

Calcium-Induced Expression of *Period1* in SK-N-SH Human Neuroblastoma Cells

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ABSTRACT

In mammals, the suprachiasmatic nuclei (SCN) of the hypothalamus control endogenous circadian rhythms and entrainment to the environment. Light stimuli transiently increase the gene expression of *Period 1* (*Per1*) that is important in resetting of the mammalian central clock in the SCN. Previous studies demonstrated that glutamate and N-Methyl-D-Aspartate (NMDA) mediate the effect of light stimuli on the central clock *in vivo*. However, molecular mechanism underlying the acute induction of *Per1* by the extra stimuli remains unclear. The present study is designed to set up an *in vitro* model and to examine the effect of calcium ion on *Per1* gene expression in the SK-N-SH human neuroblastoma cells using ribonuclease protection assay. Calcium ionophore A23187 (A23187) dramatically increased *Per1* mRNA levels in a time- and dose-dependent manner although glutamate and NMDA were unable to alter *Per1* gene expression in the SK-N-SH human neuroblastoma cells. In addition, the activity of human *Per1* promoter (-3829 to +123) was significantly promoted by treatment with A23187. The transcriptional inhibitor actinomycin D completely blocked A23187-induced increases in *Per1* mRNA levels, whereas cycloheximide, a potent translational inhibitor, did not. These results suggest that calcium ion may play an important role in the activation of *Per1* gene expression at the transcriptional level probably through the calcium responsive cis-elements located in 4.0 kb promoter region of *Per1* gene.

Key words: *Period 1*(*Per1*), clock gene, calcium ion, A23187, circadian clock, SK-N-SH human neuroblastoma cells

INTRODUCTION

All living organisms on earth have unique biological

rhythms at period of approximately 24 h. That is so called circadian rhythm. In mammals, the master circadian clock resides in the suprachiasmatic nucleus (SCN) located in the hypothalamus of the brain (Johnson et al., 1988; Weaver, 1998). Light stimuli synchronize circadian oscillations of the core clock genes in SCN. Interactions among these

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"clock cells" in the whole SCN serve to synchronize individual circadian clocks to generate coordinated circadian outputs. These outputs ultimately control a vast array of circadian rhythms in physiology and behavior.

A clear view for the mammalian molecular clock within SCN neurons is beginning to emerge (Ikeda and Nomura, 1997; Tei et al., 1997; Reppert, 1998). The central clockwork appears to involve the transcriptional/translational feedback loop, which is similar to that in *Drosophila melanogaster*. In the fly, the positive transcriptional regulation of *Period* (*Per*) and *timeless* (*tim*) is controlled by two basic helix-loop-helix (bHLH)/PAS proteins, dCLOCK and dBMAL1, which heterodimerize and bind to the E box (CACGTG) (Hao et al., 1997; Allada et al., 1998; Darlington et al., 1998; Rutila et al., 1998; Lee et al., 1999).

Prominent differences between mammals and the other lives so far are discovered in the anatomical and molecular pathways by which light entrains the clock. In *Drosophila*, light leads to a degradation of TIM that is independent of eyes. In mammals, light coming into the eyes is transsynaptically communicated to the SCN, where it leads to an induction of *Per* mRNA. Photoc signals are communicated via the retinohypothalamic tract (RHT), a direct neural projection from the retina to the SCN (Van den Polan, 1991; Gannon and Rea, 1993; Johnson et al., 1988). The master clock restricts its own sensitivity to stimulation so that only nocturnal light adjusts its timing, causing phase delays in early night and phase advances in late night (DeCousery, 1960; Takahashi and Zatz, 1982; Summers et al., 1984). This ensures that daily behavioral rhythms are synchronized appropriately to phases of the environmental cycle of darkness and light.

