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## Signaling Pathways Mediating Beta 3 Adrenergic Receptor-induced Production of Interleukin-6 in Adipocytes

Inna E. Tchivileva<sup>a,1,\*</sup>, Kai Soo Tan<sup>b,1</sup>, Maria Gambarian<sup>d</sup>, Andrea G. Nackley<sup>a</sup>, Alexander V. Medvedev<sup>d</sup>, Sergei Romanov<sup>d</sup>, Patrick M. Flood<sup>b,c</sup>, William Maixner<sup>a</sup>, Sergei S. Makarov<sup>d</sup>, and Luda Diatchenko<sup>a</sup>

<sup>a</sup>The Center for Neurosensory Disorders, School of Dentistry, University of North Carolina, Chapel Hill, NC 27599, USA

<sup>b</sup>The Comprehensive Center for Inflammatory Disorders, School of Dentistry, University of North Carolina, Chapel Hill, NC 27599, USA

<sup>c</sup>Department of Periodontology, School of Dentistry, and Department of Microbiology and Immunology, School of Medicine, University of North Carolina, Chapel Hill, NC 27599, USA

<sup>d</sup>Attagene, Inc., P.O. Box 12054, Research Triangle Park, NC 27709, USA

### Abstract

The  $\beta_3$ -adrenergic receptor ( $\beta_3$ AR) is an essential regulator of metabolic and endocrine functions. A major cellular and clinically significant consequence of  $\beta_3$ AR activation is the substantial elevation in interleukin-6 (IL-6) levels. Although the  $\beta_3$ AR-dependent regulation of IL-6 expression is well established, the cellular pathways underlying this regulation have not been characterized. Using a novel method of homogenous reporters, we assessed the pattern of activation of 43 transcription factors in response to the specific  $\beta_3$ AR agonist CL316243 in adipocytes, cells that exhibit the highest expression of  $\beta_3$ ARs. We observed a unique and robust activation of the CRE-response element, suggesting that IL-6 transcription is regulated *via* the  $G_s$ -protein/cAMP/protein kinase A (PKA) but not nuclear factor kappa B (NF- $\kappa$ B) pathway. However, pretreatment of adipocytes with pharmacologic inhibitors of PKA pathway failed to block  $\beta_3$ AR-mediated IL-6 up-regulation. Additionally, stimulation of adipocytes with the exchange protein directly activated by cAMP (Epac) agonist did not induce IL-6 expression. Instead, the  $\beta_3$ AR-mediated transcription of IL-6 required activation of both the p38 and PKC pathways. Western blot analysis further showed that transcription factors CREB and ATF-2 but not ATF-1 were activated in a p38- and PKC-dependent manner. Collectively, our results suggest that while stimulation of the  $\beta_3$ AR leads to a specific activation of CRE-dependent transcription, there are several independent cellular pathways that converge at the level of CRE-response element activation, and in the case of IL-6 this activation is mediated by p38 and PKC but not PKA pathways.

### Keywords

Cytokine; p38; PKC; obesity; inflammation

\*Corresponding author. tchivilei@dentistry.unc.edu, Tel.: +1-919-966-2953, Fax: +1-919- 966-5339.

<sup>1</sup>Both authors contributed equally in this work

## 1. Introduction

Interleukin-6 (IL-6) is a pleiotropic cytokine that modulates immune response, inflammation, nervous, hematopoietic, and endocrine systems (Kamimura et al., 2003; Kishimoto, 2005). Although IL-6 has a widespread tissue distribution, previous reports have indicated that up to 30% of circulating IL-6 is derived from adipose tissue (Fried et al., 1998; Mohamed-Ali et al., 1997). Consistent with these reports, plasma IL-6 levels are markedly elevated in obese subjects (Vgontzas et al., 1997). Obesity is closely associated with chronic low-grade inflammation characterized by abnormal production of cytokines and acute phase proteins (Hotamisligil, 2006; Wellen and Hotamisligil, 2005). The inflammatory response that emerges in the presence of obesity seems to be triggered by and predominantly reside in adipose tissue. Additionally, the production of IL-6 is well documented in different adipose tissues and adipocyte models (Burysek and Houstek, 1997; Fain et al., 2004; Hoch et al., 2008; Mohamed-Ali et al., 2001; Path et al., 2001; Skurk et al., 2007; Vicennati et al., 2002).

Both white (WAT) and brown (BAT) adipose tissues are innervated by sympathetic nervous system (SNS) (Fliers et al., 2003; Slavin and Ballard, 1978; Wirsén and Hamberger, 1967). Although sympathetic innervation is less abundant in WAT than in BAT, norepinephrine turnover in WAT can be significant in the immediate vicinity of nerve terminals, particularly in response to stress (Bamshad et al., 1998; Collins et al., 2004). In rodents SNS activation during stress has been associated with elevated plasma IL-6 levels (Takaki et al., 1994; Zhou et al., 1993). Similarly, in humans a positive correlation has been observed between exercise-induced peak plasma epinephrine or norepinephrine and IL-6 levels (Papanicolaou et al., 1996). Epinephrine and norepinephrine regulate IL-6 release from adipocytes predominantly *via* activation of  $\beta$ -adrenergic receptors ( $\beta$ ARs). *In vitro*, administration of norepinephrine, the general  $\beta$ AR agonist isoproterenol, or the selective  $\beta_3$ AR agonist CGP-12117 stimulated IL-6 expression in murine brown adipocytes (Burysek and Houstek, 1997), and isoproterenol elevated IL-6 production in human breast adipocytes (Path et al., 2001). *In vivo*, administration of  $\beta_2$  or  $\beta_3$  agonists in mice or infusion of isoproterenol in human volunteers resulted in heightened plasma IL-6 concentrations (Mohamed-Ali et al., 2001). Although  $\beta_3$ AR-dependent regulation of IL-6 expression is well established, the intracellular signaling pathways underlying this regulation have not been characterized.

Activation of the  $\beta_3$ AR by epinephrine, norepinephrine, or specific agonists typically results in the  $G_s$ -dependent activation of adenylate cyclase, increases in intracellular cAMP, and stimulation of protein kinase A (PKA) (Guan et al., 1995; Lindquist et al., 2000; Soeder et al., 1999). PKA, in turn, regulates expression of various genes *via* phosphorylation of the transcription factor cAMP-responsive element binding protein (CREB) which binds to cAMP-responsive element (CRE) sites in the promoter region of cAMP-responsive genes (Rockman et al., 2002). Recently, cAMP has been shown to activate not only PKA but also a class of cyclic nucleotide-gated (CNG) cation channels and a small family of guanine nucleotide exchange factors (GEFs) known as exchange proteins directly activated by cAMP (Epacs) (de Rooij et al., 1998; Kawasaki et al., 1998b).

New layers of complexity have been added to the field of  $\beta_3$ AR signaling with the discovery that  $\beta_3$ ARs couple to  $G_i$  as well as  $G_s$ . In adipocytes, stimulation of the  $\beta_3$ AR activates the extracellular signal-regulated kinases 1 and 2 (ERK1/2) *via* the  $G_i$ -dependent pathway (Cao et al., 2000; Gerhardt et al., 1999; Soeder et al., 1999). However, discrepant reports from other groups suggest that  $\beta_3$ AR-dependent ERK1/2 activation is mediated *via* the  $G_s$ /PKA pathway (Lindquist et al., 2000; Mizuno et al., 1999). In addition to ERK1/2, activation of  $\beta_3$ ARs in adipocytes has been shown to stimulate another mitogen-activated protein kinase (MAPK) p38 through the classical  $G_s$ - and PKA-dependent pathway (Cao et al., 2001; Moule and Denton, 1998), although an obligatory role of PKA in p38 phosphorylation was not confirmed in another

work (Mizuno et al., 2002). Finally, activation of  $\beta_3$ ARs leads to stimulation of one more major family of signaling enzymes- protein kinases C (PKCs). It has been demonstrated that  $\beta_3$ AR agonists increase glucose uptake in brown adipocytes stimulating conventional and novel PKCs (Chernogubova et al., 2004). Thus,  $\beta_3$ ARs exhibit a dynamic capacity to stimulate divergent signaling pathways.

