq-Mediated Phospholipase C β_1 Signaling

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The microtubule protein tubulin regulates adenylyl cyclase and phospholipase C β_1 (PLC β_1) signaling via transactivation of the G-protein subunits G α_s , G α_i1 , and G α_q . Because most tubulin is not membrane associated, this study investigates whether tubulin translocates to the membrane in response to an agonist so that it might regulate G-protein signaling. This was studied in SK-N-SH neuroblastoma cells, which possess a muscarinic receptor-regulated PLC β_1 -signaling pathway. Tubulin, at nanomolar concentrations, transactivated G α_q by the direct transfer of a GTP analog and potentiated carbachol-activated PLC β_1 . A specific and time-dependent association of tubulin with plasma membranes was observed when SK-N-SH cells were treated with carbachol. The same phenomenon was observed with membranes from Sf9 cells, expressing a recombinant PLC β_1 ,

cascade. The time course of this event was concordant both with transactivation of G α_q by the direct transfer of [³²P]P³(4-azidoanilido)-P¹-5'-GTP from tubulin as well as with the activation of PLC β_1 . In SK-N-SH cells, carbachol induced a rapid and transient translocation of tubulin to the plasma membrane, microtubule reorganization, and a change in cell shape as demonstrated by confocal immunofluorescence microscopy. These observations presented a spatial and temporal resolution of the sequence of events underlying receptor-evoked involvement of tubulin in G-protein-mediated signaling. It is suggested that G-protein-coupled receptors might modulate cytoskeletal dynamics, intracellular traffic, and cellular architecture.

Key words: tubulin; microtubules; cytoskeleton; G-protein; phospholipase C; muscarinic receptor

Cytoskeletal proteins appear to be involved in the control of intracellular signaling. The microtubule protein tubulin regulates adenylyl cyclase and phospholipase C β_1 (PLC β_1) signal transduction (Wang et al., 1990; Roychowdhury et al., 1993; Popova et al., 1994, 1997; Roychowdhury and Rasenick, 1994; Yan et al., 1996). The microtubule cytoskeleton is also suggested to be involved in the regulation of voltage-gated calcium channel activity (Unno et al., 1999). Certain isoforms of the microtubule-associated protein tau appear to regulate PLC γ (Hwang et al., 1996). Actin binds and inhibits the G-protein-coupled receptor kinase 5 (GRK5) (Freeman et al., 1998), and actin regulatory proteins, like profilin, gelsolin, and CapG, bind to the phospholipase C substrate phosphatidylinositol 4,5-bisphosphate (PIP₂) and inhibit PLC isoenzymes (Goldschmidt-Clermont et al., 1990, 1991; Banno et al., 1992; Sun et al., 1997). Although cytoskeletal elements affect growth factor-directed phospholipid metabolism (Payrastre et al., 1991; Rhee, 1991; Banno et al., 1992), the phosphorylation of PLC γ_1 by the epidermal growth factor receptor tyrosine kinase

overcomes profilin-induced inhibition of this signaling pathway (Goldschmidt-Clermont et al., 1991).

Tubulin regulates adenylyl cyclase and PLC β_1 signaling via specific interactions with the α subunits of the regulatory G-proteins, G α_s , G α_i1 , and G α_q (Wang et al., 1990; Roychowdhury et al., 1993; Popova et al., 1994, 1997). The interaction of tubulin with these polypeptides involves a GTP transfer from the exchangeable GTP-binding site (E site) of tubulin to G α , which activates the G-protein (transactivation) (Roychowdhury and Rasenick, 1994). Tubulin binds specifically to G α_i1 , G α_s , and G α_q with a K_d of ~130 nM (Wang et al., 1990). Complexes of tubulin and G α_i1 or G α_s have been immunoprecipitated from brain extracts and were suggested to represent functional assemblies, responsible for the observed G-protein activation by tubulin (Yan et al., 1996).

PLC β_1 , which is regulated by G α_q (Blank et al., 1992; Boyer et al., 1992; Park et al., 1993), evokes Ca²⁺-dependent hydrolysis of PIP₂ and generates two second messengers, inositol 1,4,5-trisphosphate (IP₃) and diacylglycerol (DAG) (Rhee and Choi, 1992). DAG is the physiological activator of protein kinase C (PKC), whereas IP₃ mobilizes stored calcium and promotes external calcium influx to activate Ca²⁺-dependent protein kinases ((Rhee and Choi, 1992; Berridge, 1993; Noh et al., 1995). A biphasic pattern of response of PLC β_1 to dimeric tubulin with guanosine 5'-(β,γ -imido)triphosphate (GppNHp) bound (tubulin-GppNHp) was detected in membranes from Sf9 cells expressing m₁ muscarinic receptors, G α_q , and PLC β_1 , as well as in a purified reconstituted system (Popova et al., 1997). At lower (nanomolar) concentrations, tubulin activated, whereas at higher (micromolar) concentrations, tubulin inhibited, PLC β_1 . The stimulatory effect of tubulin-GppNHp was more efficacious than that

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of GppNHp alone (Popova et al., 1997). The muscarinic receptor agonist carbachol significantly potentiated PLC β_1 activation by tubulin-GppNHp and was able to counteract partially the inhibitory effect seen at higher tubulin concentrations (Popova et al., 1997). Transactivation by tubulin appeared responsible for Gq and, subsequently, PLC β_1 activation, because nucleotide-free tubulin inhibited PLC β_1 at any tested concentration (Popova et al., 1997). The observed interaction of tubulin with the PLC β_1 substrate PIP $_2$ was thought to be responsible for enzyme inhibition at high concentrations of tubulin (Popova et al., 1997).

Most of the above-mentioned studies were performed with exogenous tubulin added to membranes or in reconstituted systems. The origin of the tubulin that might normally be involved in this process has not been investigated. It is possible that specific signals initiate a redistribution and translocation of cytosolic tubulin to the cell membrane (Popova et al., 1997; Panagiotou et al., 1999).

The present study investigates how cytoskeletal tubulin is involved in the regulation of PLC β_1 signaling. SK-N-SH neuroblastoma cells, which naturally maintain a PLC β_1 -signaling cascade (Fisher and Snider, 1987; Fisher, 1988; Fisher and Heacock, 1988), were used to test the hypothesis that agonist stimulation evokes a redistribution of cellular tubulin, tubulin membrane association, and engagement in G α_q -PLC β_1 regulation. Demonstration of agonist-induced reorganization of the microtubule cytoskeleton suggests that the interface between tubulin and heterotrimeric G-proteins may regulate both cellular signaling and cell shape or form.

MATERIALS AND METHODS

Cell culture. SK-N-SH neuroblastoma cells were maintained in DMEM supplemented with 10% fetal bovine serum according to standard procedures (Fisher and Snider, 1987). Sf9 cells were grown in Sf-900 II SFM media as described (Popova et al., 1997).

