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Journal of Chromatography B, 789 (2003) 323–335

**IOURNAL OF CHROMATOGRAPHY B** 

www.elsevier.com/locate/chromb

## A pplication of multiplexed capillary electrophoresis with laser-induced fluorescence (MCE–LIF) detection for the rapid measurement of endogenous extracellular signal-regulated protein kinase (ERK) levels in cell extracts

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Received 8 November 2002; received in revised form 20 January 2003; accepted 20 January 2003

## **Abstract**

Multiplexed (96-lane) capillary electrophoresis with laser-induced fluorescence (MCE–LIF) detection was used for the rapid analysis of extracellular signal-regulated protein kinase (ERK) levels from in vitro cell extracts. The levels of ERK enzyme in cell extracts were determined by monitoring the conversion of a fluorescent-labeled peptide substrate to a phosphorylated fluorescent-labeled peptide product using MCE–LIF. The incorporation of a fluorescent internal standard was found to improve the precision of the analysis. The enzyme assay conditions including substrate concentration, reaction time and enzyme linear range were rapidly optimized using the MCE–LIF approach for both direct and immunoprecipitationbased ERK assays. The levels of ERK from in vitro cell extracts stimulated with angiopoietin 1 (Ang1\*) were determined using the MCE–LIF approach. The advantages of MCE–LIF for developing and applying enzyme assays, as well as the figures of merit for the direct and immunoprecipitation ERK assays, are discussed. 2003 Elsevier Science B.V. All rights reserved.

## *Keywords*: Multiplexed capillary electrophoresis; Protein kinase; Enzymes

super family of serine/threonine kinases that play a  $MAPK$  subfamilies  $[1-3]$ . One subfamily, the exkey role in intracellular signal transduction in all tracellular signal-regulated protein kinases (ERKs)

**1. Introduction** tyrosine kinases and G protein-coupled receptors by a variety of extracellular signals, such as growth Mitogen-activated protein kinases (MAPK) are a factors, cytokines and stress, activate the various eukaryotic cells [1–3]. Activation of receptor are activated by a variety of growth factors including vascular endothelial growth factor (VEGF) and <sup>\*</sup>Corresponding author. Tel.: +1-513-622-2149; fax: +1-512- angiopoietins (Ang) [4,5]. Two isoforms, ERK1 (also 622-1196. known as p44 MAPK) and ERK2 (also known as *E*-*mail address*: [wehmeyer.kr@pg.com](mailto:wehmeyer.kr@pg.com) (K.R. Wehmeyer). p42 MAPK), are activated via dual phosphorylation

<sup>1570-0232/03/\$ –</sup> see front matter  $\circ$  2003 Elsevier Science B.V. All rights reserved. doi:10.1016/S1570-0232(03)00102-8

catalytic activation loop by upstream kinases [2,3]. [19]. Two immunoassays were developed for The activated ERKs subsequently phosphorylate a tyrosine kinases that employed fluorescence polarizavariety of intracellular proteins including cytosolic tion for detection [20,21]. 70/90-kDa ribosomal S6 kinase, c-Raf, EGF re- Capillary electrophoresis (CE) can be used to ceptor, cytosolic phospolipase A2, myelin basic assay active kinase enzymes since phosphorylated protein and transcription factors c-*myc*, c-*jun* and and nonphosphorylated peptides are easily separable c-*fos* [6–9]. Modulation of transcription factors by based on differences in charge-to-size ratio imparted the ERK pathway regulates expression of genes by the addition of the phosphate group. CE provides essential for cellular processes such as growth, inherent advantages such as reduced sample condifferentiation, and proliferation. Abnormalities in sumption, separation of background interferents and signal transduction via the ERK pathway may lead to on-line detection. Several CE applications have been diseases and proliferative disorders, most notably developed for assaying a variety of protein kinases cancer [10]. Elucidating the role of the ERK path- [22–28]. Conventional CE and microchip-based CE way, as well as other MAPK pathways, in cellular assays for measuring protein kinase A levels using response and disease requires methodology for moni- fluorescent-labeled peptide substrates have been retoring cellular levels of the active kinases. ported [22,23]. Another microchip-based assay was

