

Product datasheet for MR218661

Pde4dip (NM_001110163) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Pde4dip (NM_001110163) Mouse Tagged ORF Clone
Tag:	Myc-DDK
Symbol:	Pde4dip
Synonyms:	4732458A06Rik; 9430063L05Rik; C87016; D3Bwg1078e; D130016K21Rik; mKIAA0454; Usmg4
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
ORF Nucleotide Sequence:	>MR218661 representing NM_001110163 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGTCTAATGGATATCGCACTCTGTCCCAGCACCTCAATGACTTGAAGAAGGAGAACTTCAGCCTCAAGC
TGCGCATCTACTTCTGGAGGAGCGCATGCAACAGAAGTATGAAGTCAGCCGGGAGGATGTCTACAAGCG
GAACATCGAGCTGAAGGTTGAAGTGGAGAGCCTGAAACGAGAAGTCCAGGACAGGAAACAGCATCTAGAT
AAAACATGGGCCGATGCAGAGGATCTCAACAGCCAGAATGAGGCAGAGCTCCGGCCAGGTTGAAGAAC
GGCAGCAGGAGACAGAACAGTATGAGCTCCTAGGGAACAAGATCCAGCTGCTGCAGGAGGAACCCAG
GCTAGCAAAGAATGAAGCCACAGAGATGGAGACTCTGGTGGAGGCAGAGAAGAGGTGCAATCTGGAGCTC
TCAGAGAGGTGGACGAATGCTGCCAAGAACAGGGAAGATGCAGCAGGAGACCAGGAGAAGCCTGACCAAT
ATTCTGAGGCACTGGCTCAGAGGGACAGGAGAATTGAAGAGCTGAGGCAGAGCTTGGCTGCTCAGGAGGG
GCTTGTGGAACAGCTGTCTCAAGAGAAACGACAACCTGTTACATCTGCTGGAGGAGCCAGCGAGCATGGAA
GTGCAGCCCGTGCTAAAGGGTTACCCACGCAACAGAAGCCGACTTGCATGAGACCCCTACAACCCAGC
CACCTGTGTCTGAGTCCCACCTGGCAGAAGTCCAGGACAAAATCCAGCAAACAGAGGCCACCAACAAGAT
TCTTCAAGAGAACTGAATGACCTGAGCTGTGAGCTGAAATCTGCACAGGAGTCATCTCAGAAGCAAGAT
ACGACAATCCAGAGCCTCAAGGAAATGCTTAAAGCAGGGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGA
TCGAAGGACAAAATGACACAATGGCAAAGCTTCGGGAAATGCTGCACCAGAGCCAGCTCGGACAGCTCCA
CAGCTCAGAGGGCATTGCCCTGCTCAGCAACAGGTAGCCCTGCTTGACCTTCAGAGCGCTCTATTCTGC
AGCCAGCTTGAATACAGAGGCTCCAGAGGCTGGTCCGGCAGAAAGAAGCCAGCTGGCGGATGGCAAGC
GATGTGTGCAGTTAGTGGAGGCTGCAGCCAGGAGAGAGACCAGAAGGAAGCTGCTTGGAAACATAA
CCAGGAATTACGAAAGGCTTTACAGCACCTCCAAGGAGAAGTGCACAGCAAGAGCCAGCAGCTCCATGTT
CTGGAGGCGGAAAATACAATGAGATTCGAACCCAGGGACAAAACATCCAACACCTAAGTACAGTCTGA
GTCACAAAAGAGCAGCTAATTCAGGAACTCCAGGAGCTCCTACAGTATCGTGACAACGCAGACAAAACCT
AGACACAAAACGAAGTGTTCCTTGAGAAATTCGGCAACGAATACAAGACCAGGCTGTTGCTCTAGAGCGG



[View online »](#)