Baculovirus-directed protein expression in Sf9 cells. Sf9 cells were infected with baculoviruses bearing the m $_1$ muscarinic receptor, G α_q , or PLC β_1 cDNAs, as described previously (Popova et al., 1997). The construction of recombinant baculoviruses was described elsewhere (Parker et al., 1991; Graber et al., 1992; Boguslavsky et al., 1994). Membranes were prepared from cells collected 60 hr after infection.

Membrane preparation and Western blotting. Cells were sonicated in ice-cold 20 mM HEPES, pH 7.4, 1 mM MgCl $_2$, 100 mM NaCl, 1 mM dithiothreitol (DTT), and 0.3 mM PMSF (three times for 30 sec each; Branson sonifier; output control, 4). Membrane pellets were prepared as described (Popova et al., 1997). Protein concentrations were determined by the Bradford (1976) dye-binding assay with bovine serum albumin as a standard. Expression of receptors, G-proteins, and PLC β_1 was determined by immunoblotting. Membrane proteins transferred to nitrocellulose were probed with antibodies specific for the m $_1$ muscarinic receptor (#71; from G. Luthin, Philadelphia, PA), G $\alpha_q/11$ (#0945; from D. Manning, Philadelphia, PA), or PLC β_1 (K-32-3, monoclonal; from S. G. Rhee, Bethesda, MD) at a dilution of 1:500 (polyclonal) or 1:5000 (monoclonal). Biotinylated goat anti-rabbit IgG or anti-mouse IgG was used as a secondary antibody, accordingly. Either streptavidin-alkaline phosphatase or -horseradish peroxidase conjugates were used for detection via colorimetric or chemiluminescent techniques, as described. Expression levels were estimated by densitometry of the corresponding protein bands (Storm 840; Molecular Dynamics, Sunnyvale, CA). They varied by no >10% for a given recombinant protein. Receptor-binding studies using [3 H]-quinuclidinyl [phenyl-4(*n*)]benzilate ([3 H]QNB) as a ligand were also performed to monitor m $_1$ muscarinic receptor expression (Popova et al., 1997). When coexpressed with G α_q and PLC β_1 in the Sf9 cells, m $_1$ muscarinic receptor density was estimated at 240 fmol/mg of membrane protein (Popova et al., 1997). SK-N-SH cells have a high density of m $_3$ muscarinic receptors (500 fmol/mg of membrane protein) that are coupled to phosphoinositide turnover (Fisher and Snider, 1987; Fisher, 1988; Fisher and Heacock, 1988).

Tubulin preparations. Microtubules were isolated as described (Shelanski et al., 1973). Microtubule-associated proteins were removed by phos-

phocellulose chromatography, and the remaining pure tubulin fraction was termed PC-tubulin (Wang and Rasenick, 1991). Replacement of GTP in β -tubulin was performed as described previously (Wang et al., 1990). GTP was removed from PC-tubulin by charcoal pretreatment, and tubulin was incubated with 150 μ M guanine nucleotide [guanosine 5'-O-(3-thiotriphosphate) (GTP γ S), GppNHp, GDP, or [32 P]P 3 (4-azido-anilido)-P 1 -5'-GTP (AAGTP)] for 30 min on ice. Before use, these samples were passed through P6-DG columns twice to remove the unbound nucleotide. This procedure yields 0.4–0.6 moles of guanine nucleotide bound per mole of tubulin dimer. Tubulin-guanine nucleotide concentrations used throughout the study were based on the protein concentration.

PLC β_1 assays. Forty micrograms of SK-N-SH or 20 μ g of Sf9 membrane protein was incubated with a [3 H]PIP $_2$ substrate mixture (30 μ M final concentration, unless stated otherwise) as described (Popova et al., 1997). Ten microliters of GppNHp or tubulin-GppNHp and carbachol were added at appropriate concentrations to a final volume of 80 μ l. The tubes were incubated for 10 min (unless stated otherwise) at 37°C with constant shaking, as described (Popova et al., 1997). [3 H]IP $_3$ production was quantified by liquid scintillation counting (Popova and Dubocovich, 1995; Popova et al., 1997).

Photoaffinity labeling and nucleotide transfer. Cell membranes were incubated with the indicated concentrations of dimeric tubulin with [32 P]AAGTP bound (tubulin-[32 P]AAGTP) in the presence or absence of carbachol in 100 mM 1,4-piperazinediethanesulfonic acid (PIPES) buffer, pH 6.9, 2 mM EGTA, and 1 mM MgCl $_2$ (buffer A) as described (Rasenick et al., 1994). The tubes were UV irradiated, and the reaction was quenched with ice-cold PIPES buffer, 1 mM MgCl $_2$, and 4 mM DTT. After centrifugation at 20,000 \times g for 15 min, the membrane pellets were washed with buffer and dissolved in SDS Laemmli sample buffer with 50 mM DTT as described (Laemmli, 1970). SDS-PAGE of the samples was performed (Popova et al., 1997), and the gels were either stained (Coomassie blue) or subjected to Western blotting, followed by autoradiography (Kodak XAR-5 film) or phosphorimage analysis of the bands (Storm 840; Molecular Dynamics).

Immunoprecipitation. Membrane preparations were extracted with 1% sodium cholate in buffer A for 1 hr at 4°C with constant stirring. The tubes were centrifuged at 20,000 \times g for 15 min at 4°C. Membrane extracts (0.5 mg/ml membrane protein) were incubated with guanine nucleotide-bound tubulin (1 μ M), as described above. After UV irradiation and preclearing (Pansorbin; Calbiochem, La Jolla, CA), each membrane extract was incubated overnight with the appropriate specific antiserum or preimmune serum (1:20 dilution for polyclonal and 1:500 dilution for monoclonal antibodies) at 4°C with constant stirring. Immune complexes were precipitated with Pansorbin, and each immunoprecipitate was subjected to SDS-PAGE and autoradiography, as described (Popova et al., 1997). Where indicated, Western blotting followed by an ECL detection of the bands was performed.

Membrane association of tubulin in SK-N-SH cells. SK-N-SH cells were collected and washed three times with PBS, and aliquots of 1×10^7 cells were distributed in plastic tubes on ice. Carbachol (1 mM) was immediately added, and the samples were incubated for the indicated time periods at 37°C with constant shaking. When tested, atropine (100 μ M) was added before carbachol. Samples were transferred on ice and immediately sonicated, as described. Each sample was centrifuged at 600 \times g at 4°C (P $_1$ pellets). Supernatants were centrifuged at 20,000 \times g at 4°C (P $_2$ pellets). SDS-PAGE of the fractions was followed by immunoblotting, using polyclonal anti-tubulin antiserum (raised against the C-terminal 422–431 amino acid region of β -tubulin) and ECL detection (Amersham, Arlington Heights, IL). The results were analyzed in a Storm 840 image system (Molecular Dynamics).