active MAPK levels are radioenzymatic and Western single run [24]. CE-based assays were developed assays. In the radioenzymatic method, the amount of using fluorescent-labeled substrates for calcium/calactive MAPK is determined by measuring the trans-<br>fer of a radioactive  $\gamma$ -phosphate ( $^{33}P$  or  $^{32}P$ ) group kinase C [27]. Additionally, an assay based on the from adenosine-5'-triphosphate (ATP) to a protein or use of CE with UV detection for GST-tyrosine peptide target [11–17]. Myelin basic protein (MBP), kinase [28] has been reported. All of these CE-based or a synthetic peptide representing the consensus approaches were performed serially using a single sequence of MBP, is most commonly used as a capillary. substrate for assaying the ERK isoforms. Quantita- In the early 1990s, parallel or multiplexed capiltion of the phosphorylated peptide product via lary electrophoresis (MCE) was introduced by the scintillation counting or phosphoimaging is done Mathies [29,30], Yeung [31–33] and Kambara after separation of the product from excess radio- [34,35] groups. The use of multiplexed capillary labeled ATP using phosphocellulose P81 paper or arrays offer the potential of increasing the sample gel electrophoresis. The radioenzymatic-based ap- throughput without a tremendous increase in instruproach suffers from drawbacks such as environmen- mentation complexity or cost. Initial efforts with tal concerns with radioisotope use, multiple washing multiplexed arrays focused on capillary gel electrosteps, and off-line detection. In the Western ap- phoresis (CGE) for DNA sequencing due to the proach, anti-phospho-MAPK antibodies are used to interest in the Human Genome Project [29–46]. A probe the levels of active MAPKs either directly in few reports have also addressed the development, the gel or after transfer of the MAPKs to nitro- characterization and application of MCE for enzyme cellulose films. The Western approach is limited by analysis, combinatorial screening of catalysis rerelatively long incubation times, the need for multi- action conditions, the analysis of small organic ple antibodies, multiple wash steps and quantitative molecules and peptide mapping [47–52]. We report accuracy. on the application of MCE with laser-induced fluo-

recently to overcome the shortcomings of conven- precise and rapid determination of active ERK tional assay techniques. An enzyme-linked immuno- enzyme levels. Both direct and immunoprecipitation sorbent assay was done for several kinases [18] and MCE–LIF-based assays were developed and applied an immunoassay using europium-chelate labeled to the analysis of ERK levels from human umbilical

of threonine and tyrosine residues contained in their antibodies was developed for protein tyrosine kinases

Two widely employed approaches for measuring developed to monitor multiple kinase reactions in a

Alternative assay methods have been pursued rescence (MCE–LIF) detection for the accurate,

vein endothelial cells (HUVEC) following stimula- 2 .2. *Buffers and reagents* tion with angiopoietins 1 (Ang1\*), a ligand for the TIE family of receptor tyrosine kinases. 2.2.1. *Buffers* 

 $MgCl<sub>2</sub>$ , adenosine 5'-triphosphate (ATP), borax (so- 50 ml of the ADB. dium tetraborate decahydrate, 99.0–105.0%), bglycerophosphate and bovine serum albumin (BSA, 2 .2.1.2. *Cell lysis buffer*. Triton lysis buffer con-98.5%) were from Sigma (St. Louis, MO, USA). sisted of 20 m*M* Tris–HCl, pH 8.0, 137 m*M* NaCl, Ethylene-glycol-bis(2-aminoethyl)-*N*,*N*,*N'*,*N'*-tetra- 2 m*M* EDTA, 1 m*M* Na<sub>3</sub>OV<sub>4</sub>, 1 m*M* sodium acetic acid (EGTA) was purchased from Fluck fluoride, 1 m*M* phenylmethylsulfonyl fluoride, 1  $\mu$ g/ acetic acid (EGTA) was purchased from Fluck Chemie (Buchs, Switzerland). Sodium *ortho*-van- ml leupeptin, 1 mg/ml pepstatin A, 10% glycerol and adate (99.98%) and dithiothreitol (DTT) were ob- 1% Triton X-100. tained from Aldrich (Milwaukee, WI, USA). Reagent grade sodium hydroxide (0.1 and 1 N) and HPLC- 2 .2.1.3. *Water*–*BSA* (*H O*–*BSA*) *diluent*. Bovine <sup>2</sup> grade methanol were purchased from J.T. Baker serum albumin (BSA) was dissolved in distilled- (Phillipsburg, NJ, USA). Protease inhibitor cocktail deionized water at the 1 mg/ml level. tablet (complete, EDTA-free) was from Roche Diagnostics (Mannheim, Germany). Peptides labeled on 2 .2.2. *Growth media* the N-terminus with fluorescein isothiocyanate Endothelial growth media (EGM, Clonetics, San (FITC) included APRTPGGRR-COOH, APRT- Diego, CA, USA) consisted of 10 <sup>m</sup>g/ml human <sup>2</sup><sup>2</sup> <sup>2</sup><sup>2</sup> (PO )PGGRR-COOH and FITC-LRRAS(PO )- recombinant epidermal growth factor, 1.0 mg/ml 3 3 USA). Active EKR2 (100  $\mu$ g/ml) was from Upstate amphotericin B, 3 mg/ml bovine brain extract, 2% Biotechnology (Lake Placid, NY, USA). Polyclonal fetal bovine serum (Clonetics) and 1% antibiotic/ phospho-p44/p42 MAP kinase antibody (p-Ab) di- antimycotic from Gibco (Grand Island, NY, USA). rected against the phosphorylated forms of ERK1 and ERK2 was obtained from Cell Signaling Technology (Beverly, MA, USA). Distilled-deionized 2.2.3. *Combined ATP/MgCl<sub>2</sub>* stock water (dd-H<sub>2</sub>O) was from a Milli-Q system (Milli-<br>pore, Bedford, MA, USA). Magnetic Porous Glass stock solution (ATP/MgCl<sub>2</sub>) was prepared by dispore, Bedford, MA, USA). Magnetic Porous Glass (MPG) beads coated with protein A and a 3-in-1 magnetic particle separator (MPS) were obtained ADB. from CPG (Lincoln Park, NJ, USA). Ang1\* was a generous gift from Regeneron Pharmaceuticals (Tarrytown, NY). Ang1\* is a genetically engineered 2.2.4. Internal standard (I.S.) stock solution<br>variant of angiopoietin-1 generated by replacing the FITC-LRRAS( $PO_3^2$ )LG was dissolved in H<sub>2</sub>O–<br>first 77 residues of ang first 77 residues of angiopoietin-1 with the first  $73$ residues of angiopoietin-2 (Ang-2). In addition, a the I.S. stock solution. I.S. spiking solutions were non-conserved cysteine residue in the Ang-2 se- prepared from this stock solution by appropriate quence is mutated to serine.  $\ddot{\text{d}}$  dilution in H<sub>2</sub>O–BSA diluent.

