

The cloning and developmental regulation of murine diacylglycerol kinase ζ

Li Ding^a, Thomas M. McIntyre^b, Guy A. Zimmerman^b, Stephen M. Prescott^{a,*}

^aHuntsman Cancer Institute, Eccles Program in Human Molecular Biology and Genetics, Salt Lake City, UT, USA

^bNora Eccles Treadwell Cardiovascular Research and Training Institute, Suite 4220, Building 533, University of Utah, Salt Lake City, UT 84112, USA

Received 16 April 1998

Abstract Diacylglycerol kinases (DGKs) regulate the key signaling intermediates diacylglycerol (DAG) and phosphatidic acid (PA). We isolated cDNA clones of mouse diacylglycerol kinase ζ (mDGK ζ) and found that it shares 88% identity at the nucleic acid level and 95.5% identity at the amino acid level with human DGK ζ (hDGK ζ). Murine DGK ζ protein rose gradually during embryonic development, and was abundant in newborn and adult brains. By RNA whole-mount in situ hybridization, mDGK ζ was shown to be expressed in spinal ganglia and limb buds at low level in E11.5 embryos and at higher level in E12.5 embryos. In E13.5 embryos, DGK ζ mRNA was highly expressed in vibrissa follicles, in spinal ganglia, and in the interdigital regions of the developing limbs. Northern blotting showed that DGK ζ expression was limited to specific anatomical regions of the brain. Thus, the expression of DGK ζ is regulated temporally and spatially during mammalian development and correlates with the development of sensory neurons and regions undergoing apoptosis.

© 1998 Federation of European Biochemical Societies.

Key words: Diacylglycerol kinase; Embryonic development; Distribution

1. Introduction

Diacylglycerol (DAG) is a key intermediate in the synthesis of complex lipids and serves as an important second messenger. DAG exerts its signaling function by activating protein kinase C (PKC), which in turn regulates many cellular responses including growth, differentiation and apoptosis [1]. In addition, the level of DAG is essential for cellular events such as protein export from the Golgi complex in yeast [2]. The signals from DAG are terminated by its conversion to phosphatidic acid (PA), a reaction that is catalyzed by diacylglycerol kinase(s) (DGK). The role of DGK in regulating DAG levels was established by the observation that overexpression of DGK α decreased the elevated DAG level in *ras*-transformed fibroblasts [3]. In ending the signal(s) from DAG, the DGK reaction generates PA as the product which has been implicated in the regulation of DNA synthesis, in the induction of *c-myc*, *c-fos*, platelet-derived growth factor, in cAMP formation [4–6], and in modulating the activity of *n*-chimaerin [7] and NF1 [8]. Therefore, the DGK reaction occupies an interesting niche – it removes one lipid messenger but creates another. Besides its effect on DAG signaling, the DGK reaction is the first step in the recycling of phosphatidylinositol species following their hydrolysis.

Disruption of this pathway can have severe consequences as demonstrated by the *Drosophila rdgA* (DGK2) mutant in which photoreceptor cells degenerate within a week after eclosion [9].

The known eight mammalian DGKs can be divided into five structural subgroups and differ in their expression patterns, activators, and substrate specificity. Type I includes DGK α [10,11], β [12], γ [13] and is defined by the presence of E-F hand motifs at the N-termini, which bind Ca²⁺ with different affinities [14]. DGK α was originally found in lymphocytes and oligodendrocytes, while DGK β was cloned from brain and was found to be expressed in the olfactory tubercles, nucleus accumbens, caudate and putamen [12]. DGK γ , cloned from a HepG2 library, was found to be enriched in retina [13]. DGK δ [15] and DGK η [16] are type II DGKs, which have a pleckstrin domain (PH domain) at their N-termini. The PH domain has been found in a number of proteins involved in signal transduction and can serve as a site of protein-protein and protein-phospholipid interactions [17–19]. DGK δ was cloned from testis and HepG2 cells, and is primarily expressed in skeletal muscle and testis [15]. The expression of DGK η is found in a broad range of tissues and is regulated by glucocorticoids [16]. The third type of DGK, DGK ϵ [20], has the simplest structure and shows substrate specificity for DAG with an arachidonoyl residue at the *sn*-2 position. DGK ϵ mRNA has been found in retina, brain, testis, heart, spleen and lung [21]. Type IV is typified by DGK ζ [22] which has two zinc fingers, four ankyrin repeats at its C-terminus and a unique region homologous to MARCKS phosphorylation site domain. DGK ζ is highly expressed in the brain and skeletal muscle. RNA in situ hybridization by using rat brain section showed that DGK ζ is expressed in cerebral cortex, hippocampus and olfactory bulb [23]. The *Drosophila* DGK2, *rdgA* gene also belongs to this type IV group and is expressed almost exclusively in the retina [9]. Proteins containing ankyrin repeats are involved in a variety of cellular processes such as gene regulation [24–26] and cell cycle control [27]. The type V DGK, DGK θ , is mostly expressed in brain and small intestine. It contains three cysteine-rich repeats – rather than two – and a *ras*-binding domain [28].

Here, we report the cloning and characterization of murine DGK ζ . We also studied the developmental expression of mDGK ζ , and detected the protein as early as on day 10.5 of embryogenesis with a marked change at later stages of development. The spatial distribution of the mDGK ζ message was determined by RNA whole-mount in situ hybridization in mouse embryos at various stages of development. We found that DGK ζ expression is temporally and spatially restricted, and is highly expressed in tissues involved in sensory function.