GTCATAGATGAGAAGTTCTCTGCTCTAGAAGAAAAGGACAAGGAACTGCCGCAGCTTCGCCTAGCTGTGA
 GGGACCGAGACCATGACTTAGAGAGACTGCGTTGTGCCTGTCCGCAATGAAGCTACCATGCAGAGCAT
 GGAGAGTCTCCTGAGGGCCAGAGGCCTGGAAGTGGAGCAGTAACTGCCACCTGCCAAAACCTCCAGTGG
 CTGAAAGAAGAACTGGAAACCAATTTGGCCATTGGCAGAAGGAACAGGAGAGCATCATTAGCAATTGC
 AGACATCTCTACACGACAGGAACAAAGAAGTAGAGGATCTCAGCGCAACTCTGCTCTGTAACCTGGACC
 GGGTCAGAGTGAAGTAGCTGAGGAACTGTGCCAGCGCTTGCCAGCGAAAGGAACGGATGCTGCAGGACCTT
 CTGAGCGATCGGAACAAACAAGCCGTGGAGCAGAGATGGAGATCCAGGGGCTGCCACATCAGAACTGAGCA
 CCAGGGAGCAGGAAGACAGGCTGCTGCAGAAAAAATGGTCCAGGCCTTCATGGAAAGGAACTCAGAACT
 GCAGGCCCTGCGCCAGTATTTAGGGGGGAAGAACTAATGACATCGTCTCAGACGTTTCATCTCTAACCCAG
 CCAGCTGGAGTGACGTCCATCGGGCCTCACCACGGAGAGCAAACCGATCAAGTTCTATGCAGATGCCCT
 CTCGAGATGATAGCACCTCACTGACTGCTAGAGAGGAGGCCAGCATACCCCGGTCCACATTAGGAGACTC
 GGACACAGTTGCAGGGCTGGAGAAAGAACTGAGCAACGCCAAGGAGGAGCTCGAGCTCATGGCCAAAAAA
 GAAAGAGAAAGCCAGATGGAAGTGTCTGCCCTGCAGTCCATGATGGCCATGCAAGAGGAAGAGCTGCAGG
 TGCAGGCTGCTGACTTGGAGTCCCTGACCAGGAATGTGCAGATAAAAGAAGATCTCATAAAGGACCTGCA
 GATGCAACTGGTGCACCCTGAAGATATACCAGCCATGGAGCGTCTTACCAAGAGGTCTTACTTCTTCGG
 GAAAAAGTTGCTTCCGTGGAACCCAGGGTCAGGAAGTATCAGGGAACAAGAGACAGCAGTTGCTGCTGA
 TGTTAGAAGGACTAGTGATGAACGGAGTCGGCTCAACGAGGCCCTGCAAGCTGAGAGGCAACTCTACAG
 CAGCCTGGTCAAGTTCATGCCAGCCAGAGAACTCTGAGAGAGACGGAACTCTGCAGGTGGAAGTGGAA
 GGGGCCAGGTGTTACGCACTCGACTAGAAGAAGTTCCTTGAAGAAGCCTGGAGCGTTTAAAGCAGGCTGG
 AGAGCCTGGCCGCCATTGGAGGTGCTACTGCAGGCAATGAGACTGAAGATAACAAGCACGGAGTTCACAGA
 CAGCATTGAGGAGGAGGCTGCACACACCAGCCACCAGCAACTCATCAAGGTGGCTTTGGAGAAAAGCCTG
 GCAACCATGGAGACCCAGAACATATGTCTTCAGCCCCCTCCCCAGTAGGAGAGGACAGTAACAGGTGTC
 TTCAGGAGGAAATGCTCCACCTGAGGGCTGAAATCCACCAGCACTTAGAGGAGAAGAAAAGCTGAGGT
 GGAATCAAGGAGCTAAAGGCTCAAATTGAGGAAGCAGGATTCTCCTCTGTGTCCACATCAGGAACACC
 ATGCTGAGCCTTTGCCTTGAGAACGCAGAGCTGAAGGAGCAGATGGGAGAAGCAATGTCTGATGGATGGG
 AGGTCGAGGAGGACAAGGAGAAGGGCGAGGTGATGCTGGAGACAGTGGTGCCAAAGGGTGTCTGAATGA
