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Erk1/2 dependent Phosphorylation of PKCα at Threonine 638 in Hippocampal 5-HT1A Receptor Mediated Signaling

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Abstract

Stimulation of the serotonin 1A receptor $(5-HT_{1A}-R)$ causes activation of extracellular signalregulated protein kinase (Erk) and protein kinase C alpha (PKCα) in both hippocampal HN2-5 cells and cultured hippocampal slices from postnatal day-15 (P15) mice. Our earlier studies demonstrated that PKC α is coimmunoprecipitated with Erk and the phosphorylation of PKC α in this Erk–PKC α complex is dependent on the Erk pathway. Furthermore, the T⁶³⁸ residue, which must be phosphorylated for the complete activation of $PKC\alpha$, is within an authentic MAP kinase consensus domain (S/TP), and the PKCα protein also contains two docking sites for Erk such as KRGRIYL and KRGIIYRDLKL. Using Föster Resonance Energy Transfer (FRET) we have confirmed an association between Erk and $PKC\alpha$. Employing $PKC\alpha$ and Erk mutants we next demonstrated that Erk causes direct phosphorylation and activation of PKCα. By mutating the phosphoinositide dependent kinase-1 (PDK-1)-promoted phosphorylation site (S^{497}) and the kinase site (K^{368}) in PKC α , we observed that both of these autophosphorylation-deficient mutants are phosphorylated at T638 in an Erk-dependent manner. To confirm that Erk indeed catalyzes phosphorylation of PKC α at T⁶³⁸, we used a mutant Erk construct in which a bulky amino acid residue in the ATP binding site (Q^{103}) had been replaced with glycine, enabling this mutant to utilize a bulky analog of ATP, cyclopentyl ATP. An *in vitro* kinase assay using this mutant Erk protein, radiolabeled cyclopentyl ATP, and a synthetic oligopeptide containing the S/TP site of $PKC\alpha$ showed phosphorylation of the peptide by Erk1/2. These results confirm the novel possibility that PKC α is a direct substrate of Erk1/2 in neuronal cells and help link two important signaling molecules that regulate maturation and protection of hippocampal neurons as well as many other cell types.

Keywords

Hippocampus; PKC α ; Erk; 5-HT_{1A} receptor

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Introduction

The serotonin 1A receptor (5-HT_{1A}-R) is a heptahelical G_i - and G_o -coupled receptor, which functions by inhibiting adenylyl cyclase, stimulating inwardly rectifying K^+ channels, or modestly stimulating phospholipase $C[1]$. In the brain, this receptor is present in the raphé neurons as their target postsynaptic neurons in the front brain [1;2]. The receptor at the presynaptic site (autoreceptor) mainly regulates serotonin release and at the postsynaptic site (heteroreceptor) regulates synaptic activities by stimulating inwardly rectifying K^+ channels or directly inhibiting Ca^{2+} channels. Functionally, the 5-HT_{1A}-R receptor is crucial in various aspects such as synaptic plasticity, neurogenesis, neuroprotection, and also in anxiety, depression, learning and memory [2]. In our earlier studies, we have shown that stimulation of the receptor with 8-OH-DPAT (a 5 -HT_{1A} agonist) causes activation of mitogen-activated protein kinase (MAPK) isozymes Erk1 and Erk2) and also Protein kinase C alpha (PKCα) in both in HN2-5 cells (a mouse hippocampal neuron-derived cell line [3]), and organotypic hippocampal slice cultures from postnatal day 15 (P15) mice [4]. Additonally, PKCαis also co-immunoprecipitated with Erk1/2 [4]. The phosphorylation of PKC α is dependent on Erk1/2 as MEK (Erk kinase) inhibitors PD98059 and U0126 inhibit PKC α activation but the general PKC inhibitor GFX does not abrogate activation of PKC α or ERk1/2. A similar Erk1/2 dependent activation of PKCαhas been reported in parathyroid induced opossum kidney cells [5], although in many studies, the $PKC\alpha$ activation is shown to be upstream of Erk1/2 [6;7;8]. The PKC α isoenzyme is synthesized as an inactive membrane bound precursor and undergoes phosphorylation at three sites T^{497} (Activation loop), T^{638} (Turn motif), and S^{657} (hydrophobic domain) for its activation. Phosphorylation of the activation loop is promoted by the enzyme phosphoinositide dependent kinase 1 (PDK-1), which is reported to be an essential event in the activation of PKC isoenzymes as well as other members of ABC family members of protein kinases [9;10]. This activation loop phosphorylation stimulates autophosphorylation of both "turn motif" and "hydrophobic domain", turning it into a catalytically competent and stable mature PKC [10]. Dephosphorylation of the "turn motif" abolishes PKC enzyme activity [11]. However the PKC molecule remains in an inactive state *via* association of the pseudo-substrate, which is a part of the PKC molecule itself, to the active site [12]. The final activation of PKC requires removal of the pseudo-substrate from its active site, which is carried out upon binding of DAG and calcium.