*Corresponding author. Fax: (1) (801) 585-6345.
E-mail: steve.prescott@genetics.utah.edu

2. Materials and methods

2.1. Isolation and characterization of murine cDNAs

Hybridizations were performed in $5\times$ SSPE, $5\times$ Denhardt's, 0.2% SDS, and 0.1% $\text{Na}_2\text{P}_4\text{O}_7$ at 52.5°C for low stringency and 65°C for normal stringency. Primers for genomic PCR screening are: LD121-2, 5' CAG CAT CTT AGC TAC ACA GTG CTG 3' and LD121-3, 5' ATC ACC AAG TCG GGC CTC CAG 3'.

2.2. Western immunoblots for murine DGK ζ protein expression

Mouse embryos of different ages, newborn mouse brain and adult mouse brain were homogenized in 50 mM Tris-HCl pH 8.0, 2 $\mu\text{g}/\text{ml}$ pepstatin A, 20 $\mu\text{g}/\text{ml}$ leupeptin, 40 $\mu\text{g}/\text{ml}$ TLCK, 20 $\mu\text{g}/\text{ml}$ aprotinin, 20 μM BH_4 , 3 mM DTT, 10 mM CHAPS and 1 mM PMSF. All homogenates were frozen and stored at -70°C until assayed. Samples with ~ 100 μg protein were loaded on a 7.5% SDS-PAGE gel and Western blotting was performed as previously described [22]. The protein concentration of each sample was determined by standard BCA protein assay. The β -actin antibody was used as a control to ensure that an equal amount of protein was loaded in each lane. The Western blotting protocol with ECL detection is capable of detecting as little as 1 pg protein (Amersham). For the peptide competition experiment, the primary antibody was incubated with 4 volumes of peptide (0.1 mg/ml) at 4°C for 2 h prior to incubation with the membrane.

2.3. RNA whole-mount in situ hybridization

The RNA whole-mount in situ hybridization was performed as previously described [29]. The mouse strain used was Swiss Webster. The day that a copulation plug was observed was considered embryonic day 0.5 (E0.5). Embryos were collected from timed pregnant females. Embryos were dissected free of extraembryonic membranes in phosphate-buffered saline (PBS).

Two mDGK ζ antisense RNA probes were used. The first probe was a 716-bp 3' *Pst*I/*Eco*RI fragment. It was subcloned into pBluescript II and used to make digoxigenin-labeled RNA probe by in vitro transcription (Boehringer Mannheim). The second one was from the catalytic domain region of the mDGK ζ gene. A pair of primers with T7 or T3 promoters attached was used to amplify a 558-bp mDGK ζ cDNA fragment. Amplified PCR products were used as templates to make digoxigenin-labeled RNA probes by using an in vitro transcription kit (Boehringer Mannheim). The sequences of the primers were:

5' CAGAGATGCAATTAACCCCTCACTAAAGGGAGAGTCTCGAGAAGCCAACCCA

T3

GAG 3',

5' CCAAGCTTCTAATACGACTCACTATAGGGAGACCTGCTCACCTGGATCCTC

T7

AG 3'.

The probe concentration was 0.5 $\mu\text{g}/\text{ml}$.

2.4. Northern blotting

Human brain Northern blots I and II were purchased from Clontech. Two filters with 2 μg mRNA from different regions of the human brain were probed with a digoxigenin-labeled 761-bp fragment of hDGK ζ as described previously [22]. The β -actin cDNA probe was used to ensure that an equal amount of mRNA was loaded in each lane. The detection limit for digoxigenin-labeled riboprobe is around 0.1 pg according to manufacturer estimation (Boehringer Mannheim).

3. Results

3.1. Isolation and characterization of murine cDNA and genomic clones

We screened a mouse brain cDNA library with a *Sma*I-*Hind*III probe derived from the hDGK ζ cDNA by low stringency hybridization. The first identified clone (3-2) differed from any known mouse DGKs and showed a high percentage of sequence identity with hDGK ζ , suggesting it was the murine

homolog of hDGK ζ . The translation initiation site was not found either in this clone or others (7-1 and 8-1) isolated from another two rounds of screening. A 500-bp mouse genomic fragment was amplified by polymerase chain reaction (PCR) and used to isolate four P1 genomic clones. Splicing patterns were conserved between hDGK ζ [30] and mDGK ζ as demonstrated by reverse transcriptase-PCR. As summarized in Fig. 1A, the composite mDGK ζ sequence was obtained from mouse cDNA clone 3-2, 7-1, 8-1 and the mouse P1 genomic clones. The initiation ATG of the open reading frame was identified as the first ATG sequence following with an in-frame stop codon 9-bp upstream. The first zinc finger of the mDGK ζ protein includes the sequence $\text{HX}_{11}\text{CX}_6\text{CX}_{12}\text{CX}_2\text{CX}_4\text{HX}_2\text{CX}_{10}\text{C}$ and the second includes $\text{HX}_{11}\text{CX}_2\text{CX}_{19}\text{CX}_2\text{CX}_4\text{HX}_4\text{CX}_9\text{C}$. The cysteine and histidine residues of the two zinc fingers are completely conserved between hDGK ζ and mDGK ζ proteins. The MARCKS phosphorylation site is conserved between hDGK ζ and mDGK ζ proteins except for a (m)Arg²⁷⁰ to (h)Lys²⁶⁹ replacement in the mDGK ζ protein. In addition, the catalytic domain of the mDGK ζ protein is similar to all known DGKs. The main feature that distinguishes DGK ζ is the ankyrin repeats, and the deduced mDGK ζ protein, like the human protein, has four such repeats (Fig. 1B). Mouse DGK ζ shares 88% and 95.5% identity with the human sequence at the nucleotide and amino acid level, respectively.

