

Opposing effects of protein kinase C and protein kinase A on metabotropic glutamate receptor signaling: Selective desensitization of the inositol trisphosphate/Ca²⁺ pathway by phosphorylation of the receptor-G protein-coupling domain

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Signaling by the metabotropic glutamate receptor 1 α (mGluR1 α) can lead to the accumulation of inositol 1,4,5-trisphosphate (InsP₃) and cAMP and to the modulation of K⁺ and Ca²⁺ channel opening. At present, very little is known about how these different actions are integrated and eventually turned off. Unraveling the molecular mechanisms underlying these functions is crucial for understanding mGluR-mediated regulation of synaptic transmission. It has been shown that receptor-induced activation of the InsP₃ pathway is subject to feedback inhibition mediated by protein kinase C (PKC). In this study, we provide evidence for a differential regulation by PKC and protein kinase A of two distinct mGluR1 α -dependent signaling pathways. PKC activation selectively inhibits agonist-dependent stimulation of the InsP₃ pathway but does not affect receptor signaling via cAMP. In contrast, protein kinase A potentiates agonist-independent signaling of the receptor via InsP₃. Furthermore, we demonstrate that the selectivity of PKC action on receptor signaling rests on phosphorylation of a threonine residue located in the G protein-interacting domain of the receptor. Modification at Thr⁶⁹⁵ selectively disrupts mGluR1 α -G_{q/11} interaction without affecting signaling through G_s. Together, these data provide insight on the mechanisms by which selective down-regulation of a specific receptor-dependent signaling pathway can be achieved and on how cross-talk between different second messenger cascades may contribute to fine-tune short- and long-term receptor activity.

The excitatory actions of glutamate in the central nervous system are mediated by two distinct classes of receptors: ionotropic and metabotropic. While ionotropic receptors drive fast neurotransmission, the stimulation of metabotropic glutamate receptors generates slower and longer lasting changes in the signaling cascades activated in neuronal and glial cells. Metabotropic glutamate receptors (mGluRs; mGluR1 through mGluR8) are classified into three distinct groups according to their sequence homology, pharmacological profile, and signaling properties; group I receptors are linked to phosphoinositide metabolism while groups II and III inhibit adenylyl cyclase (1, 2). Many functions have been ascribed to these receptors, such as regulation of neurotransmitter release, direct mediation of glutamatergic synaptic transmission, modulation of membrane excitability, and regulation of *N*-methyl-D-aspartate, α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid, and γ -aminobutyric acid type A receptor activity (1). This vast array of effects underscores the role of mGluRs in modulating synaptic plasticity and their neurotoxic as well as neuroprotective action (1, 3).

As for other G protein-coupled receptors (GPCRs), mGluRs induce diverse and long-lasting cellular responses; these actions depend on the effectors regulated by each second messenger molecule and on the cross-talk between signaling pathways. In

the past several years, it has become increasingly clear that many GPCRs can initiate simultaneously multiple second messenger cascades because of their ability to couple to more than one G protein α subunit (4) and to the functional properties of the released $\beta\gamma$ subunits (5). In addition, several new effector molecules interacting with GPCRs have been uncovered; often these functional associations appear to be direct and not secondary to G protein activation, thus disclosing unexpected physiological properties of the receptor superfamily (6). In the case of mGluR1 α , the receptor has been shown to induce InsP₃ accumulation and consequent release of Ca²⁺ from intracellular stores (7), accumulation of cAMP (7–9), and possibly activation of the mitogen-activated protein kinase pathway (10, 11). In addition, mGluR1 α appears to regulate K⁺ (12) and Ca²⁺ (13, 14) channel opening and *N*-methyl-D-aspartate receptor activity (15), although the molecular mechanisms of these functional interactions are not clear at present.

Generally, GPCR responses desensitize rapidly, a process which is largely dependent on phosphorylation of either the receptor itself or other molecules engaged in the signaling cascade; in particular, for the β_2 -adrenergic receptor the molecular basis of homologous and heterologous desensitization have been partially deciphered (16–18). However, most of these studies have dealt with the analysis of linear signaling cascades, while fewer have addressed the question of how multiple receptor-dependent pathways are coordinately regulated (19–21).

