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Rapid CB₁ cannabinoid receptor desensitization defines the time course of ERK1/2 MAP kinase signaling

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Summary

Molecular mechanisms regulating the development of physiological and behavioral tolerance to cannabinoids are not well understood. Two cellular correlates implicated in the development and maintenance of tolerance are CB₁ cannabinoid receptor internalization and uncoupling of receptor signal transduction. Both processes have been proposed as mediators of tolerance because of observations that chronic Δ⁹-tetrahydrocannabinol (THC) treatment causes both region-specific decreases in CB₁ receptors and G-protein coupling in the brain. To determine the balance of these two processes in regulating CB₁ receptor signaling during sustained receptor stimulation, we evaluated the parameters affecting ERK1/2 MAP kinase activity in HEK293 cells stably expressing CB₁ receptors. CB₁ receptor stimulation by the potent CB₁ receptor agonist, CP 55,940 transiently activated ERK1/2. To determine if CB₁ receptor desensitization or internalization was responsible for the transient nature of ERK1/2 activation, we evaluated ERK1/2 phosphorylation in HEK293 cells expressing a desensitization-deficient CB₁ receptor (S426A/S430A CB₁). Here, the duration of S426A/S430A CB₁ receptor-mediated activation of ERK1/2 was markedly prolonged relative to wild-type receptors, and was dynamically reversed by SR141716A. Interestingly, the S426A/S430A CB₁ receptor was still able to recruit βarrestin-2, a key mediator of receptor desensitization, to the cell surface following agonist activation. In contrast to a central role for desensitization, pharmacological and genetic approaches suggested CB₁ receptor internalization is dispensable in the transient activation of ERK1/2. This study indicates that the duration of ERK1/2 activation by CB₁ receptors is regulated by receptor desensitization and underscores the importance of G-protein uncoupling in the regulation of CB₁ receptor signaling.

Keywords

cannabinoids; MAP kinase; desensitization; internalization; phosphorylation; βarrestin

1. Introduction

Most CNS effects of cannabinoids, the principal psychoactive ingredients of marijuana and their endogenous counterparts, are mediated by the CB₁ cannabinoid receptor. It is well established that the CB₁ receptor couples in an inhibitory manner to voltage-dependent Ca²⁺

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channels and adenylyl cyclase and activates G-protein-regulated inwardly rectifying K⁺ (GIRK) channels (Howlett et al., 2002; Howlett and Fleming, 1984; Mackie and Hille, 1992; Mackie et al., 1995). *In vitro* studies have demonstrated that cannabinoids can also induce the activation of mitogen-activated protein (MAP) kinases of the extracellular signal-regulated kinase (ERK) superfamily (Bouaboula et al., 1995). Consistent with these findings, acute THC administration has been reported to increase CB₁ receptor-mediated ERK1/2 activation in the dorsal striatum, nucleus accumbens, and hippocampus (Derkinderen et al., 2003; Valjent et al., 2001), while chronic THC treatment has been shown to increase ERK1/2 protein levels in the prefrontal cortex and hippocampus *in vivo* (Rubino et al., 2004).

Chronic administration of cannabinoids leads to the rapid development of tolerance. This tolerance is accompanied by changes in CB₁ receptor number and/or function (Bass et al., 2004; Breivogel et al., 2003; Gonzalez et al., 2005; Sim-Selley et al., 2003) and the inhibition of cannabinoid-mediated plasticity in different brain regions (Hoffman et al., 2003; Mato et al., 2004). The uncoupling of G-proteins from their cognate receptor, or desensitization, terminates receptor signaling and has been implicated as one of the cellular adaptations underlying tolerance (Martin et al., 2004). G-protein-coupled receptors (GPCRs) can be desensitized following agonist activation through phosphorylation by GPCR kinases (GRKs) and subsequent β arrestin binding (Gainetdinov et al., 2004). The physical interaction of β arrestin with the phosphorylated receptor reduces the affinity of the GPCR for G-proteins, likely via a mechanism involving steric hindrance, thereby attenuating signaling (Dewire et al., 2007). Previous studies have found that the β arrestin-2 isoform of β arrestin and GRK3 are capable of desensitizing CB₁ receptor-mediated activation of GIRK channels and inhibition of neurotransmission (Jin et al., 1999; Kouznetsova et al., 2002). Furthermore, serines 426 and 430 in the proximal carboxy-terminus of CB₁ are the likely GRK3 phosphorylation sites underlying β arrestin-2 mediated CB₁ receptor desensitization of GIRK activation (Jin et al., 1999).

Following binding of agonist and activation, CB₁ receptors internalize via clathrin-coated pits. However, the details and regulation of this process are now only emerging (Coutts et al., 2001; Hsieh et al., 1999; Leterrier et al., 2004; McDonald et al., 2007; Rinaldi-Carmona et al., 1998). Specifically, while β arrestin-2 is a multi-functional protein (Dewire et al., 2007), whose binding has also been proposed to regulate many forms of receptor endocytosis (Claing et al., 2002), its involvement in CB₁ receptor internalization remains unknown.

In these studies we used HEK293 cells stably expressing CB₁ to investigate the relative contributions of CB₁ receptor desensitization and internalization to the time course of ERK1/2 phosphorylation following CB₁ receptor activation. We found that CB₁ receptor desensitization primarily determines the time course of ERK1/2 activation. In contrast, CB₁ internalization is required for neither the onset nor the decay of ERK1/2 phosphorylation.

2. Materials and Experimental Methods

2.1 Reagents and drugs

CP 55,940 ((-)-*cis*-3-[2-hydroxy-4-(1,1-dimethylheptyl)phenyl]-*trans*-4-(3-hydroxypropyl)cyclohexanol) and SR141716A (4-Chlorophenyl)-1-(2,4-dichloro-phenyl)-4-methyl-*N*-(piperidin-1-yl)-1*H*-pyrazole-3-carboxamide) were provided by the National Institute of Drug Abuse Drug Supply Program (RTI, Research Triangle Park, NC). Concanavalin A, was purchased from Vector Laboratories (Burlingame, CA). Pertussis toxin and cholera toxin were purchased from List Biological Laboratories, Inc. (USA).

2.2 Generation of mutant CB₁ receptor constructs

The CB₁ receptor mutant, S426A/S430A, was made using a modified QuikChange PCR strategy (Stratagene, La Jolla, CA) to introduce serine to alanine point mutations. Full-length rat CB₁, with an amino-terminal pplsHA (prolactin signal sequence and hemagglutinin epitope) tag (Andersson et al., 2003) in pcDNA3.0 (Invitrogen, Carlsbad, CA), was used as a template with the following primers: forward 5'

ccgcacagcctcttgacaacgccatgggggacgcagactgcctgc-3' and reverse 5'-

gcaggcagtctgcgtccccatggcgttgcaagaggctgtgcgg-3'. A silent mutation was inserted (the 14th nucleotide) to remove the XbaI site for screening purposes. Integrity of the coding region for all constructs was confirmed by sequencing. The D164N CB₁ receptor mutant was generated by overlapping extension PCR using full-length rat CB₁ in pcDNA3.0 as template (Roche et al., 1999).