Confocal immunofluorescence microscopy. SK-N-SH cells were plated onto glass coverslips in 12-well culture plates at a density of 1×10^5 . After ~24 hr, treatment with 1 mM carbachol, 10 μ M atropine, or both was performed for the indicated times at 37°C. After washing with PBS buffer, the cells were fixed in -20°C methanol for 6 min. Cells were washed three times in PBS and blocked at room temperature for 30 min in 5% NGS-containing PBS. After a PBS wash, cells were incubated for 2 hr with the above-mentioned polyclonal anti- β -tubulin antiserum (1:100 dilution). After washing three times in PBS, secondary fluorescein isothiocyanate (FITC)-conjugated goat anti-rabbit antibody (EY Laboratories; 1:100 dilution) was applied for 1 hr, followed by washing and mounting. Images were acquired using a Carl Zeiss laser-scanning confocal microscope LSM 510 equipped with a 40 \times immersion objective. A single 488 nm beam from an argon-krypton laser was used for excitation.

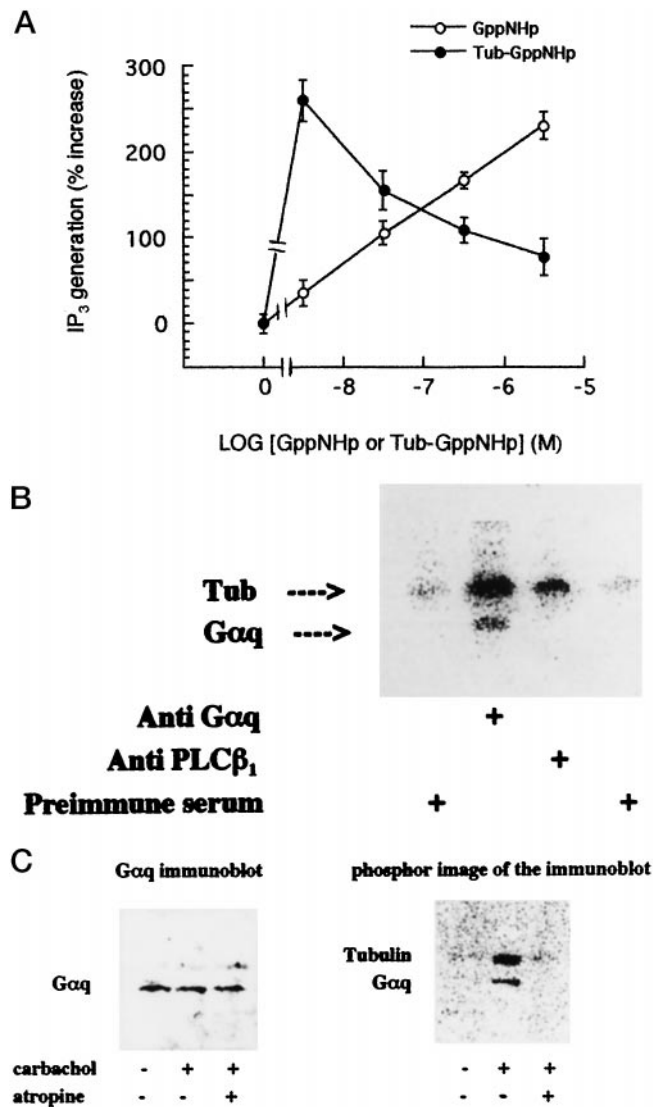


Figure 1. Tubulin regulates PLC β signaling in SK-N-SH neuroblastoma cell membranes. *A*, Dual regulation of PLC β_1 by tubulin-GppNHp. SK-N-SH membranes (50 μ g of membrane protein) were assayed for PLC β_1 activity in the presence of 1 mM carbachol and the indicated concentrations of GppNHp or tubulin-GppNHp. Control activity was 0.70 ± 0.08 nmol of IP $_3 \cdot \text{min}^{-1} \cdot \text{mg of protein}^{-1}$. A representative of three experiments with similar results is shown. *B*, Coimmunoprecipitation of tubulin with G α_q and PLC β_1 . Extracted SK-N-SH membranes (0.5 mg/ml) and 1 μ M tubulin-[32 P]AAGTP were tested as described. Tubulin was not precipitated when preimmune serum replaced the anti-G α_q antibody (*left lane*) or the anti-PLC β_1 antibody (*right lane*). A phosphorimage of one of three independent experiments with similar results is shown. *C*, Carbachol-triggered membrane association of tubulin and guanine nucleotide transfer from tubulin to G α_q . SK-N-SH membranes were incubated with 100 nM tubulin-[32 P]AAGTP, 10 μ M carbachol, and 100 nM atropine (as indicated) for 5 min at 23°C. UV irradiation, SDS-PAGE (50 μ g of membrane protein in *each lane*), immunoblotting, and phosphorimage analysis of the blots were performed as described. A representative experiment of three with similar results is shown. In the absence of carbachol, no increase in tubulin membrane association or nucleotide transfer to G α_q was detected at any time point tested. *Tub*, Tubulin.

Emission from FITC was detected via an LP505 filter. Differential interference contrast (DIC) images of the cells were also acquired. Images of computer-generated cross sections of cells were collected as well (*x-z* and *y-z* planes). Coverslips were examined at random. At each time point a total of 300 randomly selected cells over four consecutive experiments were evaluated for tubulin redistribution, changes in micro-

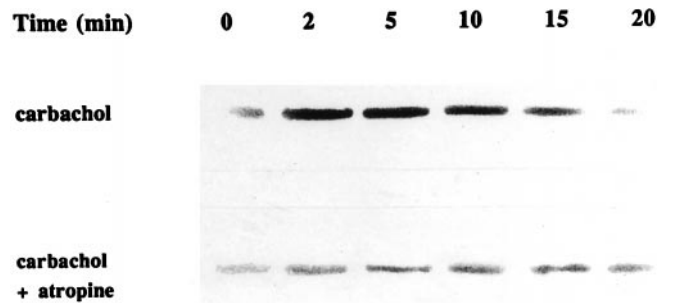


Figure 2. The activation of muscarinic receptors in SK-N-SH cells triggers the membrane association of tubulin. SK-N-SH cells were incubated for the indicated times with 1 mM carbachol with or without 100 μ M atropine, as described. Membrane pellets were subjected to SDS-PAGE (50 μ g of membrane protein in *each lane*) and immunoblotting with anti- β -tubulin antiserum, as described. A representative experiment of four with similar results is shown.