Altogether, very little is known on how mGluR signaling is turned off, despite the fact that this is a critical step *in vivo* for receptor functions such as regulation of glutamate release (22). Very recently, it has been shown that overexpressed RGS proteins alter mGluR-mediated modulation of Ca²⁺ and K⁺ channels, a mechanism that may play a role in the regulation of receptor activity in pathological conditions (14, 23). On the other hand, in physiological conditions, mGluR-mediated stimulation of the InsP₃ pathway desensitizes upon prolonged exposure to specific agonists (24–27) or following repeated stimulation (28–30). In these cases, protein kinase C (PKC) activation appears to mediate these effects.

Abbreviations: mGluR, metabotropic glutamate receptor; GPCR, G protein-coupled receptor; InsP₃, inositol-1,4,5-trisphosphate; PKC, protein kinase C; PKA, protein kinase A; GST, glutathione S-transferase; i2, second intracellular loop; InsP, inositol phosphate.

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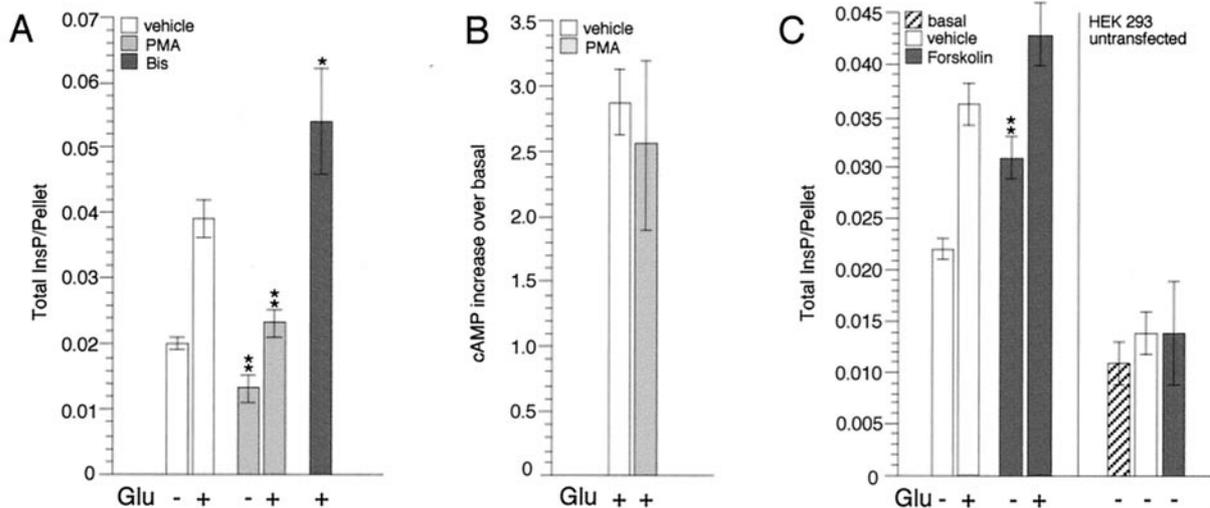


Fig. 1. PKC and PKA regulate mGluR1 α signaling. (A) Effect of PMA and bisindolylmaleimide I (Bis) on mGluR1 α -dependent InsP accumulation. For each experiment, HEK 293 cells were transfected with 10 μ g of mGluR1 α and 10 μ g of carrier DNA and seeded into six wells of a 24-well cluster. Stimulation was carried out in Hanks' saline solution (containing divalents and glucose) after clearing glutamate from the media with glutamic-pyruvic transaminase/pyruvate (9). The cells were treated with either Me₂SO (vehicle) or test reagents and then stimulated with Glu. Results represent means \pm SEM of duplicate or triplicate determinations obtained from at least two independent experiments (*, $P \leq 0.05$; **, $P \leq 0.01$; two-tailed t test). (B) Effect of PMA on mGluR1 α -dependent cAMP accumulation. HEK 293 cells were transfected with 5 μ g of mGluR1 α along with 5 μ g of G_s and 10 μ g of carrier DNA and seeded into four wells of a 24-well cluster. The cells were preincubated with either Me₂SO or PMA for 30 min and then stimulated with Glu for 20 min in the continued presence of the test reagents. Results represent means \pm SEM of duplicate determinations obtained from three independent experiments. (C) Effect of forskolin on mGluR1 α -dependent InsP accumulation. Cell transfection and treatment were conducted as described in A; in control experiments, 4×10^5 untransfected cells were plated for each well. Results represent means \pm SEM of triplicate determinations obtained from three independent experiments; for untransfected cells, results shown are means \pm SD of triplicate determinations, representative of two independent experiments.