It is of notice that glutamate is the primary neurotransmitter transmitting light signals to the SCN (Johnson et al., 1988; Castel et al., 1993; Ding et al., 1994; Hannibal et al., 2000). Throughout night, a phase-resetting light stimulus evokes release of glutamate from the RHT and activation of multiple glutamate receptor types, of which NMDA receptors are critical (Colwell and Menaker, 1992; Ding et al., 1994; Mintz et al., 1999). The consequent influx of Ca^{2+} activates nitric oxide synthase to produce nitric oxide (Ding et al., 1994; Amir et al., 1995).

After liberation of nitric oxide, the light signaling pathways diverge (Gillette, 1997). Light stimulus induce phase delay of the master clock through activation of neuronal ryanodine receptors in early night (Ding et al., 1998), whereas it promote the phase advance through activation of cGMP/PKG-dependent signal transduction cascade in late night (Weber et al., 1995; Ding et al., 1998). In both case, light-induced signaling cascades eventually induce phosphorylation of Ca^{2+} /cAMP response element binding protein (CREB) and CRE-mediated transcriptional activation (Ginty et al., 1993; Ding et al., 1997; Obrietan et al., 1998, 1999) and expression of a number of immediate-early genes including *c-fos* and *junB* in the retinorecipient zones of the SCN. These data strongly support the possibility that transcriptional regulation may play a critical role on resetting of the mater clock in the SCN.

Rapid and transient induction of *Period 1* expression is a prominent result of light-induced signaling cascades in both early and late night (Albrecht et al., 1997; Shearman et al., 1997; Shygeyoshi et al., 1997; Takumi et al., 1998; Zylka et al., 1998) and this increase is most critical for the entrainment of circadian behavior. However, the molecular mechanism underlying resetting of the master clock is largely unknown manly because of the lack of *in vitro* model. The present study is designed to set up an *in vitro* model and to examine the direct effect of calcium ion on the expression of *Per1* gene was examined in SK-N-SH human neuroblastoma cells.

MATERIALS AND METHODS

Materials

The human neuroblastoma cell line SK-N-SH (HTB-11) was obtained from the American Type Culture Collection (Rockville, MD, U.S.A.) and cell culture reagents were from Gibco (NY, USA). A23187 were from Calbiochem (La Jolla, CA, USA). pGL3-luciferase vectors were from Promega (Madison, W1, USA) and pCMV- β gal that consists of cytomegaloviral promoter fused to the β -galactosidase gene was obtained from Stratagene (La Jolla, CA, USA). Other chemicals were purchased from Sigma.

Cell culture and drug treatment

SK-N-SH cells were maintained in Dulbecco's modified Eagle's medium (DMEM) with 4 mM glutamine, 1 mM sodium pyruvate, 100 U/ml penicillin, 100 µg/ml streptomycin, and 10% fetal bovine serum under a humidifying atmosphere containing 5% CO₂ at 37°C. Medium was refreshed every 2~3 d. For drug treatment, medium was changed to serum-free DMEM and incubated for 24 h. After incubation, medium was changed to 0.5 µM A23187 containing medium. Actinomycin D (5 µg/ml) and cycloheximide (10 µM) were treated 3 h and 2 h before A23187 treatment, respectively.

Ribonuclease protection assay

Total RNAs were isolated by an acid guanidinium thiocyanate-phenol-chloroform method (Chomczynski and Sacchi, 1987). Total RNA from SK-N-SH cells was reverse-transcribed into cDNA using random hexamer. A 220 bp *hPer1* (position: 599-818) fragment was generated by reverse transcription-polymerase chain reaction (RT-PCR) with primers of 5'-ctcatgacagcacttcgaga-3' and 5'-cagacgtgatgtgctcagc-3' using cDNA as a template. This *hPer1* cDNA fragment was first cloned into pGEM-T easy vector (Promega) to generate *hPer1-T* and the sequence was confirmed by sequencing. Antisense strand riboprobe was generated by *in vitro* transcription using T7 polymerase (Promega) in the presence of ³²P-UTP using Sal-linearized *hPer1-T* as a template.