2 .2.1.1. *Assay dilution buffer* (*ADB*). ADB contained 20 m*M* MOPS, 25 m*M* b-glycerophosphate, 5 m*M* **2. Experimental** EGTA, 1 m*M* sodium *ortho*-vanadate and 1 m*M* DTT and was adjusted to pH 7.2 using 1 N sodium 2 .1. *Chemicals and materials* hydroxide. Following the pH adjustment, BSA was added at the 1 mg/ml level. Immediately before use, 3-[*N*-Morpholino] propanesulfonic acid (MOPS), one protease inhibitor cocktail tablet was added per

hydrocortisone, 50 mg/ml gentamicin, 50  $\mu$ g/ml

solving the appropriate amount of each compound in

for immunoprecipitation experiments were prepared added to each microtube, the contents of each tube following the instructions provided by the vendor to mixed and incubated at  $37^{\circ}$ C. The final concengive a series of 1.6-ml microtubes each containing trations of ERK2 substrate in the incubation mixtures 0.25 mg of MPG protein A beads. were 7.2, 15, 30, 60, 120, 240, 480 and 960 mg/ml

dissolving 38.14 g of borax in 2 l of distilled- 310 and 440 s) and added to individual polywas filtered through a 0.22-µm filter and degassed 230 ng/ml I.S. solution. The dilution step effectively under vacuum for 1 h. stopped the enzyme reaction. Each of the resulting

The stock solution was used to prepare product standard solutions covering a concentration range action time was generated for each substrate confrom 40 to 8000 ng/ml by serial dilution with ADB. centration and linear regression of each plot was An aliquot  $(10 \mu l)$  of each product standard solution used to obtain the initial reaction rate velocity for was added to a series of 1.6-ml polypropylene each substrate concentration. Only data collected microtubes already containing 10  $\mu$ l of a 2.1  $\mu$ g/ml before 10% substrate conversion was used for linear I.S. solution,  $5 \mu$  of a 108  $\mu$ g/ml substrate peptide regression. The initial rates were plotted versus the (FITC-APRTPGGRR) solution and 75  $\mu$ l of the peptide concentration to generate the Micheals–Men- $BSA-H<sub>2</sub>O$  solution. The working ERK2 product ton plot using GraFit Version 4.0 (Erithacus Soft-<br>standards covered a concentration range from 4 to ware, Staines, UK). standards covered a concentration range from 4 to 800 ng/ml. Aliquots (10  $\mu$ l) of each ERK2 working standard solution were placed into 11 consecutive wells of a 96-well plate and analyzed by MCE–LIF 2 .3.3. *Enzyme progression plot* as described below. For each capillary, the peak Aliquots (60  $\mu$ l) of a series ERK2 enzyme stock height for the product peptide was divided by the solutions (10, 100 and 250 ng/ml) in ADB were peak height of the I.S. to generate a peak height ratio added to a series of polypropylene microtubes al- (PHR) for that standard. The average PHR for each ready containing  $180 \mu$  of peptide substrate solution standard was calculated and plotted versus the (333  $\mu$ g/ml) and 60  $\mu$ l of the ATP–MgCl<sub>2</sub> solution. product concentration to yield the product standard The resulting solutions were mixed and incubated at curve.  $37^{\circ}$ C. At given time intervals, a  $10$ -µl aliquot of

An aliquot (60  $\mu$ l) of each substrate solution was substrate concentration was 200  $\mu$ g/ml. Each soluready containing 30  $\mu$ l of the ATP–MgCl, solution described below. The average PHR was determined