Selective Desensitization of the InsP₃ Pathway Proceeds from PKC-Mediated Phosphorylation of Thr⁶⁹⁵ in the G Protein-Coupling Domain of mGluR1 α .

Prompted by the observation that PKC activation selectively desensitizes the InsP₃ pathway following receptor stimulation, we sought to elucidate the molecular mechanisms of this effect. It has been shown that mGluR1 α itself is substrate for PKC and that its level of phosphorylation increases upon stimulation with agonist (34). We reasoned that if PKC-mediated phosphorylation targeted residues within the receptor involved in the interaction with G_{q/11} but not G_s, this could lead to the selective uncoupling from one signaling pathway. We have previously shown that Thr⁶⁹⁵ in the second intracellular loop of mGluR1 α takes part in the interaction with G_{q/11} but not with G_s (9). Thr⁶⁹⁵ is located within a hinge region that connects two putative α -helices and that is present only in group I mGluRs. This region is rich in basic amino acids and Thr⁶⁹⁵ falls within a consensus site for PKC-mediated phosphorylation (ref. 35; Fig. 2A). Together, these observations suggested that Thr⁶⁹⁵ could be a potential regulatory site of receptor activity, possibly involved in the selective desensitization of the InsP₃/Ca²⁺ pathway.

We first tested whether a mutant receptor in which Thr⁶⁹⁵ is substituted with an alanine (Thr⁶⁹⁵Ala), a residue that cannot be phosphorylated, differed from the wild-type receptor in its susceptibility to PKC in transfected HEK 293 cells. Following PMA treatment, both the constitutive and Glu-stimulated activity of the mutant receptor were decreased as for the wild-type (Fig. 2B). Consistently, incubation of the transfected cells with bisindolylmaleimide I caused an increase in total InsP accumulation (Fig. 2B). From this analysis, mutation of Thr⁶⁹⁵ did not appear to affect the receptor-mediated stimulation of the InsP₃ pathway and the PKC-dependent desensitization. However, in this experimental paradigm, receptor activity is monitored over a long period and therefore rapid and transient changes in receptor responses that happen in neurons cannot be detected. In addition, constitutive activation along with the prolonged