A 248 bp cyclophilin (position: 160~407) fragment was generated by RT-PCR with primers of 5'-agcactggggagaaaggatt-3' and 5'-agccactcagtcttg-cagt-3' using cDNA as a template. This cyclophilin cDNA fragment was first cloned into pGEM-7G vector (Promega) to generate hcyclophilin-7G using SmaI and the sequence was confirmed by sequencing. Antisense strand riboprobe was generated by *in vitro* transcription using T7 polymerase in the presence of ³²P-UTP using SpeI linearized cyclophilin-4G as template.

All reagent used in the ribonuclease protection assay (RPA) are from the Ambion RPA II (Ambion), unless otherwise specified. Total RNA (10 µg) was hybridized with *hPer1* (100000 cpm) and cyclophilin (10000 cpm) riboprobes at 42°C overnight. The hybridized mixture was then treated with RNase A/T1

mix (1 : 200) at 37°C for 1 h. The ribonuclease digestion was terminated by addition of 300 µl of RNase inactivation and precipitation solution and protected fragment was precipitated at 20°C for 30 min. The pellets were dissolved in loading buffer and subjected to electrophoresis in the denaturing gel (8 M urea, 5% polyacrylamide) at 250 V for 2 h. The gel was dried and exposed to x-ray film for 24~48 h using intensifying screens.

Transfection

phPer1-Luc vector containing 4 kb *hPer1* promoter (-3829 to +123) was generated by RT-PCR using primers (5'-aagctagcgggatcatgcaaatacgtga-3' and 5'-aa ctcgaggtagccatcagattggaag-3'). For the transfection experiments, cells were plated in 60 mm dishes and grown to 30~40% confluence for 2~3 d. Cells were washed twice with 1X Dulbecco's phosphate-buffered saline (D-PBS) and medium was changed to DMEM with 10% fetal bovine serum. After two days, cells were washed twice with 1X D-PBS, received 5 ml of fresh medium, and incubated at 37°C for 2 h before transfection. Reporter plasmids for transfection experiments were purified by Qiagen columns (Qiagen GmbH, Hilden, Germany) according to manufacturer's instructions and dissolved in 0.1X TE buffer (1 mM Tris (pH 8.0), 0.1 mM EDTA) at a concentration of 0.5~1 mg/ml. DNA-calcium phosphate mixture was prepared by mixing one part of 2X BES-buffered saline (50 mM BES (pH 6.96), 280 mM NaCl, and 1.5 mM Na₂HPO₄) with one part of DNA-calcium chloride solution (60 µg/ml DNA, 250 mM CaCl₂) and incubated at room temperature for 15 min before transfection. Each dish received 0.5 ml of resulting DNA-calcium phosphate mixture containing 5 µg β-galactosidase reporter vector (pCMV-βgal) drop wise and was incubated at 37°C under 5% CO₂ tension for 24 h. Excess DNA-calcium phosphate precipitates were washed out by rinsing twice with calcium and magnesium-free D-PBS. Then cells were cultivated in an appropriate drug-added medium and further incubated for 24~48 h before making cell extracts. Cell extracts were prepared by incubating cells in 0.4 ml of 1X reporter lysis buffer (Promega) for 15 min at room temperature. After centrifugation, supernatants were kept at

-70°C till assay. Protein contents were measured by commercial protein assay kit (Bio-Rad laboratories, Hercules, CA, USA). Beta-galactosidase assay and luciferase assay were performed using β -galactosidase assay kit and luciferase assay kit (Promega), respectively. *phPer1*-Luc exhibited high luciferase activity (3×10^4 to 1×10^5 light unit per 30 seconds depending on experimental conditions). Untransfected SK-N-SH cell extracts showed no background luciferase activity and endogenous β -galactosidase activity was negligible under the assay condition used. Cytomegaloviral promoter-driven β -galactosidase activity was used to compensate transfection efficiency.