2 .2.5. *Preparation of MPG protein a beads* and 30 ml of ADB and mixed. An aliquot (30 ml) of MPG protein A beads (50 mg in 10 ml solution) a 200-ng/ml ERK2 enzyme solution in ADB was while the final ERK2 enzyme concentration was 40 2.2.6. *MCE background electrolyte* ng/ml. During the incubation, aliquots (10 μl) were A 50 m*M* borate buffer (pH 9.3) was prepared by removed from each tube at several time points (140, deioinzed water without adjusting pH. The solution propylene microtubes already containing 90  $\mu$ l of a diluted incubation mixtures was analyzed by MCE– 2 .3. *Methods* LIF in triplet as described below. The average PHR for the product peptide was obtained for each 2.3.1. *ERK2 product standard curve* substrate concentration and was used to calculate the A 0.8-mg/ml stock solution of product peptide concentration of the product peptide by interpolation (FITC-APRT(PO $_3^2$ )PGGRR) was prepared in ADB. from a product standard curve prepared as described 3 The stock solution

The resulting solutions were mixed and incubated at each incubation mixture was removed and pipetted 2 .3.2. *Michaelis*–*Menton curve* into a plastic microtube already containing 90 ml of a ERK2 substrate peptide (FITC-APRTPGGRR) 230 ng/ml I.S. solution to stop the reaction. The solutions were prepared in H<sub>2</sub>O–BSA solution over final concentrations of ERK2 in the incubation a concentration range from 18  $\mu$ g/ml to 2.4 mg/ml. mixtures were 2, 20 and 50 ng/ml and the peptide mixtures were 2, 20 and 50 ng/ml and the peptide added to a series of polypropylene microtubes al- tion was analyzed in triplicate by MCE–LIF as

for the generated peptide product for each ERK2 below. The average PHR for the generated product enzyme concentration and the PHR was plotted was obtained for each enzyme standard and plotted versus incubation time to generate the enzyme versus enzyme concentration to generate a linear progression curves. The regression standard curve. Similarly, an ERK2 en-

standards or cell lysate samples, were added to polypropylem microtubes containing 170  $\mu$ l of ADB<br>and 4 2.3.5.2. Immunoprecipitation ERK2 standard curve.<br>and 45 pl of polyclonal ERK antiboty, vortexted and incurve cov-<br>containing (2.5 mg MPG protection), the solution

## 2 .3.5. *ERK*<sup>2</sup> *enzyme standard curves cell* (*HUVEC*) *studies*

the reaction mixtures ranged from 1.6 to 50 ng/ml. Prior to stimulation, cultured HUVEC were serum-After incubation, the reaction was stopped by adding starved for 2 h by replacing EGM with phosphateanalyzed in triplicate by MCE–LIF as described Ang1\* in PBS–0.2% BSA for 7 min. Cells were

zyme standard curve ranging from 0.25 to 10 ng/ml 2.3.4. *Immunoprecipitation of ERK2* was prepared and the incubation time increased to Aliquots (100 μl) of ERK2 solutions, either 60 min.

# 2 .3.6. *In vitro human umbilical vein endothelial*

HUVECs were obtained as primary cells from 2.3.5.1. *Direct assay ERK2 standard*. Aliquots (15 Clonetics. HUVECs (passage <4) were maintained  $\mu$ l) of ERK2 stock solutions (7.8–250 ng/ml) in in endothelial growth medium (EGM) at 37 °C in a ADB were added to separate polypropylene micro-<br>tubes already containing 45  $\mu$ l of a 333 mg/ml Cells were handled in accordance with manufactur-Cells were handled in accordance with manufacturpeptide substrate solution and 15  $\mu$ l of the ATP– er's recommendations. HUVECs were plated in MgCl<sub>2</sub> solution, vortexed and incubated at 37 °C for tissue culture dishes at  $1 \times 10^6$  cells per 100-mm dish 15 min. The f and grown to 90% confluency in EGM (Clonetics). 10  $\mu$ l of the enzyme reaction mixture to 90  $\mu$ l of a buffered saline to reduce basal ERK activity. ERK 230 ng/ml I.S. solution. The resulting solutions were was activated by adding various concentrations of lysed directly in the plates by the addition of 1 ml of a linear regression standard curve prepared by the cell lysis buffer (see above). The plates were then immunoprecipitation approach as described above. incubated at  $4^{\circ}$ C for 20 min with shaking. The supernatants were transferred to a clean poly- 2.3.8. Accuracy and precision of ERK2 analysis in propylene tubes and stored frozen at  $-80$  °C until *the cell lysate* analysis.

