

(19)  
(12)(KR)  
(A)(51) 。 Int. Cl. <sup>7</sup>  
C12N 15/55(11)  
(43)2002 - 0076562  
2002 10 11(21) 10 - 2001 - 0016517  
(22) 2001 03 29

(71) 100 104 206

(72) 100 104 206

2가43 - 1,1

111 1703

B 202

가 414 - 401

(74)

:

(54)

ERK , G<sub>1</sub> S , 가 , ERK

6a

1 a DMKP - 3 Drosophila MAPK cDNA , 1b ,  
1c .

2 DERK, DJNK, Dp38 MAPK DMKP - 3 . 2a  
DMKP - 3  
, 2b 2c LPS NaCl 가 - DJNK - Dp38  
, 2c Cys - 302 (DMKP - 3 - CA)  
- DERK .

3 MAPK - ERK DMKP - 3 . 3a DMK  
P - 3 Elk - 1 -  
, 3a c - Jun - (JNK ) DMKP - 3  
(Dp38 ) DMKP - 3  
3b GST - DMKP - 3 GST - DMKP - 3 가  
- ERK , - ERK .

4 DMKP - 3 DERK . 4a B42 - DMKP - 3 LexA - DERK  
DMKP - 3 ERK가 가 , B  
42 - DMKP - 3 LexA - DJNK LexA - Dp38 가  
DMKP - 3가 DERK , 4b B42 - DMKP - 3 - R56A/R57  
가 18 DMKP - 3  
, DMKP - 3가 N - ( R56 R57 ) ERK .

5 DMKP - 3 . 5a Schneider DMKP - 3  
, 5b 8 - 12  
DMKP - 3 mRNA

6 DMKP - 3 G<sub>1</sub> S ( ) .  
6a G<sub>1</sub> S , G<sub>0</sub>/G<sub>1</sub> , S  
DMKP - 3 DMKP - 3 - CA , G<sub>1</sub> S 가  
, 6b BrdU % , BrdU D  
MKP - 3 ( 6b, ), DMKP - 3  
19% 12% BrdU ( 6b, ) .

7 (Forward light scatter) Schneider  
, 가 가 , DMKP - 3 DMKP - 3 - CA(  
) .

Extra - cellular regulated kinase( , " ERK" )

가

ERK, SAPK p38

가

가

5,420,245

(Ras)

가

(RYK)

( , " ERK  
DMKP - 3" )

1

42AD - DMKP - 3

pOT2 - DMKP - 3, pMT/V5 - DMKP - 3, pPacPL - DMKP - 3, pcDNA3.1 - DMKP - 3, pB

1

가

2

Drosophila ERK

, G<sub>1</sub> S

(Mitogen - activated protein kinase, " MAPK" )

ERK, JNK, p38 MAPK MAPK 가  
(Blumer and Johnson, 1994. TIBS 19:232 - 240; Waskiewicz and Cooper, 1995. Curr.Opin.Cell Biol.7:798 - 805). MAPK  
Drosophila MAPK  
MAPKs (Haneda et al., 1999. Eur. J. Pharmacol.365:1 - 7). MAPKs  
Thr - X - Tyr  
MAPK (MAPK phosphatase, " MKPs" ) MAPKs MAPK MAPK  
MKPs 가 MAPKs MAPK  
MKPs MAPKs MKPs가  
MKP - 1 (CL100) MKP - 4 ERK (Extra - cellular regulated kinase), JNK (c - Jun N - terminal kinase) p38 , PAC1 ERK p38 MAPK  
MKP , MKP - 3 VHR ERK  
(Camps et al., 1998. FASEB J.14:6 - 16).

MAPKs , MKP MAPK Drosophila  
(Keyse, 1995. Nature. 359:644 - 647). MAPKs MKP  
s Drosophila Drosophila ERK(DERK)  
가 (Lee et al., 2000. Biochem. J.349:821 - 828). Drosophila MAPK  
(Duffy and Perrimon, 1996. Curr. Opin. Cell. Biol. 8:231 - 238). ERK가  
/ (Kerkhoff and Rapp, 1998. Oncogene 17:1457 - 1462), Drosophila DERK

in vitro in vivo ERK Drosophila MKP DMKP -  
3 DMKP - 3 , DE  
RK가 , G<sub>1</sub> S

1: DMKP - 3 MKPs

1) Drosophila MAPK cDNA

(dual) ( " MKP" ) Drosophila dbEST  
MKP cDNA . C - cDNA (GM13896)  
Research Genetics, Inc.(Huntsville, AL) , N cDNA pBlueScript SK(Stratagene)  
T3 5 - AATTAACCCTCACTAAAGGG - 3 Drosophila cDNA 5 - G  
TAGCATTCGCCAGATGG - 3 PCR

2)