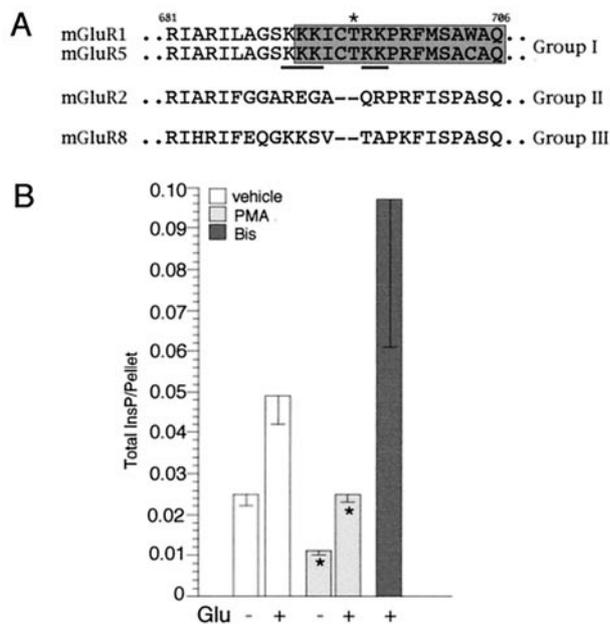


Fig. 2. (A) Sequence alignment of the second intracellular loop of mGluRs; the region involved in selective coupling of mGluR1 α with G_{q/11} is boxed in gray. An asterisk highlights Thr⁶⁹⁵, which we propose to be substrate for PKC-mediated phosphorylation. Basic amino acid residues which can constitute the consensus for PKC are underlined. (B) Effect of PMA and bisindolylmaleimide I (Bis) on the Thr⁶⁹⁵Ala mutant receptor activity. Cell transfection and treatment were conducted as described for the wild-type receptor (Fig. 1A). Results represent means \pm SEM of duplicate or triplicate determinations obtained from at least two independent experiments (*, $P \leq 0.05$; two-tailed t test).

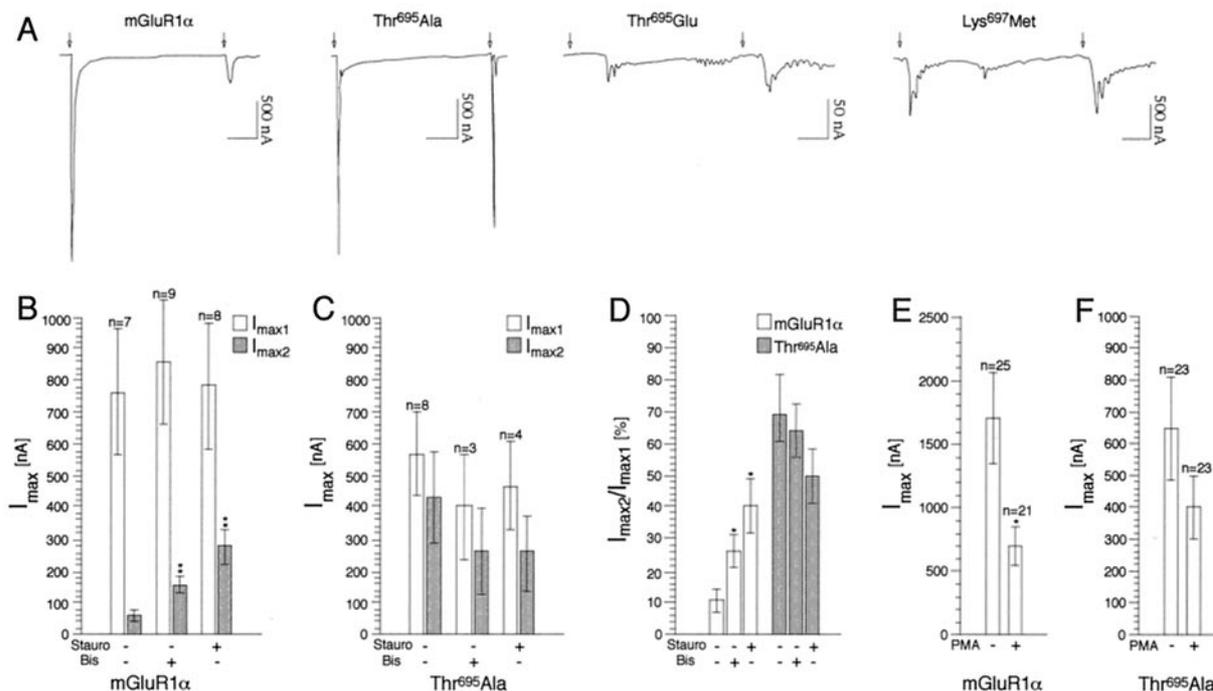


Fig. 3. Rapid desensitization of mGluR1 α depends on the phosphorylation state of Thr⁶⁹⁵. (A) Representative tracings of currents generated by wild-type and mutant receptors expressed in oocytes upon repeated application of Glu (1 mM); arrows indicate beginning of Glu application which lasted for 1 min. Time scale: 1 min. (B–F) PKC dependence of wild-type and mutant receptor responses. Injected oocytes were incubated with either staurosporin (100 nM) or bisindolylmaleimide I (2 μ M) or PMA (100 nM) before recordings. Results are means \pm SEM of values obtained from two independent experiments (*, $P \leq 0.05$; **, $P \leq 0.01$; two-tailed t test).

exposure to agonist are likely to induce a down-regulation of receptor activity that may mask transient alterations in the kinetic of the responses.