added into separate polypropylene microtubes al- added into polypropylene microtubes already conready containing 45  $\mu$ l of 333  $\mu$ g/ml ERK2 sub- taining 7.5  $\mu$ l of ADB, 45  $\mu$ l of a 333  $\mu$ g/ml peptide strate and 15  $\mu$ l of ATP–MgCl<sub>2</sub> solution and substrate solution and 15  $\mu$ l of ATP–MgCl<sub>2</sub> solution incubated at 37 °C for 15 min. A 10- $\mu$ l aliquot of and mixed by vortexing. The resulting solutions were incubated at 37 °C for 15 min. A 10- $\mu$ l aliquot of each incubation mixture was removed and added to incubated at  $37^{\circ}$ C for 15 min. Following incubation, polypropylene microtubes already containing 90  $\mu$ l 10  $\mu$ l of each reaction mixture was added into of a 230 ng/ml I.S. solution and analyzed in polypropylene microtubes already containing 90  $\mu$ l triplicate by MCE–LIF as described below. The of a 230 ng/ml I.S. solution, vortexed and analyzed average PHR for the product obtained for each in triplicate by CE–LIF as described below. The sample was used to determine the concentration of average PHR of the product for each spiked sample ERK2 in the cell-lysate sample by interpolation from was used to calculate the concentration of the ERK2 a linear regression standard curve prepared by the enzyme by interpolation from a linear regression direct assay approach described previously. ERK2 standard curve prepared by the direct assay

Aliquots (100  $\mu$ ) of the cell lysate samples were dividing the ERK2 concentration found in the samadded to microtubes containing 170  $\mu$ l of ADB. The ples by the expected ERK2 concentration and multiresulting solutions were immunoprecipitated with plying by 100. ERK2 polyclonal antibody and MPG protein A beads as described previously. The isolated beads for each 2 .3.8.2. *Immunoprecipitation spiking studies*. sample were resuspended in 15-µl aliquots of ADB Aliquots (100 µl) of spiked cell lysates prepared and added to separate polypropylene microtubes above were studied by immunoprecipitation as dealready containing 45  $\mu$ l of 333  $\mu$ g/ml ERK2 scribed previously. The isolated beads containing the substrate and 15  $\mu$ l of ATP–MgCl<sub>2</sub> solution. The bound ERK2 enzyme were suspended in 15- $\mu$ l resulting mixtures were vortexed and incubated at aliquots of ADB and added to separate poly-37 °C for 15 min with vortexing every 5 min to propylene microtubes already containing 45  $\mu$ l of resuspend the beads. Following incubation, the beads 333  $\mu$ g/ml ERK2 substrate and 15  $\mu$ l of ATP– were removed using the MPS device and aliquots  $MgCl_2$  solution. The resulting mixtures were vortex- (10  $\mu$ l) of each incubation mixture were added to ed and incubated at 37 °C for 15 min with vortexing polypropylene microtubes already containing  $90 \mu l$  every 5 min to suspend the beads. Then aliquots (10) of a 230 ng/ml I.S. solution and analyzed in  $\mu$ ) of each mixture were added to polypropylene triplicate by MCE–LIF as described below. The microtubes already containing 90  $\mu$ l of a 230 ng/ml average PHR for the product obtained for each I.S. solution and analyzed in triplicate by MCE–LIF sample was used to determine the concentration of as described below. The average PHR for the ERK2 in the cell-lysate sample by interpolation from product of each sample was used to determine the

2 .3.8.1. *Direct assay spiking studies*. Aliquots (180  $\mu$ l) of unstimulated HUVEC cell lysate were spiked 2 .3.7. *Analysis of cell lysate samples by MCE*–*LIF* with 20 ml of either ADB or various ERK2 standard stock solutions (0.125, 1.25 and 5  $\mu$ g/ml) to give 2 .3.7.1. *Direct analysis of cell lysates*. Aliquots (15 lysates with 0, 12.5, 125 and 500 ng/ml ERK2  $\mu$ l) of the HUVEC cell lysates described above were enzyme. Aliquots (7.5  $\mu$ l) of each spiked lysate were approach as described previously. Recoveries of 2 .3.7.2. *Immunoprecipitation analysis of cell lysates*. ERK2 from the spiked lysates were calculated by

> aliquots of ADB and added to separate polyed and incubated at 37 °C for 15 min with vortexing

concentration of ERK2 in the spiked cell-lysate sample by interpolation from a linear regression ERK2 standard curve constructed using the immunoprecipitation approach described previously. Recoveries of ERK2 from the spiked lysates were calculated by dividing the ERK2 concentration found in each sample by the expected ERK2 concentration and multiplying by 100.