To elucidate the signaling pathways controlling IL-6 production in white adipocytes, we employed a novel method of homogenous reporters (Romanov et al., 2008) and assessed the activation pattern of 43 transcription factors in response to the  $\beta_3$ AR-specific agonist CL316243. We observed a unique and robust activation of the CRE-response element, but not NF- $\kappa$ B which is a pivotal regulator of pro-inflammatory cytokine expression (Baldwin, 1996). CRE activation suggested regulation of IL-6 transcription *via*  $G_s$ /cAMP/PKA activity. However, subsequent experiments demonstrated that IL-6 expression is not mediated through PKA or NF- $\kappa$ B pathways, but instead requires activation of p38- and PKC-dependent signaling mechanisms.

## 2. Materials and Methods

### 2.1. Cell culture

The C3H10T1/2, 3T3-L1, and HEK 293 cells were obtained from American Tissue Culture Collection Center (Rockville, MD). Cells were grown in DMEM (Sigma, St Louis, MO) supplemented with 10% heat-inactivated FBS (Sigma), 2mM L-glutamine (Gibco, Carlsbad, CA), and 1x penicillin/streptomycin (Gibco) under a humidified atmosphere with 5% CO<sub>2</sub> at 37°C. 3T3-L1 fibroblast cells were treated with 0.5 mM IBMX (Sigma), 1  $\mu$ M dexamethasone (Sigma), and 10  $\mu$ g/ml insulin (Sigma) to initiate adipogenesis as described previously (Mizuno et al., 1999). C3H10T1/2 adipogenesis was induced by incubating cells in growth media containing 1  $\mu$ M dexamethasone, 0.5 mM IBMX, 1  $\mu$ M rosiglitazone, and 10  $\mu$ g/ml insulin for 2 days, after which cells were allowed to differentiate in growth media with two more boosts of 1  $\mu$ M rosiglitazone and 10  $\mu$ g/ml insulin. After more than 90% of cells became differentiated, the media containing inducers of differentiation was replaced with growth medium without inducers, and cells were maintained for two more days in culture. Then adipocytes were treated with various reagents.

### 2.2. Factorial homogenous reporter system

The mouse pluripotent fibroblasts C3H10T1/2 were plated at subconfluent density in 6-well plates and then transfected the next day with the Factorial reporter library as described previously (Romanov et al., 2008). The C3H10T1/2 cells were selected because we found that they provide a unique fibroblast cell system that will continue to differentiate into adipocytes after transfection, a critical step required for the Factorial application. After appropriate adipocyte differentiation, the cells were stimulated with various reagents, total cellular RNA was extracted and processed according to the Factorial detection protocol and quantified as described previously (Romanov et al., 2008).

### 2.3. Drugs and treatment

CL316243, ICI 118,551, Betaxolol, SR59230A, RO31-8220, PTX, and 8CPT-2'-O-Me-cAMP were obtained from Tocris Cookson, Ellisville, MI. NBD peptide, MG-132, U0126, and SB203580 were purchased from Biomol, Plymouth Meeting, PA. CTX was bought from Calbiochem, La Jolla, CA. An aliquot of each drug solution was added to the medium, and the final concentration of the vehicle in the medium was adjusted to 0.1% (v/v). The control medium contained the same amount of the vehicle. For inhibitor experiments, cells were pretreated with the indicated drug for 1 h prior. Dose-response curves were established for the agonist and antagonist. Doses within one log unit of the ED<sub>50</sub> or ID<sub>50</sub> were used for all *in*

*vitro* experiments. The effectiveness of the chosen inhibitor doses was confirmed in our previous report (Tan et al., 2007).

#### 2.4. Detection of IL-6 by real-time PCR and ELISA

Total RNA was isolated using the RNeasy Mini Kit (Qiagen, Valencia, CA). Prior to reverse transcription, RNA was treated with 2 units of TURBO DNase I (Ambion, Austin, TX) at 37°C for 30 min. Reverse transcription was performed using Superscript III (Carlsbad, CA, Invitrogen) according to the manufacturer's instructions. IL-6 transcripts were quantified by real-time PCR using TaqMan Gene Expression Assays (Applied Biosystems, Foster City, CA) in an ABI PRISM 7000 Sequence Detection System. The expression of cytokine mRNAs was normalized to the relative abundance of glyceraldehyde-3-phosphate dehydrogenase (GAPDH). Fold induction was calculated using the  $2^{-\Delta\Delta Ct}$  method (Livak and Schmittgen, 2001). Experiments were performed in quadruplicate. IL-6 protein was determined in the culture supernatant of adipocytes using ELISA kits (R&D Systems, Minneapolis, MN). ELISA was performed according to the manufacturer's instructions. Experiments were performed in triplicate.

#### 2.5. Plasmids and transfections

Reporter vectors containing specific *cis*-acting DNA sequence fused to secreted alkaline phosphatase (SEAP), pCRE-SEAP, pAP-1-SEAP, pC/EBP $\beta$ -SEAP, pNF- $\kappa$ B-SEAP and pETS-SEAP, were purchased from Clontech, Inc. (Palo Alto, CA). Mouse  $\beta_3$ -AR clone (GB#BC132000) was purchased from I.M.A.G.E. Consortium Collection (Open Biosystems, Huntsville, AL) and subcloned into pCMV-SPORT6 expression vector (Invitrogen, Carlsbad, CA). The 3<sup>rd</sup> exon of the clone (nucleotides 1239 to 1327) has been replaced with the PCR amplified cDNA fragment corresponding to 3<sup>rd</sup> exon of the major  $\beta_3$ -AR isoform NM\_013462 (nucleotides 1975 to 2042). The functional activity of the clone has been confirmed in transfection experiments. HEK 293 cells were transfected with Effectene transfection reagent (Qiagen) according to the manufacturer's instructions. For normalization of transfection efficiency, cells were co-transfected with a  $\beta$ -galactosidase plasmid (Clontech). Cell stimulation experiments were performed 24 h after transfection. Experiments were performed in triplicate.

#### 2.6. SEAP activity assay

The culture medium was changed to remove the accumulated SEAP prior to treatment of cells with CL316243. The cells were incubated with CL316243 for 6, 12 or 24 h. Culture medium was collected and subjected to SEAP activity assays. The SEAP activity in the culture medium was measured with a Phospha-Light assay kit (Tropix, Foster City, CA), according to the instructions of the manufacturer. Experiments were performed in triplicate.

#### 2.7. Western blot analyses

Cells were lysed using cell lysis buffer containing phosphatase inhibitors (Cell Signaling Technologies, Danvers, MA). The amount of protein in the cell lysates were quantified using the BCA Protein Assay Reagent (Pierce, Rockford, IL). Proteins were resolved on 12% sodium dodecyl sulfate (SDS) NuPAGE Novex Gels (Invitrogen, Carlsbad, CA). Following gel electrophoresis, proteins were transferred to nitrocellulose membranes (Amersham, Piscataway, NJ). Membranes were blocked with 5% non-fat milk at room temperature for 1 h and then probed with the specific antibodies at 4°C overnight. Protein expression was determined with the specific primary antibodies according to the manufacturer's instructions. Antibodies to p-p38, pPKCs, pATF-2 and pCREB were obtained from Cell Signaling Technologies, Danvers, MA, and antibodies to p-PKC $\epsilon$  were received from Santa Cruz Biotechnology, Santa Cruz, CA. The pCREB antibody also detects the pATF-1 at a different size.  $\beta$ -actin was used as the

loading control. Antibodies were detected with corresponding horseradish peroxidase-linked secondary antibodies. Blots were developed using Enhanced Chemiluminescent (ECL) reagent (Perkin Elmer, Waltham, MA) and signals captured using the ImageQuant ECL system (Amersham). Experiments were performed in triplicate.

## 2.8. Statistical analysis

Statistical significance was determined by performing an analysis of variance (ANOVA) followed by Bonferroni test. Data are expressed as mean  $\pm$  SEM. Differences were considered significant if  $P < 0.05$ .