 GAACAGCCTTCAGGCTGAGTTCAGGAAAGTCCAGGGGAACTCAAGAGTGCCTACAACATCATCAACCTC
 CTCAAAGAGCAGCTGCTCCTGAGGAGCTCGAAGGGAACAGTAAAGAGATGCCAGAGCTCCTTGTGCGCC
 TGGCCAGGAGGTGGACAGAATGAACACGGGTCTGCCTCCCTGGGGAAGCATCAGCACCAAGAGCAGGA
 GAATACGACCACCGAAGCCTGGCTCGAGACCCAGAGCCTCCCCTTGGGGCAGCCCTCTCAGTGGAT
 GGCTACCAACTGGAGAACAAGTCTCAGGCCAAGACTCTGGACACCAGCCAGAATTTAGCCTCCCGGGCT
 CCACCAAACACCTGCGTTCCAGCTGGCTCAGTGCAGACAACGATACCAAGATCTCCAGGAGAAGTTGCT
 CATCTCAGAAGCTACCGTGTTCGCCAGGCAAACAGCTGGAGAAGTACAGAGCCGATTTCAGTGAATCC
 CTGGTGAAACAGGACAGCAAGCAGATCCAGGTGGACCTTCAAGACCTGGGCTACGAGACTTGTGGCCGAA
 GTGAGAAATGAAGCTGAACGTGAGGAGACCACCAGCCCCGAGTGTGAGGAACACAATAACCTGAGGCCCGT
 GGTGCTCATGGAGGGGCTGTGCTCTGAGCAGGGGTACCTGGACCTGTCTTGGTTCAGCCACCTGCGAAG
 AAGCCCTTGGAGAACAAGCCGGGAAAGCAAGAGGAGTTCGTCACATGGAACCTCCGGACGACAGCTCTC
 TCCTGAGGAAGGACATCCGAGACCTGAAAGCCCAGCTACAGAATGCCAACAAAGGTATTGAGAACCTGAG
 GAGCCGGGTCCGGTCCCTGTCTGCCACAAGCGACTACTCATCAAGTCTGGAGAGACCCCGCAAGCTGAGA
 GCCGTGGCGACCCCTTGGGGGGCTTACCCACAGCGTGACTGATGAAGATGAGGGGTGGCTGTCAGATG
 GCACTGGGGCTTTTTACCTCCAGGACTCCAGGCCAAAAAAGATCTAGAGAGTCTCATCCAGCGAGTATC
 CCAACTGGAGGCCAGCTCCCCAAAAGTGGACTAGAAGGGAAGCTGGCCGAGGAGCTGAGGTGTGCCTCG
 TGGCCTGGAAAAATGATTCCTTGATTGAGGATCAGGCCCGAGAATCATATCTGCGTCAAAAGATAC
 GAGAAGGGAGGGGATATGTTATCTTCCACCAACATGCAAAAGATACTGTAATAATCTTTGAGGACCT
 CCTTAGGAGCAACGACATTGACTACTACCTGGGCCAGAGCTTCCGGGAGCAACTAGCCAGGAGGTCAG
 CTGACGGAGAGGCTCACCAGCAAACCTCAGCACAAGGATCATAAGAGTAAAAGGAAGAAGCTGGGCTTG
 AGCCACTGGCCCTCAGGCTCAGCAGGAACTACAGGAGAAAGAGAAAGTAATTGAGTCTCGAGGCCAA
 GCTGGATACCCGGTCTCTCTACCCCAAGCAGCCAGCCGTGTCTGACTCCACCGCTCCGCCAGCACC
 ACATCCTTCTGTGCGATGACATAGAAGCCTGCTCTGACATGGACGTAGCCAGCGAGTACACACACTACG
 ACGAGAAGAAACCTCACCCAGTCACTCAGCAGCCAGTGCATCTCAGGGGCTTAAGGGCGAGTCCAGCAG
 CAGCCCCATCAGCTTGCCAACTCCCAGAATCCCCCTAAGGAGGCCAGCCAGGCCATCCAGGCTTTCAC
 TTTCACTCCATACCAAGCCGGCTAGCCTTCCAGACACCAATGCACTCCGCTCTGCCAGCTTTGTGC