## 2 .4. *MCE conditions*

The MCE–LIF analysis was done with an HTS9610 electrophoresis system from SpectruMedix (State College, PA, USA). The system was equipped with 96 uncoated fused-silica capillaries (52 cm $\times$ 50 mm I.D., 35 cm effective length), a 25-A three-phase air-cooled all-line argon ion laser with output power (300 mV) for the wavelength from 458 to 514 nm (457.9 nm, 4%; 476.5 nm, 12%; 488 nm, 22%; 496.5 nm, 13%; 501.7 nm, 7%; and 514.5 nm, 42%) and a CCD camera. A standard 96-well plate format was used at the inlet end with 96 individual inlet electrodes. At the detection window, capillaries were arranged in a parallel format. The CCD camera was used for signal detection with an f4 stop, bin 3 and 570 ms exposure time. Data was collected using Checkmate software and processed with SpectroPro1.80 software (SepectruMedix).

For all studies, the capillaries were conditioned prior to each run by rinsing with 0.1 N NaOH, Fig. 1. CE separation of ERK2 product standards: (A) a charge-<br>distilled-deionized water and running buffer for 2 coupled device image collected for the analysis of a typical E distilled-deionized water and running buffer for 2<br>min each at 5 ml/min flow-rate. All electrophoresis<br>was performed at ambient temperature. Samples<br>were injected by pressure  $(-0.35 \text{ p.s.i})$ . for 10 s) and<br>were injected b separation was conducted at 2.8 kV for 45 min. The peptide standard. running buffer was 50 m*M* sodium tetraborate (pH 9.3). consecutive capillaries with the product concentra-

1A. The standards were injected sequentially into 11 constant. Additionally, impurities from the substrate



Time, s

a representative electropherogram for the 160 ng/ml product

tion increasing from the top to the bottom of the array. A water sample was injected into a single **3. Results and discussion** capillary between each set of product concentrations. The dots within a given capillary correspond to 3 .1. *Separation of substrate*, *product and internal* individual components and the intensity of the dots is *standard by MCE–LIF* proportional to the concentration of the components. The intensity of the ERK2 product can be seen to A typical CCD image of the 96-capillary array is increase as you move down the array, while the shown for the ERK2 product standard curve in Fig. concentration of the substrate and the I.S. remain

streaking observed in some of the capillaries is due instrumentation has been reported previously [52]. A to a slight tailing of the analytes being accentuated plot of the PHR for the product peptide was linear by the high contrast used to generate the picture. over two orders of magnitude with a correlation High contrast was employed to allow the lowest coefficient of 0.999 (data not shown). standards to be easily visualized. A representative electropherogram constructed from the CCD image 3 .2. *Michaelis*–*Menton plot and enzyme* is shown in Fig. 1B. *progression curve for ERK*<sup>2</sup> *enzyme*

The migration time of each component can be seen to be relatively constant across the array and the In developing the ERK2 enzyme assay conditions intensity for the components within a product stan- it was important to determine the optimal substrate dard concentration are relatively constant. Slight concentration for the enzyme reaction. Ideally, the variations in the migration time are expected due to substrate concentration should be on the flat part of slight differences in the capillary surface chemistry the Michaelis–Menton plot to avoid dramatic between capillaries and the potential for thermal changes in the reaction rate due to small errors in the gradients across the capillaries. Similarly, slight substrate concentration and to provide for the maxivariations in the intensities within a standard con- mum reaction rate. A Michaelis–Menton plot was centration are expected due to variation in camera constructed by holding the ERK2 enzyme level digitization error, minor differences in capillary constant and varying the substrate concentration diameters, variations in laser intensity across the from 7 to 960  $\mu$ g/ml. The reaction rate was decapillary and differences in the vacuum injection termined for each concentration of substrate and the pressure experienced by individual capillaries during resulting reaction rates were plotted versus the the injection. The incorporation of the I.S. allows for corresponding substrate concentrations to obtain the correction of the non-uniformities introduced by the Michaelis–Menton plot (Fig. 2). A  $K_{\text{max}}$  value of capillary array. The migration time and peak height for the product in a given capillary are corrected by dividing these values by the corresponding migration time and peak height obtained for the I.S. to give the corrected migration time and the PHR. The percent relative standard deviation (%RSD) obtained withinday for the migration time, corrected migration time, peak height and the PHR are shown in Table 1 for several ERK2 product standards and similar results were obtained across days (data not shown). The %RSD values were significantly improved by the incorporation of the I.S. correction. The long-term

Table 1 Injection precision of ERK2 product standards

[Product] standard] (ng/ml)	Percent relative standard deviation $(n=6)$			
	Migration time(s)	Corrected migration time(s)	Peak height	Peak height ratio
4	1.19	0.26	17	8.8
40	0.33	0.07	10	3.0
400	3.79	0.91	11	5.1

and I.S. can also be seen in the CCD image. The performance and reproducibility of the MCE–LIF



Fig. 2. Michaelis–Menten plot. Plots of reaction rates versus FITC-labeled peptide substrate concentration. Incubations performed using 40 ng/ml ERK2 with various FITC-labeled peptide substrate concentrations (7.5–960  $\mu$ g/ml) at 37 °C for 15 min (*n* = 6 for each substrate concent