## 3. Results

### 3.1. $\beta_3$ AR activation leads to IL-6 increase in white adipocytes

We first sought to confirm that  $\beta_3$ AR stimulation can modulate IL-6 mRNA expression in the 3T3-L1 adipocyte cell model (Green and Kehinde, 1976). 3T3-L1 adipocytes were treated with CL316243, a specific  $\beta_3$  agonist (Bloom et al., 1992), and real-time quantitative PCR was used to quantify the level of mRNA transcripts present. Treatment of 3T3-L1 cells with CL316243 (1  $\mu$ M) resulted in about 30-fold increase in IL-6 transcripts, when compared to untreated cells (Fig. 1a). In line with previously reported correlation between up-regulation of IL-6 mRNA and protein secretion (Burysek and Houstek, 1997; Path et al., 2001), this increase in IL-6 mRNA expression was accompanied by a 3.5- and 4.8-fold increase in IL-6 protein secretion by 24 and 48 h, respectively (Fig. 1b). Similarly, an up-regulation of IL-6 expression was also observed in the murine white adipocytes C3H10T1/2 (results not shown). The CL316243-induced up-regulation of IL-6 transcripts was blocked by the specific  $\beta_3$ AR antagonist SR59230A (1  $\mu$ M), but not by the specific  $\beta_1$ AR antagonist betaxolol (0.5  $\mu$ M) or the specific  $\beta_2$ AR antagonist ICI118,551 (0.5  $\mu$ M) (Fig. 1a). These results indicate that  $\beta_3$ AR stimulation leads to a significant up-regulation of IL-6 expression in white adipocytes.

### 3.2. $\beta_3$ AR-mediated IL-6 expression is PKA- and Epac- independent

To identify the regulatory pathways involved in  $\beta_3$ AR-mediated signaling, we used a recently described homogeneous reporter system termed Factorial (Romanov et al., 2008). The system comprises 43 *cis*-regulatory reporter transcription units enabling simultaneous functional profiling of multiple transcription factors present in the cell. We found that of all the Factorial reporters, the CRE reporter was the only one whose activity was strongly induced (12-fold induction) by CL316243 (1  $\mu$ M) in C3H10T1/2 adipocytes (Fig. 2a). Activation of the CRE reporter in the homogenous reporter experiment together with the presence of a functional CRE element in the promoter region of IL-6 (Ammit et al., 2002; Edwards et al., 2007; Ray et al., 1988) led to the expectation that the cAMP/PKA/CREB cascade underlies  $\beta_3$ AR-mediated IL-6 regulation. Surprisingly, however, stimulation of the CRE reporter in adipocytes by CL316243 was not inhibited by the PKA inhibitor H89 (10  $\mu$ M) (Fig. 2a). Furthermore, IL-6 mRNA expression was not blocked in 3T3-L1 cells pretreated with the PKA inhibitors H89 or KT5720 (Fig. 2b), suggesting that the  $\beta_3$ AR-dependent signaling pathway(s) leading to IL-6 up-regulation are distinct from the cAMP/PKA regulatory pathway.

We further examined the possibility that cAMP could activate IL-6 production through the PKA-independent Epac (guanine nucleotide exchange factors directly activated by cAMP) pathway (de Rooij et al., 1998; Kawasaki et al., 1998b). Epac proteins bind cAMP with high affinity and activate the Ras superfamily small GPTases Rap1 and Rap2. To determine if Epac activation results in IL-6 production, 3T3-L1 cells were treated with the Epac agonist, 8CPT-2'-O-Me-cAMP (CPTOMe, 10  $\mu$ M), and the level of IL-6 mRNA measured by real-time quantitative PCR (Fig. 2c). Treatment of cells with CPTOMe failed to promote increased IL-6 mRNA levels. However, similar to previous reports, treatment of 3T3-L1 cells with cholera

toxin (CTX), an activator of G<sub>s</sub> proteins (Cassel and Selinger, 1977), did lead to a 16-fold increase in IL-6 mRNA levels (Fig. 2c). Together, these data indicate that  $\beta_3$ AR-mediated IL-6 production in white adipocytes depends on CRE activation but not PKA or Epac activation.

### 3.3. $\beta_3$ AR-mediated IL-6 expression is NF- $\kappa$ B-independent

Transcription factor NF- $\kappa$ B is a pivotal regulator of inflammation and has been shown to play a central role in the transcriptional regulation of pro-inflammatory cytokines (Baldwin, 1996). Therefore, we sought to investigate whether NF- $\kappa$ B stimulation is required for  $\beta_3$ AR-mediated IL-6 expression. In a homogenous reporter system, NF- $\kappa$ B reporter activity was not affected by CL316243 (1  $\mu$ M) treatment, though it was strongly induced by TNF $\alpha$  (10 ng/ml) which served as a positive control for NF- $\kappa$ B activation (Fig. 3a).

As the homogenous reporter assay results did not suggest a contribution of NF- $\kappa$ B in CL316243-dependent IL-6 up-regulation, we confirmed this conclusion by independent approaches. NF- $\kappa$ B activity can be inhibited using either NEMO binding peptide (NBD), which binds to and inactivates IKK $\gamma$  thereby blocking the phosphorylation and degradation of I $\kappa$ B (May et al., 2000; Yamaoka et al., 1998), or MG-132, a proteasome inhibitor (Fenteany et al., 1995; Myung et al., 2001). Pre-treatment of 3T3-L1 cells with NBD (100  $\mu$ M) or MG-132 (25 and 50  $\mu$ M) prior to  $\beta_3$ AR stimulation with CL316243 (1  $\mu$ M) did not block IL-6 up-regulation (Fig. 3b) confirming the mechanism is NF- $\kappa$ B-independent.

In agreement with these results and data from homogenous reporter assay, human embryonic kidney cell line 293 (HEK 293) co-transfected with the pNF- $\kappa$ B-SEAP reporter vector and  $\beta_3$ AR expression construct did not show significant changes in SEAP reporter activity following CL316243 (1  $\mu$ M) treatment (Fig. 3c). Thus, all these experiments strongly indicate that  $\beta_3$ AR stimulation does not result in NF- $\kappa$ B activation.

### 3.4. Role of MAPK pathways in $\beta_3$ AR-mediated IL-6 expression

It has been reported that  $\beta_3$ ARs can activate ERK1/2 and p38 MAPK cascades through interchangeable coupling to both G<sub>i</sub> and G<sub>s</sub> (Collins et al., 2004). Although findings from various studies differ,  $\beta_3$ AR coupling with G<sub>i</sub> proteins has been implicated in phosphorylation of ERK1/2 (Cao et al., 2000; Gerhardt et al., 1999; Robidoux et al., 2006; Soeder et al., 1999). We used pertussis toxin (PTX), which ADP-ribosylates and inactivates G<sub>i</sub> proteins, to test the plausible contribution of this G<sub>i</sub>-mediated pathway to  $\beta_3$ AR-mediated IL-6 production. The CL316243-induced IL-6 mRNA up-regulation was not inhibited in cells pretreated with the PTX (Fig. 4a), excluding the role of G<sub>i</sub>-dependent signaling in the regulation of IL-6 expression.

We next examined whether inhibition of the MAPK pathways could block the  $\beta_3$ AR-dependent IL-6 production. Pretreatment of 3T3-L1 cells with the p38 inhibitor SB203580 (5-10  $\mu$ M) blocked the CL316243-induced increase in IL-6 mRNA level (Fig. 4b). In contrast, pretreatment with ERK inhibitor U0126 (10  $\mu$ M) had no effect on CL316243-induced IL-6 expression (Fig. 4b). In the homogenous reporter assay, activation of the CRE reporter by CL316243 was also blocked by pretreatment with SB203580 (10  $\mu$ M; Fig. 4c). Additionally, p38 phosphorylation following CL316243 treatment was observed in cell lysates by Western blot analyses (Fig. 4d). Taken together, these results indicate that while no contribution of G<sub>i</sub>-dependent ERK1/2 stimulation was observed, p38 activation is required for  $\beta_3$ AR-mediated IL-6 production.