2.3 Cell culture and transfection

HEK293 cells were grown in DMEM containing 10% fetal bovine serum and penicillin/streptomycin (GIBCO, Carlsbad, CA) at 37°C in 5% CO₂. Transfections were performed in 35-mm dishes with 2 µg of the appropriate plasmid and Lipofectamine 2000 (Invitrogen, Carlsbad, CA), following the manufacturer's protocol. Stable cell lines were generated by selection in Geneticin (G418; Invitrogen, Carlsbad, CA). G418-resistant colonies were evaluated for the surface expression of CB₁ by live cell immunostaining using an antibody (Covance, Berkeley, CA) directed towards the amino-terminal, extracellular HA epitope tag and a fluorescein isothiocyanate (FITC) secondary antibody (Jackson ImmunoResearch Laboratories Inc., West Grove, PA). Clones expressing uniform and moderate to high levels of CB₁ were expanded and used for the subsequent experiments as previously described (Brown et al., 2002; Hsieh et al., 1999).

2.4 Quantitative internalization assay

HEK293 cells stably expressing rat CB₁ receptors (wild-type or with S426A/S430A or D164N mutations) were seeded onto poly-D-lysine-coated 96-well plates (Corning, Corning, NY) and grown until 90% confluent in DMEM containing 10% fetal bovine serum and penicillin/streptomycin at 37°C in 5% CO₂. Prior to drug treatment, plates were washed once in HEPES-buffered saline (HBS; 130 mM NaCl, 5.4 mM KCl, 1.8 mM MgCl₂, and 10 mM HEPES, pH 7.5) containing 1.0 mg/ml bovine serum albumin. Cells were then incubated with treatment supplied in HBS containing 1.0 mg/ml bovine serum albumin and incubated at 37°C for the specified times. In all cases, the matching concentration of vehicle (DMSO) was included in control experiments. At the end of the drug treatment, the plate was placed on ice, the buffer was removed, and cells were immediately fixed with 4% paraformaldehyde for 30 min at room temperature. The integrity of the plasma membrane was evaluated following fixation using antibodies directed against both the amino- and carboxy-terminus of CB₁ (with and without the addition of detergent). Paraformaldehyde itself did not cause substantial permeabilization in this cell model (data not shown). Following fixation, cells were washed 5X for 30 min in phosphate-buffered saline (PBS; 137 mM NaCl, 10 mM NaH₂ PO₄, 2.7 mM KCl, pH 7.4) (without detergent to avoid membrane permeabilization) and blocked for 90 min in LI-COR Odyssey Blocking Buffer® (LI-COR Biosciences, Lincoln, Nebraska) at room temperature with gentle rocking. After blocking, cells were incubated overnight at 4°C with a monoclonal anti-HA antibody (1:100; Covance, Berkeley, CA). The cells were washed 5X the following day in Tris-buffered saline containing 0.05% Tween-20 (TBST; 137 mM NaCl, 10 mM Tris, 0.05% Tween-20, pH 7.4) for 30 min. Bound primary antibody was detected with an IRDye 800 conjugated anti-mouse IgG (1:800; Rockland Immunochemicals, Inc., Gilbertsville, PA). Plates were washed 5X in TBST, dried, and the immunocomplex was quantified on a LI-COR Odyssey following the manufacturer's recommendations.

2.5 Quantitative Measurement of ERK1/2 Phosphorylation

HEK293 cells stably expressing either wild-type or modified CB₁ receptors were seeded onto poly-D-lysine-coated 96-well plates, grown until 90% confluent at 37°C in 5% CO₂. The cells were serum-starved overnight in DMEM containing only penicillin/streptomycin prior to the experiment. Drug treatments were performed as described in the preceding section. Following drug treatment, cells were fixed with 4% paraformaldehyde on ice for 15 min and then at room temperature for an additional 45 min. Cell membranes were permeabilized with 100% methanol at -20°C for ≥ 20 min and wells were blocked for 1.5 hr in Tris-buffered saline (TBS; 137 mM NaCl, 10 mM Tris, pH 7.4) containing 5.0 mg/ml bovine serum albumin. Following blocking, cells were incubated overnight at 4°C with a phospho-ERK1/2-specific antibody (1:200; Cell Signaling Technologies Inc., Danvers, MA). The primary antibody was removed with 5X washes over 1.5 hr in TBST and detected with an IRDye 800 conjugated anti-rabbit IgG (1:1000; Rockland, Gilbertsville, PA). Plates were washed 5X for 1.5 hr in TBST, dried and immunofluorescence was quantified on the LI-COR Odyssey.

2.6 Immunocytochemistry and fluorescent imaging

Immunostaining of cells stably expressing either the wild-type, S426A/S430A, or D164N CB₁ receptors was performed using a previously described method to detect surface and intracellular receptors (Hsieh et al., 1999; Kearn et al., 2005). Briefly, cells were grown on poly-D-lysine-coated glass coverslips, washed in HBS and incubated with the specified drug diluted in HBS solution containing 0.2 mg/ml bovine serum albumin. After drug treatment, cells were fixed with 4% paraformaldehyde for 20 min, washed 2X with phosphate-buffered saline (PB; 10 mM NaH₂PO₄, 2.7 mM KCl, pH 7.4), and 3X with PBS. Cells were blocked for 1 hr in PBS containing 5% donkey serum and 0.1% saponin (to permeabilize cell membranes) at room temperature. After blocking, cells were incubated overnight at 4°C with a monoclonal mouse anti-HA antibody (1:1000; Covance, Berkeley, CA). Following 5X washes in PBS, immunoreactivity was detected with a FITC anti-mouse IgG secondary antibody (1:150; Jackson ImmunoResearch Laboratories Inc., West Grove, PA). The coverslips were washed 2X in PBS, 2X PB, 1X in water, dried, mounted in Vectashield (Vector Laboratories, Burlingame, CA), and imaged on a Leica SP1/MP confocal microscope. All images (1024 X 1024 pixels) were obtained by the use of a 100X objective.

For immunocytochemical phospho-ERK1/2 detection, HEK293 cells were treated the same as above through the fixation step. However, cells were then permeabilized with 100% methanol for ≥ 30 minutes at -20°C and then incubated overnight with a rabbit polyclonal p-ERK1/2 antibody (1:500; Cell Signaling Technology, Danvers, MA) diluted in a TBS solution containing 5.0 mg/ml BSA. A FITC anti-rabbit IgG antibody (1:150; Jackson ImmunoResearch Laboratories Inc., West Grove, PA) was used to detect the immunocomplex the following day. All confocal images of the phospho-ERK1/2 immunocomplex were acquired with the same instrument settings.