 $247\pm17$   $\mu$ g/ml (168 $\pm$ 2  $\mu$ *M*) was found for the For routine analysis, the incubation was performed substrate. However, a substrate concentration of  $200$  for 15 min at 37 °C.  $\mu$ g/ml was chosen as a compromise between rapid enzyme reaction rate without excessive use of sub- 3 .3. *Direct ERK*<sup>2</sup> *enzyme standard curve* strate during the enzyme assays. The high throughput capability of the MCE–LIF system allowed the data In the direct assay, the standard curve for ERK2 the unlabeled substrate, however, for our quantitative

enzyme concentrations for at least 20 min. At longer the standards ranged from 3 to 15%. incubation points there was a deviation from linearity at all ERK2 concentrations, however, the 2 ng/ml 3 .4. *Immunoprecipitation ERK*<sup>2</sup> *standard curve* standard provided linear results for at least 60 min.

## **ERK2 Enzyme Progression Curve**



taken at various time points. Peak height represents average control samples maintained a constant level of ERK2  $(n=4)$  for replicate injections. over both incubation periods indicating the enzyme

for the Michaelis–Menton plot to be generated in a was constructed in ADB by adding the peptide single run. It should be noted that the introduction of substrate to a series of tubes containing various a fluorescent group into a substrate peptide sequence ERK2 concentrations, incubating at  $37^{\circ}$ C for 15 can change the  $K_{\text{max}}$  and reaction kinetics relative to min, diluting an aliquot of the incubated standard the unlabeled substrate, however, for our quantitative solution with the I.S. solution and analyzing the purposes this was not of concern. amount of product formed by multiplexed MCE– The time period over which the enzyme incubation LIF. The standard curve for the direct assay was yields a linear increase in product concentration was found to be linear over ERK2 concentrations from 4 determined in a single run at three ERK2 enzyme to 250 ng/ml using a 15-min incubation period (Fig. levels  $(2, 25 \text{ and } 50 \text{ ng/ml})$  using a fixed substrate 4A). The linear range could be lowered to 0.2–10 concentration of 200  $\mu$ g/ml (Fig. 3). The generation ng/ml using a 60-min incubation interval (Fig. 4B). of product was a linear function of time at all In general, the RSD for replicate injections  $(n=3)$  of

Direct assay of ERK2 levels in cell lysate has the advantage of convenience and speed. However, the HUVEC cell lysate may contain additional kinases that could utilize the FITC-labeled peptides as substrates. To evaluate this possibility an immunoprecipitation assay was developed to isolate ERK2 from the cell lysates prior to analysis. Lysates from HUVEC cells stimulated with Ang1\* were treated with a polyclonal antibody selective for the active form of ERK2. The resulting ERK2-antibody complex was isolated from the solution using MPG protein A beads. The efficiency of the isolation step was evaluated by performing two consecutive immunoprecipitation reactions on the same cell lysate. Additionally, control and nonspecific binding samples were obtained by doing no treatment to the lysate and by treating the lysate with protein A beads in the absence of antibody, respectively. The amount of ERK2 remaining in the cell lysate samples and attached to the beads after each treatment was determined by incubation with the peptide substrate followed by MCE–LIF analysis. The PHR obtained Fig. 3. ERK2 product progression curve. Various ERK2 enzyme<br>concentrations (2, 20 and 50 ng/ml) were incubated with FITC-<br>labeled peptide substrate (200  $\mu$ g/ml) at 37 °C and samples were<br>samples are shown in Fig. 5A,B,



(A) ERK2 standards with 200  $\mu$ g/ml FITC-labeled peptide 112% and 106 to 120% for the direct and immuno-<br>substrate at 37 °C for 15 min and (B) ERK2 standards incubated precipitation assays respectively. The %RSD for substrate at 37 °C for 15 min and (B) ERK2 standards incubated<br>with 200 µg/ml peptide substrate at 37 °C for 60 min. An<br>FITC-labeled peptide internal standard was spiked into each<br>standard prior to analysis. All standards  $(n=4)$  and the average peak height was plotted versus ERK2 tion approaches demonstrate the direct approach can

slight decrease in the ERK2 enzyme level in the cell HUVECs. lysate (Fig. 5A) and the increase in ERK2 levels on the beads (Fig. 5B) following treatment of the lysate 3 .6. *Analysis of ERK*<sup>2</sup> *levels in HUVECs* with only beads indicates a small amount of non- *stimulated with Ang*1\* specific binding. The level of ERK2 was markedly decreased in the cell lysate following the immuno- HUVECs stimulated with various concentrations precipitation treatments (Fig. 5A) and this is re- of Ang1\* were analyzed for ERK2 levels by the flected by an increased amount of ERK2 found on direct assay approach. Prior to conducting the direct the beads (Fig. 5B). assay, the FITC-labeled substrate and product were