### 3.5. Role of the PKC pathway in $\beta_3$ AR-mediated IL-6 expression

Previous work has demonstrated that protein kinases C (PKCs) are essential for  $\beta_3$ AR-mediated glucose uptake regulation (Chernogubova et al., 2004). Thus, we sought to determine the effect

of PKC inhibition on  $\beta_3$ AR-mediated IL-6 production using the specific PKC inhibitor RO31-8220 which blocks conventional ( $\alpha$ ,  $\beta$ ,  $\gamma$ ) and novel ( $\epsilon$  and  $\theta$ ) PKC isoforms (Wilkinson et al., 1993; Yuan et al., 2002). Pretreatment with RO31-8220 (1 and 10  $\mu$ M) completely inhibited CL316243-induced IL-6 mRNA expression in 3T3-L1 cells (Fig. 5a) and significantly inhibited CL316243-mediated activation of the CRE reporter in the homogenous reporter assay (Fig. 5b). Additionally, CL316243-mediated phosphorylation of several PKC subtypes was detected applying an antibody specific for a panel of PKCs ( $\alpha$ ,  $\beta_I$ ,  $\beta_{II}$ ,  $\delta$ ,  $\epsilon$ , and  $\eta$ ) phosphorylated at C-terminal residue homologous to serine 660 of PKC $\beta_{II}$ . In a subsequent attempt to identify individual PKC isoforms activated by CL316243 treatment, phosphorylation of PKC  $\alpha/\beta_{II}$ ,  $\theta$ , and  $\epsilon$  was observed with the use of isoform-specific antibodies (Fig. 5c). These findings indicate that conventional and novel isoforms of PKC play a significant role in  $\beta_3$ AR-mediated IL-6 expression.

### 3.6. Stimulation of $\beta_3$ AR leads to the phosphorylation of transcription factors CREB and ATF-2

To determine transcription factors that mediate  $\beta_3$ AR-dependent IL-6 expression, the transcription factors which bind to the CRE-response element were investigated. First, to confirm the results of the homogeneous reporter assay, a pCRE-SEAP reporter vector with the CRE-response element was co-transfected into HEK293 cells with a  $\beta_3$ AR expression construct (Fig. 6a). Cells were treated with CL316243, and SEAP activity was measured over a 24-hour period. In agreement with the initial screening results, we observed a 5- to 6-fold increase in CRE-dependent SEAP expression (Fig. 6a). To confirm that other transcription factors most commonly associated with activation of PKC or p38 pathways are not activated in response to  $\beta_3$ AR stimulation (Buchwalter et al., 2004; Isakov and Altman, 2002; Trautwein et al., 1993), analogous experiments were conducted with pAP-1-SEAP, pC/EBP $\beta$ -SEAP, and pETS-SEAP reporter vectors. Again, in full agreement with the homogeneous reporter assay, no significant differences in AP-1, C/EBP $\beta$  or ETS activities were found (Fig. 6b-d).

Transcription factors that specifically recognize the CRE-response element include the CREB/ATF family (Hai and Hartman, 2001). We examined the phosphorylation states of CREB, ATF-1, and ATF-2 transcription factors in 3T3-L1 adipocytes. An increased phosphorylation of CREB, ATF-1, and ATF-2 was observed following CL316243 treatment (Fig. 7a). Pretreatment of cells with the p38 inhibitor SB203580 (10  $\mu$ M) remarkably reduced the phosphorylation of both CREB and ATF-2, while administration of the PKC inhibitor RO31-8220 (10  $\mu$ M) decreased reduced phosphorylation of ATF-2 only (Fig. 7b). In contrast, the phosphorylation of ATF-1 was unaffected by either SB203580 or RO31-8220. Thus, our results suggest that the  $\beta_3$ AR-dependent up-regulation of IL-6 transcription in adipocytes is mediated by the transcription factors CREB and ATF-2.

## 4. Discussion

Activation of  $\beta_3$ ARs in adipocytes and adipose tissue results in increased production of the proinflammatory cytokine IL-6 (Burysek and Houstek, 1997; Mohamed-Ali et al., 2001; Path et al., 2001). In the present study, we elucidated the signaling pathways underlying  $\beta_3$ AR-mediated regulation of IL-6 production in white adipocytes. In view of previous reports that circulating IL-6 concentrations are elevated in obesity and that adipose tissue releases up to 30% of total IL-6 in the circulation (Fried et al., 1998; Mohamed-Ali et al., 1997; Vgontzas et al., 1997), the molecular mechanisms contributing to enhanced production of this pro-inflammatory cytokine may play an active role in generating low grade chronic inflammation in obesity.

Here, we demonstrated that  $\beta_3$ AR stimulation in white adipocytes dramatically increases both mRNA and protein levels of IL-6 (30- and 5-fold, respectively). Adrenergic regulation of IL-6

release has been reported in several other cell types, including macrophages (Tan et al., 2007), hepatocytes (Jung et al., 2000), astrocytes (Norris and Benveniste, 1993), cardiac fibroblasts (Yin et al., 2006), and endothelial cells (Gornikiewicz et al., 2000), although the relative contribution of these tissues to circulating IL-6 concentrations is unknown. Adipose tissue is innervated by postganglionic sympathetic nerves releasing norepinephrine, which promote increased IL-6 levels in humans and animal models (Papanicolaou et al., 1996; Takaki et al., 1994; Zhou et al., 1993). Thus, elevated systemic concentrations of IL-6 associated with obesity could be related to higher SNS activity as well as increased adipose tissue mass in obese subjects (Tentolouris et al., 2006).

To examine the transcriptional pathways involved in  $\beta_3$ AR-mediated signaling, we utilized a novel homogeneous reporter system (Romanov et al., 2008) enabling quantitative assessment of multiple transcription factor activities in a eukaryotic cell. The robust activation of a single CRE reporter along with the presence of a functional CRE element in the promoter region of IL-6 (Ammit et al., 2002; Edwards et al., 2007; Ray et al., 1988), suggested that the cAMP/PKA/CREB cascade underlies  $\beta_3$ AR-mediated IL-6 production. Nevertheless, specific PKA inhibitors failed to block either the IL-6 mRNA up-regulation induced by CL316243 in adipocytes or the CRE activation in the homogenous reporter assay. Thus, our findings suggest that the signaling mechanisms required for  $\beta_3$ AR-mediated IL-6 production do not involve the PKA pathway.

PKAs were considered to be the major effectors of cAMP until the recent discovery of Epacs, exchange proteins directly activated by cAMP, which have been shown to activate the small G proteins Rap1 and Rap2 (de Rooij et al., 1998; Kawasaki et al., 1998b). Stimulation of the  $\beta_2$ AR has been reported to induce PKA-independent cell adhesion through Epac and Rap1 (de Rooij et al., 1998). Epac activation produces an inhibitory effect on PKB phosphorylation in rat white adipocytes (Zmuda-Trzebiatowska et al., 2007) while does not affect glucose uptake in brown adipocytes (Chernogubova et al., 2004). Using the Epac-specific agonist CPTOMe, we found no evidence for involvement of an Epac pathway in activation of IL-6 release in our experiments, although CTX treatment significantly elevated IL-6 mRNA expression. These results suggest that  $\beta_3$ AR-induced IL-6 production could be mediated by a novel cAMP pathway through PKA- and Epac-independent mechanism. Indeed, several atypical cAMP binding proteins have been described, including the cAMP receptor which regulates development in *Dictyostelium discoideum* (Klein et al., 1988) and cyclic nucleotide gated channels from olfactory neurons (Goulding et al., 1992). Whether atypical cAMP pathways are involved in IL-6 secretion in adipocytes still needs further investigation. Interestingly,  $\beta_3$ AR-mediated IL-6 production was also independent from another pivotal regulator of cytokine expression, NF- $\kappa$ B (Baldwin, 1996). The PKA- and NF- $\kappa$ B-independent nature of  $\beta_3$ AR-mediated IL-6 expression in adipocytes is in line with similar reports which have been recently published regarding  $\beta_2$ AR-mediated cytokine production. Secretion of IL-6 after  $\beta_2$ AR stimulation is PKA-independent and p38-driven in mouse cardiac fibroblasts (Yin et al., 2006), while  $\beta_2$ AR-mediated release of IL-6 and IL-1 $\beta$  is PKA- and NF- $\kappa$ B-independent in macrophages (Tan et al., 2007).