CTTTCAGCCCCTCCGGGCTCCCTTCTGGGTTGCTGTGAGACACCGATGGTGTCTTGGCTGAGGCTCA
ACAAGAGCTGCAGATGCTGCAGAAGCAGCTGGGAGAAAGTGTAGCATTGCCCTCCCGCTCCACATCC
ACGCTGCTCAGCAACCAGACTGAAGCTAGCTCTCCCCACTACATCAACCTGCCAGCCCCACACTCCGA
CAAGGAGCACCATAGAGCTGGGACGAATCTGGAGCTGGATACCTGGGAAGCAGCGGCCAGTGGGACAT
GATGAGGCCCCAGAAAGGGAGCGTCTCTGGGAGCTGTCTCAGGCTCCTCGATGTACCAGCTTAACTCC
AAGCCCACTGGCGTGACCTGTTGGAAGAGCATTTAGGTGAGATCCGGAACCTGCCCAGCGCTGGAGG
AGTCCATATGTGTCAATGACCGGCTGCGGGAGCAGCTGCAGCACAGGCTCAGTCCACTGCCCGAGAAAA
CGTTCCACCTCTCACTTCTACAGTCAGGGCTGGAGTCCATGCCTCAGCTCTACAATGAGAACAGAGCC
CTAAGGGAAGAAAACCAAGCCTGCAGACACGGCTCAGCCATGCTTCCAGAGGACTCCAGGAAGTGG
ATCACCTGCGGGAGGCTCTGCTCTCCTCCAGATCCCAGCTTCCAGGAGCTGGAGAAGGAGCTGGAGCAGCA
GAAGGCTGAGCGGCAGCAGCTCCTGGAAGACTTGCAAGGAGAAGCAGGATGAGATCGTGCAGTTCCGAGAG
GAGAGGCTGTGCTCCAGGAAAACAACCTCAGGCTGCAGCACAAGCTGGCCCTCTGCAACAACAGTGG
AAGAGAAACAGCAGCTCTCCCTGTCCCTGCAGTCAGAGCTCCAGATCTACGAGTCCCTCTGTGAAAATCC
CAAGAAGCCCTTAAAGCTTTTAGCCTAGATTCCTGTACCAAGTCCCGGTGAGTTAAGCTGCCTGGT
GCAGAGATTCGAGCTCTGAGAGGACAGTTGGAGCAGAGCATTGAAGTGAACAACCGTCTTCGGCTACAGC
TGGAACAGCAGATGGATCGTGGTGTGGCAAAGCCAGTCTCGGCCCATCGCTGTTGGCCAGAGCTTCCC
AGACAAGGCAGAGCCAGCAAACCTGCACCAAGGTTCCGCTGCTTCCCCTCCAGTGCGGGACGTTGGCTTG
AATTCCCCAGCCATGGTCTCCCAACTCTTCTGCTCCGCTCCTGGCTCAGACCATGCCATTGTCACCA
GGACAAACAACGAGCTAAGTTCAGATGATTCTGCAGCAATGAAGAACCCTCCCAAGCTGGAGGTCGATGC
TACCGATGGCCATTTGCCAACAACACGGAAGACACGTCATCGGCCATGTTGATGACTACGATGCTCTG
CAACAGCAGATTTGGGAAGGCAAGCTGCTGATTCAAAAGATACTGTCTCTCATGAGGTCAGCACGCAGCA
TCCCTGGGCAAGAAGCTCAGGACACAGAGGCACCAGGTAACATAAGTGCCATGAGCTTCGGAGCAGCGC
CAAGGCTCTGAACCACGCCCTAGAAGAGTCAACATCCCTACTCAACATGTTCTGGAGAGCAGCCTTGCCA
AACACTCATGGTCTGTACTGGTAGGCAAAGAGGGACAACCTGATGGAGAAAGAAGCTTGGACCTGCGAG
CCCAAGTATCCCAACAGGAACAGATCCTTCAGAACACTGCTGCACGCTCTGAAGAGGGCAACCAGAGGAA
GAAAAGCATGGAACAATTCATCGTGTAGCCATTTGACTAGGACCCATGATGTCTTGAAGAAAGCGGGACT
AATTTAGAGATGAAATCCTTCAGGGCTCTGACGTGCACTCCGGCCTTG