N Drosophila MKP 1.3kb PCR pBlueScript SK(Stratagene) EcoRV  
 pBS - N - DMKP - 3 cDNA cDNA C cDNA  
 (GM13896) pOT2 Smal/SphI - DMKP - 3 pGST - DM  
 KP - 3 pGEX2TK(Pharmacia) EcoRI pOT2 - DMKP - 3 1.6kb . Droso  
 phila DMKP - 3 pPacPL - DMKP - 3 pPacPL XbaI - NotI 5 - GGAATTCGGCTCTA  
 GACCATGGCAGAAACGGAGCACGA - 3 , 5 - GGCAACGGCGATGTGGCGGCCGCTGCAAATGGGAT  
 CTC - 3 pOT2 - DMKP - 3 PCR DMKP - 3 pcDNA3.  
 1 - DMKP - 3 pcDNA3.1(Invitrogen) EcoRI 1.6kb pOT2 - DMKP - 3  
 pMT/V5 - DMKP - 3 pMT/V5 - C(Invitrogen) EcoRI 1.6kb pOT2 - DMKP - 3  
 5 - CGGCACGAATTCATGCCAGAAACGGAGCAC - 3 5 - GCCACTCTCGAGTCATTTAA  
 GACCCGTGTCCG - 3 , pOT2 - DMKP - 3 pB42AD - DMKP - 3  
 pB42AD(Clonetech) EcoRI - XhoI . pB42AD - DMKP - 3(1 - 239) pB42AD EcoRI - XhoI  
 5 - CGGCACGAATTCATGCCAGAAACGGAGCACG - 3 5 - ATACTTCTCGAGTCACTTCTT  
 CAACGCTTCCGAG - 3 pOT2 - DMKP - 3 PCR DMKP - 3 . pB42AD - DM  
 KP - 3(210 - 411) pB42AD EcoRI - XhoI 5 - CACAGTGAATTCAATTACAACGAGGC  
 GCCCG - 3 5 - GCCACTCTCGAGTCATTTAAGACCCGTGTCCG - 3 pOT2 - DMKP - 3  
 PCR DMKP - 3 . pLexA - DERK pLexA EcoRI 5 - GAAACGGAATTCAT  
 GGAGGAATTTAATTTCGAGCG - 3 5 - TACAGCGAATTCTTAAGGCGCATTGTCTGGTTGTC - 3  
 pPacPL - His - DERK PCR DERK . pLexA - DJNK pLexA BamHI - XhoI  
 5 - ATCAGTGGATCCTGACGACAGCTCAGCACCAACAC - 3 5 - AAAAGTCTCGAGCTACCGC  
 GTTCTATTATTTGTATTG - 3 pPacPL - His - DJNK PCR DJNK . pLexA  
 - Dp38 pLexA BamHI - XhoI 5 - TCAAGCGAATTCATGTCAGTGTCCATTACAAAAA  
 G - 3 5 - GATGGTCTCGAGTCACTTTACATCCTTTAGAACC - 3 pPacPL - His - Dp38  
 PCR Dp38 .

3)

DNA Sanger ,  
 NTI 6.0.1 (InforMax Inc, MD) .

DMKP - 3 Drosophila MAPK cDNA 45.7kDa  
 411 ( 1a). DMKP - 3 ERK  
 MKP - 3 ( 1b), 33.3%(44.8%)  
 ) . DMKP - 3 N Cdc25 (CH2)  
 ( 1 c). DMKP - 3 MKPs N  
 , MKP - 3 24.6% ERK  
 MKP - 3 " IMLRR" DMKP - 3 " IVLRR" ( 1c).

2: MAKPs DMKP - 3

1)

DMKP - 3 PCR (Stratagene) . DMKP - 3 - CA  
 5' - GAAACTCCGGCCAGGGCGTGGACCAGC - 3' , DMKP - 3 - CS 5'  
 - GGTGCTGGTCCACTCCCTGGCCGGAG - 3' , DMKP - 3 - A56R/A57R 5' - CCCAGC  
 ATCGTCCTCGCGGCCCTGGCGGTTGGC - 3'  
 가 . DMKP - 3 .

2)

GST - DMKP - 3 pGST - DMKP - 3 가 (Sigma)  
 . DMKP - 3 Lee (Biochem.J.349:821 - 829) , GST - DMKP - 3

3) DMKP - 3

DMKP - 3, DMKP - CA DMKP - 3 - CS Schneider Lee (Biochem.J.349:821 - 829)  
 , pMT/V5 - DMKP - 3( ) pCoHygro  
 . Schneider Schneider's Insect Media(Sigma) , Schneider DMKP - 3  
 DJNK Dp38 Lee (Biochem.J.349:821 - 829) . DERK 10 $\mu$ g/ml  
 5 (Clemens et al., P.N.A.S. 97:6449 - 6503).