It has been reported that  $\beta_3$ ARs can interchangeably couple to both  $G_s$  and  $G_i$  and activate ERK1/2 via  $G_i$ -dependent mechanism in adipocytes (Gerhardt et al., 1999; Soeder et al., 1999). Unlike  $\beta_2$ AR, the  $\beta_3$ AR doesn't require receptor phosphorylation and, instead, recruits c-Src kinases for ERK activation in PKA-independent manner (Cao et al., 2000; Robidoux et al., 2006). In contrast, results from other groups argue that  $\beta_3$ AR-mediated phosphorylation of ERK1/2 is  $G_s$ - and PKA-dependent (Lindquist et al., 2000; Mizuno et al., 1999; Mizuno et al., 2000).  $\beta_3$ AR signaling is further complicated by alternative splicing of the gene. Two splice variants,  $\beta_{3a}$  and  $\beta_{3b}$ ARs that differ only in their C-termini (Evans et al., 1999), also display differences in their signaling properties. The  $\beta_{3a}$ AR couples only with  $G_s$ , probably due to



localization in caveolae (Sato et al., 2007), whereas the  $\beta_{3b}$ AR couples with  $G_s$  and  $G_i$  (Hutchinson et al., 2002). Together, these reports indicate that  $\beta_3$ AR may activate different signaling mechanisms in different cell systems (white or brown adipocytes, cell lines or primary cultures). We observed in white adipocyte cell lines that  $\beta_3$ AR-mediated IL-6 expression is  $G_i$ - and ERK1/2-independent, but instead is up-regulated by p38 MAPK and PKC pathways (Fig. 8).

Next, we found that conventional PKCs ( $\alpha/\beta_{II}$ ) and novel PKCs ( $\theta$  and  $\epsilon$ ) are activated by CL316243 in adipocytes. Our results are in line with previous work showing that  $\beta_3$ AR agonists increase glucose uptake in brown adipocytes *via* conventional and novel PKCs (Chernogubova et al., 2004). Furthermore, up-regulation of IL-6 in a PKA-independent but PKC- and p38-dependent manner has been previously reported for various G-protein-coupled receptors (GPCR) in other types of cells, such as P2Y receptors in microglia (Shigemoto-Mogami et al., 2001) and corticotropin-releasing factor receptor 2 $\beta$  in smooth muscle cells (Kageyama and Suda, 2003) as well as under inflammatory conditions in astrocytes (Norris et al., 1994). Further studies are needed to elucidate the exact mechanism of PKC activation by  $\beta_3$ AR and its relation to p38. As PKCs are known to be downstream effectors of phosphatidylinositol 3-kinase (PI3K) (Standaert et al., 1997), it is plausible that PKCs activated by  $\beta_3$ ARs in adipocytes are downstream of PI3K and not directly activated by elevated cAMP. Interestingly, while p38 and PKC pathways appear to be distinct, our findings reveal that inhibition of either one of them can completely abolish IL-6 expression, suggesting an interaction of these two pathways in IL-6 induction, most likely at the level of transcriptional regulation and/or mRNA stability.

The p38 and PKC pathways activate multiple transcription factors including CREB, ATF-1, ATF-2, AP-1, C/EBP $\beta$ , and ETS (Buchwalter et al., 2004; Deak et al., 1998; Isakov and Altman, 2002; Kawasaki et al., 1998a; Li et al., 2006; Lim et al., 2005; Morton et al., 2004; Tan et al., 1996; Togo, 2004; Trautwein et al., 1993). Besides, the promoter region of IL-6 gene contains several functional binding sites such as CRE, AP-1, C/EBP, and NF- $\kappa$ B (Ammit et al., 2002; Edwards et al., 2007; Eickelberg et al., 1999; Zhu et al., 1996). Searching for transcription factors activated by  $\beta_3$ AR agonist CL316243, we found that  $\beta_3$ AR stimulation leads to increased expression of reporter constructs containing CRE, but not AP-1, C/EBP, ETS or NF- $\kappa$ B binding sites, which is in accord with our results from the homogenous reporter assay. Furthermore, significant increases in phosphorylation of CREB, ATF-1, and ATF-2 transcription factors were observed in response to  $\beta_3$ AR-agonist treatment. These transcription factors belong to a large ATF/CREB family of the basic region-leucine zipper (bZip) proteins which are defined by their ability to bind to the consensus CRE site (Hai and Hartman, 2001). We next found that pretreatment with a p38 inhibitor remarkably decreased CREB and ATF-2 phosphorylation, and that pretreatment with a PKC inhibitor diminished ATF-2 activation alone. Overwhelming evidence indicates that ATF/CREB proteins form selective homo- and heterodimers (De Cesare and Sassone-Corsi, 2000). Thus, it is plausible that activation of both CREB and ATF-2 transcription factors is required for IL-6 production, and that these two factors may work synergistically to regulate cytokine expression in adipocytes. The exact mechanism of their interaction merits further research. Nevertheless, our results strongly suggest that the CRE site in the IL-6 promoter is the main target for  $\beta_3$ AR-induced transcription, and that this gene activation is mediated by the actions of CREB and ATF-2 transcription factors.

Our findings have important clinical implications. Converging lines of evidence have revealed that inflammation is a key feature of obesity and type 2 diabetes (Wellen and Hotamisligil, 2005). Although a causal role of IL-6 in diabetes still needs to be clarified, IL-6 levels correlate with the degree of insulin resistance in human subjects (Bastard et al., 2002; Kern et al., 2001) and predict the development of type 2 diabetes (Hu et al., 2004; Thorand et al., 2007). Additionally, circulating levels of IL-6 are elevated in patients with painful inflammatory and

musculoskeletal conditions (Liebregts et al., 2007). Selective  $\beta_3$ AR antagonists have been shown to reduce circulating levels of IL-6 and corresponding pain behavior in an animal pain model (Nackley et al., 2005; Nackley et al., 2007). On the other hand, IL-6 plays protective role improving adenosinergic signaling and increasing neuronal survival under various pathological conditions (Biber et al., 2008). Furthermore, experiments with hepatectomy and hepatic warm ischemia/reperfusion injury demonstrated that IL-6 is a critical component for hepatocyte proliferation and liver regeneration (Camargo et al., 1997; Cressman et al., 1996). In pursuit of novel therapeutic approaches a series of human  $\beta_3$ AR agonists and antagonists has recently been generated (Brockunier et al., 2001; Candelore et al., 1999; Mathvink et al., 2000; Parmee et al., 2000). Thus, this work may promote the development of new therapeutic avenues for the treatment of obesity, inflammatory pain conditions, brain pathology, and liver diseases.

## 5. Conclusions

The present work provides novel insights into signal transduction cascades mediating the  $\beta_3$ AR-dependent release of pro-inflammatory cytokine IL-6 in adipocytes. Here, we report that this IL-6 increase is  $G_i$ -independent and doesn't rely on ERK1/2, PKA, or NF- $\kappa$ B pathways. Instead, it's mediated by p38 and PKC signaling through activation of transcription factors CREB and ATF-2. Elucidation of cellular network regulating IL-6 expression improves our understanding of molecular mechanisms in obesity, inflammatory pain, and other pathological conditions where a significant role for IL-6 is implicated.