Statistical analysis

Data for *hPer1* mRNA induction by ionophore and promoter activity were statistically evaluated using Student's *t*-test. Statistical significance was set at $p < 0.05$.

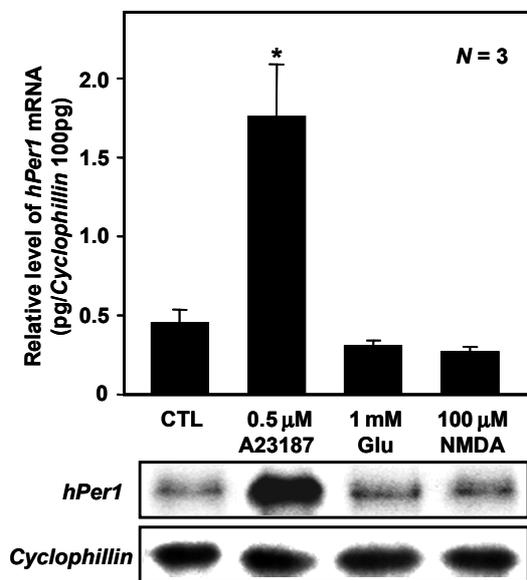


Fig. 1. Effects of calcium ionophore A23187, glutamate and NMDA on *Per1* mRNA levels. SK-N-SH cells cultured for 24 h in the growth medium were incubated for 24 h in the serum-free medium. After incubation, SK-N-SH cells were treated with 0.5 μ M A23187, 1 mM glutamate and 100 μ M NMDA, respectively. Two hours after the treatment, total RNA was isolated and mRNA levels of *Per1* were determined by RPA (* $p < 0.05$).

RESULTS

Glutamate and NMDA failed to regulate *hPer1* mRNA levels in SK-N-SH cells

SK-N-SH cells are derived from human neuroblastoma, and their neuronal origin has been extensively characterized (Biedler et al., 1973). To investigate whether glutamatergic neurotransmitters influence *Per1* gene expression, 0.5 μ M A23187, 1 mM glutamate and 100 μ M NMDA were treated for 2 h (Fig. 1). A23187 significantly increased *hPer1* mRNA levels in the SK-N-SH cells. At this dose and time window, glutamate and NMDA did not increase *Per1* mRNA levels. Similarly, excess amount of NMDA (1 mM and 10 mM) also had no alteration in *hPer1* mRNA levels (data not shown).

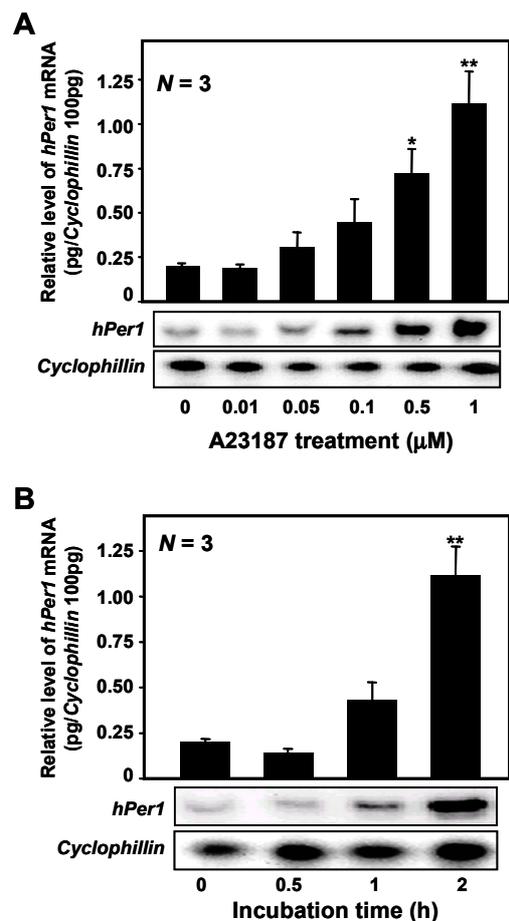


Fig. 2. Dose- and time-dependent increase in *Per1* mRNA levels by A23187. SK-N-SH cells were treated with increasing amounts of A23187. Two hours after the treatment, total RNA was isolated and subjected to RPA (A). SK-N-SH cells were treated with 0.5 μ M A23187 and incubated for indicated time (B) (* $p < 0.5$, ** $p < 0.01$).