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAATGATATCCTGGATT
ACAAGGATGACGACGATAAGGTTTAA

Protein Sequence: >MR218661 representing NM_001110163
 Red=Cloning site Green=Tags(s)

```

MSNGYRTLSQHLNDLKKENFSLKLRIFYLEERMQQKYEVSRQDVEYKRNIELKVEVESLKRELQDRKQHL
KTWADAEDLNSQNEAELRRQVEERQQE TEHVYELLGNKIQLLQEEPR LAKNEATEMETLVEAEKRCNLEL
SERWTNAAKNREDAAGDQEKPDQYSEALAQDRRIEELRQSLAAQEGLEQLS QEKRQLLHLLLEEPASME
VQPVPKGLPTQQKPDHETPTTQPPVSESHLAELQDKIQQTEATNKILQEKLNDL SCELKSAQESSQKQD
TTIQSLKEMLKSRSESETEEL YQVIEGQNDTMAKLREMLHQSQLGQLHSSEGIAPAQQQVALLDLQSALFC
SQLEIQRLQRLVRQKERQLADGKRCVQLVEAAAQEREHQEA AWKHNQELRKALQHLQGELH SSKSQQLHV
LEAEKYNEIRTQGQNIQHL SHSLSHKEQLIQELQELLQYRDNADKTLDTNEVFLEKLRQRIQDRAVALER
VIDEKFSALEEKDKELRQLRLAVRDRDHLERLRCVLSANEATMQSMESLLRARGLEVEQLTATCQNLQW
LKEELETKFGHWQKEQESI IQQLQTSLHNRKEVEDLSATLLCKL GPGQSEVAEELCQRLQRKERMQLDL
LSDRNKQAVEHEMIEI QGLLQSMGTREQERQAAA EKMQVAFMERNSELQALROYLGGKELMTSSQTFISNQ
PAGVTSIGPHHGEQTDQGSMMQPSRDDSTSLTAREEASIPRSTLGDSDTVAGLEKEL SNAKEEELMAKK
ERESQMELSAQSMAMQEEELQVQAADLES LTRNVQIKEDLIKDLQMQLVDPEDIPAMERLTQEVLLL R
EKVASVEPQGEVSGNKRQQLLLMLEGLVDERSRLNEALQAERQLYSSLVKFHAQPENSERDGT LQVELE
GAQVLRTRLEEVLRSLERL SRIE SLAAIGGATAGNETEDTSTEF TDSIEEEAAHTSHQQLIKVALEKSL
ATMETQNICLQPPSPVGEDSNRCLQEEMHLRAEIHQHLEEKRAEVELKELKAQIEEAGFSSVSHIRNT
MLSLCLENAE LKEQMGEAMSDGWEVEEDKEKGEVMLETVVAKGC L NENSLQAEFRKVQGGKLSAYNIINL
LKEQLLLRSSEGNKEMPELLVRLAREVDRMNTGLPSL GKHQHQEQENTTARPGSRPQSLPLGAAL SVD
GYQLENKSAQDQSGHQPEFSLPGSTKHLRSQLAQCRQRYQDLQEKLLI SEATVFAQANQLEKYRAVFS ES
LVKQDSKQIQVDLQDLGYETCGRSENEAEREETT SPECIEHNLRPVVLM EGLCSEQGYLDPVLVSPPAK
KPLENKPGKQEEFRAHGT PDDSSLLRKDIRDLKAQLQNANKVIQNLRSRVRSL SATSDYSSSLERPRKLR