4)

Lee (Biochem.J.349:821 - 829) DERK, DJNK  
 Dp38 - - ERK, JNK p38 MAPK (New England Bio Labs)  
 - - (Calbiochem) horseradish pero  
 xidase(HRP) - - IgG(Bio Rad) - IgG(Promega) , e  
 enhanced chemiluminescence(ECL) (Genepia) , X - .

DMKP - 3 - DERK (2a).  
 Schneider DERK DPI3K - DAkt 가 (Clemens, 2000).  
 Schneider DMKP - 3 DERK 가 가  
 , DMKP - 3 2A , Schneider DERK 2 - 3 가 가  
 , DMKP - 3 - DERK .

- DERK , LPS 가 - DJNK (2b). DMKP - 3  
 2b). DMKP - 3 NaCl 가 - Dp38 (2b). DMKP - 3  
 Cys - 302 MKP , Cys - 302 (DMKP - 3 - CA) (DMKP -  
 . DERK , DMKP - 3 , DERK  
 3 - CS) , - DERK DMKP - 3 - CA

## DMKP - 3 - CS

( 2c).

3: MAPK - ERK DMKP - 3

## 1) PathDetect Trans - Reporting System

in vivo MAPK (Stratagene), Pathfind  
 er Trans - reporting system CV - 1 100 /ml 100 µg/ml  
 10%(v/v)FBS DMEM . CV - 1 37 , 5% CO<sub>2</sub>  
 18 , 3x 10<sup>7</sup> pFR - Luc - (pFA - Elk - 1, pFA2 - J  
 un, pFA - CHOP) CMV - - gal . Lipofectamin plus (Gibco BRL)

2)

In vitro DMKP - 3 GST - DMKP - 3 5 ng  
 - ERK (Stratagene) 25µl (50mM Tris pH 7.5, 1mM EDTA, 10mM  
 DTT) 30 30 , 가 , - ERK  
 - ERK .

DMKP - 3 in vivo , , Elk - 1, c - Jun c - CHOP MAPK  
 Elk - 1 - 3a , DMKP - 3 , Elk - 1 -  
 DMKP - 3 - CS , DERK  
 ( 2c). Elk - 1 , JNK c - Jun -  
 DMKP - 3 ( 3a, ), p38 MAPK  
 c - CHOP - DMKP - 3 ( 3a, ).  
 DMKP - 3 ERK , GST (GST - DMKP - 3)  
 DMKP - 3 , 가 GST - D  
 MKP - 3 ERK( - ERK) 가 . 3b  
 , GST - DMKP - 3 GST - DMKP - 3 가  
 - ERK , - ERK

4: DMKP - 3 DERK ( )

EGY48(MAT his3 trp1 ura3 lexAop(x6) - LEU2) p8op - lacZ (Clonetech) pLe  
 xA pB42AD , (SC - Ura His Trp)  
 2% 1% , 가 30  
 1/6 , - 4  
 30 5 .

DERK DMKP - 3 , -  
 DERK, DJNK Dp38 LexA DNA - p8op - LacZ  
 (" B42" ) DMKP - 3 B42 - DMKP - 3 LexA - DE  
 RK - 가 , B42 - DMKP - 3 LexA - DJNK LexA - Dp3  
 8 - 가 ( 4a). DERK  
 DMKP - 3 , DMKP - 3 N (1 - 239) C (210 - 411) B42  
 , DERK N C LexA - DERK  
 , B42 - DMKP - 3(1 - 239 ) 가 ,  
 LexA - DERK B42 - DMKP - 3(210 - 411 ) 가  
 MKP - 3 N - (64 65 ) 가 ERK ,  
 DMKP - 3 " IVLRR" Arg - 56 - 57 Ala  
 B42 - DMKP - 3 - R56A/R57 LexA - DERK  
 4b , B42 - DMKP - 3 - R56A/R57  
 가 18 .

## 5: DMKP - 3

### 1) (Immunocytochemistry)

DMKP - 3 CuSO<sub>4</sub> 1mM  
 24 100% - 20 10 1% BSA 5%  
 PBS 30 1:100 DMKP - 3  
 2 PBS 3 - IgG (1:100 ;J  
 ackson) 50% 1 . PBS , Radiance2000/M  
 P(Bio - Rad, UK) .

### 2) mRNA

Drosophila mRNA DNA cDNA Drosophila rapid - S  
 can<sup>TM</sup> Gene Expression panel(Origene) , DMKP - 3 5 - GCAAGGAGTGGCTGCAG  
 TCC - 3 , 5 - GGGATTATCTCTACGGGCGC - 3 DMKP - 3  
 PCR . 15 µl PCR 0.5 µg/ml EtBr 가

MKP - 3 , - DMKP - 3 DMKP - 3  
 Schneider DMKP - 3 , ( 5a).  
 DMKP - 3 DMKP - 3 - CA ( 5a).  
 8 - 12 , DMKP - 3 mRNA ( 5b).  
 RP49 mRNA .