A23187 increased *Per1* mRNA levels in a time- and dose-dependent fashion

To examine effective time and dose-response, SK-N-SH cells were exposed to A23187. As shown in Fig 2A and B, the increases in *hPer1* mRNA levels by A23187 were time and dose dependent. In particular, increases in *hPer1* mRNA levels were peak after 2 h incubation with 0.5µM A23187. Also, these effects were inhibited by treatment with EGTA, a calcium chelator, indicating that calcium influx by A23187 appears to increase *Per1* mRNA

levels (data not shown).

Actinomycin D, but not cycloheximide blocked A23187-mediated increase of *hPer1* mRNA

To elucidate classical transcriptional aspect, SK-N-SH cells were preincubated with 5µg/ml of actinomycin D (Act D), a polymerase II inhibitor, 3 h before A23187 treatment and then 0.5µM A23187 was treated. After 2 h incubation, *Per1* mRNA levels were examined. Fiveµg/ml of Act D completely blocked A23187-mediated induction of *Per1* mRNA (Fig. 3A). Time course experiment showed that *Per1* mRNA levels remained until 2 h after Act D treatment, and then declined almost at the basal levels (Fig. 3B). These results indicate that A23187 regulates *Per1* mRNA level at transcription level.

CHX had no effect on A23187-mediated increase of *hPer1* mRNA

To delineate whether protein synthesis is involved in A23187-mediated *Per1* induction, 10µM cyclohexamide, a protein synthesis inhibitor, was treated to SK-N-SH cells. While CHX did not alter A23187-induced increase in *hPer1* mRNA levels (Fig. 4), pretreatment of CHX alone for 2 h increased *hPer1* mRNA levels by 3 times. These results suggest