The MAPK enzymes require a very specific phosphorylation sequence where a serine or threonine is followed by proline (S/TP) [13]. In addition, a proline at position -2 is favorable but not absolutely required [14]. The Turn Motif of $PKC\alpha$ at T^{638} is within the consensus Erk phosphorylation sequence PVXTP. A very similar Erk phosphorylation site is reported in peroxiredoxin-6 where PVATP is the phosphorylation motif [15]. Additionally the MAPK enzymes require docking sites in the substrates for high affinity interactions [14]. The PKCαmolecule also contains two Erk1/2 docking domains such as KRGRIYL and KRGIIYRDLKL. So, there is a possibility that the "turn motif" phosphorylation may be catalyzed by Erk1/2. We have used Föster Resonance Energy Transfer (FRET) to demonstrate the interaction of PKC α with Erk1/2. Furthermore, we have demonstrated that autophosphorylation-deficient mutant forms of $PKC\alpha$ are phosphorylated at T^{638} in an Erk1/2-dependent manner upon stimulation of the $5-HT_{1A}$ -R in the HN2-5 cells. Finally, in order to validate Erk1/2 mediated phosphorylation of PKC α at T⁶³⁸, we have used a mutant Erk1/2 construct [16] in which a bulky amino acid residue in the ATP binding site (glutamine 103) has been replaced with glycine, enabling this mutant to utilize a bulky analog of ATP (cyclopentyl ATP). A kinase reaction assay using this mutant Erk1/2 protein in its activated state, radiolabeled cyclopentyl ATP, and a synthetic peptide from the PKCαprotein containing the S/TP site in a reaction buffer have been used to demonstrate mutant Erk1/2 mediated phosphorylation of this peptide.

Materials and Methods

Cell culture, transfection, and immunoprecipitation

HN2-5 cells were grown in DMEM media with 10% FBS, 1% (v/v) penicillin-streptomycin solution in a 5% -CO₂, humidified incubator. Transfection of different vector constructs was carried out in six-well plates (3 μg DNA/well) using Turbofect Transfection Reagent (MBI, Fermentas). Overnight-transfected cells were differentiated by treatment with 5 μM retinoic acid in 1% FBS containing medium. After 24 h, cells were serum starved for 1 h then treated with different drugs with or without 1 μ M 8-OH-DPAT (D) for 1 h (for the stimulation of PKC). The cells were washed once with ice cold PBS and were lysed in a buffer (0.5 ml) containing 20 mM HEPES pH 7.4, 2mm EGTA, 2 mM $MgCl₂$, 200 mM $Na₃VO₄$, 1% Triton X-100, 2 mM PMSF and protease inhibitor cocktail (Roche, Mannheim, Germany). All manipulations were carried out on ice. The cell lysates were incubated overnight at 4 °C with a Myc antibody (Cell Signaling) at 1:200 dilution. Next, a 40 μL aliquot of a slurry of previously washed A/G agarose beads was added to each lysate sample and the mixture was again incubated with rocking overnight at 4° C. Following this, the Protein A/G agaroseimmunoprecipitated complex was centrifuged, the pellet washed eight times with the same lysis buffer (1 ml at a time). Finally each pellet was reconstituted in 30-μL 2X Laemmli treatment buffer, heated in boiling water for 10 min, centrifuged, and the supernatant resolved using SDS PAGE. Resolved proteins were then transferred to a nitrocellulose membrane, which was probed successively with different antibodies. The membranes were incubated in blocking buffer [5% bovine serum albumin in TBS-T (20 mmol/L Tris–HCl, pH 7.4, 0.8% NaCl, 0.1% Tween 20)], and then probed with different primary antibodies, followed by treatment with HRP-linked respective secondary antibodies. Each nitrocellulose membrane was first probed with a rabbit polyclonal PPKC α /ß II (T^{638/641}) antibody (1:1000). This was followed by stripping of the blot in 0.2 M glycine (pH 2.5), blocking and reprobing with the same Myc antibody (1:1000) as used for immunoprecipitation. Detection was carried out with the Supersignal luminol kit and bands were densitometrically quantified using a Fluorchem FC2 imaging system (Alpha Innotech, San Leandro, CA). The intensities of the luminescent bands were digitized using SPOT DENSO software, and the P- $PKC\alpha$ band intensities were normalized to the corresponding Myc band intensities. All the experiments were repeated 3 times and statistical analysis was performed using student's ttest and ANOVA.