tubule network, and cell shape. When analyzed, cellular processes were defined as projections of $>2 \mu\text{m}$ in length. Final image composites were created using Adobe Photoshop 5.0. No specific FITC labeling was observed in cells treated with preimmune serum instead of anti- β -tubulin antiserum. FITC labeling was not observed when the anti-tubulin antiserum was preincubated overnight at 4°C with PC-tubulin (1:1 ratio) and tested at the same antibody dilution (1:100) afterward. Although the anti- β -tubulin antiserum did not cross-react with actin in immunoblotting, SK-N-SH cells were tested for tubulin and actin distribution throughout the cell. Cells were double stained with the above-described polyclonal anti- β -tubulin antibody (1:100 dilution) and an anti-actin antibody (monoclonal; kindly provided by Dr. J. Lessard, University of Cincinnati, Cincinnati, OH; 1:1000 dilution). Secondary FITC-conjugated goat anti-rabbit antibody was used to detect tubulin (1:100 dilution), and Texas Red dye-conjugated AffiniPure goat anti-mouse antibody (Jackson ImmunoResearch, West Grove, PA; 1:100 dilution) was used to detect actin. Distinct protein structures were recognized by the anti-tubulin and anti-actin antibodies.

Materials. [α - 32 P]GTP was from ICN Biomedicals (Cleveland, OH). [32 P]AAGTP and AAGTP were synthesized as described (Rasenick et al., 1994). *p*-Azidoaniline was synthesized by Dr. William Dunn (University of Illinois, Chicago, IL). [^3H]PIP $_2$ and myo-[^3H]inositol were from American Radiolabeled Chemicals (St. Louis, MO). [^3H]QNB was from Amersham. GppNHp, GTP, GDP, and GTP γ S were from Boehringer Mannheim (Indianapolis, IN). Carbachol, atropine sulfate, and PIP $_2$ were from Sigma (St. Louis, MO). The P6-DG desalting gel was from Bio-Rad (Hercules, CA). P11 cellulose phosphate was from Whatman (Maidstone, UK). All other reagents were of analytical grade.

RESULTS

PLC β_1 signaling is regulated by tubulin in neuroblastoma SK-N-SH cells

Both reconstitution studies with purified proteins and studies using membranes from Sf9 cells with recombinant m_1 muscarinic receptors, G α_q , and PLC β_1 indicated that tubulin regulated PLC β_1 signaling (Popova et al., 1997).

To test whether tubulin would regulate PLC β_1 in a cell with endogenous expression of a PLC β_1 cascade, membranes were prepared from SK-N-SH neuroblastoma cells, and carbachol-evoked activation of PLC β_1 was studied. Tubulin-GppNHp evoked a biphasic regulation of PLC β_1 in SK-N-SH cells similar to that seen in the Sf9 system. Enzyme activation at lower (nanomolar) and inhibition at higher (micromolar) concentrations of tubulin-GppNHp were observed (Fig. 1*A*). Tubulin-GppNHp potentiated carbachol-evoked phosphoinositide hydrolysis more than did GppNHp. Thus it appears that in SK-N-SH cells, tubulin might participate in the regulation of PLC β_1 via a receptor-Gq-signaling cascade.

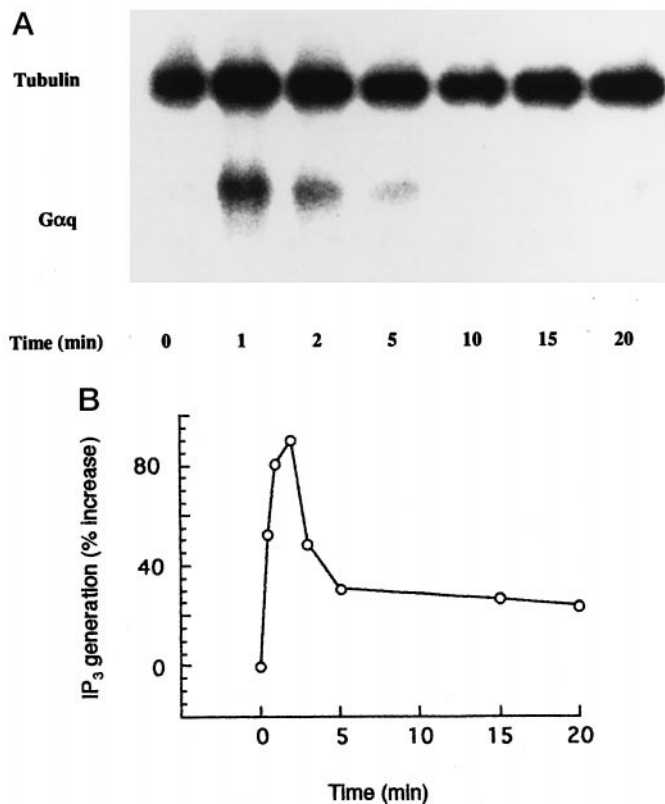


Figure 3. Tubulin membrane association, nucleotide transfer to G α q, and PLC β_1 activation follow a similar time course in membranes from Sf9 cells that contain a recombinant PLC β_1 cascade. *A*, Time course of the carbachol-evoked tubulin-[³²P]AAGTP association with the membrane and the nucleotide transfer to G α q. Membranes were incubated for the indicated times with 1 μ M tubulin-[³²P]AAGTP and 10 μ M carbachol at 23°C, followed by UV irradiation, SDS-PAGE (70 μ g of membrane protein in each lane), and autoradiography. The experiment shown is representative of three with similar results. In the absence of carbachol, no increase in tubulin membrane association or nucleotide transfer to G α q was detected at any time point tested. Both tubulin membrane association and G α q transactivation, triggered by carbachol, were inhibited by 100 nM atropine. *B*, Carbachol-evoked activation of PLC β_1 by tubulin-GppNHp. Twenty micrograms of Sf9 cell membranes containing the indicated recombinant proteins were incubated with 100 μ M PIP₂ substrate as described. Carbachol (10 μ M) and tubulin-GppNHp (30 nM) were added and incubated for the indicated times at 37°C as described. The experiment shown is representative of three with similar results. Control activity was 0.61 ± 0.09 nmol of IP₃ · min⁻¹ · mg of protein⁻¹. When GppNHp was tested under these experimental conditions, a linear increase in PLC β_1 activation was observed. Tubulin-GppNHp activation of PLC β_1 in the presence of carbachol was atropine-sensitive.

Tubulin transactivates G α q in SK-N-SH cell membranes

To investigate whether tubulin interacted with G α q in SK-N-SH membranes, these membranes were extracted with sodium cholate and incubated with tubulin-[³²P]AAGTP, and the extracts were immunoprecipitated with anti-G α q or anti-PLC β_1 antisera (Fig. 1*B*). Tubulin coimmunoprecipitated with endogenous G α q and, to a lesser extent, with PLC β_1 .