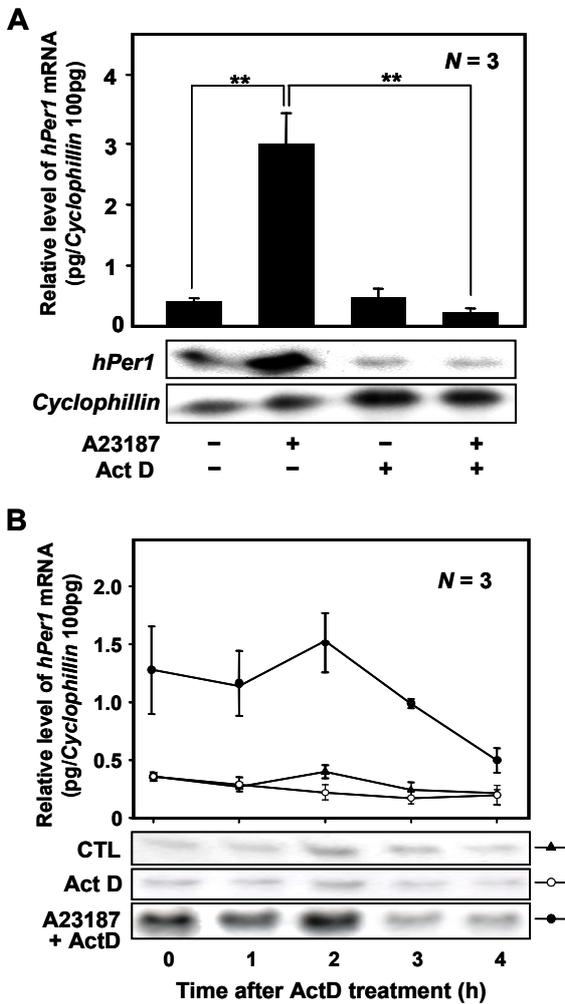


Fig. 3. Effect of actinomycin D on A23187-mediated *Per1* induction. (A) SK-N-SH cells were incubated for 3 h in the presence (+) or absence (-) of 5µg/ml actinomycin D. After incubation, they were treated with 0.5µM A23187 and further incubated for 2 h. (B) the cells were pretreated with actinomycin D (5µg/ml) for indicated periods followed by co-treatment with 0.5µM A23187 for 2 h. Total RNA was isolated and subjected to RPA (**p<0.01).

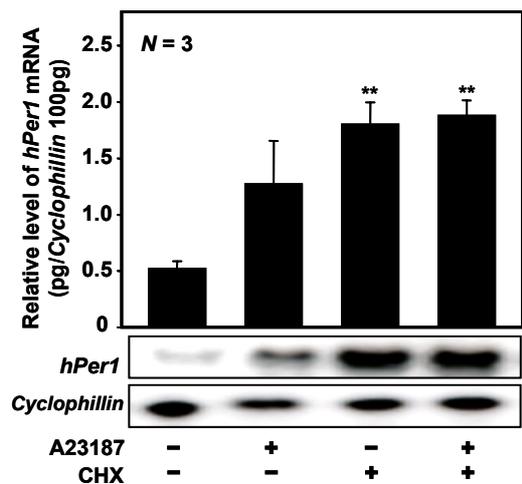


Fig. 4. Effect of cycloheximide on the A23187-mediated *Per1* induction. The SK-N-SH cells were incubated for 3 h in the presence (+) or absence (-) of 10µM cycloheximide (CHX). After incubation, they were treated with 0.5 µM A23187 and further incubated for 2 h. Total RNA was isolated and subjected to RPA (**p<0.01).

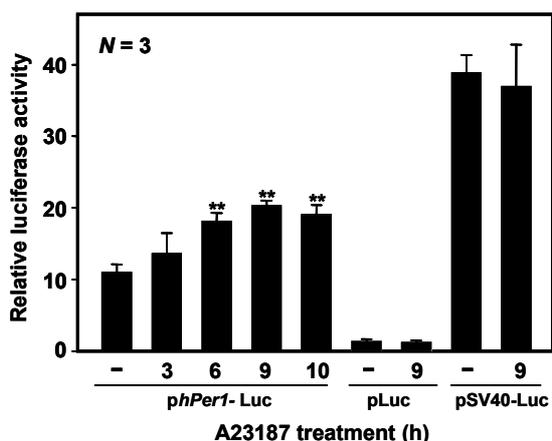


Fig. 5. Time course change of A23187-mediated activation of *hPer1* promoter. SK-N-SH cells were transfected with *phPer1*-4.0 Luc or pGL3-basic by the calcium phosphate co-precipitation method. Twenty-four hours after the transfection, cells were incubated for 9 h in the presence (+) or absence (-) of 0.5 μ M A23187. Cells were then harvested and luciferase activity was determined (* $p < 0.05$).

that de novo protein synthesis is not required for *Per1* induction by A23187, rather is needed for rapid *Per1* degradation.

A23187 increased the human *Per1* promoter activity

To test whether the changes in *hPer1* mRNA levels by A23187 are coupled with enhanced gene transcription, *hPer1* promoter activity was determined by transient transfection assay. SK-N-SH cells were transfected with *phPer1*-4.0Luc containing 4kb (-3829 to +123) *hPer1* promoter-driven luciferase reporter. Treatment of SK-N-SH cells with 0.5 μ M A23187 increased the promoter activity by about 1.8 folds (Fig. 5). As expected, treatment of A23187 did not alter the basic reporter vector and SV40 promoter-driven luciferase activity as controls.