3.2. Immunodetection of the murine DGK ζ protein: the DGK ζ protein accumulates during late embryogenesis and post-natal development

Human DGK ζ and murine DGK ζ proteins have an identical peptide sequences at their C-termini, thus, we were able to use a previously described antibody [22]. This antibody recognized a ~ 120 -kDa protein in homogenates of the mouse brain, and pre-incubation with the peptide antigen block the recognition indicating specific binding (Fig. 2A). We next examined the expression of mDGK ζ during embryonic development by performing Western immunoblots of homogenates from whole embryos and from brains of newborn and adult mice. The mDGK ζ protein was detected in the homogenates of E10.5 embryos, and the protein level increased gradually during development (Fig. 2B). A band (indicated by an ar-

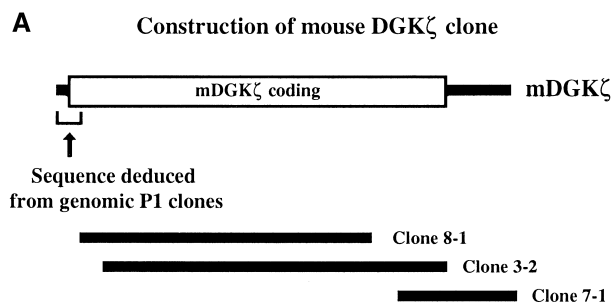


Fig. 1. Sequence of the murine DGK ζ cDNA. A: The overlapping map of representative clones. The cDNA and genomic clones were isolated as described (Section 3). The clones were sequenced in an automated ABI system. B: The nucleic acid sequence and deduced amino acid sequence of mDGK ζ . The zinc fingers are underlined, and conserved cysteine and histidine residues are marked with *. Serine residues within the MARCKS homology region are marked by +. The residues within the ATP binding motif are double underlined. The ankyrin motifs are displayed within boxes.

B

CGGTGCGGAGCCGCGTGTGAGCCCCGGCCGGCCCGGCATGGGCGTCTCCCGCGG 60
 CCCGCCCGCGCGGGCTAGGGCTGATGGAGCCGGGACCCAGCCCGAGGGCCGG 120
 M E P R D P S P E G R