Previous studies suggested that tubulin activated G α q by the direct transfer of GTP (Popova et al., 1997). This was tested in SK-N-SH cells using the photoaffinity GTP analog AAGTP. Carbachol increased the association of tubulin-[³²P]AAGTP with SK-N-SH membranes (Fig. 1*C*). Although the amount of G α q was

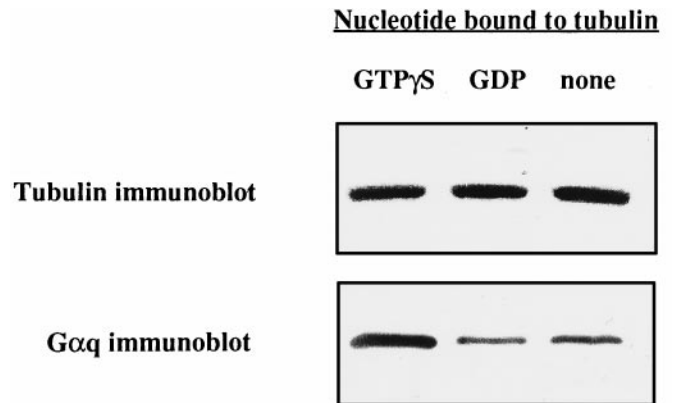


Figure 4. Preferential association of G α q with the GTP γ S-bound form of tubulin. Membrane extracts (0.5 mg/ml membrane protein) from Sf9 cells infected with baculoviruses carrying G α q cDNA were incubated at 4°C with 1 μ M tubulin-GTP γ S, tubulin-GDP, or nucleotide-free tubulin. Immunoprecipitation with anti- β -tubulin antibody (polyclonal) was followed by SDS-PAGE (10% gels) and immunoblotting with anti-G α q and anti- α -tubulin antibodies (monoclonal DM 1A). ECL detection and densitometry of the bands were used to estimate the G α q bound to different tubulin states. No tubulin or G α q immunoprecipitated when preimmune serum replaced the anti- β -tubulin antiserum in the immunoprecipitation. G α q-containing Sf9 membranes were run in each experiment to ensure that the immunoprecipitated protein was G α q. The image of one of three independent experiments with similar results is shown.

Table 1. Coimmunoprecipitation of G α q with different tubulin-guanine nucleotide bound species

Tubulin species	Arbitrary units (% change)
Tubulin-GTP γ S	100.0 \pm 8.2
Tubulin-GppNHp	89.0 \pm 7.4
Tubulin-GDP	31.0 \pm 5.1*
Tubulin stripped of nucleotide	43.2 \pm 9.3*

Membranes were extracted with sodium cholate, incubated with the indicated tubulin species, and immunoprecipitated with anti-tubulin antiserum as described in Figure 4. The results presented are the means \pm SD of three independent experiments.

*Significantly less than tubulin-GTP γ S ($p < 0.05$), Student's t test.

similar under these experimental conditions, [³²P]AAGTP transfer from tubulin to G α q (transactivation) was observed only after muscarinic receptor stimulation. Thus, an activated receptor was required to initiate the process of G α q activation by tubulin. This effect was specific, because it was blocked by atropine.

Carbachol triggers the membrane association of tubulin in SK-N-SH cells

In the studies described above, exogenous tubulin was shown to bind to isolated membranes and regulate PLC β_1 signaling. To clarify whether endogenous tubulin would gravitate toward the membrane in response to agonist stimulation, we incubated SK-N-SH cells with carbachol and quantified membrane-associated tubulin by immunoblotting (Fig. 2). Carbachol evoked a rapid and time-dependent increase in the tubulin recruited to membranes of the SK-N-SH cells. Tubulin association with membranes (post-nuclear fraction) increased by ~ 2.5 -fold [$244.8 \pm 31.3\%$ (\pm SD)] after 2 min of carbachol treatment and gradually declined afterward. The process was atropine sensitive. A decrease in cytosolic tubulin was also detected [$35.9 \pm 10.5\%$ (\pm SD)]. Carbachol did

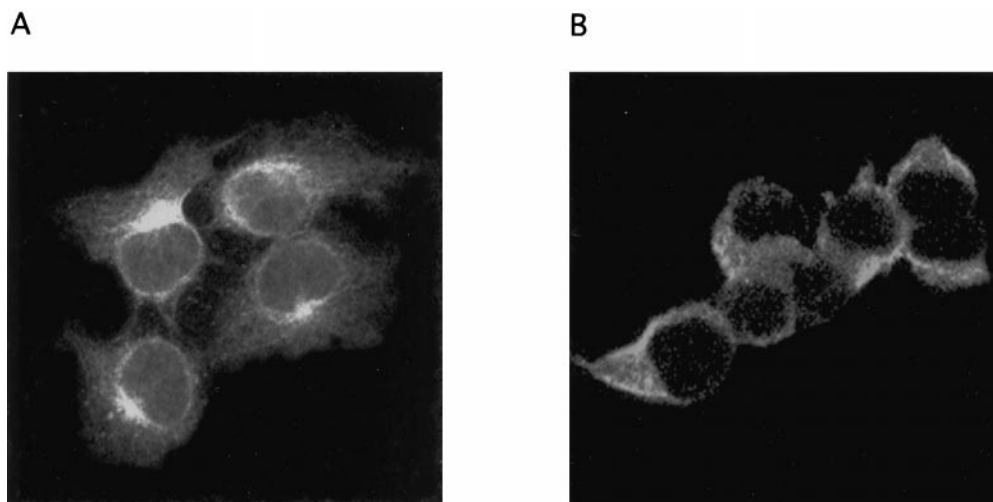


Figure 5. Carbachol stimulation causes microtubule reorganization and translocation of tubulin to the membrane in the SK-N-SH cells. Cells were untreated (*A*) or treated with 1 mM carbachol for 2 min (*B*) before fixation and immunofluorescence labeling, as described. A representative image of cells obtained in one of four independent experiments with similar results is shown.

not increase the amount of tubulin associated with the nuclear fraction.