was prepared in ADB using the polyclonal antibody mixture that contained protease and phosphatase and the MPG protein A beads as described above. inhibitors. An electropherogram obtained from The ERK2 isolated on the beads was determined by Ang1\*-stimulated HUVEC cells is shown in Fig. 7B. incubating the beads with the peptide substrate and In addition to the substrate, product and I.S. peaks a analyzing by MCE–LIF. The immunoprecipitation number of additional peaks are present due to the

standard curve was linear over ERK2 concentrations from 1 to 318 ng/ml using a 15-min incubation period (Fig. 6). The RSD for replicate injections  $(n=3)$  of the standards ranged, in general, from 3 to 15%. As stated previously, the direct assay has the advantage of simplicity and speed while the immunoprecipitation assay offers a higher degree of selectivity by first isolating the ERK2 enzyme from potentially interfering kinases.

## 3 .5. *Accuracy and precision of ERK*<sup>2</sup> *assay*

The accuracy and precision of measuring ERK2 in cell lysate was evaluated using the direct and immunoprecipitation assay approaches. Cell lysate from unstimulated HUVEC cells was spiked with various amounts of ERK2 (0, 6.25, 62.5 and 250 ng ERK2/ ml lysate) and the spiked lysates were analyzed by the direct and the immunoprecipitation approaches. The concentration of ERK2 in the spiked lysates was determined by interpolation from standard curves constructed by the direct and immunoprecipitation approaches. The percent recovery, corrected for the basal levels, for the spiked cell lysates are shown in Fig. 4. ERK2 direct standard curves obtained by incubation of: Table 2. The ERK2 recovery ranged from 103 to concentration. be employed for assaying ERK2 levels in HUVEC cells. Therefore, the direct approach was applied to was stable under the study conditions (Fig. 5A). The measuring the levels of ERK2 in stimulated

An ERK2 immunoprecipitation standard curve shown to be stable in the cell lysate incubation



## **ERK2 Immunoprecipitate Analysis**

B



Fig. 5. Immunoprecipitation of ERK2. Plot of peak height ratio obtained for the incubation of FITC-labeled peptide substrate with (A) cell lysates before and after immunoprecipitation with a polyclonal antibody against active ERK2 and (B) GMP beads obtained following immunoprecipitation with polyclonal antibody against active ERK2. Incubations were performed using with 200 µg/ml FITC-labeled peptide substrate at 37 °C for 15 min.



Fig. 6. ERK2 immunoprecipitation-based standard curve obtained by incubating GMP beads obtained from the immunoprecipitation of ERK2 standards with a polyclonal antibody against active ERK2 with 200  $\mu$ g/ml FITC-labeled peptide substrate for 15 min at  $37^{\circ}$ C.

presence of trace impurities in the substrate. A plot<br>of the PHR for the product generated in the stimu-<br>(HUVEC) with angiopoietin (Ang 1<sup>\*</sup>): (A) plot of the average increased, as indicated by the increasing PHR values, with  $1000 \text{ ng/ml Ang1*}$ . with increasing levels of Ang1\* used for the stimulation. The MCE–LIF methodology allowed the analysis of 40 samples, in duplicate, to be completed in less than 2 h. Typically, the capillary array was **4. Conclusion** usable for  $\sim$ 100 runs before performance deterioration was noted, usually peak tailing and decreased The determination of active kinase enzymes levels sensitivity were the major symptoms. from in vitro and in vivo settings can provide

Spiked [ERK2]	% Recovery (% RSD)			
(ng/ml)	<b>Direct</b> assay	Immuno- precipitation assay		
250 62.5 6.25	103(3.5) 102(9.0) 112 (20)	106(7.5) 120(22) 112(10)		



lated HUVEC samples versus Ang1\* concentration peak height ratio versus Ang1\* concentration,  $n=3$  replicate is shown in Fig. 7A. The amount of active ERK2 HUVEC samples for Ang 1\* concentration and (B) a representa-<br>increased as indicated by the increasing PHP values tive electropherogram obtained for a HUVEC sample stimulated

important information for determining intervention points in disease processes, for monitoring the effect Table 2 contract the mechanism of t Accuracy and precision of ERK2 analysis nism of drug action. We have shown MCE–LIF can provide a rapid approach for monitoring the levels of ERK2 in HUVEC lysates by employing a fluorescently labeled peptide substrate. The methodology was shown to provide accurate and precise quantitation of ERK2 levels using both direct and immunoprecipitation assay formats. The direct assay format has the advantage of simplicity and speed and allows for the analysis of 40 samples in duplicate in less Iy applicable for the determination of variety of  $\begin{array}{c} \text{process } 22 \text{ (2001) } 4000. \\ \text{125} \text{ S. Jeong, T.T. Nikiforov, BioTechniques } 27 \text{ (1999) } 1232. \\ \text{126} \text{ B. Liu, L. Zhang, Y. Lu, J. Chromatogr. A } 918 \text{ (2001) } 401. \end{array}$ 

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