For β arrestin-2 recruitment studies, HEK293 cells were transiently transfected with constructs encoding either wild type or S426A/S430A CB₁ receptors and an RFP-tagged β arrestin-2 fusion protein. GFP-tagged β arrestin-2 (a gift from Dr. Marc Caron) was modified using standard molecular biology techniques to exchange the GFP for RFP (Campbell et al., 2002). The cells were plated onto coverslips the same day and the experiment was performed 24 hours post-transfection. CB₁ receptor expression was detected using the above-mentioned HA antibody and protocol.

2.7 Statistical analysis

In the internalization studies, the percentage of cell surface receptors remaining was determined by first calculating the mean integrated intensity values for each drug concentration.

Background subtraction was not taken into account for individual experiments. However, non-specific binding was determined to be 5–10% for antibodies tested during the assay development phase. Final values were normalized to the mean integrated intensity values for the no drug treated controls and are represented as the mean \pm SEM. Extent of internalization curves were fit by nonlinear regression. The half-life of the decay ($t_{1/2}$) was calculated by fitting the data to a single-phase exponential decay model. In the ERK1/2 phosphorylation studies, percent maximal response for ERK1/2 activation was calculated by defining basal as the value at time zero and setting the maximal peak to 100%. The statistical significance with respect to the extent of internalization and maximal ERK1/2 activation was determined with an unpaired student's t-test (** $p < 0.01$, * $p < 0.05$). To quantify β arrestin-2 recruitment, the membrane to cytosolic ratio of fluorescence in the 561 nm channel (obtained using the 100X objective with 2X zoom), was calculated by region-specific analysis of averaged integrated intensity values using MetaMorph software (Molecular Devices, Downingtown, PA). Co-detection of RFP and FITC fluorescence was done by excitation with 488 and 561 nm lasers. These data are represented as the mean AFU \pm SEM. Statistical significance with respect to the membrane:cytosolic ratios was determined with an unpaired student's t-test (** $p < 0.01$, * $p < 0.05$). All graphs and statistical analyses were generated using GraphPad Prism 4.0 software (Hearne Scientific Software, Chicago, IL).

3. Results

3.1 Mutation of serines 426 and 430 of the CB₁ receptor prolong the time course of ERK1/2 activation in a SR141716A reversible manner

To test the hypothesis that putative GRK phosphorylation sites (serines 426 and 430) (Jin et al., 1999) in the proximal carboxy-terminus of the CB₁ receptor determine the rate of ERK1/2 inactivation, the kinetics of ERK1/2 phosphorylation were evaluated following agonist stimulation of wild-type or S426A/S430A CB₁ receptors stably expressed in HEK293 cells. A maximally efficacious concentration (100 nM) of the CB₁ agonist, CP 55,940, was used for subsequent experiments ($EC_{50} = 0.97 \text{ nM} \pm 0.06$ for ERK1/2 activation at 7 min; data not shown). CP 55,940 stimulation of wild-type CB₁ resulted in the transient activation of ERK1/2 with peak activation at 5 min (Fig. 1A). ERK1/2 phosphorylation then rapidly decayed ($t_{1/2} = 4.9 \text{ min}$) to near basal levels. While both wild-type and S426A/S430A CB₁ receptors activated ERK1/2 to a similar extent (2–3 fold activation over basal; data not shown), there was little decay ($71 \pm 4\%$ of peak at 15 min) in ERK1/2 phosphorylation with S426A/S430A CB₁ receptor stimulation (Fig. 1A, Bf) relative to the wild-type receptor ($33 \pm 5\%$ of peak at 15 min). Significantly, the sustained ERK1/2 activation was very long lasting in S426A/S430A CB₁ expressing cells ($71 \pm 4\%$ of peak at 45 min for S426A/S430A CB₁ versus $20 \pm 3\%$ at 45 min for wild-type CB₁). This prolonged ERK1/2 activation was observed in eight different monoclonal cell populations expressing S426A/S430A CB₁ receptors (data not shown) and is not due to an increase in the constitutive activity of the modified receptors. Treatment with an inverse agonist, 1 μM SR141716A, does not decrease basal ERK1/2 activation in cells expressing either wild-type or S426A/S430A CB₁ receptors ($107 \pm 1\%$ of peak for S426A/S430A CB₁ versus $100 \pm 3\%$ for wild-type CB₁ following a 30 min incubation with SR1; data not shown). Additionally, the time course of ERK1/2 activation following 3 μM Δ^9 -THC treatment is similar in cells expressing either wild-type or S426A/S430A CB₁ receptors (data not shown). Since these residues are required for homologous desensitization of GIRK activation by CB₁ (Jin et al., 1999), this result suggests that CB₁ receptor desensitization modulates the rapid decay of ERK1/2 phosphorylation. To evaluate if ERK1/2 phosphorylation was dependent upon CB₁ receptor activation, we pretreated cells expressing wild-type or S426A/S430A CB₁ receptors with 1 μM of the CB₁ antagonist, SR141716A and then applied 100 nM of CP 55,940 in the presence of 1 μM SR141716A (Fig. 1Bg,h). SR141716A treatment prevented ERK1/2 phosphorylation indicating it is a CB₁ receptor-dependent process.

To determine if the prolonged phosphorylation of ERK1/2 reflects sustained S426A/S430A CB₁ receptor activation, cells were first treated with 10 nM CP 55,940 for 35 min. Then 1 μM of the CB₁ receptor antagonist, SR141716A was added at 2.5 min intervals in the continued presence of 10 nM CP 55,940 until 45 min had elapsed (Fig. 1C). Phosphorylation of ERK1/2 by S426A/S430A CB₁ was rapidly ($t_{1/2}$ = 2.97 min) reversed by addition of SR141716A, suggesting sustained ERK1/2 phosphorylation was due to the continued signaling by the mutant S426A/S430A CB₁ receptor.

3.2 CB₁ receptor internalization is not necessary for ERK1/2 activation and does not contribute to the decay of ERK1/2 phosphorylation

Agonist-induced internalization of G-protein coupled receptors is a widely observed phenomenon. Receptor internalization has been suggested to be necessary for the activation of ERK1/2, as well as for receptor resensitization (Daaka et al., 1998). In both heterologous and endogenous expression systems, wild-type CB₁ receptors are internalized following agonist treatment (Coutts et al., 2001; Hsieh et al., 1999; Rinaldi-Carmona et al., 1998). Previously, we have shown using confocal microscopy that serines 426 and 430 are not required for rapid (< 30 min), agonist-induced CB₁ receptor internalization in AtT20 cells (Jin et al., 1999). In the present study we extended this observation to CB₁ receptors expressed in HEK293 cells in a more quantitative fashion. Consistent with previous findings, after 15 minutes of 100 nM CP 55,940 treatment (a time window that includes the period of peak ERK1/2 activation) S426A/S430A CB₁ and wild-type CB₁ receptors are internalized to the same extent ($78 \pm 2\%$ versus $78 \pm 0.6\%$ of receptors on the cell surface, respectively) (Fig. 2A). However, for agonist treatments >15 min, the extent of internalization for the mutant CB₁ receptor is significantly attenuated compared to wild-type receptors (for example, $77 \pm 2\%$ versus $58 \pm 0.4\%$ of receptors remain at the cell surface after 120 minutes of agonist treatment) (Fig. 2A).