FRET Analysis

HN2-5 cells, grown on poly-L-lysine-coated coverslips to 50-70% confluence, were cotransfected with Erk-CFP and $PKC\alpha$ -YFP vectors (1 µg each). The control cells were similarly cotransfected with the empty vectors, pECFP and pEYFP (1 μg each). After 24 h, cells were treated for 1 h with carrier or 8-OH-DPAT, fixed in 4% paraformaldehyde (30 min at room temperature), washed with PBS, and mounted on glass slides. Images were acquired using a Leica Confocal Scanning System (Exton, PA) TCS SP2. ECFP (Donor) and EYFP (Acceptor) fluorescence were excited with 458 and 514 nm lasers, respectively. We used acceptor photo bleaching method in which a small region of interest expressing both the flourophores of which the EYFP fluorescence channel was photobleached using a 514 nm laser to 30% of the original intensity and an increase in the intensity of the donor Erk-CFP was recorded. ECFP and EYFP images were taken both before and after acceptor photobleaching [17]. FRET efficiency (*E*) was calculated using Leica software using the

following equation: $FRETeff = \frac{Dpost - Dpre}{Dpost}$ for all $Dpost > Dpre$, where *Dpre* and *Dpost* are the ECFP emission spectra before and after regional photobleaching.

Cloning and site-directed mutagenesis

The $PKC\alpha$ coding sequence, without the stop codon, was cloned into a pGEM-T vector (Promega, USA) and subcloned into a pcDNA6A vector in frame with Myc and His tags (Invitrogen). The cloned DNA was used as template for PCR based site-directed mutagenesis using primers: Muta-497-Fwd-5′TCACGACCAGG*G*CCTTCTGTG3′ and Muta-497-Rvs-5′ CACAGAAGG*C*CCTGGTCGTGA3′. The mutated bases (italicized) yield T ⁴⁹⁷**→**A mutant PKCα (ACC**→** GCC). Overlapping PCR products were generated using primer pairs (EcoR1-Fwdα and mutα-497-rvs) and (muta-497-Fwdα and XHO-Rvsα). The two PCR products were gel purified and mixed together. 10 ng of the mixed DNA was used as template in a PCR reaction containing the primers ECoR1-Fwdα and XHO-Rvsα. Sequencing was performed by Genewiz laboratory (NJ, USA). A Flag-tagged mutant ErkpCDNA3 vector used in this study has been described previously [16]. The full length mouse Erk and PKC cDNAs were cloned into pECFP and pEYFP respectively by standard procedures.