To examine whether phosphorylation at Thr⁶⁹⁵ could play a role in the rapid desensitization of receptor responses, we expressed wild-type and mutant receptors in *Xenopus* oocytes. In these cells, the release of intracellular Ca²⁺ induced by receptor stimulation causes the opening of Ca²⁺-activated Cl⁻ channels, thus generating an inward current which is a measure of receptor activity. A 1-min pulse of Glu induced in mGluR1 α -injected oocytes a large and transient current which rapidly goes back to baseline (Fig. 3A). Repeated applications of Glu after 1-, 2-, or 3-min intervals do not elicit responses (data not shown). After a 5-min washout with bath solution, a second application of Glu gave rise to a transient current of largely decreased amplitude compared with the first response ($I_{2\max}/I_{1\max} = 0.17 \pm 0.036$, $n = 23$; Fig. 3A, B, and D). To test whether desensitization of the receptor response was due to PKC activation, the injected oocytes were incubated with two different PKC inhibitors, staurosporin and bisindolylmaleimide I, for ≥ 20 min before stimulation with Glu. Both compounds caused a significant increase in the amplitude of the second response (Fig. 3B and D), although they were not capable to fully prevent desensitization. Conversely, incubation with PMA at submaximal doses (100 nM for 10–20 min) before stimulation with Glu induced about 60% inhibition in current amplitude (Fig. 3E, control $I_{\max} = 1694 \pm 360$, $n = 25$; PMA-treated $I_{\max} = 686 \pm 151$, $n = 21$), confirming the role of PKC in regulating receptor desensitization.

We then analyzed the responses evoked by two different mutant receptors in which Thr⁶⁹⁵ was replaced with either a neutral amino acid (Thr⁶⁹⁵Ala) or with a negatively charged residue (Thr⁶⁹⁵Glu), a mutation which mimics the effect produced by the addition of a phosphate group. A mutant was also

tested in which the putative consensus for PKC-mediated phosphorylation is disrupted (Lys⁶⁹⁷Met; Fig. 2A). The Thr⁶⁹⁵Ala mutant evoked robust and transient currents comparable to the wild-type receptor (Fig. 3A). Interestingly, after a second challenge with agonist, the receptor showed only partial desensitization ($I_{2\max}/I_{1\max} = 0.52 \pm 0.08$, $n = 17$; Fig. 3A, C, and D). Neither staurosporin nor bisindolylmaleimide I have any significant effect on the amplitude of the second response (Fig. 3C and D). In addition, incubation with PMA was much less effective (about 30%) in blocking receptor activity compared with wild type (Fig. 3F, control $I_{\max} = 646 \pm 161$, $n = 23$; PMA-treated $I_{\max} = 401 \pm 95$, $n = 23$). Together, these results demonstrate that the Thr⁶⁹⁵ residue is indeed critical for the Glu-induced receptor desensitization mediated by PKC. Mutant receptors harboring the Thr⁶⁹⁵Glu substitution gave rise to currents decreased in amplitude by an order of magnitude, slower in arising and returning to baseline. In addition, their waveform differed from wild type, being very often oscillatory: a representative current tracing obtained from this mutant is shown in Fig. 3A. These responses do not desensitize ($I_{2\max}/I_{1\max} = 1.23 \pm 0.28$, $n = 8$). Finally, disruption of the PKC consensus sequence by the Lys⁶⁹⁷Met mutation generated currents slightly smaller in amplitude than those in wild type, partially oscillatory and with largely reduced desensitization ($I_{2\max}/I_{1\max} = 0.65 \pm 0.13$, $n = 15$; Fig. 3A). For all receptors, a certain variability in the amplitude of the responses was observed when comparing experiments conducted with different batches of oocytes. However, the extent of desensitization was not correlated with the amplitude of the first response.

Thr⁶⁹⁵ Is a Substrate for PKC-Dependent Phosphorylation *In Vitro*. To test whether Thr⁶⁹⁵ can be a substrate for PKC-dependent phosphorylation, we generated a chimeric protein carrying GST fused to the second intracellular loop of mGluR1 α (GST-i2).

