

Product datasheet for SC323670

AKT3 (NM_005465) Human Untagged Clone

Product data:

OriGene Technologies, Inc.

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| Product Type: | Expression Plasmids |
|------------------------------|---|
| Product Name: | AKT3 (NM_005465) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | AKT3 |
| Synonyms: | MPPH; MPPH2; PKB-GAMMA; PKBG; PRKBG; RAC-gamma; RAC-PK-gamma; STK-2 |
| Mammalian Cell Selection: | None |
| Vector: | pCMV6-XL4 |
| E. coli Selection: | Ampicillin (100 ug/mL) |



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| Fully Sequenced ORF: | >OriGene ORF within SC323670 sequence for NM_005465 edited (data generated by NextGen Sequencing) |
|----------------------|---|
| | ATGAGCGATGTTACCATTGTGAAAGAAGGTTGGGTTCAGAAGAGGGGAGAATATATAAAA |
| | |
| | AACTGGAGGCCAAGATACTTCCTTTTGAAGACAGATGGCTCATTCAT |
| | |
| | TTAATGAAAACAGAACGACCAAAGCCAAACACACTTTATAATCAGATGTCTCCAGTGGACT |
| | ACTGTTATAGAGAGAACATTTCATGTAGATACTCCAGAGGAAAGGGAAGAATGGACAGAA |
| | GCTATCCAGGCTGTAGCAGACAGACTGCAGAGGCAAGAAGAGAGAG |
| | CCAACTTCACAAATTGATAATATAGGAGAGGAGGAGGAGGATGGAT |
| | AAAAGAAAGACAATGAATGATTTTGACTATTTGAAACTACTAGGTAAAGGCACTTTTGGG |
| | AAAGTTATTTTGGTTCGAGAGAAGGCAAGTGGAAAATACTATGCTATGATGATTCTGAAG |
| | AAAGAAGTCATTATTGCAAAGGATGAAGTGGCACACACTCTAACTGAAAGCAGAGTATTA |
| | AAGAACACTAGACATCCCTTTTAACATCCTTGAAATATTCCTTCC |
| | TTGTGTTTTGTGATGGAATATGTTAATGGGGGGCGAGCTGTTTTTCCATTTGTCGAGAGAG |
| | CGGGTGTTCTCTGAGGACCGCACACGTTTCTATGGTGCAGAAATTGTCTCTGCCTTGGAC |
| | TATCTACATTCCGGAAAGATTGTGTACCGTGATCTCAAGTTGGAGAATCTAATGCTGGAC |
| | AAAGATGGCCACATAAAAATTACAGATTTTGGACTTTGCAAAGAAGGGATCACAGATGCA |
| | GCCACCATGAAGACATTCTGTGGCACTCCAGAATATCTGGCACCAGAGGTGTTAGAAGAT |
| | AATGACTATGGCCGAGCAGTAGACTGGTGGGGCCTAGGGGTTGTCATGTATGAAATGATG |
| | TGTGGGAGGTTACCTTTCTACAACCAGGACCATGAGAAACTTTTTGAATTAATATTAATG |
| | GAAGACATTAAATTTCCTCGAACACTCTCTTCAGATGCAAAATCATTGCTTTCAGGGCTC |
| | TTGATAAAGGATCCAAATAAACGCCTTGGTGGAGGACCAGATGATGCAAAAGAAATTATG |
| | AGACACAGTTTCTTCTCTGGAGTAAACTGGCAAGATGTATATGATAAAAAGCTTGTACCT |
| | CCTTTTAAACCTCAAGTAACATCTGAGACAGATACTAGATATTTTGATGAAGAATTTACA |
| | GCTCAGACTATTACAATAACACCACCTGAAAAATATGATGAGGATGGTATGGACTGCATG |
| | GACAATGAGAGGCGGCCGCATTTCCCTAATTTTCCTACTCTGCAAGTGGACGAGAATAA |
| | Clone variation with respect to NM_005465.4 530 a=>t |
| 5' Read Nucleotide | >OriGene 5' read for mutant NM_005465 unedited |
| | CCGCCCGTTGTAGCAATGGGCGGTAGGCGTGTACGGTGGGAGGTCTATATAAGCAGAGCTCGTTTAGTGA |
| Sequence: | ACCGTCAGAATTTTGTAATACGACTCACTATAGGGCGGCCGCGAATTCGGCACGAGGCCAAACCCTAAAG |
| | CTGATATCACAAAGTACCATTTCTCCAAGTTGGGGGGCTCAGAGGGGAGTCATCATGAGCGATGTTACCAT |
| | TGTGAAAGAAGGTTGGGTTCAGAAGAGGGGAGAATATATAAAAAACTGGAGGCCAAGATACTTCCTTTTG |
| | AAGACAGATGGCTCATTCATAGGGATATAAAGAGAAAACCTCAAGATGTGGATTTACCTTATCCCCCTCAA |
| | CAAACTTTTTCAGTGGCAAAAATGCCCAGGTTAATGAAAACCAGACCGACC |
| | TAATCCAAATGTTCTCCATTGGACTACTGGTTATAGAAGAAAAACATTCATGGGAAAATACTCCAAAGGA |
| | AAGGGAAAATGGGGACAAAACCTTTCCAGGCCTGTACCAAAACAAAC |
| | TGATTTGTATCCCAACTTCAAAATTTGAAAATATAGGAAAGAAA |
| | ATAAGGAGAGCACATGTGAGATTTAGAATATTTGAAAACTATAGGAAAGCGCCCTTTTGGGGAAGTTTTT |
| | TTGGTTTCAGAAGGGGCTGGTGGAAAACTCTTTGTGCTAGGGAGTTTTCAGAAAAAAATGTTTATTTGG |
| | GCAAGAGGAGAGAGAGTGTGCACACCTCTCCTGTGAGACCGCGAGATTTATAGACACACCTCCACCCCTC |
| | TGTTTAACACCCTGCGAGAATTCCTCCACACAACAACACACAC |
| | TGGCGAGCCAGTTTTCTATCTTTGAAGAAGAGGCGTGTGTTCTTAGACGACACCACCACTATATAGAG |
| | GGCTGCTGATACTACTTACTCGCGAAGTAGTAGTGTTCTGT |
| | |