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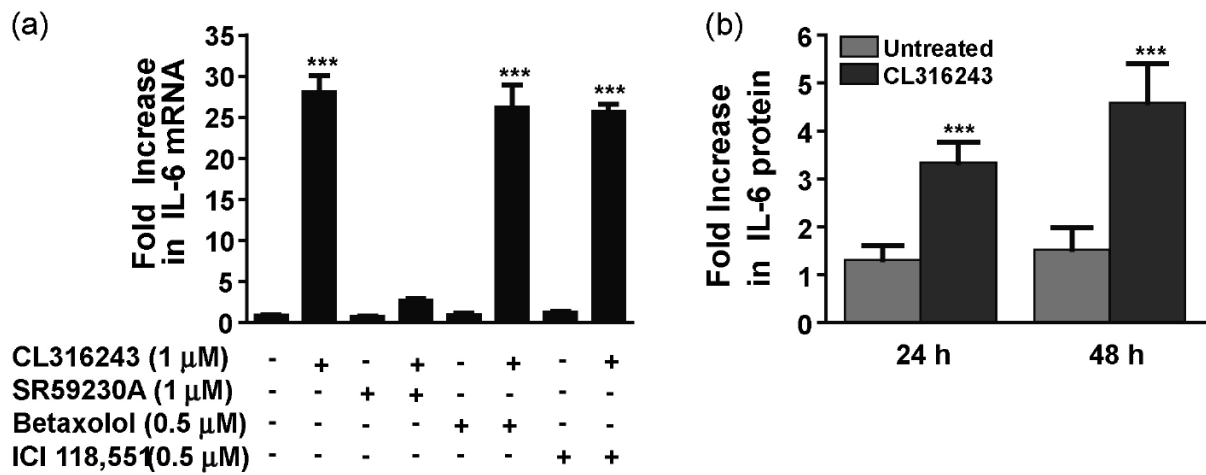
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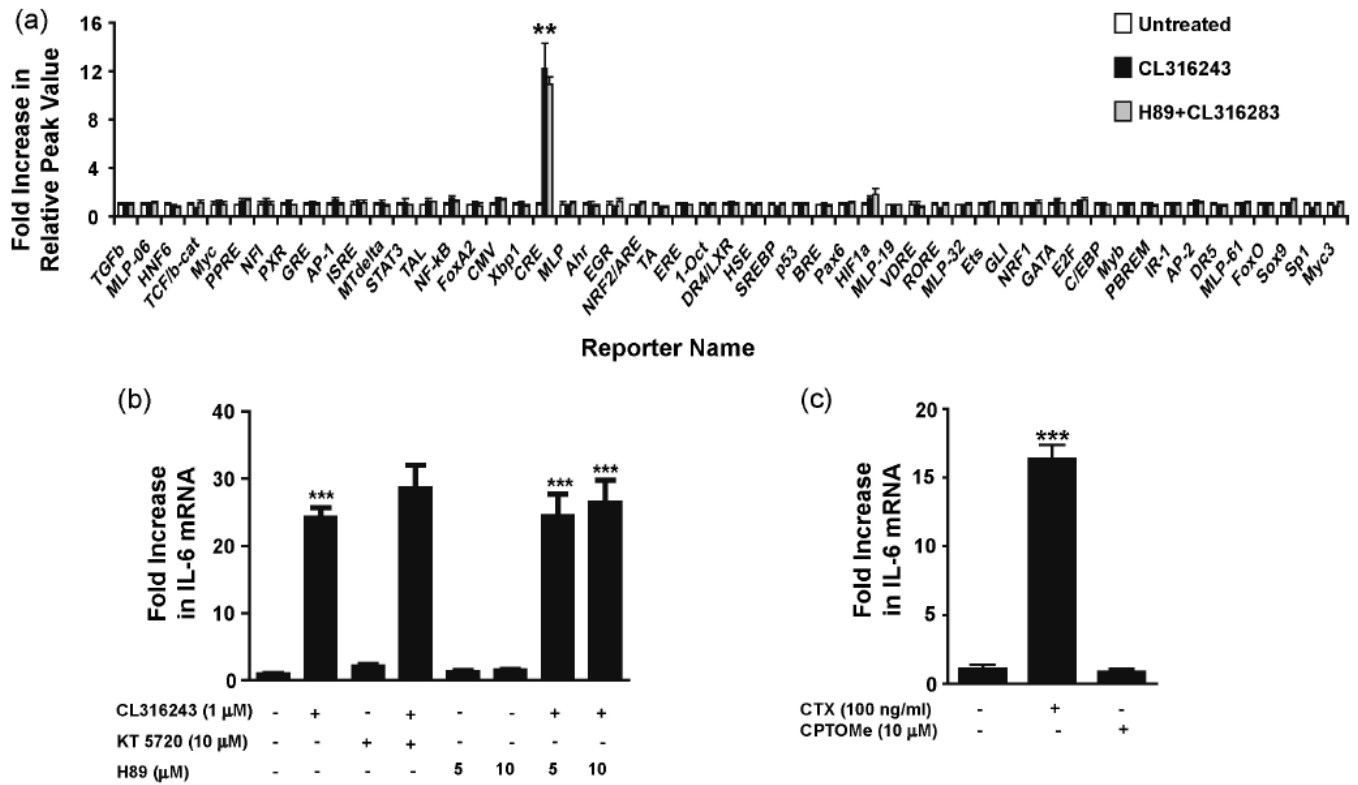
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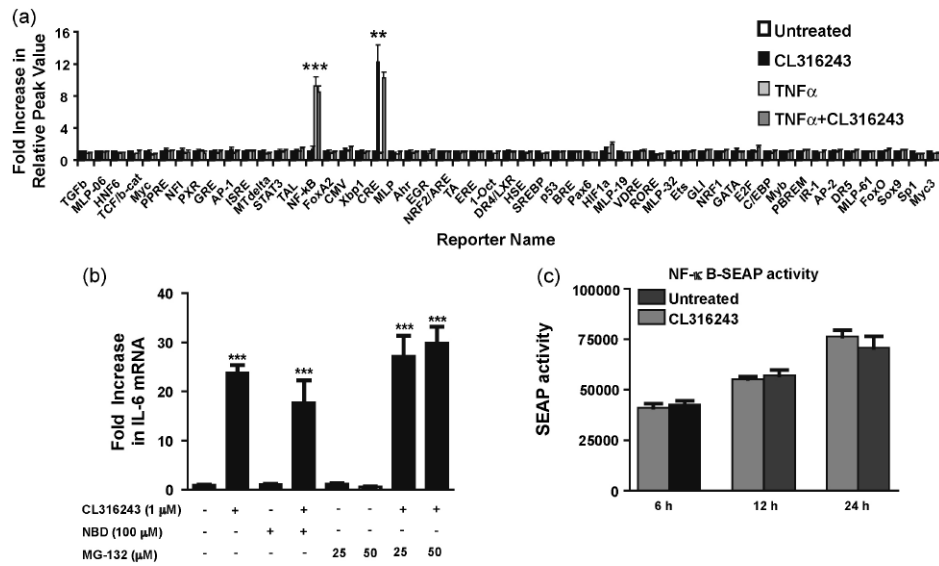
**Fig. 1.**

$\beta_3$ AR stimulation up-regulates IL-6 production in adipocytes. 3T3-L1 cells treated with the  $\beta_3$ AR-agonist CL316243 (1  $\mu$ M for 45 min) showed a significant increase in IL-6 mRNA. The CL316243-induced increase in cytokine transcripts was completely blocked by pre-treatment with SR59230A (1  $\mu$ M for 1 h) but not with betaxolol (0.5  $\mu$ M for 1 h) or ICI118,551 (0.5  $\mu$ M for 1 h) (a). 3T3-L1 cells treated with the  $\beta_3$ AR-agonist CL316243 (1  $\mu$ M for 24 and 48 h) showed a significant increase in IL-6 protein measured in cell supernatant (b). \*\*\* $P$ <0.001 different from untreated.