Mutant Erk (Erk*) mediated in-vitro kinase assay

Mutant Erk mediated *in vitro* kinase assay was followed, as previously described [16]. In the Flag-tagged mutant Erk construct, the normal amino acid residue (glutamine 103) is changed to a glycine residue (glycine 103), which causes the ATP binding site to be bulky and therefore, allows the utilization of an analog of ATP (cyclopentyl ATP) that cannot be used by wild-type Erk2 or other cellular kinases. The empty vector and the mutant FLAG-Erk vector were transiently transfected into 70-80% confluent HN2-5 cells using EXGen500 (MBI Fermentas, USA) transfection reagent. After 24 h, the cells were differentiated using 5 μM retinoic acid in 1% serum for 18 h, serum starved for 1 h, and then treated with 1 μM 8-OH-DPAT for 30 min (for the stimulation of Erk1/2). The cells were lysed in a hypotonic lysis buffer (20 mM HEPES pH 7.4, 2 mM EGTA, 2 mM $MgCl₂$, 200 mM $Na₃VO₄$, 2 mM PMSF and protease inhibitor cocktail). The expressed mutant FLAG-Erk protein was immunoprecipitated using anti-FLAG-M2- agarose beads (Sigma). Both the immunoprecipitated products from the empty vector and mutant Erk vector transfected cells were used for a kinase reaction assay by mixing ${}^{32}P$ -labeled cyclopentyl ATP, a synthetic, 17-amino acid oligopeptide from the putative Erk phosphorylation site of $PKC\alpha$ protein (FFTRGQPVLTPPDQLVIA), in a reaction buffer and incubating for 20 min at 30°C. The reaction mixture was heat inactivated and spotted onto a high performance thin layer chromatography (HPTLC) plate. The labeled and unlabeled oligopeptides were separated by developing the plate in n-butanol: acetic acid: water: 4:2:2. The resolved, radiolabeled peptide was visualized by autoradiography.

Results and Discussion

Physical interactions between Erk and PKCα

We employed the FRET microscopy method to monitor the in-cell physical association of Erk and PKC α . Insignificant FRET efficiency (<0.4%) was observed in the empty vectortransfected cells. Representative confocal images for FRET are shown in Figure 1. From five independent experiments, we obtained a mean FRET efficiency of 7.19±0.75%. This strongly indicated that Erk and PKCα protein molecules are held in a close physical association. It should be noted, that activated Erk distributes in both cytosol as well as nucleus (Fig. 1, upper panel, cyan fluorescence) whereas $PKC\alpha$ always remains in the cytoplasm (Fig. 1, lower panel, yellow fluorescence). Non-specific association accounted for <0.4% FRET efficiency.

Previously we have demonstrated that stimulation of $5-HT_{1A}-R$ causes Erk1/2-dependent phosphorylation of $PKC\alpha$ in both mouse hippocampus (P15) and hippocampal derived HN2-5 cells [4]. Here we show that transiently expressed wild type Myc-tagged $PKC\alpha$ protein in HN2-5 cells also undergoes phosphorylation at T^{638} in an Erk1/2 dependent manner. To achieve this, we transfected HN2-5 cells with a wild type PKCα construct and after 18 h, the cells were differentiated with retinoic acid for 16 h, then serum starved for 1 h, followed by treatment for 1 h with carrier or 8-OH-DPAT (a 5-HT_{1A} agonist) (1 μ M) in the absence or presence of Way100635 (a 5-HT_{1A} antagonist), U0126 (an MEK inhibitor), and GFX (a general PKC inhibitor) for 1 h. The protein products were immunoprecipitated with a Myc antibody and analyzed by Western blotting as described in methods. As shown in Figure 2, 8-OH-DPAT treatment caused an induction in Myc-tagged P- T^{638} -PKC α , which was eliminated in the presence Way100635 or U0126, but not GFX. Thus, $5-HT_{1A}-R$ -linked phosphorylation of $PKC\alpha$ was Erk-dependent, but not PKC-dependent in the hippocampal neuron-derived HN2-5 cells.

Phosphorylation of PKCα at T638 is not PDK-1 dependent

It has been reported that the activation loop of $PKC\alpha$ is phosphorylated by a phosphoinositide dependent kinase 1 enzyme (PDK-1). The PDK-1 enzyme binds to the Cterminal hydrophobic loop of the newly synthesized $PKC\alpha$ molecule, which serves as the docking site [18], and then phosphorylates the T^{497} residue of PKC α . Once this T^{497} is phosphorylated, the PKC α is reported to undergo autophosphorylation of T^{638} . We mutated the PDK-1 phosphorylation site of PKC α (T⁴⁹⁷ to A). This mutant, Myc-tagged PKC α cDNA construct was transfected into HN2-5 cells, the cells were differentiated and treated with carrier or 8-OH-DPAT (1 μ M) in the absence or presence of Way100635 (4 μ M), or U0126 (10 μM). The cells were lysed and immunoprecipitated using a Myc antibody. Western blotting analysis of the immunoprecipitated proteins revealed $5-HT_{1A}-R$ and Erkdependent phosphorylation of PKC α at T⁶³⁸ (Fig. 3).