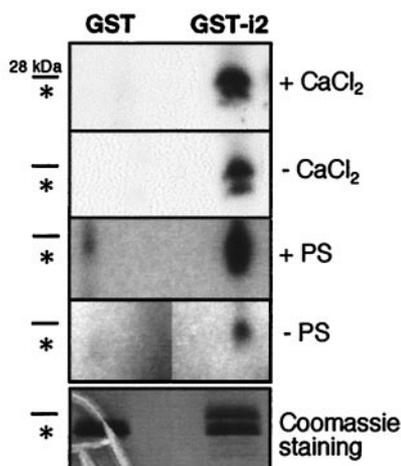


Fig. 4. Thr⁶⁹⁵ is phosphorylated *in vitro*. Recombinant fusion protein (GST-i2) and GST were used as substrate for phosphorylation by PKC purified from brain. Data shown are representative of three independent experiments. To verify for the specificity of the phosphorylation, the effects of lipids [phosphatidyl serine (PS)] and calcium on enzyme activity were tested. GST-i2 predicted molecular mass is 28 kDa; GST alone is 26 kDa and its position on the gel is indicated by an asterisk. The gels were exposed for autoradiography at -70°C for 12 h (with or without CaCl_2) and 8 h (with or without PS), respectively.

Point mutations were introduced at Ser⁶⁸⁹ and Ser⁷⁰² of i2, so that the Thr⁶⁹⁵ residue was the only one available as substrate for PKC-mediated phosphorylation. The construct was expressed in bacteria and the corresponding recombinant protein was purified by affinity chromatography and incubated with PKC holoenzyme purified from brain. The results shown in Fig. 4 confirm that Thr⁶⁹⁵ is a substrate for PKC and the specificity of the reaction is further illustrated by its lipid and calcium dependence.

Discussion

In recent years, our appreciation of the importance of intracellular signaling networks in shaping cellular responses has considerably increased (36). In analogy with other systems, we reasoned that the interplay between different signaling cascades is likely to have a critical role in determining the final output of mGluR activation. As an experimental paradigm to explore the possible cross-talk between different mGluR effectors, we selected two second messenger pathways activated in response to glutamate, $\text{InsP}_3/\text{Ca}^{2+}$ and cAMP, and examined their regulation by the respective downstream targets, PKC and PKA. Activation of the $\text{InsP}_3/\text{Ca}^{2+}$ pathway by group I mGluRs in neurons has been extensively characterized (1). In addition, stimulation of these receptors has been shown to increase cAMP levels during early developmental stages in the hippocampus (37), during the critical period in the visual cortex (38), and in neuronal cultures (39), although the molecular mechanisms underlying this effect are less well understood.

We found that PKC activation drastically curtails both the basal and Glu-stimulated activity of mGluR1 α ; however, under the same conditions, receptor-mediated stimulation of the cAMP pathway was not significantly altered. Feedback inhibition by PKC appears therefore to target selectively only mGluR1 α signaling via InsP_3 . This opens the interesting possibility that receptor stimulation in neurons following neurotransmitter release could lead to the selective desensitization of this signaling pathway while leaving intact other receptor functions. Remarkably, such dual regulation of mGluR activity has been reported to take place both at presynaptic (30, 40) and postsyn-

aptic sites (41). For example, activation of a phosphoinositide-linked mGluR at presynaptic terminals caused both facilitation and inhibition of glutamate release; of these two independent actions, only the signaling pathway that leads to facilitation desensitizes rapidly following PKC activation (40).

The effect of PKA on mGluR activity was more complex; treatment with forskolin potentiated rather than decreased receptor signaling via the $\text{InsP}_3/\text{Ca}^{2+}$ pathway, affecting in particular its basal activity. PKA appears therefore to have an opposing effect to PKC by inducing a sustained coupling of mGluR1 α to the InsP_3 pathway even in the absence of agonist. Several considerations arise from this observation: first, PKA action may target directly the receptor itself and/or indirectly other molecules involved in the InsP_3 pathway. However, the fact that we did not observe a forskolin-induced increase in total inositols in control cells seems to point to a direct effect of the kinase on the receptor. Second, in neurons, simultaneous activation of other GPCRs linked to PKA activation may also lead to potentiation of mGluR1 responses by strengthening its signaling via InsP_3 .