To determine if CB₁ receptor internalization is required for ERK1/2 activation and/or the decay of phosphorylation in the continued presence of agonist, we examined the time course under conditions where internalization was blocked. We used a pharmacological inhibitor, concanavalin A (Con A), to prevent clathrin-dependent endocytosis following agonist treatment, a strategy utilized in previous studies (Arttamangkul et al., 2006). HEK293 cells stably expressing wild-type CB₁ receptors were treated with Con A (100 μg/ml) and stimulated with 10 nM CP 55,940 (Fig. 2B). Con A completely inhibited CB₁ receptor internalization during peak ERK1/2 activation (7 min) and greatly inhibited endocytosis following prolonged drug treatment (Fig. 2Bb, c). In the absence of Con A, a significant fraction of CB₁ receptors were endocytosed (Fig. 2Bd). To examine whether CB₁ receptor internalization was required for the transient activation of ERK1/2, we stimulated HEK293 cells expressing wild-type CB₁ receptors in the presence of Con A (Fig. 2C). Blockade of endocytosis by Con A did not significantly alter the time course of receptor-mediated ERK1/2 phosphorylation. Additionally, 10 nM CP 55,940-induced ERK1/2 phosphorylation by S426A/S430A CB₁ receptors was unaffected by Con A treatment (Fig. 2C).

A complementary approach was taken to further determine if CB₁ receptor internalization contributes to the decay of ERK1/2 phosphorylation. We evaluated the ability of a previously characterized mutant CB₁ receptor, D164N CB₁ (aspartate to asparagine at residue 164), (Roche et al., 1999) to phosphorylate ERK1/2. CB₁ receptors with a D164N modification do not undergo WIN 55,212-2-induced internalization in AtT20 cells but remain partially functional – in that they can inhibit Ca²⁺ channels and adenylyl cyclase, and activate MAP kinase, yet do not activate GIRK channels (Roche et al., 1999). We first verified the internalization-deficient finding in a quantitative manner by assessing agonist-induced internalization of D164N CB₁ receptors expressed in HEK293 cells by CP 55,940 (100 nM). Consistent with previous work, no significant internalization of D164N CB₁ receptors was

observed ($101 \pm 3\%$ surface receptors at 120 min) relative to wild-type CB₁ receptors ($58 \pm 0.4\%$ surface receptors at 120 min) following treatment with 100 nM CP 55,940 at time points ≥ 30 min (Fig. 3A). To preclude the possibility of agonist-specific trafficking, the extent of internalization of D164N CB₁ receptors was evaluated using a different CB₁ agonist, WIN 55,212-2 (1 μ M). In agreement with the CP 55,940 results, no internalization of D164N CB₁ receptors was observed with WIN 55,212-2 stimulation ($99 \pm 0.8\%$ surface receptors at 120 min for D164N versus $50 \pm 0.1\%$ for wild-type; data not shown). We then examined the time course of D164N CB₁ receptor-mediated activation of ERK1/2. Both wild-type and D164N CB₁ receptors activated ERK1/2 to a similar extent (2-fold over basal; data not shown) and with the same temporal pattern of activation (Fig. 3B). Taken together, these results indicate that CB₁ receptor internalization is not required for ERK1/2 phosphorylation nor does it modulate the time course of receptor-mediated activation.

3.3 Wild-type and S426A/S430A CB₁ receptors activate ERK1/2 independent of G_{i/o} G-protein signaling

Agonist stimulation of CB₁ receptors has been reported to activate ERK1/2 via pertussis toxin (PTX) sensitive G-proteins in heterologous expression systems (Bouaboula et al., 1999; Bouaboula et al., 1995; Kearm et al., 2005). Thus, we expected ERK1/2 activation in HEK293 cells to be PTX-sensitive. However, overnight treatment with 500 ng/ml PTX followed by 100 nM CP 55,940, did not alter the time course of wild-type or S426A/S430A CB₁ receptor-mediated phosphorylation of ERK1/2 (Fig. 4). However, the same treatment did inhibit depolarization-induced suppression of excitation in cultured hippocampal neurons (data not shown), attesting to the activity of the toxin. CB₁ receptors have been shown to functionally couple to stimulatory G_{as} G-proteins under certain conditions (Glass and Felder, 1997; Kearm et al., 2005). However, selective down-regulation of G_{as} with overnight cholera toxin treatment (100 ng/ml) did not prevent ERK1/2 activation (data not shown). Collectively these results suggest that CB₁ receptors expressed in HEK293 cells can mediate ERK1/2 activation through G-proteins other than those of the G_{ai/o} and G_{as} families.

3.4 Wild-type and S426A/S430A CB₁ receptors recruit β arrestin-2-RFP to the plasma membrane following agonist stimulation

Phosphorylation of serine and threonine residues in the carboxy-terminus of many GPCRs by GRKs and subsequent β arrestin binding is thought to be a common mechanism for both receptor internalization and desensitization of GPCRs (Gainetdinov et al., 2004). β arrestin-2 has been implicated as a key regulator of desensitization of CB₁ receptor-mediated coupling to GIRK currents and excitatory glutamatergic transmission in *Xenopus* oocytes and cultured hippocampal neurons, respectively (Jin et al., 1999; Kouznetsova et al., 2002). These studies determined the necessity of β arrestin-2 for GPCR desensitization using dominant negative arrestin constructs or direct protein over-expression. If β arrestin binding is necessary for receptor desensitization, this work indirectly suggests that β arrestin-2 is recruited to CB₁ receptors on the cell surface following agonist activation. To test this hypothesis, we evaluated β arrestin-2 translocation in HEK293 cells transiently expressing RFP-tagged β arrestin-2 and wild-type CB₁ receptors by confocal imaging (Fig. 5A). In the absence of receptor stimulation, β arrestin-2-RFP is uniformly expressed within the cytosol (Fig. 5Ac-d). Agonist activation of wild-type CB₁ receptors with 100 nM CP 55,940 causes a rapid redistribution of β arrestin-2-RFP (Fig. 5Ag). Clusters of this protein are found at the surface in close proximity to CB₁ receptors following CP 55,940 treatment (highlighted by arrows) (Fig. 5Ag). Quantitative analysis of β arrestin-2-RFP recruitment to wild-type CB₁ receptors shows a 2-fold increase in the membrane/cytosol fluorescence ratio within 2 min (0.8 ± 0.05 at time 0 versus 1.6 ± 0.1 at 2 min) indicative of an increase in membrane associated β arrestin-2 (Fig. 5B). Interestingly, the membrane/cytosol ratio remains constant (1.5 ± 0.1 at 1 min versus 1.5 ± 0.2 at 15 min)

indicating a sustained association of β arrestin-2-RFP with the plasma membrane and perhaps with activated CB₁ receptors.