PVLTP is a MAPK phosphorylation site and Erk1/2 causes phosphorylation of PKCα oligopeptide

All MAPK phosphorylates their substrates to a minimal consensus sequence Ser/Thr-Pro (S/ TP) and many potential substrates contains this motif [19]. The T^{638} residue of PKC α is located within an authentic MAPK consensus phosphorylation site (PVXTP). A very similar phosphorylation site (PVATP) has been identified in the peroxiredoxin-6 protein [15]. In order to establish that PVXTP is an Erk1/2 phosphorylation site we performed an *in vitro* kinase assay using a mutant Erk construct in which a bulky amino acid residue in the ATP binding site (Q^{103}) was replaced with glycine, enabling this mutant to utilize a bulky analog of ATP, cyclopentyl ATP. An 18-aa segment of PKCα protein sequence harboring the putative Erk target site (FFTRGQ*PVLTP*PDQLVIA) was used as a substrate in an *in vitro* reaction. (4a) Flag-tagged mutant Erk vector or the empty vector were transfected into hippocampal neuron-derived HN2-5 cells. The transfected cells were differentiated and then treated with 1 μM 8-OH-DPAT for 30 min. The mutant Erk was immunoprecipitated using a Flag-antibody and the IP resuspended in PKC kinase buffer, supplemented with ³²P-labeled cyclopentyl ATP and the 18-aa oligopeptide, followed by incubation at 30 °C for 20 min. HPTLC analysis of the product showed Erk-catalyzed phosphorylation of the $PKC\alpha$ peptide (Fig. 4b). Neither PKC α nor wild-type Erk can use cyclopentyl- $\left[3^2P\right]ATP$, therefore this phosphorylation must have been catalyzed by the mutant Erk molecule expressed in the differentiated HN2-5 cells.

A kinase site mutant form of PKCα(K368→A) is also phosphorylated at T638 in an Erkdependent manner

As an additional evidence beyond the observation that $5-HT_{1A}-R$ stimulation in the HN2-5 cells causes Erk-dependent phosphorylation of the PDK-1 site mutant version of PKC α , we transfected the HN2-5 cells with the cDNA for a Myc-tagged kinase site mutant form of PKC α . As observed for the PDK-1 site mutant, 5-HT^{1A}-R stimulation also caused increased T⁶³⁸ phosphorylation of the kinase site mutant in an Erk-dependent manner (Supplemental figure 1).

PKCα has two Erk1/2 docking sites

MAPKs recognize their substrates by initially binding to a motif called docking domain. The docking domain contains an amino acid sequence "K/R-K/R-(X)2–6-L-X-L", which harbors a cluster of at least two positively charged amino acids followed by a spacer of 2-to-6 residues from a hydrophobic-X-hydrophobic sequence containing long-chain aliphatic amino acids such as Leu and Ile. Pro, Asn, and/or Gly, which form turns and break helices, occur at a high rate in the spacer and in the sequence immediately up- stream of the hydrophobic-X-hydrophobic element [20,21]. Another type of docking motif is a short peptide containing the sequence FXFP that is usually downstream of the phosphorylation site. This motif was originally found to bind the ERK subfamily but lately it has been shown to promote interactions with p38 alpha and p38 beta2 [22,23]. Substrates can contain the classical docking motif, or the FXFP sequence, or both [19]. Interestingly, the $PKC\alpha$ protein, like other MAPK substrates, also contains two putative docking domains such as RGRIYL and KRGIIYRDLKL (Fig. 4C). This provides further evidence that PKC α is a direct substrate of Erk1/2 in the 5-HT1A-R-mediated signaling cascade.

An earlier study by Kobe and coworkers used FRET to analyze and assess association between 5-HT_{1A}-R and 5-HT₇-R [17]. Both of these proteins mainly reside in the plasma membrane. Because of their restricted ability to move perpendicular to the plasma membrane, these two proteins will then be restricted to lateral movement, which may enhance the possibility of intermolecular interactions between these two molecules. This may have been the reason for a high FRET efficiency observed in this study. Even the FRET efficiency of non-specific interactions was relatively high $(\sim 5\%)$. In contrast, P-Erk is distributed in the entire cell, whereas $PKC\alpha$ and $P-PKC\alpha$ remain in the cytosol and the cytoplasmic membranes. Consequently, we observed insignificant non-specific FRET efficiency and 7% FRET efficiency between Erk-CFP and $PKC\alpha$ -YFP in our experiments.