DISCUSSION

It is well known that light pulse or NMDA can elicit rapid induction of *Per1* mRNA in the SCN. For example, a single brief exposure to light induces *mPer1* RNA within 1 h and shifts phase of the oscillation of its own expression (Shigeyoshi et al 1997). In this regard, anti-sense *Per1* nucleotide blocks phase shifts induced by light, glutamate, and NMDA (Akiyama et al., 1999).

In the present study, we shows that Ca^{2+} influx induced by A23187, a calcium ionophore, increased *hPer1* mRNA levels significantly within 2 h in SK-N-SH human neuroblastoma cells. Such increase in *hPer1* mRNA levels was clearly time- and dose-dependent and attenuated by chelation of extracellular Ca^{2+} using EGTA. This observation is consistent with recent studies. For instance, light-stimulated *mPer1* induction in mice is a rapid, transient event that is initiated by 15~20 min in the retinorecipient SCN peaks at 60 min, and spread throughout the SCN (Albrecht et al., 1997; Shearman et al., 1997; Shigeyoshi et al., 1997; Takumi et al., 1998; Zylka et al., 1998).

In the present study, we found that treatment with actinomycin D completely blocked the increase of *Per1* mRNA level induced by A23187. This result suggests that Ca^{2+} regulate the expression of *Per1* gene at the transcriptional level. Gene induction has been believed to be the basis on resetting the biological clock in the SCN. Similarly, light pulses that reset the clock also induce the expression of a number of immediate-early genes, including *c-fos* and *junB* in the retinorecipient zones of the SCN (Kornhauser et al., 1996; Best et al., 1999). Also, it has been reported that light-induced gene expression in the SCN is in the control of a well established signaling pathway, in which RHT synaptic glutamate acts through postsynaptic NMDA receptors to increase intracellular levels of Ca^{2+} , which in turn trigger a kinase cascade to activate transcription factors CREB and ERK (Kornhauser et al., 1996; Obrietan et al., 1998). Recently, it is demonstrated that there is a link between light signaling and CRE activation, and these effects are mediated by glutamate, nitric oxide synthase, MAPK, and CREB phosphorylation in the mouse (Obrietan et al., 1999). In addition to *mPer1* induction, light/ glutamate stimuli promote phosphorylation of CREB and CRE-mediated transcription, either directly or indirectly (Ginty et al., 1993; Ding et al., 1997).

We observed that increased levels of *hPer1* mRNA induced by A23187 were quickly declined to control levels 4 h after treatment with actinomycin D. It is noticeable that degradation of *Per1* mRNA was faster in the A23187-treated group than in A23187-untreated group. This result suggests that stability of *hPer1* mRNA induced by A23187 is

shorter than 4 h and there may be an active degradation of *Per1* mRNA triggered by calcium ion. On the other hands, pretreatment with cycloheximide, a protein synthesis blocker did not block induction of *Per1* mRNA by A23187. This result suggests that protein synthesis is not required for A23187-mediated *Per1* induction. Treatment with cycloheximide alone also led to a 3.5 fold increase of *hPer1* mRNA, which may be resulted from prolonged half-lives (Balsalobre et al., 1998, 2000). Based on such results, we speculate that de novo protein synthesis is not required for induction of *Per1* mRNA mediated by A23187 but for rapid degradation of *Per1* mRNA.

In the present study, using whole proximal promoter (-3829 to +123, 4 kb) of *hPer1* gene containing 3 E-boxes (Hida et al., 2000; Yamaguchi et al., 2000), we also found that *hPer1* promoter-driven luciferase activity was significantly increased 6 h after treatment with A23187 in SK-N-SH cells. Thus, it is conceivable that there resides the calcium responsive element in the 4.0 kb 5' flanking region of *hPer1* gene. These results provide evidence that calcium can play an important role in transcriptional activation of *hPer1* gene.

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