AGCAGCGACTCGGAGTCGGCTCCGCTCGTCCAGCGGCTCCGAGCGGACGCGGGTCCG 180
 S S D S E S A S A S S S G S E R D A G P

GAGCCCGACAAGCGCCGCGGCGCTGACCAAGCGGCGTTCGCCGGGCTCGGCTCTTT 240
 E P D K A P R R L T C F R R F P G L R L F

GGGCACAGGAAGCCATCACCAAGTCGGGCTCCAGCACCTGGCACCCCTCTTCCACG 300
 G H R K A I T K S G L Q H L A P P L P T

CCCCGGGCCCCGTGCGTGAATCTGAGGAGCAGATCCAGAGCACTGTGGACTGGAGTGA 360
 P G A P C G E S E E Q I Q S T V D W S E

TCAGCAGTGTATGGGAGCACATCTGGTTTGAAGCAACGTGTCGGTGACTTCTGCTAT 420
 S A V Y G E H I W F E T N V S G D F C Y

GTCGGGAGCAGCACTGTGTAGCTAAGATGCTGCCGAAGTCAGCGCCAGAAAAAGTGT 480
 V G E Q H C V A K M L P K S A P R K K C

GCAGCCTGTAAAGTCGTGGTGATACCCAAATGCATTAAGCAGCTGGAAAAAGATCAATTTC 540
 A A C K I V V H T Q C I K Q L E K I N F

CGCTGTAAGCCGCTCTTCGCGAATCAGGCTCCAGGAATGTCCTGAGCCAACTTCGTA 600
 R C K P S F R E S G S R N V R E P T F V

AGACACCACTGGTCCACAGACGACGCCAGGATGGCAAGTGTGGCACTGTGGGAAGGGC 660
 R H H W V H R R R O D G K C R H C G K G

TTCACGAGAAGTTACCTTCCACAGCAAGGAGATTGAGCCATCAGCTGCTCTGGTGC 720
 F Q Q K F T F H S K E I V A I S C S W C

AAACAGGCATACCACAGCAAGGTGCTGCTTCATGATGCAACAGATTGAGGAACCTCGC 780
 K Q A Y H S K M O Q I E E P C

TCCTAGGGGTGCATGCAGCGTGGTATCCCAACCCACCTGGATCTCGGGCCCGGAGG 840
 S L G V H A A V V I P P T W I L R A R R

CCCCAGAACACCTCAAGGCCAGCAAGAAGAAAAAGAGAGCGTCTTCAAGAGGAGTGC 900
 P Q N T L K A S K K K K R A S F K R R S

AGCAAGAAAGACCTGAGGAAGCCGCTGGAGACCTTCATCATCAGACCCACCCATCC 960
 S K K G P E E G R W R P F I I R P T P S

CCCCCATGAACCCCTGCTGGTGTGTTGTGAACCCCAAGAGTGGGGCAACAGGGTGC 1020
 P L M K P L L V F V N P K S G N Q F A

AAGATCATCCAGTCTTTTGTGGTATCTGAATCCCGACAGTCTTTGATCTGAGCCAG 1080
 K I I Q S F L W Y L N P R Q V F D L S Q

GGAGGACCCAGGGAAGCGCTGGAATATGCCGCAAGATACATAATTTGAGGATTCTGGCT 1140
 G G P R E A L E M Y R K V H N L R I L A

TGCGGGGTGATGGCAGGTTGGCTGGATTCTCTCCACCTGGACAGCTGCGCTTAAAA 1200
 C G G D G T V G W I L S T L D Q L R L K

CCACCGCTCTGTAGCCATCTGCTGCTGGGCACTGGCAATGACCTGGCCCGCAACCTC 1260
 P P P P V A I L P L G T G N D L A R T L

AAGTGGGTGGGGTTACACAGATGAGCCTGTGCAAGATCCTTTCCCATGTTGAGGAG 1320
 N W G G Y T D E P V S K I L S H V E E

GGGAATGTGTACAGCTGGACGTTGGACCTCCGAGCAGGCCAACCTGAGCGGGG 1380
 G N V V Q L D R W D L R A E P N P E A G

CCTGAGGAGCAGATGATGGAGCCACTGACCGCTGCCCTGGATGTCTTTAACAACATC 1440
 P E E R D D G A T D R L P L D V F N N Y

TTACGCTGGGCTTCGATGCCACGTCACCTGGAAATTCATGAGTCTCGAAGGCCAAC 1500
 F S L G F D A H V T L E F H E S R E A N

CCAGAGAAGTTCAACAGCCGCTTCGGAATAAGATGTTCTATGTGGGACGCGCTTCTCT 1560
 P E K F N S R F R N K M F Y A G T A F S

GACTTCCTGATGGGACGCTCCAAGGACTTAGCCAAGCACATCCGAGTAGTGTGTACGGA 1620
 D F L M G S S K D L A K H I R V V C D G

ATGGACCTAACCCCCAAGATTACAGGACCTGAAACCGCAGTGACATGCTTCTGAATATC 1680
 M D L T P K I Q D L K P Q C I V F L N I

CCCAGGTACTGTGACGACCATGCGCTGGGGCCACCCGGGAGCACCATGACTTCGAG 1740
 P R Y C A G T M P W G H P G E H H D F E

CCCCAGAGGCATGATGAGTACCTGGAGGTTCATCGGCTTACCATGACATCCTTGGCA 1800
 P Q R H D D G Y L E V I G F T M T S L A

GCATGCGAGTGGGTGGGCGGCGAGCGATTGACGAGTCCGAGAAGTGTCTCTACC 1860
 A L Q V G G H G E R L T Q C R E V L L T

ACTGCCAAGGCCATCCCTGTGACGTTGGACGTTGAGCCCTGCAAGCTTTACGATCACGG 1920
 T A K A I P V Q V D G E P C K L S A S R

ATTGCAATTGCCCTGCGCAACCAGGCCACTATGGTGCAGAAGGCCAAGCGCGGAGTACT 1980
 I R I A L R N Q A T M V Q K A K R R S T

GCCCCATGCACAGCGACCCAGCAGCTTCTGAGCAGCTGAGGATCCAGGTGAGCAGG 2040
 A P L H S D Q Q P V P E Q L R I Q V S R

GTCAGTATGCAGCACTACGAGGCTCTACATTATGACAAGGAGCAGCTCAAGGAAGCTCT 2100
 V S M H D Y E A L H Y D K E Q L K E A S

GTGCCTCTGGGCACTGTGGTGGTCCCCGGAGACAGTACCTGGAGCTCTGCGCGCCAC 2160
 V P L G T V V V P G D S D L E L C R A H

ATTGAGAGACTGCAGCGGAGCCGATGGGGCTGGAGCCAAGTCCCGATGTCACCAAA 2220
 I E R L Q R E P D G A G A K S P M C H Q