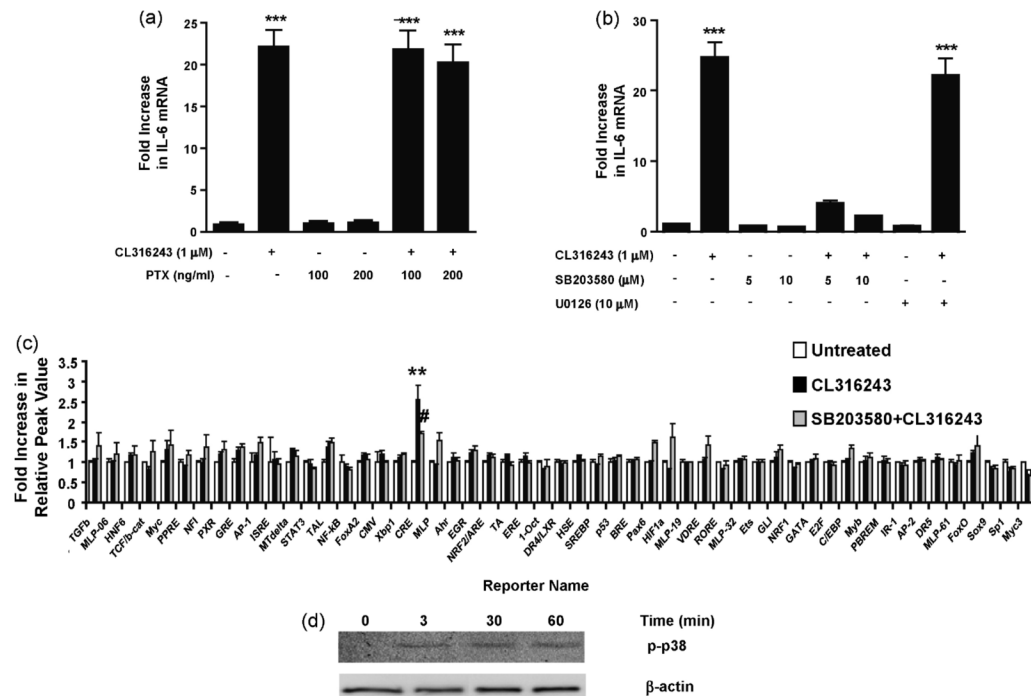




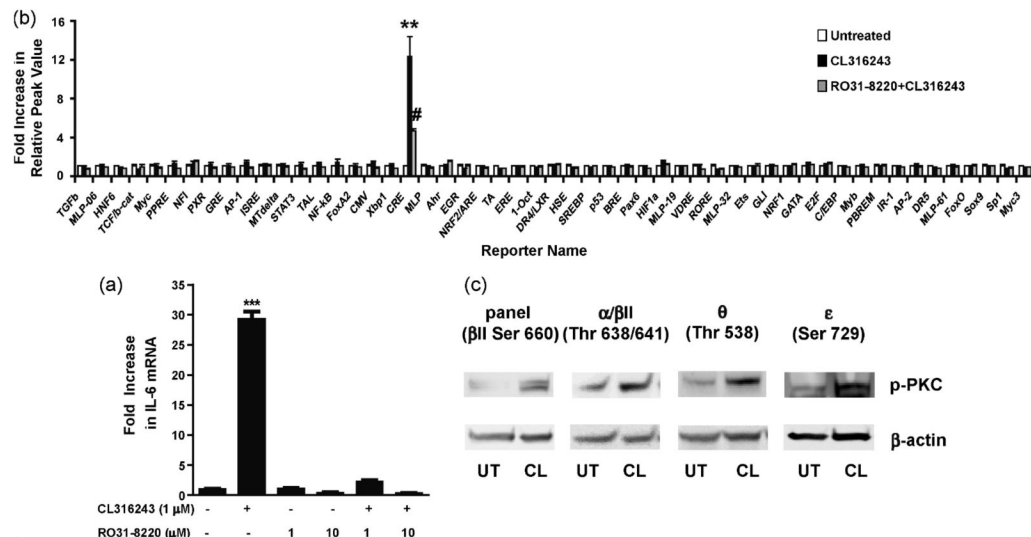
**Fig. 2.**  $\beta_3$ AR-induced IL-6 expression is PKA- and Epac-independent. C3H10T1/2 cells were pretreated with H89 (10  $\mu$ M for 1 h) prior to stimulation with CL316243 (1  $\mu$ M for 2 h). The CL316243-induced activation of the CRE reporter was not inhibited by H89 in the homogenous reporter assay (a). 3T3-L1 cells were treated with KT5720 (10  $\mu$ M) or H89 (5-10  $\mu$ M) for 1 h prior to treatment with CL316243 (1  $\mu$ M for 45 min). The CL316243-induced up-regulation of IL-6 mRNA was not inhibited by KT5720 or H89 (b). Treatment of 3T3-L1 with CTX (100 ng/ml for 4 h) led to increases in IL-6 transcript, while treatment with the Epac agonist 8CPT-2'-O-Me-cAMP (CPTOMe) (10  $\mu$ M for 45 min) did not enhance IL-6 expression (c). \*\*\* $P$ <0.001 and \*\* $P$ <0.01 different from untreated.

**Fig. 3.**

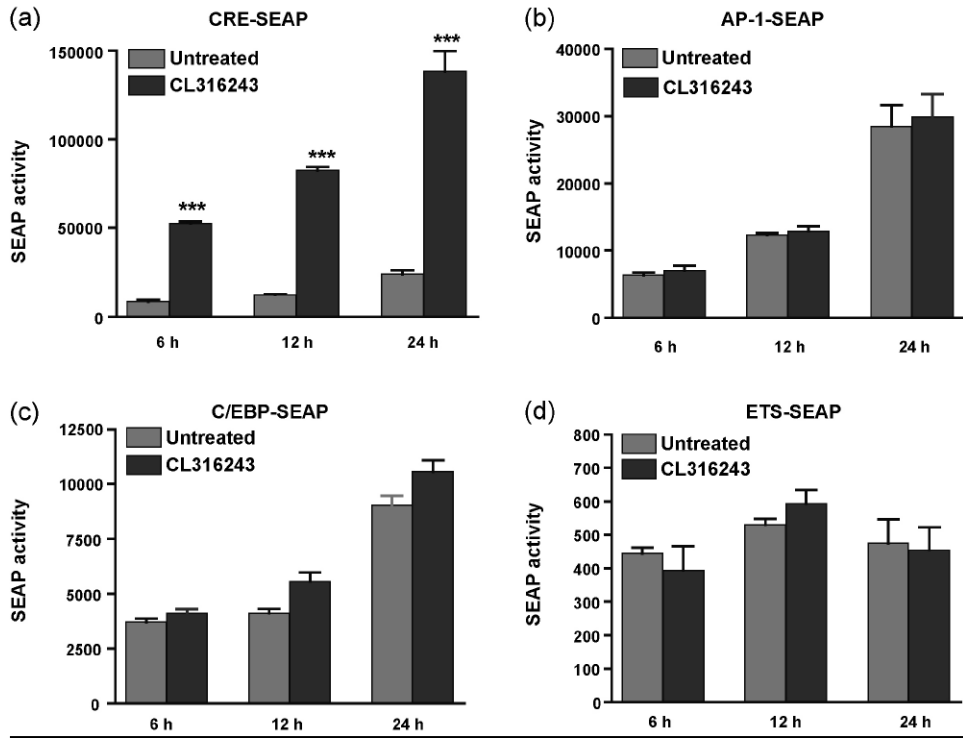
$\beta_3$ AR-induced IL-6 expression is not mediated by NF- $\kappa$ B pathway. C3H10T1/2 cells were treated with CL316243 (1  $\mu$ M) and TNF $\alpha$  (10 ng/ml) for 2 h. In the homogenous reporter assay, CL316243 didn't activate the NF- $\kappa$ B reporter, while TNF $\alpha$  did (a). 3T3-L1 cells were pre-treated with NBD (100  $\mu$ M) or MG-132 (25-50  $\mu$ M) for 1 h prior to treatment with CL316243 (1  $\mu$ M for 45 min). The CL316243-induced up-regulation of IL-6 mRNA was not inhibited by NBD or MG-132 (b). SEAP activity was assayed from the culture media of HEK 293 cells co-transfected with the pNF- $\kappa$ B-SEAP reporter construct and  $\beta_3$ AR expression vector. No significant increase in SEAP activity was observed at 6, 12 and 24 h post-treatment with CL316243 (1  $\mu$ M) (c). \*\*\* $P$ <0.001 and \*\* $P$ <0.01 different from untreated.