## 6: DMKP - 3 G1<sub>1</sub>/S



, DMKP - 3 DMKP - 3 - CA 10%FBS Schneider 50  
 % 가 . FACS(fluorescent activating cell sorting)  
 DMKP - 3 DMKP - 3 - CA , 1 mM CuSO<sub>4</sub> , 69 .  
 10 µg/ml 24 . 6 - PBS 2 70%  
 가 . 1% FCS Becton Dickinson FACS Caliber ,  
 4 30 . WinMDI2.8(Scripps Research Institute, CA)  
 ModFit LT 2.0(Verity Software House, Inc., ME) . BrdU , Schneider 6 - 60% 가  
 23 . 7 µg pPacPL , pPacPL - DMKP - 3 pPacPL - DMKP - 3 - CA 가 ,  
 Han (J.Biol.Chem. 273: 369 - 374, 1998) . 24 , 10  
 % FBS 10 µg/ml 가 24 0.2%  
 . BrdU 4 , / (99:1) - 20 15 ,  
 Triton - X100 PBS 5 . - DMKP - 3 (1:100) , PBS ,  
 - (1:100) , . 10 3.7% , PBS ,  
 BrdU 2M HCl 30 . PBS 5 1% BSA 5% P  
 BS . - BrdU (Jackson) , Cy2 (Jacks  
 on) . DNA , 1 µg/ml 4',6' - Diamidine - 2 - phenylindole dihyd  
 rochloride(Boehringer Mannheim) . 3 .

Schneider 가 .  
 , DERK가 Drosophila Schneider 가 ,  
 , G1 S , DMKP - 3  
 . G1 S , S  
 DMKP - 3 DMKP - 3 - CA , S  
 CuSO<sub>4</sub> DMKP - 3 ( 6a). DMKP - 3 - CA DERK 18.5% 가 S  
 , G1 S , G1 S 가  
 , 9.9% ( 6a). CuSO<sub>4</sub> DMKP - 3  
 , G<sub>1</sub> S , S  
 33% 가 BrdU , BrdU % 60% 가  
 ( 6b), DNA BrdU DMKP - 3 1  
 ( 6b, ), , DMKP - 3 - CA  
 9% 12% BrdU ( 6b, ). BrdU % ,  
 ( 6b). , BrdU %  
 MKP - 3 - CA 60% 35% . DMKP - 3 - CA  
 , DMKP - 3 G<sub>1</sub> S  
 Schneider .  
 (Forward light scatter) Schneider DMKP - 3  
 DMKP - 3 - CA 가 ( 7, ).  
 가 CuSO<sub>4</sub> DMKP - 3 DMKP - 3 - CA ( 7,  
 ). CuSO<sub>4</sub> DMKP - 3 DMKP - 3 - CA ( 7).  
 DERK가 가 .

Drosophila ERK 가 Drosophila MKP DMKP - 3 ERK

(57)

1.

ERK 1

2.

1 ,

3.

2 ,

pOT2 - DMKP - 3, pMT/V5 - DMKP - 3, pPacPL - DMKP - 3, pcDNA3.1 - DMKP - 3, pB42A

D - DMKP - 3 1

4.

1 2

5.

1 ,

ERK Drosophila

6.