CTTTCCTCAAGTGGTGTTCCTGGATGCCACCACTGCCAGCGCTTACAGGATCGAC 2280
 L S S K W C F L D A T T A S R F Y R I D

AGAGCCAGGAGCAGCTCAACTATGTGACGAGATTGCCAGGAGAGATTACATCCTA 2340
 R A Q E H L N Y V T E I A Q D E I Y I L

GACCCAGGCTTCTGGGAGCATCAGCCCGCTGACCTCCCCACCTTACCTCCCCACTC 2400
 D P E L G T G A S A R P D L P T P T S P L

CCTGCTTCCCCCTGCTCCCTACACCCGGTCAATGCAGGGGACACTGCACTGCCCCAA 2460
 P A S P C S P T P G S M Q G D T A L P Q

GGTGAAGAACTGATTGAAGCAGCCAAGAGGAATGACTGTGCAAGCTCCAGGAGTACAC 2520
 G E E L I E A A K R N D C C K L Q E L H

CGAGCAGGAGGTGACCTCATGCACCGTACCAGAGAGCCGACGCTTTCACCAACGG 2580
 R A G G D L M H R D Q K S R T L L H H A

GTCAGCACTGGCAAGTAAGGAAGTGGTCCGCTATCTGCTGGATCATGACCTCCAGAGATC 2640
 V S T G S K E V V R Y L L D H A P P E I

CTTGATGCTGTGGAGGAAATGGGAGACCTGTCTACACAGGACGCGCCCTGGGTGAG 2700
 L D A V E N G E T C L H Q A A A L G Q

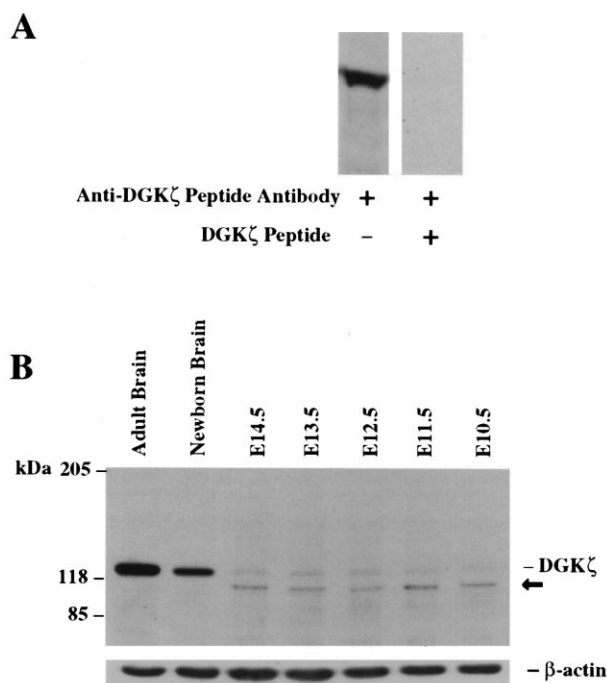
CGCACCATCTGCCACTACATTGTGGAAGCGGGGCTCCCTCATGAAGACAGATCTGCAG 2760
 R T I C H Y I V E A G A S L M K T D L Q

GGCGACACTCCCCGGCAGCGCTGAGAAGGCTCAAGACACAGAGCTCGCTGCCTACCTG 2820
 G D T P R Q R A E K A Q D T E L A A Y L

GAGAACAGACAGCATTACCAGATGATCCAGCGTGGAGACCAGGAGACAGCTGTGTAGTTA 2880
 E N R Q H Y Q M I Q R E D Q E T A V *

AGGGGACCATGGATCAACAAGGGACACTACCAGAGGACTCTCCTTGCCCATCTCACTGCC 2940
 ACATTCTGTGATGGCTATGGGGGACCTGCCACAGGGAAGGAGCCCGTGCCACCC 3000
 CCTGAGAAGCTGTTGATCTAGGGCTGGACTCTAAGGAGCTGGACTCTCACCTGTCCCT 3060
 GGTTTCATGGGGAACAGGAACAGGCTGGGCTGGCTGGTTCCTCCCTTACGGGAGCCT 3120
 CCAGCTCACCACAGCAGATGGAAGGGCTGGACAGAGCTCAGCATCGATGTGAGGAGGCC 3180
 TGCACTCAGCAGGACTTTCAAAAGCCACCTGATCCTTTGGGCTCTCTGAGGGTTCTGGG 3240
 TACCTAACCTCCTCTGTATCCACCCACCTGGGATCCAGAACTCAAGAGCCTGTCT 3300
 GTATTCACTTGCCCGCTGCCCTGCTTGGCACTACCTGGTCTTCTCTCATGATCA 3360
 GCCATTTCATTTGGGACTGTATGGCTGGGGTGGGGTGGGGTCCACGCTGACTGT 3420
 TTACAGCTGGGTTGACTCAGTAAAGTGAATTTTTTCCAAAAA

Fig. 1 (continued).



row), ~5–10 kDa smaller than the full-length protein, was detected in embryo homogenates and its recognition was also blocked by pre-incubation with the peptide antigen. We speculate that it is a proteolytic product since its intensity was

Fig. 2. The mDGK ζ protein level changes during mouse embryogenesis. A: 100 μ g of protein of mouse brain homogenate was loaded in each lane and probed with the affinity purified anti-DGK ζ antibody at a concentration of 1 μ g/ml. In the control experiment, pre-incubation with the immunogen-peptide (4 volumes 0.1 mg/ml peptide+1 volume 1 mg/ml antibody) significantly blocked the recognition of the DGK ζ protein. B: Embryos at different stages were dissected and homogenized. 100 μ g of protein from embryo homogenates, newborn brain and adult brain homogenates was loaded in each lane. They were probed with the affinity purified anti-DGK ζ antibody. The β -actin antibody was used as a control to show that an equal amount of protein was loaded in each lane. This is a representative result from five independent experiments.

variable and a PEST sequence was found in the N-terminus of DGK ζ [22]. In five independent experiments, we found a high level of DGK ζ protein in the brain of newborn mice, and a further two-fold increase at the DGK ζ protein level in adult mouse brain. The DGK ζ signal was normalized to β -actin (Fig. 2B).

3.3. Patterns of DGK ζ expression: examination of DGK ζ by RNA whole-mount in situ hybridization indicates that DGK ζ is expressed predominantly in sensory structures

After finding the change in DGK ζ expression during development, we examined its tissue distribution by in situ hybridization for mRNA on whole-mount preparations of mouse embryos. Antisense and sense probes to mDGK ζ were synthe-