The time courses of tubulin- ^{32}P AAGTP membrane association, $\text{G}\alpha\text{q}$ transactivation by tubulin, and $\text{PLC}\beta_1$ activation are concordant

The activation of $\text{PLC}\beta_1$ by tubulin shows rapid onset and decline. To understand this process, it was important to correlate the association of tubulin with the membrane and the transactivation of $\text{G}\alpha\text{q}$ with the activation of $\text{PLC}\beta_1$. This was performed in Sf9 cells in which no endogenous $\text{PLC}\beta_1$ activity is detected unless m_1 muscarinic receptors, $\text{G}\alpha\text{q}$, and $\text{PLC}\beta_1$ are expressed (Popova et al., 1997). Carbachol (10 μM) induced a rapid membrane association of tubulin- ^{32}P AAGTP with Sf9 membranes containing all three elements of the $\text{PLC}\beta_1$ cascade (Fig. 3*A*). The pattern of response was similar to that observed in the SK-N-SH membranes. The amount of membrane-associated tubulin was maximal after 1 min [an increase of $127.0 \pm 15.5\%$ (\pm SD)] and gradually decreased over 10 min. The transfer of ^{32}P AAGTP from tubulin to $\text{G}\alpha\text{q}$ followed the same pattern. All ^{32}P AAGTP bound to $\text{G}\alpha\text{q}$ originated from tubulin, because under this experimental condition ^{32}P AAGTP remains bound to tubulin unless transferred directly to $\text{G}\alpha$ (Roychowdhury and Rasenick, 1994). The association of tubulin with the membrane and $\text{G}\alpha\text{q}$ transactivation by tubulin were dependent on m_1 receptor activation and blocked by atropine (data not shown). Carbachol did not cause the association of tubulin with native Sf9 membranes or with membranes that did not contain recombinant m_1 muscarinic receptors, $\text{G}\alpha\text{q}$, or $\text{PLC}\beta_1$ (Popova et al., 1997).

The temporal relationship between the membrane association of tubulin and the activation of $\text{G}\alpha\text{q}$ and $\text{PLC}\beta_1$ was also tested (Fig. 3*B*). The time course of carbachol-induced activation of $\text{PLC}\beta_1$ in the presence of tubulin-GppNHp was strikingly parallel to the membrane association of tubulin and $\text{G}\alpha\text{q}$ transactivation. The enzyme activity reached a maximum in 2 min and declined afterward, although a 30% activation of the enzyme above basal was maintained for the duration of the experiment. The observed kinetics of tubulin membrane association, both *in vivo* and *in vitro*, supported the idea that tubulin was recruited to the membrane in response to receptor stimulation for the regulation of $\text{G}\alpha\text{q}$ -mediated $\text{PLC}\beta_1$ signaling.

Tubulin must have GTP (or a GTP analog) in the exchangeable binding site to interact with $\text{G}\alpha\text{q}$ and activate $\text{PLC}\beta_1$

Previous results suggested that GTP or a GTP analog had to occupy the exchangeable nucleotide-binding site of tubulin to activate $\text{PLC}\beta_1$, whereas $\text{PLC}\beta_1$ inhibition at high tubulin concentrations was a nucleotide-independent process. When tested, tubulin-GppNHp, but not tubulin-GDP, tubulin-GDP βS , or tubulin stripped of nucleotide, activated the enzyme (Popova et al., 1997).

To analyze further the importance of GTP in the interaction between tubulin and $\text{G}\alpha\text{q}$, tubulin with various guanine nucleotides bound was coimmunoprecipitated with $\text{G}\alpha\text{q}$. Tubulin-GTP γS , tubulin-GppNHp, tubulin-GDP, or tubulin free of nucleotide in the exchangeable GTP-binding site was incubated with detergent extracts of Sf9 cell membranes enriched in $\text{G}\alpha\text{q}$. Anti-tubulin antiserum was added, and tubulin was immunoprecipitated (Fig. 4, Table 1). Although the amounts of the tubulin species, immunoprecipitated by the anti-tubulin antiserum, were identical (this was determined by blotting with a monoclonal anti-tubulin antibody, DM 1A; Sigma), $\text{G}\alpha\text{q}$ coimmunoprecipitated with the GTP γS - and GppNHp-bound forms of tubulin. Coimmunoprecipitation of $\text{G}\alpha\text{q}$ with the GDP-bound form of tubulin or with tubulin devoid of nucleotide was identical to that seen with preimmune serum replacing the anti-tubulin antiserum (Fig. 4, Table 1). These results confirmed that a conformation of tubulin with GTP bound to the E site was required for tubulin interaction with $\text{G}\alpha\text{q}$.

Carbachol stimulation causes redistribution of intracellular tubulin

It is predicted that tubulin dimers translocate to the membrane in response to muscarinic receptor stimulation, transactivate $\text{G}\alpha\text{q}$, and promote activation of $\text{PLC}\beta_1$. A possible mechanism would be that when the intracellular calcium concentration rises in defined areas of the cell, microtubules depolymerize and more dimers are available to interact with $\text{G}\alpha\text{q}$. To clarify whether muscarinic receptor stimulation causes redistribution of intracellular tubulin, SK-N-SH cells were treated with carbachol and studied with confocal laser microscopy.

During interphase, microtubules were seen in greatest density at the centrosome, radiating out to the periphery of the cell in a fine array of threads (Fig. 5). Tubulin was not seen at the plasma