This short communication reports a fundamental discovery that the two pivotal kinases Erk and $PKC\alpha$, which regulate myriad cellular processes, physically interact to regulate signaling receptor-mediated pathways that are central to synaptogenesis in the neonatal hippocampus [4]. Although this study delineates the mechanistic details in *in vitro* studies in a hippocampal neuron-derived cell line, our prior studies had demonstrated the profile of Erk-dependent, but PKC-independent phosphorylation of PKCα in cultured hippocampal slices [4]. Our current experiments employ intra-hippocampal injections of selective agonists and inhibitors of signaling intermediates to delineate these pathways *in vivo* [24]. These studies will verify the importance of Erk-mediated phosphorylation and stimulation of $PKC\alpha$ in neonatal development of the hippocampal circuitry. Identification of such synaptogenic signaling molecules may yield appropriate therapeutic targets, which can be manipulated in an early attempt to alleviate chances of later-life affective disorders that crucially depend on neonatal synaptogenesis in the hippocampus.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1. Significant physical association occurs between Erk and PKCα molecules in the HN2-5 cells

Erk-CFP and PKC-YFP vector were cotransfected into HN2-5 cells grown on coverslips. After 24 h, the cells were fixed, mounted on slides, and analyzed by laser confocal microscopy using a Leica Confocal Microscope equipped with a FRET software package. Two small regions of interest were photobleached using YFP excitation (514 nm). The images were acquired before and after photobleaching. The florescence of the donor showed an increase in intensity in the post-photobleaching image, whereas the fluorescence of the acceptor underwent a decrease. Mean FRET efficiency determined from six discrete samples used in three discrete experiments was 7.19±0.75%.

Debata et al. Page 10

Figure 2. Transiently expressed wild type PKCαundergoes phosphorylation at T638 in an Erk dependent manner

HN2-5 cells were transfected into wild type pcDNA6-PKCα construct, differentiated by retinoic acid, treated with drugs and lysed. Lysates were subjected to immunoprecipitation with a Myc tag antibody. Immunoprecipitates were analyzed by Western blotting using a P-PKC α /B II (T^{638/641}) antibody (upper panel) and then a Myc antibody (lower panel). *p < 0.05 with respect to all other sets except GFX + D. There was no significant difference between D and GFX + D.

Debata et al. Page 11

Transfected (empty vector or T^{497} -mutant PKC α) and differentiated HN2-5 cells were treated with $1 \mu \overline{M} D$ (1 h), in the absence and presence of $1 \mu \overline{M} WAY$ or $10 \mu \overline{M} U0126$ (U). The Myc-tagged (T497**→**A) PKCα was immunoprecipitated with a Myc Ab. Western blotting analysis of the immunoprecipitate using a $P-(T^{638/641})-PKC\alpha/BI$ antibody revealed phosphorylation of PKC α at T⁶³⁸, which was abrogated by WAY and U0126 (U). $\frac{1}{\gamma}$ > 0.05 (versus all other sets).

Debata et al. Page 12

Figure 4. In vitro analysis of Erk-catalyzed phosphorylation of PKCα

(a) The scheme of the experiment, showing mutant Erk mediated phosphorylation of the PKC α oligopeptide harboring the turn motif and the T^{638} residue. **(b)** Transfected and differentiated HN2-5 cells were treated with 1 μM D (30 min), lysed, and the activated mutant Erk was immunoprecipitated (IP) using a Flag antibody (Ab) (upper panel). The IP pellet was used in a kinase assay (A) using cyclopentyl $[\gamma^{32}P]$ ATP and the PKC α peptide. The radiolabeled peptide moved faster than the free cyclopentyl $[\gamma^{32}P]$ ATP (smear) on an HPTLC plate. **(c)** PKCα harbors a putative phosphorylation site (PVLTP) and two docking site (KRGRIYL) and (KRGIIYRDLKL) for Erk1/2.