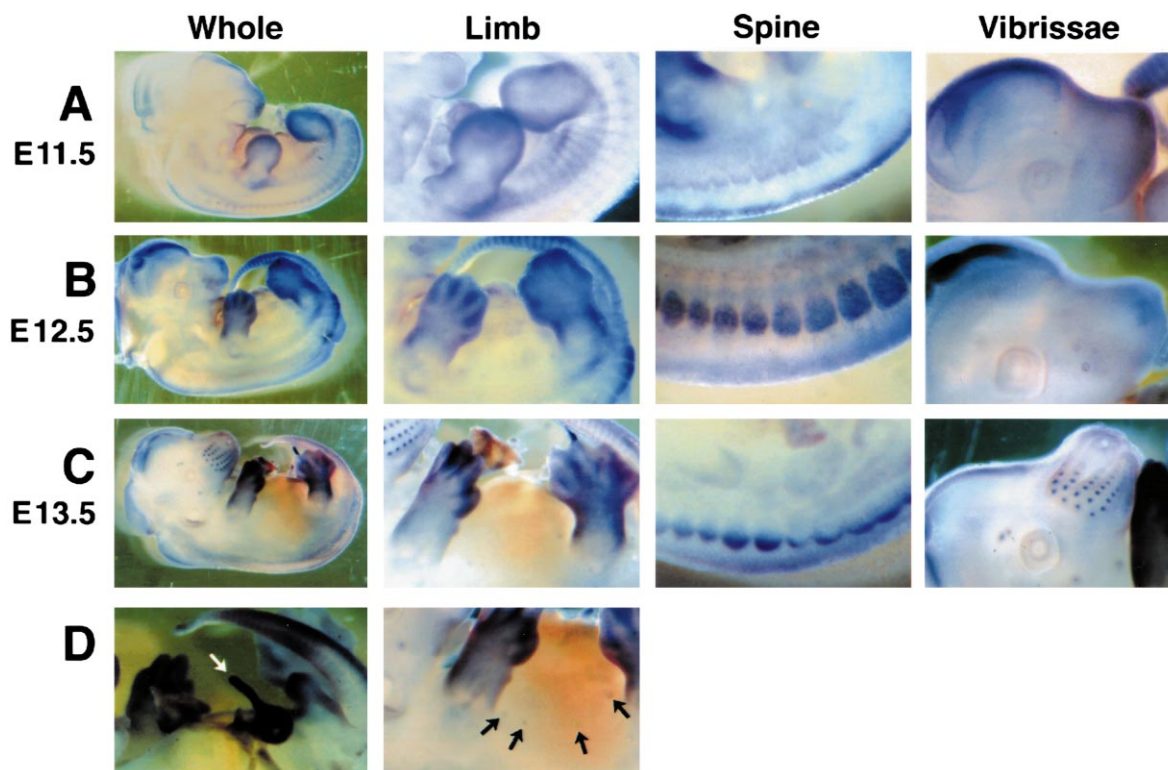


Fig. 3. Murine DGK ζ is highly expressed in the sensory nerve system during mouse embryogenesis. Mouse embryos at different stages were hybridized with DGK ζ antisense and sense digoxigenin-labeled probes as described in Section 2.3 (blue-purple staining represents the positive signals). The DGK ζ sense probe gave no specific staining. A: Low level expression of DGK ζ in somites, limb buds and spinal ganglia in E11.5 day embryos. B: Strong expression of DGK ζ in spinal ganglia, limb buds and low level expression in the follicles of vibrissae in E12.5 day embryos. C: Constitutively high level expression in limbs and spinal ganglia, and the follicles of vibrissa in E13.5 day embryos. D: Expression of DGK ζ in umbilical vessels and nipple primordia.

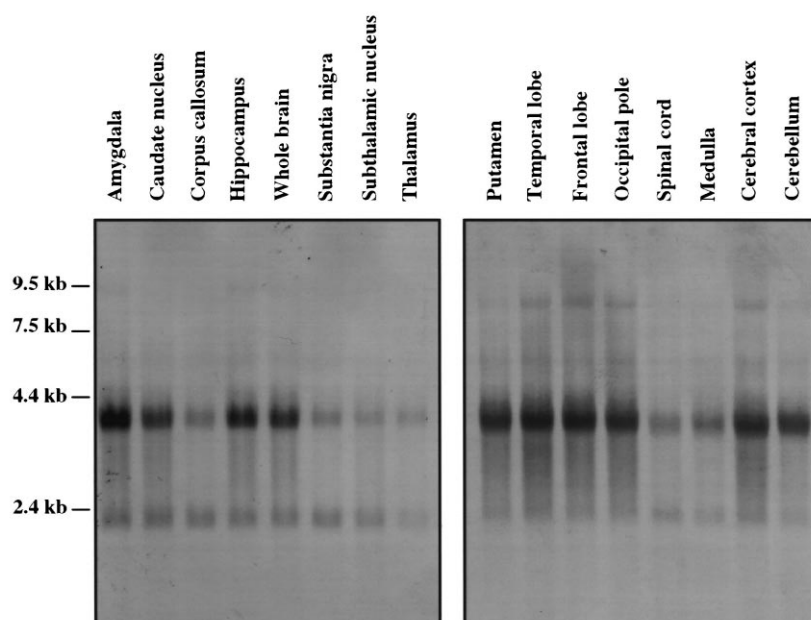


Fig. 4. Analysis of DGK ζ mRNA from different regions of the human brain. Two filters with 2 μ g mRNA from different brain regions were probed with a fragment of the hDGK ζ . A sample from whole brain shows a band at 3.7 kb. The expression level of DGK ζ varies among different regions of brain, high in cerebral cortex, cerebellum, hippocampus, occipital pole, frontal lobe, temporal lobe, putamen, amygdala and caudate nucleus, and low in spinal cord, medulla, corpus callosum, substantia nigra, subthalamic nucleus and thalamus.