Since residues 426 and 430 are putative GRK phosphorylation sites and are likely to contribute to a β arrestin-2 binding site, we examined whether β arrestin-2-RFP recruitment was reduced when these sites were mutated (S426A/S430A receptor). Surprisingly, β arrestin-2-RFP was still rapidly recruited to the surface in cells expressing S426A/S430A receptors stimulated with 100 nM CP 55,940. (Fig. 5Ah). The extent (1.6 ± 0.1 at 2 min for wild-type versus 1.4 ± 0.1 at 2 min) of β arrestin-2-RFP translocation in S426A/S430A expressing cells was not significantly different than that seen in wild-type cells (Fig. 5B). Furthermore, the time course of β arrestin-2-RFP translocation in S426A/S430A expressing cells is similar to that of the wild-type receptor. These data suggest that the receptor domain responsible for CB₁ receptor desensitization is not required for β arrestin-2 recruitment.

4. Discussion

The principal finding of this study is that CB₁ receptor desensitization dictates the kinetics of agonist-mediated ERK1/2 phosphorylation (activity). Using HEK293 cells stably expressing modified CB₁ receptors, we have shown that mutation of serines 426 and 430 of CB₁ to alanines dramatically prolongs the duration of ERK1/2 phosphorylation during protracted receptor stimulation (Fig. 1A). Importantly, the sustained phosphorylation of ERK1/2 (at time points ≤ 35 min) is a direct result of continued agonist activation of the S426A/S430A CB₁ receptor and is not a consequence of a receptor-independent phenomenon (Fig. 1B, C). The results of the present study, coupled with findings from our earlier work (Jin et al., 1999), suggest that the time course of both ERK1/2 de-phosphorylation and GIRK channel inactivation are dynamically regulated at the receptor level by a distinct domain of the CB₁ receptor – a domain that mediates G-protein uncoupling, likely by receptor phosphorylation.

In contrast to the desensitization results, mutation of serines 426 and 430 to alanine had no effect on the rate of rapid CB₁ receptor internalization (< 15 min), but did significantly attenuate the extent of internalization following prolonged agonist treatment (Fig. 2A). These results contrast observations made previously using this same mutant receptor in AtT20 cells (Jin et al., 1999). In this study, the S426A/S430A mutation did not appear to affect CB₁ receptor internalization after a 30 min application of WIN 55,212-2. The apparent discrepancy between the current and the previous findings may be due to the increased sensitivity of the quantitative approach for detecting receptor loss from the surface in the current study or may be due to a cell-type specific difference in trafficking. Whatever the cause, our present results suggest that the desensitization domain of CB₁ may have a role in the late phase of receptor endocytosis.

Our results strongly support the notion that internalization of the CB₁ receptor following agonist stimulation is not required for ERK1/2 activation nor does it dictate the time course of ERK1/2 inactivation. Pharmacological inhibition of agonist-induced CB₁ receptor internalization was without effect on the kinetics of ERK1/2 phosphorylation (Fig. 2C). Consistent with these findings, we have found that a non-internalizing, modified D164N CB₁ receptor is still capable of activating ERK1/2 with a similar temporal pattern to that of the wild-type receptor (Fig 3B). These latter data are in agreement with the initial observation in AtT20 cells that found robust D164N CB₁ receptor mediated ERK1/2 activation at a single time point (Roche et al., 1999). Taken together, these data indicate that CB₁ receptor internalization is not required for the activation of ERK1/2. An early study found that β_2 -adrenergic receptor-mediated activation of ERK1/2 was blocked by the over-expression of β arrestin-2 and dynamin dominant negative constructs in HEK293 cells (Daaka et al., 1998). This observation has caused some to conclude that receptor internalization is a necessary step in ERK1/2 activation. However, subsequent studies have suggested that agonist-induced

internalization of GPCR's, including the β_2 -adrenergic receptor, is not required for MAP kinase activation (Huang et al., 2004; Jimenez-Sainz et al., 2003; Kramer and Simon, 2000; Li et al., 1999). These results, coupled with our present findings, suggest that internalization is not a general requirement for ERK1/2 activation. Furthermore, at least one study (Huang et al., 2004) suggests that the requirement for receptor internalization in ERK1/2 activation may also be cell-type dependent, adding an additional layer of complexity.

CB₁ receptor-mediated activation of ERK1/2 did not require active G protein $\alpha_{i/o}$ or α_s subunits. Specifically, overnight treatment with PTX and/or CTX did not block agonist-induced ERK1/2 phosphorylation by either wild-type or S426A/S430A CB₁ receptors (Fig. 4). This is surprising considering that the majority of CB₁ mediated signal transduction pathways involve G_{ai/o} or G_{as} G-proteins. Possible interpretations of these results are the involvement of G_z or another G protein alpha subunit, or the involvement of a G protein-independent (potentially β arrestin-2 mediated) signaling pathway. This possibility will need to be addressed by additional experiments beyond the scope of the present study.

GPCRs are specifically phosphorylated by GRKs following agonist activation. The phosphorylated receptor then interacts with β arrestin. The physical association of β arrestin with the activated receptor has been shown to uncouple many GPCR's from their cognate G-proteins, inducing homologous desensitization (Gainetdinov et al., 2004). In *Xenopus* oocytes and cultured hippocampal neurons, homologous CB₁ receptor desensitization is mediated in part by β arrestin-2 (Jin et al., 1999; Kouznetsova et al., 2002). In these studies β arrestin over-expression or dominant negative β arrestin-2 mutants were used to reveal the necessity of this protein for the desensitization of CB₁ receptor-mediated activation of GIRK channels or inhibition of voltage-gated Ca²⁺ channels. While these results suggest a role for β arrestin-2 in receptor desensitization, β arrestin-2 recruitment to activated CB₁ receptors has not been reported. Figure 5 shows that wild-type CB₁ receptors rapidly recruit (within 2 min) β arrestin-2 to the plasma membrane during agonist treatment, where it co-localizes with CB₁. Since CB₁ serines 426 and 430 are likely candidates for mediating β arrestin-2 recruitment, we also examined β arrestin-2 recruitment by the desensitization-deficient CB₁ receptor. Surprisingly, S426A/S430A CB₁ receptors recruited β arrestin-2 to a similar extent and with the same kinetics as the wild-type receptor (Fig. 5B), indicating that the residues involved in desensitization are not required for β arrestin-2 recruitment. One interpretation of these results is that the gross recruitment of β arrestin-2 seen following CB₁ activation reflects the bulk recruitment of β arrestin-2 for its scaffolding functions (Lefkowitz and Shenoy, 2005; Shenoy and Lefkowitz, 2005) and not the much more modest quantities (i.e., equimolar with the CB₁ receptor) likely needed for desensitization.