1 4 5 ,

G<sub>1</sub> S

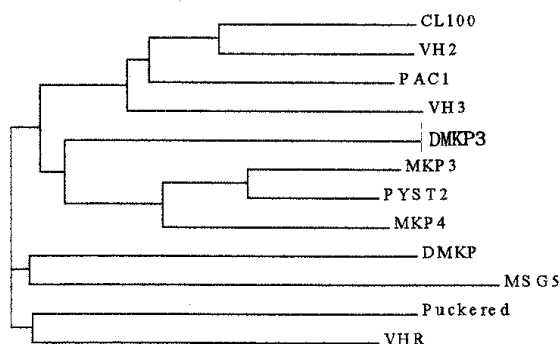
## 1a

```

1  GGCACGAGAAAATGCCAGAAACGGAGCAGCAGACCTGCAGCAAGGAGTGGCTGCAGTCCAGTTGCGATCCCTGGACTCCAGGACCTG
    M P E T E H E T C S K E W L Q S Q L R S L D S K D L
90  ATCCTGTTGGACTGCGCGGGCTCGCAGAGTACAGTGAATCGCACATCGCGGGAGCGGTCAACCTATGCTATCCAGCATCGTCCCTCGGT
    I L L D C R G S H E Y S E S H I R G A V N L C I P S U V L R
180  CGTCTGGCGGTGGCAAAATCGATCTGGCTCCACGATCAAGTCGCGCGAGCTGAAGCAAGCATCCAGTGGGCTACAGCTATGCTGG
    R L A V G K I D L A S T I K S P E L K Q R I Q S G Y K L C W
270  TTGATCCTCTACAAACGGCGAGGCGGTGCGCGGCGAGAATCAGGAGATCGCGGAGCGGATCCCTGGCGGTGCGCATGGACTCCATCATC
    F I L Y N G E G V P G Q N Q E I A G A G S L A V A M D S I I
360  AGCATCTCGACCGCTCGCTCAAGCAGGACGGCTGCGCGGTAGTTGCTTTACAGATGGCTTCAACAATTTTGGCCAGGCAATTTCCGGA
    S I L H R R L K Q D G C R V V A L Q D G F N N F R Q A F P E
450  TGGTGGAGGACGATAATCAGACCGCAGCAAGAGATCGAATCTAGTCGCAATGTTCAACCGATCAGTTAATGGGTCTTAGGTCCCTT
    W C E D D N Q T H S K E I E S S R N V Q T D Q L M G L R S L
540  CGCATTTCCACAACGCAATCCGATTCGCGTGCAGCAGTTCCGCGGAATCGTGGATTCGAGAGCTCCAGCCACCATCACCACCCAC
    R I S T T Q S D S A C S S S A B S S D C E S S S H H H H H
630  AGTCTCCCAATTAACAAGGCGCGCGTAGAGATAATCCCTGGACTACTCTTCTGGGAATGCCACACAGCTGCGACTCGGAAGCG
    S L H N Y N R A P V E I I P G L L F L G N A T H S C D S E A
720  TTGAAGAAGTACAAATATAAGTATGTTTGAATGTGACACAGATTTCGCAATAAGTTCAAGGAGTCGGGCGACATCAAGTATCTGCAG
    L K K Y N I K Y V L N V T P D L P N K F K E S G D I K Y L Q
810  ATTCGATCAAGGATCAGTACTACAAAGATTGGCCATACATTTCCCGGATGCCATACAGTTTATAGAGGAAGCGGGTCCGCAAGCTCG
    I P I T D H Y S Q D L A I H F P D A I Q F I E E A R S A S S
900  GTGTGCTGGTCCACTGCGTGGCGGAGTTTGGCGCTCGGTGACCGTGACGCTCGCTACTTGTATGACACGCGGGGCTGAGTCTCAAC
    V V L V H C L A G V S R S V T V T L A Y L M H T R G L S L N
990  GACGCTTTCGGATGTTGGGGACCGGAAGCGGATGTATGCCCACTTCCACTTCATGACAGAGCTGCTGCTCTCGAGAGCCCACTG
    D A F A M V R D R K P D V S P N F H F M Q Q L L S F E S Q L
1080  CGCGTGAGACCGCGCTCGAGTTCTCGTGAGCTGCATTGCGCGGAGTCGAATTCATGACAGACCGGGCTTTATGGCGCGCATCTG
    R L R P G S R F S C S C I A P D C N C M Q T T G F M A A H L
1170  GCGAATGCTACCGGGTCTCGCAGACTCCCGCATGAGTTGATGCTGAGACCGCTCGGACACGGGTCTTAATGAGAACAGAGTGGC
    A N A T G V S P D S G I E F D R N T P S D T G L K *
1260  GGTAAAGTTTCGTGCTACCAACAGTCAGGAGATCCCATTTGCAGCGCGGCCACATCGCGGTGCGCATGTTCTGCGGGTTAAGCGG
    G A C A A G A T G T C T C G G C C A G C A C C A G T C G T C C A C G T G A C C A C C A C C G G G G T T C C G T G T G A A T G G T C A G C A T
1350  GACAAAGTGTCTCGGCGCAGCAGCAGCTGCTCCACGTCGACACGACCCGCGGAGCGGTTTCCGTGGTGAATGGTGCAGCAT
    C G G A T C A G G A T G G C C G A A G A G A C A T A T A C C A G G G A G A T A C A T G A G G A T G G G C C T G A T G G C G A A G C T T T C T A A C T T G G
1440  CGGATCAGGAGATGGCCGAGAAGACATCATACCGAGGGAGATACGATGAGGATGGGCGCTGAATGGCGAAGCTTTTCTAATCTGG
    T T C T C G C G A A A A C C C G G A T A A C C A C A T A T A T G A G C C G T T G T T A A T A G G G G T A C G A T T C A A C T T T A T A A T G T A C A A A
1530  TTTTGGAGAAATGCCAAGGTACACATATTGATTAAATACCAAGTAAATAACTTAATGTTTAAATACACTTTTAAATGAAGTGCAGACAAA
    G T T T G G A G A A T G C C A A C G T A C A C A T A T T G A T T A A T A C C A G T A A A T A A C T T A A T G T T A A T A C A C T T T T A A G T G A A G T G C A G A C A A A
1620  ATCCAAAGTTATCCACAGCGAACCACACTACGCACTATACAAAATGAGAATGAGAATAAAATATATAAAATATATATATATATATACGC
    A T C C A A G T T A T C C A C A G C G A A C C A C T A C G C A C T A T A C A A A A T G A G A A T G A G A A T A A A A T A T A T A A A T A T A T A T A T A T A T A T A T A C G C
1710  AGTGTACGAGTGTGTTGATCTAACTCTAAGCAGTCAAGAGAATAAATATATTAAGCCGAGCGAAAAATAACAATTAACCTGAGA
    A G T G T A C G A G T G T T G T A T C T A A C T C T A A G C A G T C A A G A G A A T A A A T A T A T T A A G C C G A G C G A A A A T A A C A A T T A A C T G A G A
1800  GCAGTTTTCTAGGTATATAGTTTTAGTTTTAGTTTTAAATTTTACTAACTCGAAATCGTTTATATATGAAGTTTGTATATTCGGAG
    C G A G T T T T C T A G G T A T A T A G T T T A G T T T A A T T T A C T A A C T C G A A T C G T T T A T A T A T G A A G T T T G T T A T A T C G G A G
1890  CGATTGGCGGATAGCGCTCCCTAATATAAATATATACCACTTAATTACCGAATGCCAAGTCATATAAGTGAAGTGAACATAAATC
    C G A T T G C G G A T A G C G C T C C C T A A T A T A A A T A T A C C A C T T A A T T A C C G A A T G C C A A G T C A T A T A A G T G A A G T G A A C T A A A A T C
2070  CGGAATATCGAAGAAATTTTAAAAATCGAAAAAAGAAAAATGAACACTAAAACAAAGAAAAATAATACAAACA