sized and used to examine the expression of DGK ζ in E10.5, E11.5, E12.5, and E13.5 embryos. In E10.5 embryos, we could not define a tissue-specific pattern of expression because the signals were not sufficiently above the background. In E11.5 embryos, we observed low level expression in somites, spinal ganglia, and limb buds (Fig. 3A). We detected much stronger expression of DGK ζ in spinal ganglia and limb buds in E12.5 embryos (Fig. 3B). Additionally, very strong expression was found in the interdigital regions of the limb where cells are undergoing apoptosis to create the fully developed digits. DGK ζ staining was very strong in spinal ganglia and highlighted the segmentation of the embryo. Interestingly, the spinal cord was not highly labeled (Fig. 3B). The dorsal root ganglia are comprised of the cell bodies of sensory neurons and our results demonstrate that DGK ζ is highly expressed in them. When we examined E13.5 day embryos, we found that DGK ζ was highly expressed in the follicles of the vibrissae, in spinal ganglia, and in limb buds (Fig. 3C). The vibrissa follicle contains venous sinuses and sensory nerve fibers within a connective tissue sheath and is first observable during the 13th embryonic day. The strong DGK ζ staining in the follicles of the vibrissae of E13.5 embryos suggests that DGK ζ is expressed from the initiation of vibrissae. We also observed staining in the umbilical vessels of E13.5 embryos (Fig. 3D), which is noteworthy since hDGK ζ was originally cloned from an umbilical vein endothelial cell library [22]. Interestingly, expression of DGK ζ also was detected in the nipple primordia of E13.5 embryos (Fig. 3D). We observed DGK ζ staining in the brain of the embryos, but the precise anatomical localization of the signal was not easily revealed by this technique. Both probes mentioned in Section 2.3 gave us identical staining patterns. In all cases hybridization with the sense probe was performed to confirm that the pattern observed with the antisense probe was specific.

3.4. Northern blotting of isolated regions of brain defines specific patterns of expression

To refine the localization of DGK ζ expression in the brain, we used Northern blotting to examine mRNA samples from specific regions of human brain. The membranes were hybridized with a 761-bp digoxigenin-labeled hDGK ζ probe. The hybridization was performed as previously described [22]. We detected a band at 3.7-kb position in the whole brain sample, as expected. Interestingly, the expression level varied among the different regions with strong expression in cerebellum, cerebral cortex, and hippocampus. In contrast, the expression of DGK ζ in the spinal cord, medulla of the brain, and thalamus was low (Fig. 4). A β -actin probe was used as a control to confirm the same amount of the mRNA loaded and the integrity of the mRNA in each lane demonstrating the same amount of mRNA from different regions of brain was loaded. These results are consistent with the *in situ* hybridization experiment in which we did not detect significant staining in the spinal cord of embryos of different ages. Thus, DGK ζ expression is limited to specific regions of the human brain.

4. Discussion

To study the developmental regulation and functions of DGK ζ , we cloned the mDGK ζ and analyzed the expression of this gene during mouse embryogenesis. Murine DGK ζ has highest sequence similarity with hDGK ζ (97%) and *Drosophila* DGK2, *rdgA* (49%) at the amino acid level. The deduced mDGK ζ protein has the domain motifs that define the type IV DGKs: two cysteine-rich repeats, a conserved catalytic domain, and four ankyrin repeats at the C-terminus. Like the other members of the DGK family, mDGK ζ shares the greatest homology with other DGKs in the catalytic domain. Within that domain DGK ζ has the conserved putative ATP

binding site GXGXXG, of which a mutation in the second glycine in *Drosophila* DGK2 causes the retinal degeneration phenotype [9].

We studied mDGK ζ expression in embryos from E10.5 to E14.5 and found that the expression increases gradually during embryogenesis. We observed that brains of both newborn and adult mice have abundant DGK ζ protein suggesting that DGK ζ may be involved in neuronal development and function. We determined the location of the DGK ζ mRNA in developing embryos and found that it is highly expressed in the sensory nerve system including the dorsal root ganglia and vibrissa follicles. The expression pattern of DGK ζ shows similarity to the expression pattern of the *patched* regulatory gene at some stages [31]. *Patched* is a part of the hedgehog signaling pathway, which is down-regulated by protein kinase A. There is no known relationship between this pathway and the DGK-mediated signaling, but our findings show they have overlapping expression during development. This suggests a potential interaction between them but more experimental evidence is needed to distinguish whether this indicates functional relationship. Our whole-mount in situ hybridization method was not sufficiently sensitive to define subregions of the mouse embryonic brain, so we used samples from different anatomical regions of the human brain to test whether the DGK ζ mRNA is limited to specific regions of the brain. DGK ζ is highly expressed in the sensory nervous system, while expression is low in other areas, particularly the spinal cord.

Our data demonstrate that the expression of DGK ζ is subject to strict temporal and spatial regulation during mammalian development. Its tissue-specific expression pattern is characterized by selective expression in many sensory components of the nerve system. At the same time, the expression of DGK ζ overlaps significantly with the pattern observed with other DGK isoforms [12,21,28]. Although DGK isoforms differ in substrate specificity and cofactor usage, when two or more DGK isoforms are expressed in the same cells, they may have complementary functions, or act in concert to regulate the level of DAG and/or other important lipid messengers. They may also be functionally redundant in these cells, meaning one can compensate for the functional loss of the other. On the other hand, DGK ζ may have unique and indispensable functions especially where it is expressed alone or at a much higher level than other DGK isoforms. Taken together, it is likely that different DGK isoforms have unique functions, but also may act cooperatively to control cellular events such as growth and differentiation.

Acknowledgements: We thank Matt Topham for many suggestions and discussions, Elie Traer for the sequence analysis, and Diana Lim for the preparation of figures. Susan Mango and Diana Stafforini contributed helpful comments on the manuscript. This work was supported by Grant CA59548 from the National Cancer Institute. The DNA sequencing core facility at the University of Utah is supported by Grant CA42014 from the National Cancer Institute.