In summary, our results suggest that the rapid inactivation of ERK1/2 during stimulation of CB₁ is mediated by phosphorylation of serines 426 and 430 and desensitization of the CB₁ receptor. CB₁ receptor internalization does not seem to play a role in ERK1/2 activation (or inactivation) as its blockade did not effect on the kinetics of ERK1/2 phosphorylation. Thus, increases in the ERK1/2 cascade following chronic cannabinoid administration *in vivo* (Derkinderen et al., 2003; Valjent et al., 2001), are likely to be regulated by CB₁ receptor desensitization at the cellular level.

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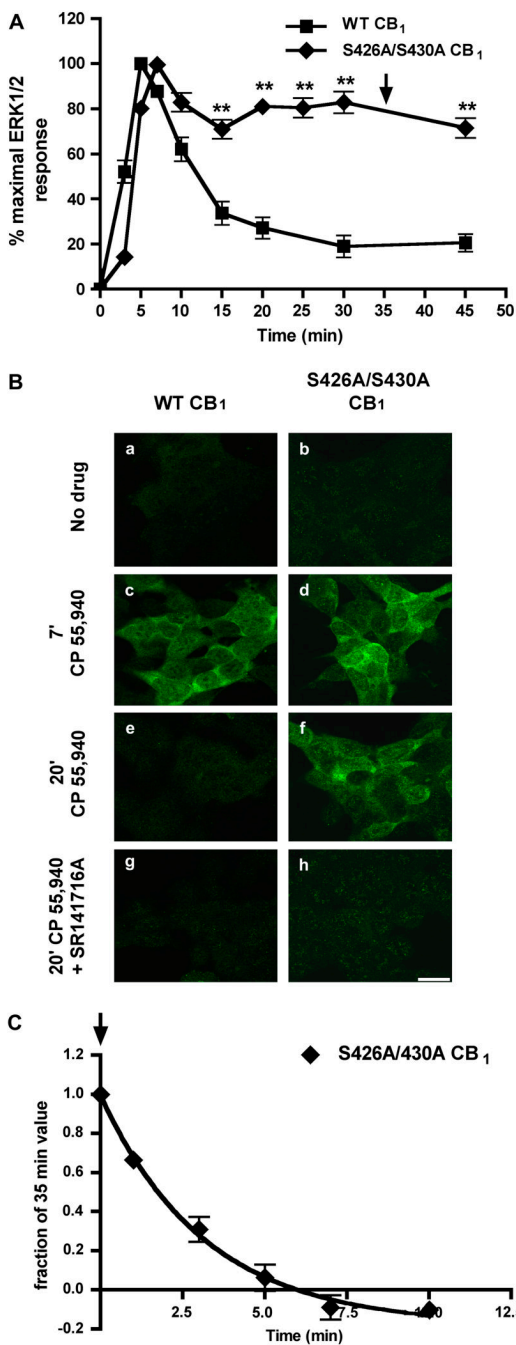


Fig. 1. The sustained ERK1/2 phosphorylation seen with S426A/S430A CB₁ is mediated by continued agonist-induced activation

(A) Quantitative detection of phospho-ERK1/2. HEK293 cells stably expressing either wild-type or S426A/S430A CB₁ receptors were treated with 100 nM CP 55,940 for the indicated times and the time course of ERK1/2 activation (measured by ERK1/2 phosphorylation) was determined (see Methods). The S426A/S430A CB₁ receptor-mediated activation of ERK1/2 is significantly prolonged. Data are mean ± SEM and were collected from 5 or more experiments performed in duplicate. **p<0.01 compared with individual wild-type time points by unpaired t-test. Arrow indicates where SR1 was added in (C). (B) Immunocytochemical detection of phospho-ERK1/2 in HEK293 cells stably expressing either wild-type or S426A/

S430A CB₁. Cells were treated with either vehicle (a–b), 100 nM CP 55,940 alone (c–f), or pretreated with 1 μM SR1 followed by application of 100 nM CP 55,940 in the presence of 1 μM SR1 (g–h). Phospho-ERK1/2 was detected immunocytochemically (see Methods). Both wild type and S426A/S430A CB₁ increased phospho-ERK1/2 immunoreactivity at 7 min. However, with S426A/S430A CB₁ substantial phospho-ERK1/2 immunoreactivity was still evident at 20 min. In contrast, with wild type CB₁ phospho-ERK1/2 immunoreactivity was transient, returning to control levels after 20 min. Pre-treatment with SR1 blocked ERK1/2 activation by both receptors. Scale bar, 20 μm. (C) Prolonged activation of ERK1/2 by the S426A/S430A CB₁ receptor was rapidly ($t_{1/2}$ = 2.97 min) reversed by addition of 1 μM SR1 after 35 min of 10 nM CP 55,940 stimulation (time of SR1 addition indicated by arrow in A and C). Data are mean ± SEM with n = 6 from three experiments performed in duplicate.

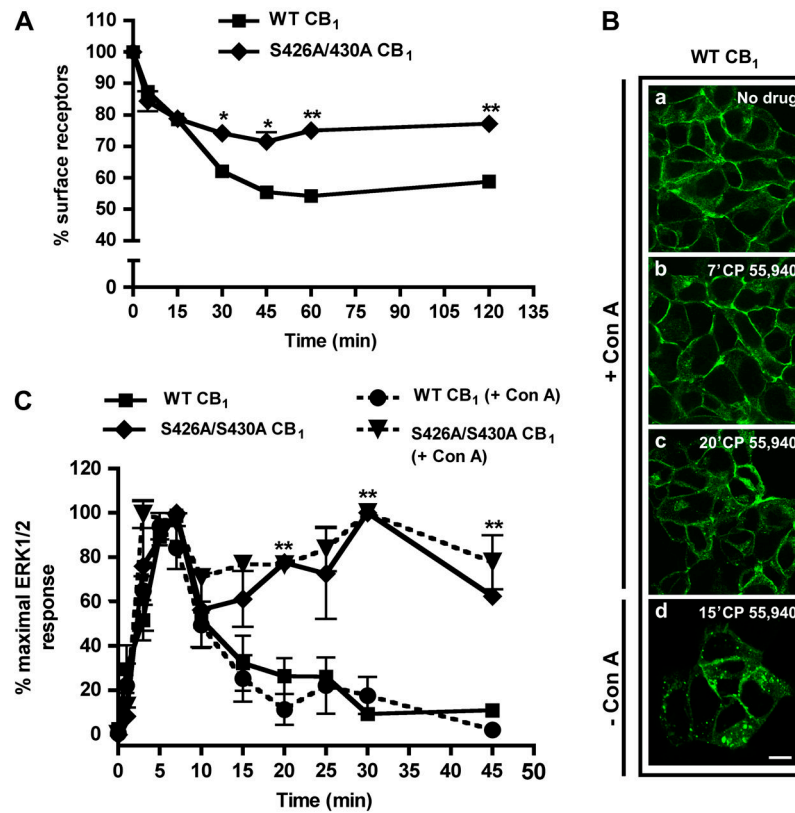


Fig. 2. S426A/S430A CB₁ receptor internalization and pharmacological blockade of CB₁ receptor internalization with Concanavalin A