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GRIGENE AKT3 (NM_005465) Human Untagged Clone – SC323670

| Kinase Domain Sequence: | >SC323670 kinase domain raw sequence. By performing <u>BLASTX</u> analysis with this sequence against NCBI refernce protein database, you can confirm the presence of the kinase- deficient mutation CYTTGMGCATGGGCGGTAGGCGTGTACGGTGGGAGGGTCTATATAAGCAGAGCTCGTTTAGTGAACCGTCA GAATTTTGTAATACGACTCACTATAGGGCGGCGCGCGAATTCGGCACGAGGCCAAACCCTAAAGCTGATAT CACAAAGTACCATTTCTCCCAAGTTGGGGGCTCAGAGGGGAGTCATCATGAGCGATGTTACCATTGTGAAA GAAGGTTGGGTTCAGAAGAGGGGAGAATATATAAAAAACTGGAGG |
|----------------------------|---|
| Restriction Sites: | Please inquire |
| ACCN: | NM_005465 |
| Insert Size: | 5000 bp |
| OTI Disclaimer: | Due to the inherent nature of this plasmid, standard methods to replicate additional amounts of DNA in E. coli are highly likely to result in mutations and/or rearrangements. Therefore, OriGene does not guarantee the capability to replicate this plasmid DNA. Additional amounts of DNA can be purchased from OriGene with batch-specific, full-sequence verification at a reduced cost. Please contact our customer care team at <u>custsupport@origene.com</u> or by calling 301.340.3188 option 3 for pricing and delivery. |
| | 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. <u>More info</u> |
| OTI Annotation: | This kinase-deficient mutant clone was generated by created by site-directed mutagenesis from the corresponding wild-type clone. See details in "Application of active and kinase-deficient kinome collection for identification of kinases regulating hedgehog signaling." <u>Cell.</u> <u>2008 May p536-548.</u> |
| 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: | Centrifuge at 5,000xg for 5min. Carefully open the tube and add 100ul of sterile water to dissolve the DNA. Close the tube and incubate for 10 minutes at room temperature. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom. 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: | <u>NM 005465.3, NP 005456.1</u> |
| RefSeq Size: | 3588 bp |
| RefSeq ORF: | 1440 bp |
| Locus ID: | 10000 |
| | |

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| UniProt ID: | <u>Q9Y243</u> |
|-------------------|--|
| Cytogenetics: | 1q43-q44 |
| Domains: | pkinase, S_TK_X, TyrKc, PH, S_TKc |
| Protein Families: | Druggable Genome, ES Cell Differentiation/IPS, Protein Kinase |
| Protein Pathways: | Acute myeloid leukemia, Adipocytokine signaling pathway, Apoptosis, B cell receptor signaling pathway, Chemokine signaling pathway, Chronic myeloid leukemia, Colorectal cancer, Endometrial cancer, ErbB signaling pathway, Fc epsilon RI signaling pathway, Fc gamma R- mediated phagocytosis, Focal adhesion, Glioma, Insulin signaling pathway, Jak-STAT signaling pathway, MAPK signaling pathway, Melanoma, mTOR signaling pathway, Neurotrophin signaling pathway, Non-small cell lung cancer, Pancreatic cancer, Pathways in cancer, Progesterone-mediated oocyte maturation, Prostate cancer, Renal cell carcinoma, Small cell lung cancer, T cell receptor signaling pathway, Tight junction, Toll-like receptor signaling pathway, VEGF signaling pathway |
| Gene Summary: | The protein encoded by this gene is a member of the AKT, also called PKB, serine/threonine protein kinase family. AKT kinases are known to be regulators of cell signaling in response to insulin and growth factors. They are involved in a wide variety of biological processes including cell proliferation, differentiation, apoptosis, tumorigenesis, as well as glycogen synthesis and glucose uptake. This kinase has been shown to be stimulated by platelet-derived growth factor (PDGF), insulin, and insulin-like growth factor 1 (IGF1). Alternatively splice transcript variants encoding distinct isoforms have been described. [provided by RefSeq, Jul 2008] Transcript Variant: This variant (1) represents the longest transcript and encodes the longer isoform (1). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments. |

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