References

- [1] Kikkawa, U., Kishimoto, A. and Nishizuka, Y. (1989) *Annu. Rev. Biochem.* 58, 31–44.
- [2] Kearns, B.G., McGee, T.P., Mayinger, P., Gedvilaite, A., Phillips, S.E., Kagiwada, S. and Bankaitis, V.A. (1997) *Nature* 387, 101–105.
- [3] Fu, T., Sugimoto, Y., Okano, Y., Kanoh, H. and Nozawa, Y. (1992) *FEBS Lett.* 307, 301–304.
- [4] Moolenaar, W.H., Kruijer, W., Tilly, B.C., Verlaan, I., Bierman, A.J. and de Laat, S.W. (1986) *Nature* 323, 171–173.
- [5] Bocchino, S.B., Wilson, P.B. and Exton, J.H. (1991) *Proc. Natl. Acad. Sci. USA* 88, 6210–6213.
- [6] van Corven, E.J., van Rijswijk, A., Jalink, K., van der Bend, R.L., van Blitterswijk, W.J. and Moolenaar, W.H. (1992) *Biochem. J.* 281, 163–169.
- [7] Ahmed, S., Lee, J., Kozma, R., Best, A., Monfries, C. and Lim, L. (1993) *J. Biol. Chem.* 268, 10709–10712.
- [8] Bollag, G. and McCormick, F. (1991) *Nature* 351, 576–579.
- [9] Masai, I., Okazaki, A., Hosoya, T. and Hotta, Y. (1993) *Proc. Natl. Acad. Sci. USA* 90, 11157–11161.
- [10] Sakane, F., Yamada, K., Kanoh, H., Yokoyama, C. and Tanabe, T. (1990) *Nature* 344, 345–348.
- [11] Schaap, D., de Widt, J., van der Wal, J., Vandekerckhove, J., van Damme, J., Gussow, D., Ploegh, H.L., van Blitterswijk, W.J. and van der Bend, R.L. (1990) *FEBS Lett.* 275, 151–158.
- [12] Goto, K. and Kondo, H. (1993) *Proc. Natl. Acad. Sci. USA* 90, 7598–7602.
- [13] Kai, M., Sakane, F., Imai, S.-i., Wada, I. and Kanoh, H. (1994) *J. Biol. Chem.* 269, 18492–18498.
- [14] Yamada, K., Sakane, F., Matsushima, N. and Kanoh, H. (1997) *Biochem. J.* 321, 59–64.
- [15] Sakane, F., Imai, S., Kai, M., Wada, I. and Kanoh, H. (1996) *J. Biol. Chem.* 271, 8394–8401.
- [16] Klauck, T.M., Xu, X., Mousseau, B. and Jaken, S. (1996) *J. Biol. Chem.* 271, 19781–19788.
- [17] Musacchio, A., Gibson, T., Rice, P., Thompson, J. and Saraste, M. (1993) *Trends Biochem. Sci.* 18, 343–348.
- [18] Lomasney, J.W., Cheng, H.F., Wang, L.P., Kuan, Y., Liu, S., Fesik, S.W. and King, K. (1996) *J. Biol. Chem.* 271, 25316–25326.
- [19] Paris, S., Beraud-Dufour, S., Robineau, S., Bigay, J., Antonny, B., Chabre, M. and Chardin, P. (1997) *J. Biol. Chem.* 272, 22221–22226.
- [20] Tang, W., Bunting, M., Zimmerman, G.A., McIntyre, T.M. and Prescott, S.M. (1996) *J. Biol. Chem.* 271, 10237–10241.
- [21] Kohyama-Kogane, A., Watanabe, M. and Hotta, Y. (1997) *FEBS Lett.* 409, 258–264.
- [22] Bunting, M., Tang, W., Zimmerman, G.A., McIntyre, T.M. and Prescott, S.M. (1996) *J. Biol. Chem.* 271, 10230–10236.
- [23] Goto, K. and Kondo, H. (1996) *Proc. Natl. Acad. Sci. USA* 93, 11196–11201.
- [24] Davis, N., Ghosh, S., Simmons, D.L., Tempst, P., Liou, H.C., Baltimore, D. and Bose Jr., H.R. (1991) *Science* 253, 1268–1271.
- [25] Watanabe, H., Sawada, J.-I., Yano, K.-I., Yamaguchi, K., Goto, M. and Handa, H. (1993) *Mol. Cell. Biol.* 13, 1385–1391.
- [26] Blank, V., Kourilsky, P. and Israel, A. (1992) *Trends Biochem. Sci.* 17, 135–140.
- [27] Andrews, B.J. and Herskowitz, I. (1989) *Nature* 342, 830–833.
- [28] Houssa, B., Schaap, D., van der Wal, J., Goto, K., Yamakawa, A., Shibata, M., Takenawa, T. and van Blitterswijk, W.J. (1997) *J. Biol. Chem.* 272, 10422–10428.
- [29] Chen, F. and Capocchi, M.R. (1997) *Dev. Biol.* 181, 186–196.
- [30] Ding, L., Bunting, M., Topham, M.K., McIntyre, T.M., Zimmerman, G.A. and Prescott, S.M. (1997) *Proc. Natl. Acad. Sci. USA* 94, 5519–5524.
- [31] Goodrich, L.V., Johnson, R.L., Milenkovic, L., McMahon, J.A. and Scott, M.P. (1996) *Genes Dev.* 10, 301–312.