(A) Agonist-induced internalization of wild-type and S426A/S430A CB₁. Cells were treated with 100 nM CP 55,940 for the indicated times at 37°C. Loss of cell surface receptors was quantified as described in Methods. Rapid (≤ 15 min) receptor endocytosis was unaffected by the S426A/S430A mutation. However, the extent of S426A/S430A CB₁ receptor internalization was significantly attenuated following prolonged drug treatment (≥ 30 min). Data are mean \pm SEM; $n = 15$ – 20 from three to five experiments performed in quadruplicate. ** $p < 0.01$ and * $p < 0.05$ compared with wild-type CB₁ receptor internalization by unpaired t-test. (B) Immunostaining of wild-type CB₁ receptors following Con A treatment. Cells were pretreated with HBS containing 100 μ g/ml Con A (a–c) or HBS alone (d) for 30 min, followed by stimulation with 100 nM CP 55,940 for the indicated times. Con A prevented significant CB₁ receptor internalization at both time points evaluated. In the absence of Con A, a significant fraction of CB₁ receptors were internalized after 15 min (d). Scale bar, 20 μ m. (C) Time course of ERK1/2 activation in the presence or absence of Con A. Cells were pretreated with either HBS alone or 100 μ g/ml Con A for 30 min (+Con A groups). Following pretreatment, the cells were stimulated with 10 nM CP 55,940 alone or in the presence of Con A (+Con A groups). Inhibition of CB₁ receptor internalization by Con A (wild-type or S426A/S430A) does not affect the kinetics of ERK1/2 activation. Data are mean \pm SEM; $n = 5$ – 15 from two to three experiments performed in duplicate. ** $p < 0.01$ compared with wild-type CB₁, no Con A by unpaired t-test.

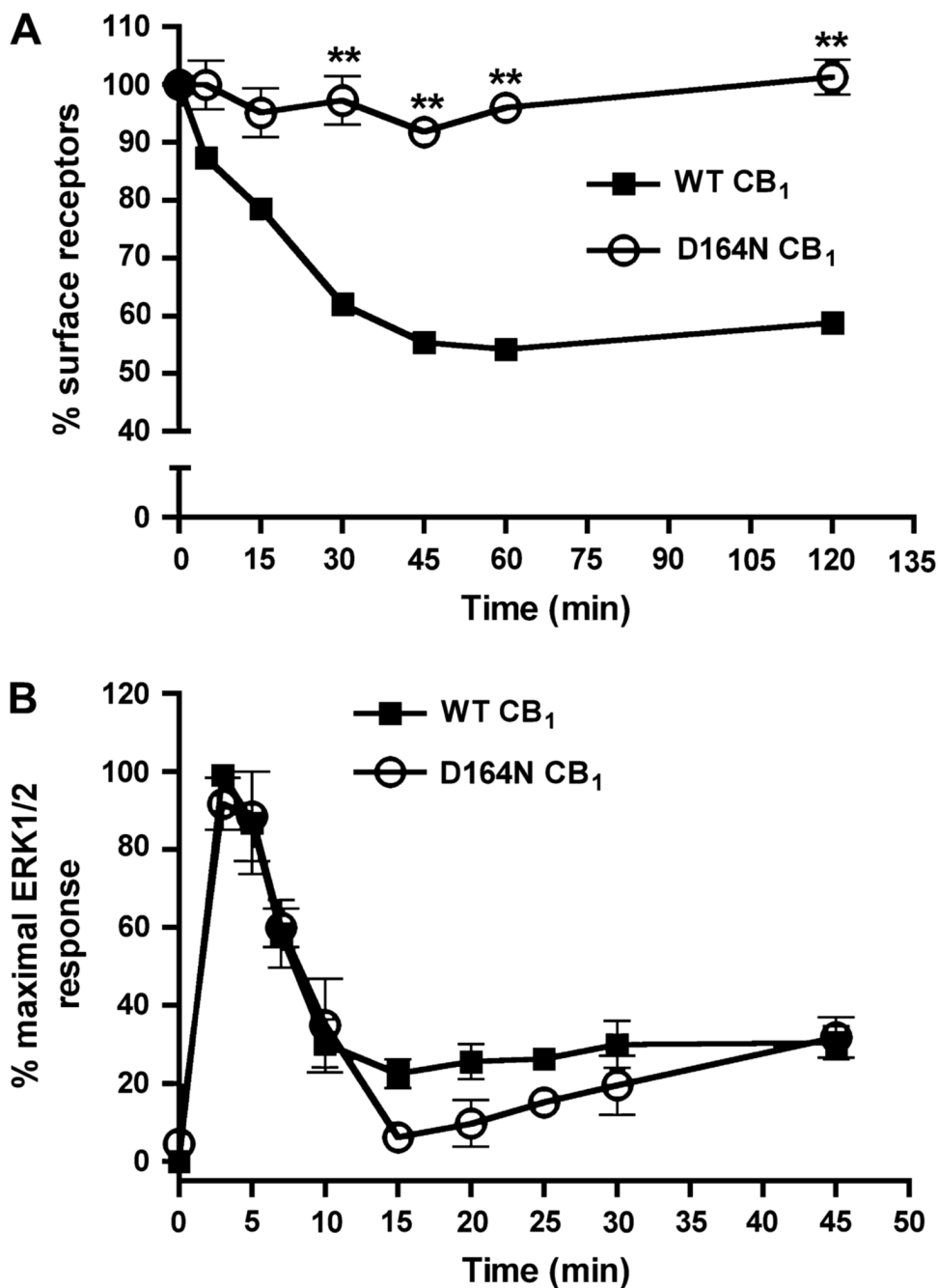


Fig. 3. The kinetics of ERK1/2 activation for an internalization-deficient mutant CB₁ receptor (D164N) is temporally similar to that of the wild-type receptor
 (A) 100 nM CP 55,940-induced internalization of wild-type and D164N CB₁ receptors. Mutation of aspartate 164 to asparagine prevented receptor internalization in HEK293 cells. Data shown are mean \pm SEM; n = 15–20 for three to four experiments. **p < 0.01 compared with the extent of wild-type CB₁ receptor internalization by unpaired t-test. (B) Activation of ERK1/2 in cells expressing wild-type or D164N CB₁ receptors. The time course of D164N CB₁ receptor-mediated activation of ERK1/2 was not significantly different from the wild-type receptor. Data shown are mean \pm SEM; n = 15–20 for three to four experiments.