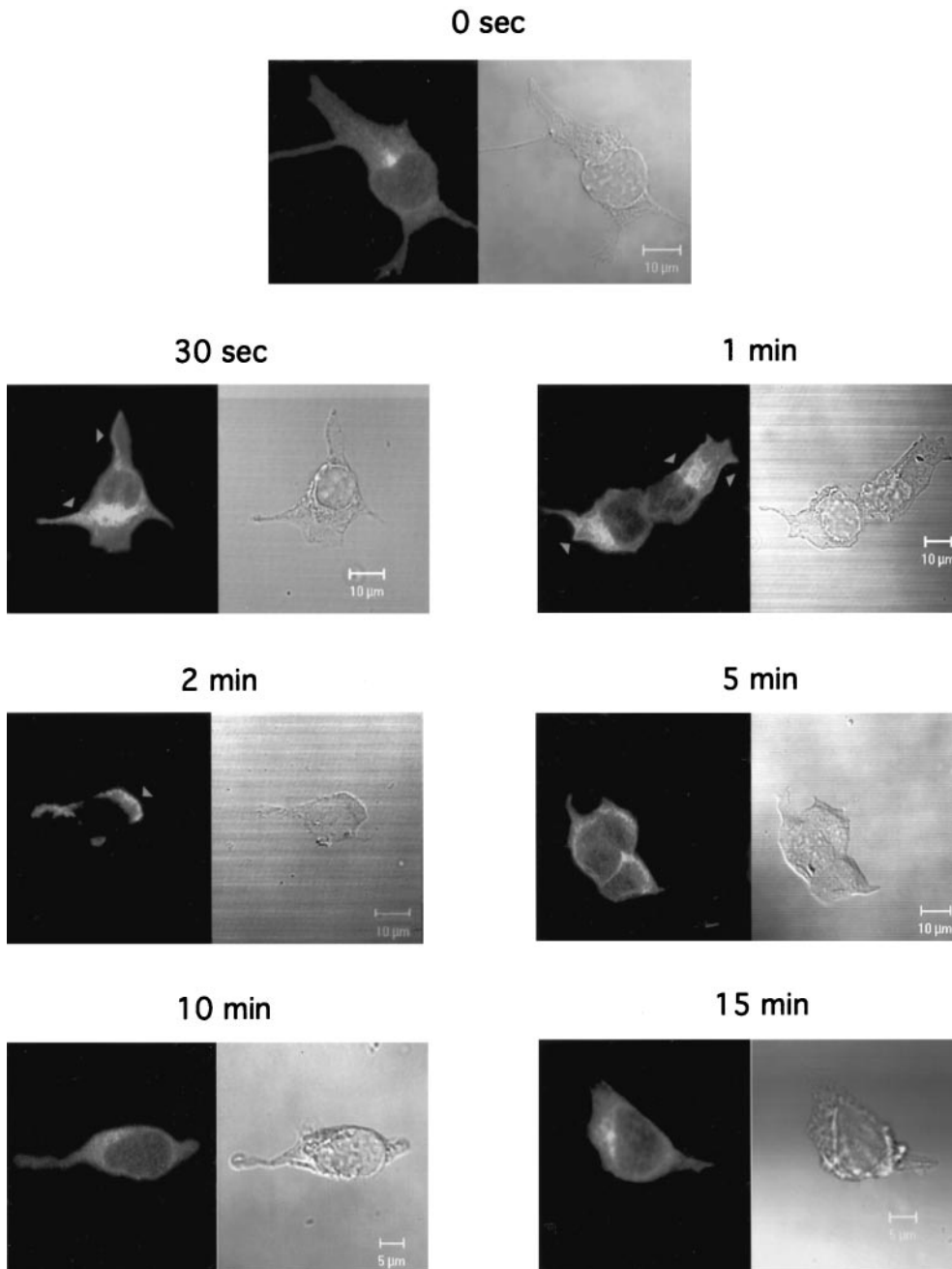


Figure 6. Time course of the redistribution and membrane translocation of tubulin during carbachol stimulation of SK-N-SH cells. Cells were treated with 1 mM carbachol for the periods indicated before fixation and immunostaining, as described. When 100 μ M atropine was applied before carbachol, the images were identical to those of control cells. Confocal micrographs of the treated cells are shown to the *left*. Shown to the *right* are the differential interference contrast micrographs of the same cells. The *arrowheads* denote areas of membrane localization of tubulin. Confocal images of 1- μ m-thick sections at the same level within the cell are presented. Four independent experiments with similar results were performed. The images shown are representative of \sim 300 cells examined at each time point.

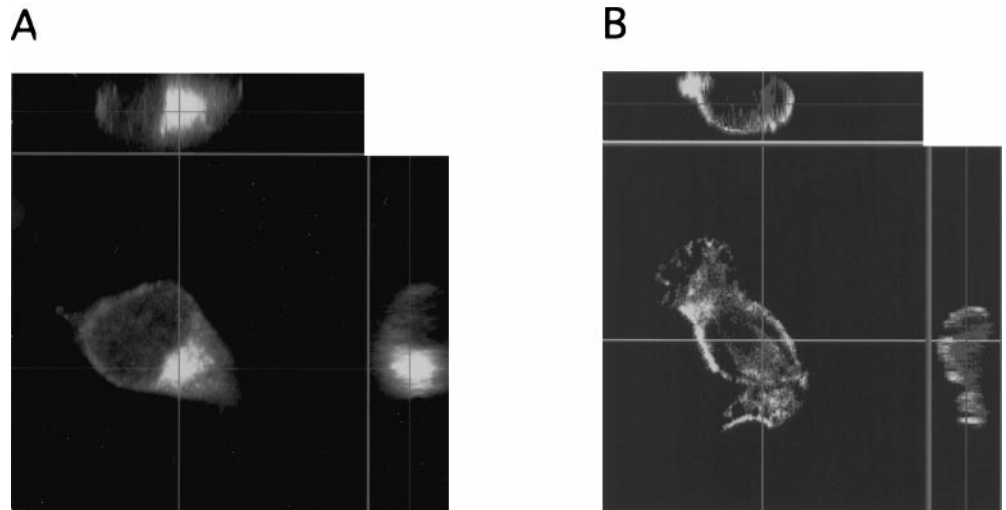
membrane. Two minutes after carbachol exposure, microtubule depolymerization and redistribution of tubulin to areas along the plasma membrane were observed. A rapid and transient translocation of tubulin to the periphery of the cell was seen as early as 30 sec after muscarinic receptor activation (Fig. 6). The membrane relocation of tubulin was most evident at 2 min after agonist stimulation. This event was accompanied by changes in the organization of the cytoskeleton, as well as cell shape. This is seen in the corresponding DIC images. Although most of the control cells (85.5%; $n = 300$) exhibited a well developed microtubule network and cellular projections, cells treated with carbachol progressively lost and regained these features. At 2 min of stimulation, microtubules in most of the cells appeared disorganized, cell projections were either lacking or shortening, and the cells had the tendency to round up (processes appeared in only 12.6% of the cells; $n = 300$).

Five minutes after agonist stimulation, the process of tubulin redistribution back to the cytosol had commenced. Distinct membrane localization of tubulin was no longer observed after 10 min of carbachol exposure of the cells. Fifteen minutes subsequent to the initiation of carbachol exposure, the resting cell shape was restored, and the microtubule network was reorganized. Cell processes became more abundant (processes were present in 61.0% of the cells; $n = 300$). The confocal z-scan images, presented in Figure 7 clearly demonstrate a normal distribution of tubulin in the untreated cell and its peripheral redistribution after 2 min of carbachol exposure.

DISCUSSION

The principal finding in this study is that muscarinic receptor activation triggers a transient redistribution and membrane association of cytosolic tubulin. The intracellular reorganization of

Figure 7. Carbachol exposure of the SK-N-SH cells translocates tubulin to the membrane of the cell. Computer-generated cross sections of the whole cell are displayed on the *top* (x - z plane) and on the *right* (y - z plane). **A**, Untreated SK-N-SH cell (0 time point, control). Note the random distribution of tubulin throughout the cell in the *top* and *right* computer-generated cross-section images. **B**, SK-N-SH cell treated with 1 mM carbachol (2 min exposure). Note the membrane localization of tubulin in the *top* and *right* computer-generated cross-section images of the cell. The images shown are representative of at least 30 cells subjected to a z -scan analysis with similar results.



this cytoskeletal protein in response to agonist stimulation was visualized in SK-N-SH neuroblastoma cells, and the carbachol-evoked association of tubulin with the plasma membrane was correlated with the regulation of Gq-mediated PLC β_1 signaling. This translocation event carries the potential of affecting a broad spectrum of cellular functions, including other signaling pathways, intracellular trafficking, cell shape, cell movement, and cell division.