```

## 1b

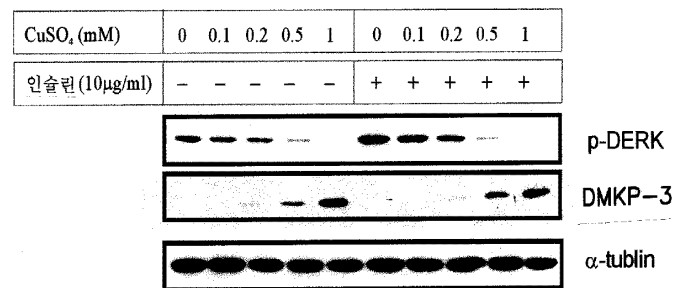


1c

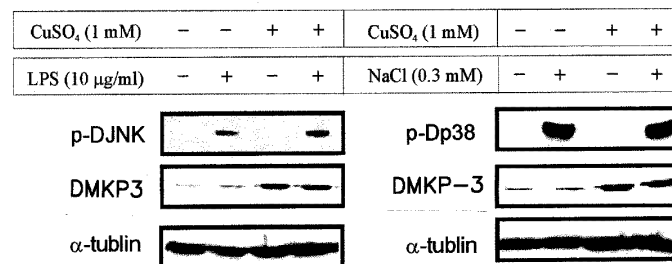
## 결합자리 모티프

DMKP3	(51)	PSIVL <b>RRL</b> AVG <b>K</b> IDLA
MKP3	(59)	PGIML <b>RRL</b> Q <b>K</b> GNLPVR
MKP4	(46)	LPALLL <b>RRLRR</b> GSLSV
PAC1	(50)	PWNALL <b>RRR</b> ARAGPP
PYST2	(1)	PGLML <b>RRLRK</b> GNLPIR
hVH2	(68)	RCNTIV <b>RRRAK</b> GSVSL
hVH3	(46)	NLNSVVL <b>RR</b> ARGGAVS
CL100	(47)	RFSTIV <b>RRRAK</b> GAKGA