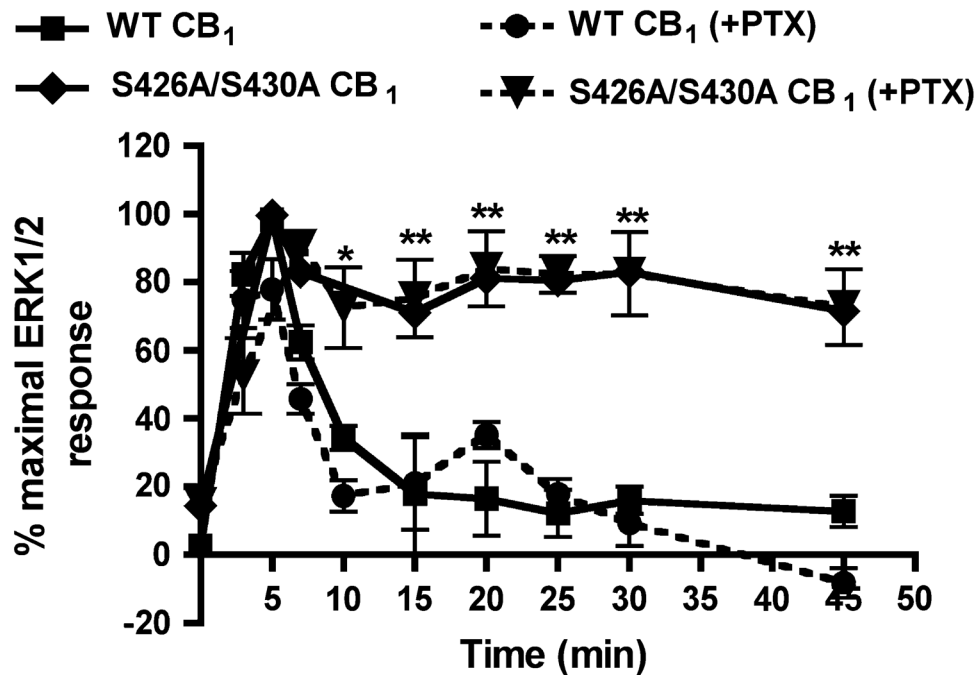


Fig. 4. ERK1/2 activation is pertussis toxin-insensitive

Time course of ERK1/2 activation following pertussis toxin treatment. Cells were treated overnight with 500 ng/ml pertussis toxin (+PTX groups) and stimulated the next day with 100 nM CP 55,940 for the indicated times. Inhibition of $G_{i/o}$ sensitive G-proteins with pertussis toxin did not attenuate agonist-activation of ERK1/2 for wild-type or mutant CB₁ receptors. Data shown are mean \pm SEM; $n = 16-20$ for three to four experiments. ** $p < 0.01$ and * $p < 0.05$ compared with wild-type CB₁ receptor without pertussis toxin treatment by unpaired t-test.

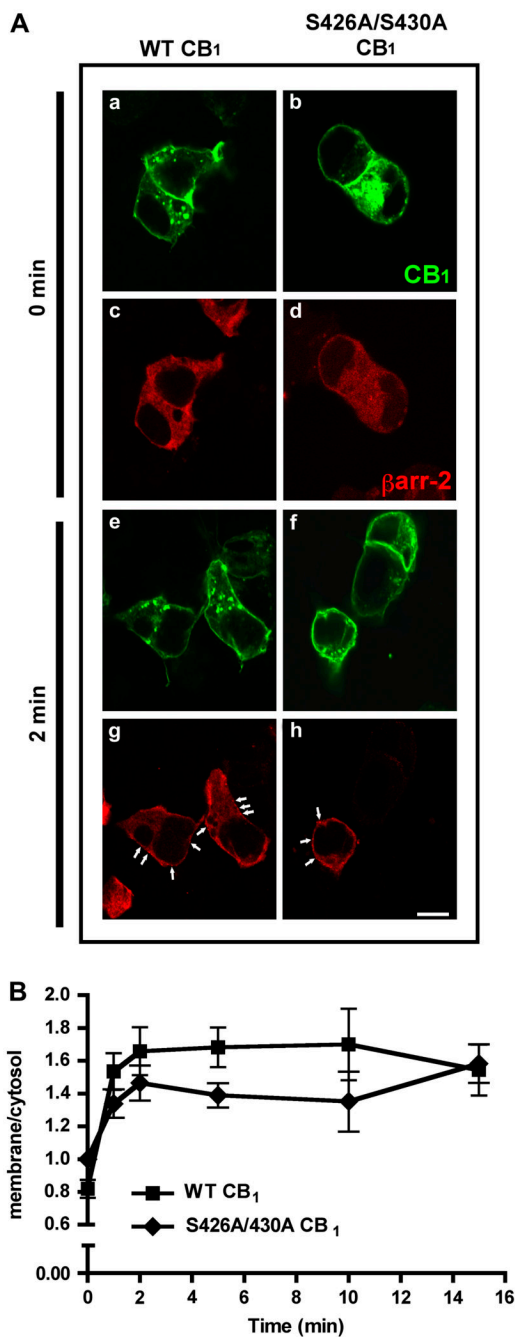


Fig. 5. β arrestin-2-RFP is rapidly recruited following CB₁ receptor activation
 (A) Confocal images of HEK293 cells transiently expressing β arrestin-2-RFP and either wild-type or S426A/S430A CB₁ receptors. CB₁ receptors were detected immunocytochemically, while β arrestin-2-RFP was detected by its red fluorescence. Transiently expressed CB₁ receptors were found both at the surface and within the cytosol (a–b). In unstimulated cells, β arrestin-2-RFP was diffusely present throughout the cytosol (c–d). CB₁ receptor activation with 100 nM CP 55,940 for 2 min caused a dramatic translocation of β arrestin-2-RFP to the surface in cells expressing both wild type and S426A/S430A CB₁ (g–h). Arrows indicate the regions of β arrestin-2-RFP enrichment at the plasma membrane. Scale bar, 5 μ m. (B) Quantitation of β arrestin-2-RFP recruitment. Cells were treated with 100 nM CP 55,940 for

the indicated times. β arrestin2-RFP translocation was quantified as described in Methods and the ratio of membrane averaged intensity values to cytosolic averaged intensity values are plotted. Data shown are mean \pm SEM; n = 5–10 cells per time point (40–80 defined regions total).