Numerous cytosolic enzymes, including phospholipases, kinases, and phosphatases, associate transiently with membrane proteins. The association of tubulin with receptors (Kirsch et al., 1991; Barsony and McKoy, 1992; Item and Sieghart, 1994), G-proteins (Wang et al., 1990; Wang and Rasenick, 1991; Roychowdhury et al., 1993; Popova et al., 1994, 1997; Rasenick et al., 1994; Roychowdhury and Rasenick, 1994; Yan et al., 1996), and regulatory enzymes (Kapeller et al., 1995; Reszka et al., 1995; Sontag et al., 1995; Popova et al., 1997; Pitcher et al., 1998) has also been reported. A direct interaction of tubulin with the α subunit of the G-proteins Gs, Gi1, and Gq is thought to mediate adenylyl cyclase and PLC β_1 signaling (Popova et al., 1994, 1997; Yan et al., 1996). The present findings extend these observations further. They raise the possibility that at least some of the tubulin associated with the cell membrane might be transiently recruited from the cytosol in response to a signal. It is suggested that previously observed tubulin-G α complexes (Yan et al., 1996) are transient dynamic formations, whose functional assembly is regulated and reversible.

The present data also reveal that tubulin dimers must be in the GTP conformation to interact with G α q. A similar functional interaction of tubulin-GTP but not tubulin-GDP with the small GTPase Rac1 has been reported (Best et al., 1996). Furthermore, tubulin-GppNHP was more potent and efficacious than GppNHP for the activation of PLC β_1 . This suggested the possibility that, after receptor activation, G α q was transactivated by tubulin-GTP more easily than activated by GTP alone. It also indicated that the effect of tubulin-GTP on PLC β_1 could be significant in cellular compartments where the local tubulin dimer concentration is comparable with or higher than that of GTP.

Although membrane-associated tubulin has been demonstrated (Bhattacharyya and Wolff, 1976; Zisapel et al., 1980; Pfeffer et al., 1983; Stephens, 1986), there is no clear evidence that a membrane-integrated tubulin isoform exists. For tubulin to transactivate G α , however, association with the membrane can be

quite transient. In fact, the evidence in Figures 3 and 5–7 is consistent with such a transient association. It is noteworthy that a redistribution of cytoskeletal components, including actin and tubulin, has been reported to occur in T47D breast cancer cells after opioid receptor stimulation (Panagiotou et al., 1999).

In addition to elements of the cytoskeleton regulating cellular signaling, cellular-signaling molecules have been shown to regulate cytoskeletal form and function. The actin cytoskeleton is regulated by the small GTP-binding proteins Rho, Rac, Cdc42, and Ras (Zigmond et al., 1997; Hall, 1998; Nobes and Hall, 1999); the heterotrimeric G-protein subunits G α i1, G α q, and G $\beta_1\gamma_2$ affect microtubule polymerization dynamics (Ravindra et al., 1996; Roychowdhury and Rasenick, 1997; Roychowdhury et al., 1999).

After a rapid increase, tubulin migration to the membrane, transactivation of G α q by tubulin, and activation of PLC β_1 declined (Figs. 2, 3, 6). The reorganization of the microtubule cytoskeleton also appeared transient and reversible. It is possible that increases in cytosolic calcium resulting from the generation of IP $_3$ might evoke localized destabilization of microtubules. As a result, an increase in tubulin dimer concentrations near regions of the membrane where receptors, Gq, and PLC β_1 are found would be expected. Because tubulin concentrations >100 nM inhibit PLC β_1 , continued activation of PLC β_1 could effect a feedback inhibition of the enzyme via tubulin.

It is possible that an increased amount of tubulin associated with the plasma membrane would interfere with the coupling between G α q and PLC β_1 . Tubulin coimmunoprecipitates with both G α q and PLC β_1 , although to a lesser extent with the later (Fig. 1B) (Popova et al., 1997). As such, a direct tubulin-PLC β_1 interaction (at high tubulin concentrations) might abolish G α q-PLC β_1 coupling. However, tubulin also binds the PLC β_1 substrate PIP $_2$ (Popova et al., 1997). Rapid hydrolysis of a readily available PIP $_2$ pool by PLC β_1 , followed by tubulin association with the remaining substrate, might also account for the observed decrease in IP $_3$ generation in the course of the experiment.

It is also possible that dissociation of tubulin from the membrane after G α q activation accounts for the observed decline in PLC β_1 activation. Such movement of tubulin from the membrane back to its original location in the cytosol could be caused by the decrease in the local membrane concentration of PIP $_2$. If G α q is the membrane anchor for tubulin, the loss of GTP from tubulin during the process of G α q transactivation would also decrease the

affinity of tubulin for Gq and release tubulin. In addition to G α q and PIP₂, other signaling proteins might enjoy a reversible association with tubulin at the plasma membrane. Both G $\beta\gamma$ (Roychowdhury and Rasenick, 1997) and the GRKs, GRK2 (Carman et al., 1998; Haga et al., 1998; Pitcher et al., 1998) and GRK5, (Carman et al., 1998) bind to tubulin. G $\beta\gamma$ assists the recruitment of GRK2 to the membrane (Pitcher et al., 1992). Thus, a possible scenario is that tubulin, GRK, and G $\beta\gamma$ are involved in a common membrane association pathway and that some interplay among these molecules is a regulatory event. Should the membrane anchor for tubulin prove to be a molecule other than those listed above, another scenario would need to be developed.

It is noteworthy that activity shapes the structure of neurons and their circuits. Synaptic activation is shown to produce rapid input-specific changes in dendritic structure (Maletic-Savatic et al., 1999). The possibility exists that the neurotransmitter-evoked recruitment of tubulin to the membrane assists with this process. In fact, it has been suggested previously that the synaptic activity-controlled balancing of monomeric, dimeric, and polymeric forms of actin and tubulin might underlie the changes in spine shape (Van Rossum and Hanisch, 1999).

In summary, the experiments described above support the notion that, in addition to its function as a structural protein that forms microtubules, tubulin serves as a signal-recruited regulator of membrane-associated signaling events. As such, tubulin is able to orchestrate the function of multiprotein signaling complexes. These results also suggest the possibility of direct cross-regulation of cellular signaling and the reorganization of the microtubule cytoskeleton. Reciprocal interactions between G-protein signaling and the cytoskeleton might channel events triggered by diverse regulatory signals to different cellular compartments. The regulation of cell division, growth, motility, and morphology, as well as the movement of intracellular components, might be coordinated along this axis.

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