AVATLEGASPHSVTDEDEGWLSDGTGAFYPPGLQAKKDLES LIQRVSQLEAQLPKTGLEGKLAEEELRCAS
WPGKYDSL IQDQARELSYL RQKIREGRGICYLLTQHAKD TVKSFEDLLRSNDIDYYL GQSFREQLAQQGQ
LTERLTSKLS TKDHKSEKEEAGLEPLALRLSRELQEKEK VIEVLQAKLDTRSLSPSSHAVSDSHRSAST
TSFLSDDIEACSDMDVASEYTHYDEKKPSPSHSAASASQGLKGESS SPSISLPTPQNPPKEASQAHPGFH
FHSIPKPASLSQTPMHSALPSFVPFSPSGPPLLGCCETPMVSLAE AQQELQMLQKQLGESVSIAPPASTS
TLLSNQTEASSPHYINPAQPHTPTRSTIELGRILEPGYL GSSGQWDMMPQKGSVSGELSSGSSMYQLNS
KPTGADLLEEHLGEIRNL RQRLEESICVNDRLREQLQHRLS STARENGSTSHFYSGGLE SMPQLYNENRA
LREENQSLQTRL SHASRGHSQEVDHLREALLSSRSQ LQELEKELEQQKAERQQLLEDLQEKQDEIVQFRE
ERLSLQENNSRLQHKLALLQQQCEEKQQLSLSLQSELQIYESL CENPKKALKAFSLDSCHQVPGEL SCLV
AEIRALRGQLEQSIEVNNRLRLQLEQQMDRGAGKASLGP IAVGQSFDPKAEPANLHQGSAASPPVRDVGL
NSPAMVLPNSSCSAPGSDHAI VTRTNNELSSDDSAAMKNPPKLEVDATDGPFANKHGRHVIGHVDDYDAL
QQQIGEGKLLIQKILSLMRSARSIPGQEAQDTEAPGNISAH ELRSSAKALNHAALEESTSLNMFWR AALP
NTHGPVLVGKEGQLMEKELLDLRAQVSQQEQILQNTAARLKRANQRKKSMEQFIVSHL TRTHDVLK KART
NLEMKSFRAL TCTPAL
  
```

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Chromatograms: https://cdn.origene.com/chromatograms/mm9041_a10.zip

Restriction Sites: Sgfl-Mlul

Cloning Scheme:



ACCN: NM_001110163

ORF Size: 6978 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001110163.1](#), [NP_001103633.1](#)

RefSeq Size: 8289 bp

RefSeq ORF: 6981 bp

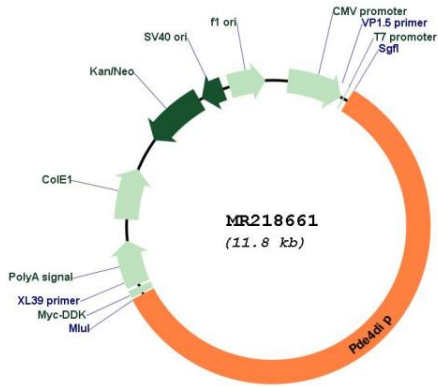
Locus ID: 83679

Cytogenetics: 3 42.28 cM

MW: 262.8 kDa

Gene Summary: Functions as an anchor sequestering components of the cAMP-dependent pathway to Golgi and/or centrosomes (By similarity).[UniProtKB/Swiss-Prot Function]

Product images:



Circular map for MR218661