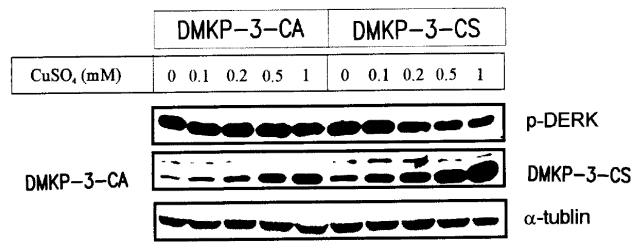
2a



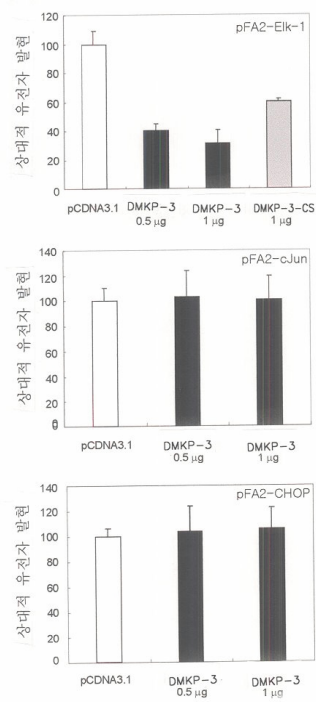
2b



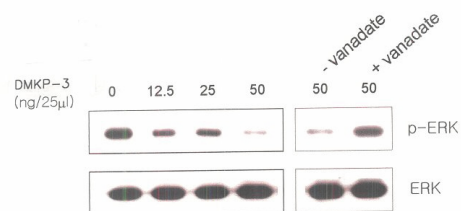
2c



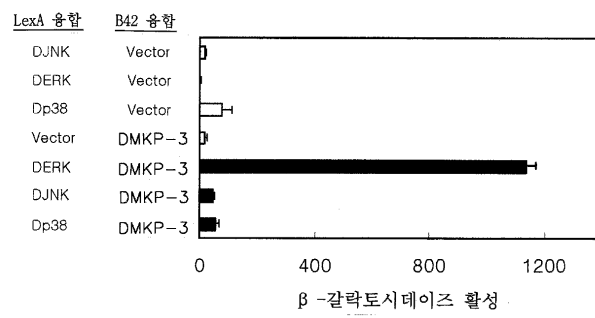
3a



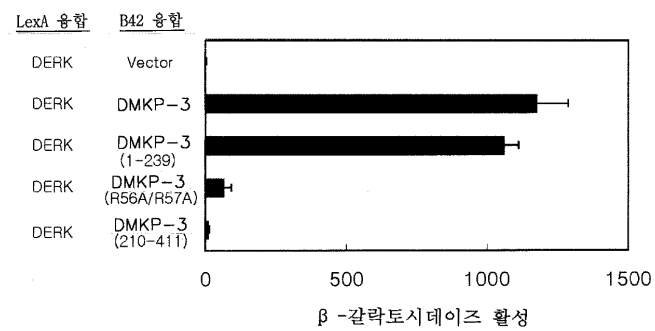
3b



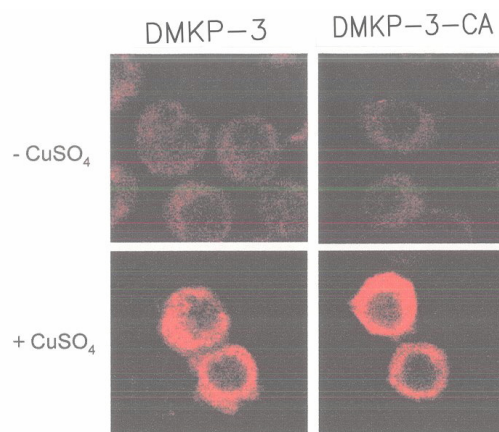
4a



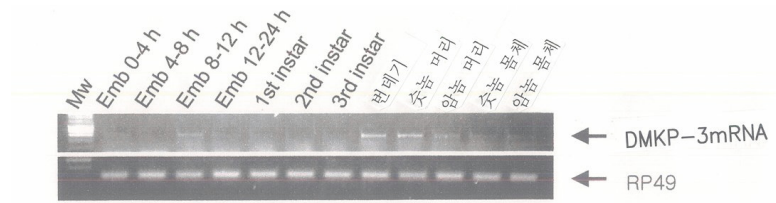
4b



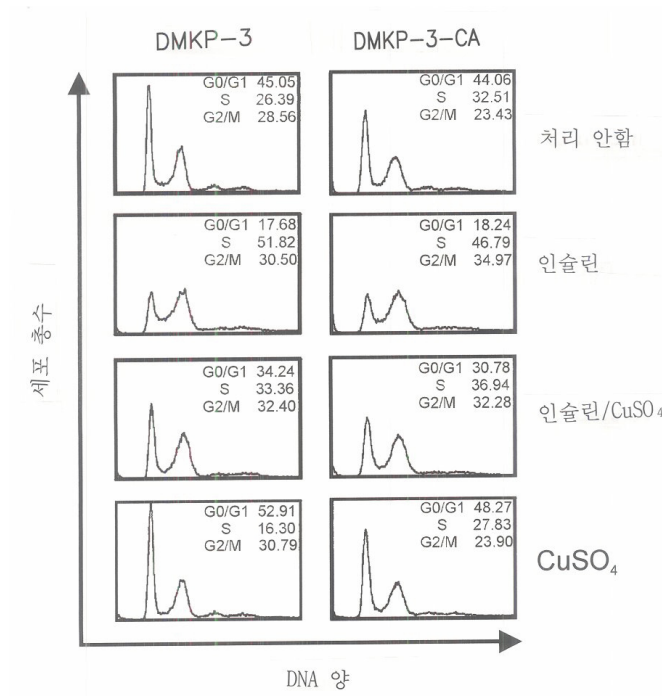
5a



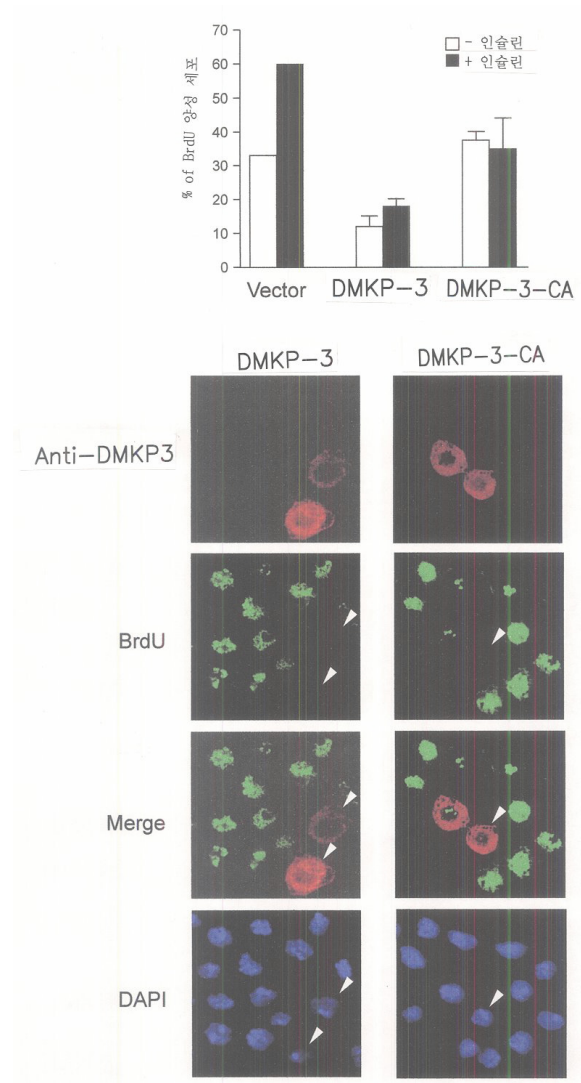
5b



6a

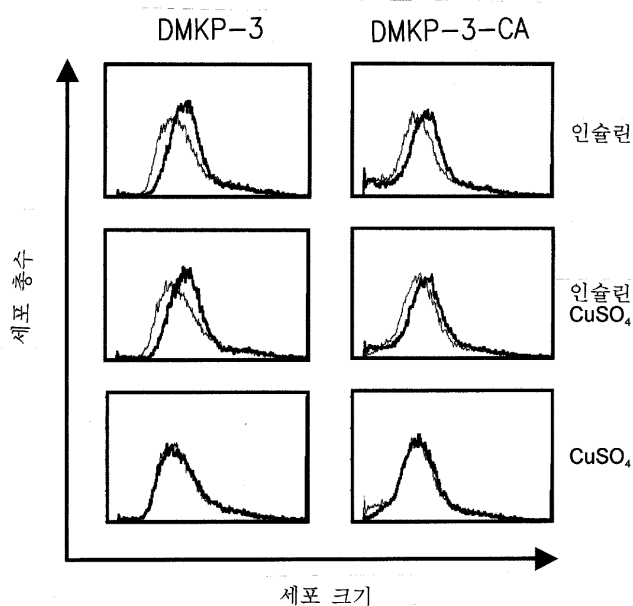


6b





7



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gacatcaagt atctgcagat tccgatcagc gatcactact cacaagattt ggccatacat      840
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 Gly Ser His Glu Tyr Ser Glu Ser His Ile Arg Gly Ala Val Asn Leu  
 35 40 45  
 Cys Ile Pro Ser Ile Val Leu Arg Arg Leu Ala Val Gly Lys Ile Asp  
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 65 70 75 80  
 Gly Tyr Lys Leu Cys Trp Phe Ile Leu Tyr Asn Gly Glu Gly Val Pro  
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 Leu Gly Asn Ala Thr His Ser Cys Asp Ser Glu Ala Leu Lys Lys Tyr  
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 245 250 255  
 Lys Glu Ser Gly Asp Ile Lys Tyr Leu Gln Ile Pro Ile Thr Asp His  
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 Tyr Ser Gln Asp Leu Ala Ile His Phe Pro Asp Ala Ile Gln Phe Ile  
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Glu Ser Gln Leu Arg Leu Arg Pro Gly Ser Arg Phe Ser Cys Ser Cys
          355          360          365
Ile Ala Pro Asp Cys Asn Cys Met Gln Thr Thr Gly Phe Met Ala Ala
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