**Fig. 4.**

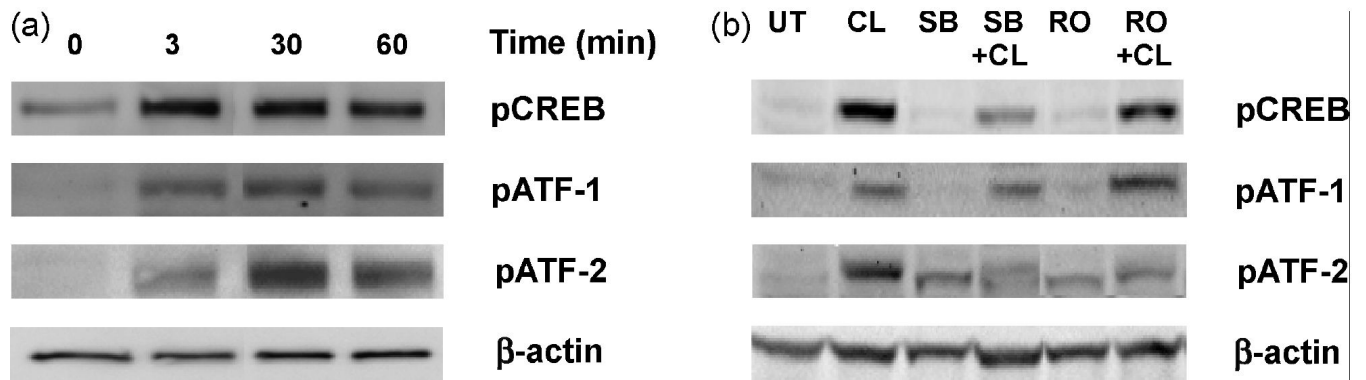
$\beta_3$ AR-induced IL-6 expression is Gi- and ERK1/2-independent and mediated by p38 signaling pathway. 3T3-L1 cells were pretreated with PTX (100 ng/ml and 200 ng/ml for 1 h) prior to stimulation with CL316243 (1  $\mu$ M for 45 min). The CL316243-induced up-regulation of IL-6 transcripts was not blocked by PTX (a). 3T3-L1 cells were pretreated with either U0126 (10  $\mu$ M) or SB203580 (5-10  $\mu$ M) for 1 h prior to stimulation with CL316243 (1  $\mu$ M for 45 min). The CL316243-induced up-regulation of IL-6 transcripts was blocked by SB203580, but not U0126 (b). C3H10T1/2 cells were pretreated with SB203580 (10  $\mu$ M for 1 h) prior to stimulation with CL316243 (1  $\mu$ M for 45 min). The CL316243-induced activation of CRE reporter was inhibited by SB203580 in homogenous reporter assay (c). The cells lysates were also subjected to Western blot analyses with antibodies to p-p38. Increased phosphorylation of p38 was observed at 3, 30 and 60 min post-treatment with CL316243.  $\beta$ -actin was used as the loading control (d). \*\*\* $P$ <0.001 and \*\* $P$ <0.01 different from untreated. # $P$ <0.05 different from CL316243-treated.

**Fig. 5.**

$\beta_3$ AR-induced IL-6 expression is dependent on the PKC signaling pathway. 3T3-L1 cells were pretreated with RO31-8220 (1-10  $\mu$ M) for 1 h prior to stimulation with CL316243 (1  $\mu$ M for 45 min). The CL316243-induced up-regulation of IL-6 transcripts was blocked by RO31-8220 (a). C3H10T1/2 cells were pretreated with RO31-8220 (10  $\mu$ M for 1 h) prior to stimulation with CL316243 (1  $\mu$ M for 2 h). The CL316243-induced activation of CRE reporter was inhibited by RO31-8220 in the homogenous reporter assay (b). The cells lysates were also subjected to Western blot analyses with antibodies to pPKCpan ( $\beta$ II Ser660), pPKC $\alpha/\beta$ II (Thr638/641), pPKC $\theta$  (Thr538), and pPKC $\epsilon$  (Ser729). Increased phosphorylation of PKC $\alpha/\beta$ II was observed at 30 sec and PKCpan, PKC $\theta$  and PKC $\epsilon$  45 min post-treatment with CL316243.  $\beta$ -actin was used as the loading control (c). \*\*\* $P$ <0.001 and \*\* $P$ <0.01 different from untreated. # $P$ <0.05 different from CL316243-treated.



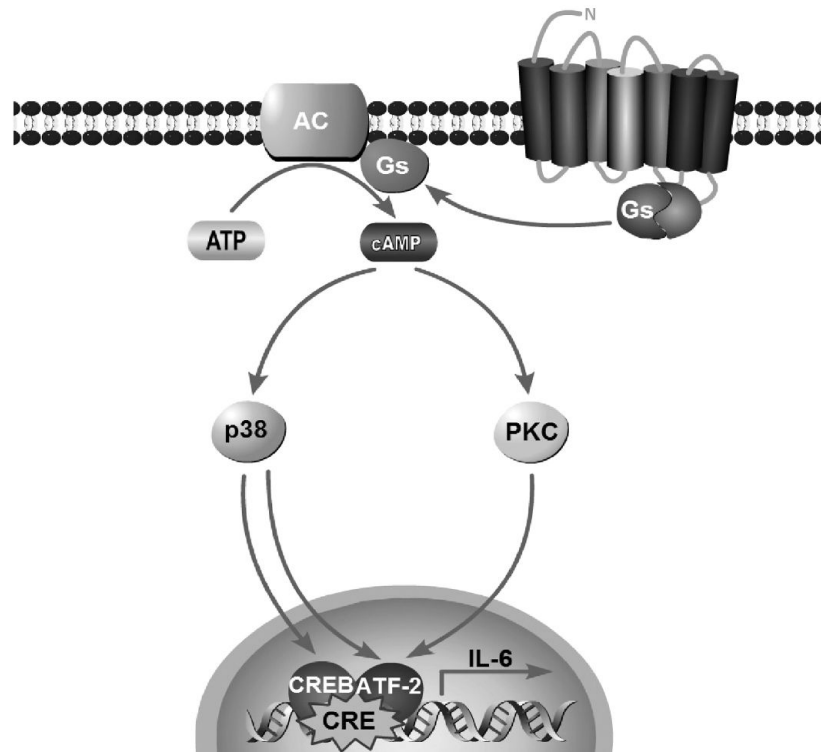
**Fig. 6.** Stimulation of  $\beta_3$ ARs results in up-regulation of the reporter construct containing the CRE site. HEK 293 cells were transiently co-transfected with pCRE-SEAP (a), pAP1-SEAP (b), pC/EBP $\beta$ -SEAP (c), or pETS-SEAP (d) reporter constructs along with the  $\beta_3$ AR expression vector. 24 h after transfection, media was changed, and cells were stimulated with CL316243 (1  $\mu$ M) for 6, 12 and 24 h. Significant increase in SEAP activity was observed with pCRE-SEAP construct only. \*\*\* $P$ <0.001 different from untreated.



**Fig. 7.**

Effect of  $\beta_3$ AR agonist stimulation on phosphorylation of CREB, ATF-1, and ATF-2. 3T3-L1 cells were stimulated with CL316243 (1  $\mu$ M) for 3, 30, and 60 min, then cell lysates were subjected to Western blot analyses with antibodies to pCREB, pATF-1, and pATF-2.

CL316243 stimulation increased phosphorylation of pCREB, pATF-1, and pATF-2 (a). 3T3-L1 cells were pretreated with SB203580 (10  $\mu$ M for 1 h) or RO31-8220 (10  $\mu$ M for 1 h) before stimulation with CL316243 (1  $\mu$ M for 45 min). CL316243-induced phosphorylation of CREB and ATF-2 was attenuated in the presence of SB203580, while ATF-2 phosphorylation was also decreased in the presence of RO31-8220, although non-specific phosphorylation of ATF-2 also has been observed in presence of inhibitors alone. ATF-1 phosphorylation was not affected by inhibitor treatment.  $\beta$ -actin was used as the loading control (b).



**Fig. 8.** Schematic diagram of the proposed signaling pathways for activation of IL-6 gene by  $\beta_3$ AR stimulation in white adipocytes.