To further understand the molecular mechanisms which govern mGluR function, we have attempted to clarify the nature of the structural determinants underlying the PKC-dependent pathway-selective desensitization. Wild-type mGluR1 α was expressed in *Xenopus* oocytes and the kinetics of the receptor response following brief and repeated stimulations with agonist was examined. In this system, mGluR1 α desensitized drastically and rapidly in response to glutamate. Desensitization ensued from PKC activation, since preincubation with specific PKC inhibitors partly rescued receptor responses. The fast and transient mGluR1 α -induced currents recorded in oocytes correspond physiologically to the Ca^{2+} transients induced in other expression systems (42, 43) where PKC-dependent desensitization has also been reported (43). Thus, having available a system in which rapid and PKC-dependent changes in receptor activity can be analyzed, we set out to examine the effect of mutations of a putative PKC substrate, Thr⁶⁹⁵. Remarkably, substitution of the residue with Ala generated a mutant receptor that shows limited desensitization. Consistently, treatment with PKC inhibitors did not affect the mutant receptor responses. Together, these data indicate that Thr⁶⁹⁵ is an important target for PKC in the regulation of Glu-induced receptor activity. Thr⁶⁹⁵ was also replaced with Glu, a residue which by itself is negatively charged and hence mimics the modification which is introduced upon phosphorylation. Such strategy has proved useful in many cases to study the effects of phosphorylation with mutant molecules resembling constitutively phosphorylated substrates. In keeping with the prediction, the mutant Thr⁶⁹⁵Glu elicited responses greatly diminished in amplitude, as would be expected for a receptor "constitutively desensitized"; in addition, this mutant does not further desensitize following repeated stimulation with Glu. It is worth noting that all of the described mutations affect selectively $\text{InsP}_3/\text{Ca}^{2+}$ signaling, since the efficacy of the mutant receptors in stimulating the cAMP pathway is unaltered compared with wild type as previously shown (9).

On the basis of these observations, we propose a model for PKC action in which Thr⁶⁹⁵ is substrate for the kinase and acts as a molecular switch for mGluR1 activity. Upon ligand binding, a conformational change in the receptor is produced which would render the Thr⁶⁹⁵ residue accessible to phosphorylation. The added phosphate group would then prevent further coupling to $\text{G}_{q/11}$ but not to G_s . Interestingly, it has been recently reported that in cortical synaptosome preparations, PKC activation inhibits an mGluR-induced increase in $\text{GTP}\gamma\text{S}$ binding (44). This observation provides further evidence for a PKC-mediated uncoupling of mGluRs from their cognate G proteins. It has been shown that mGluR1 α is phosphorylated *in vivo* both in the resting and agonist-bound conformation (34); however, the

number and identity of the phosphorylated sites is still unknown. Our results suggest that different residues could be involved in controlling the basal or Glu-stimulated activity of the receptor, with Thr⁶⁹⁵ being a target for Glu-induced PKC-mediated phosphorylation.

Phosphoinositide-linked mGluRs, mGluR1 and 5, are generally regarded as functionally homologous because of the high similarity in their primary structure. However, recent reports have pointed out some interesting differences in the mode of intracellular Ca²⁺ release induced by these receptors. While mGluR1 induces single transient peaks of Ca²⁺, mGluR5 activation elicits oscillatory responses (42, 43). Such oscillations appear to be generated by PKC-dependent phosphorylation of a threonine residue which is present only in the carboxyl tail of

mGluR5. Our work now reveals another difference in the regulation of the activity of these receptors. In fact, while in mGluR1 Thr⁶⁹⁵ plays a key role in regulating InsP₃ signaling, mutation of the corresponding residue in mGluR5 generates inactive receptors (45). The different signaling properties and mechanisms of regulation of these receptors suggest that they subserve distinct functions in the neuronal environment.

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