

Product datasheet for **RG215838**

ALPK2 (NM_052947) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	ALPK2 (NM_052947) Human Tagged ORF Clone
Tag:	TurboGFP
Symbol:	ALPK2
Synonyms:	HAK
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)
ORF Nucleotide Sequence:	>RG215838 representing NM_052947 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGAAAGACTCCGAAGGGCCCCAGAGGCCCCCGCTGTGTTTTTATCTACATTGCTTTCCAGAAGGTTCC
CTGAGAAGTCAGACGCTGTGCTTCGCTGCATAATATCTGGTCAGCCCAAGCCAGAGGTAACCTTGGTATAA
GAATGGTCAGGCCATCGATGGGAGTGGCATTATTTCCAATATGAATTCTTTGAGAATCAGTATATTCAT
GTGTTACATCTCTTGTGTACCAAAAATGATGCTGCTGTCTATCAAATCTCGGCTAAAACTCTTTT
GAATGATCTGTTGTTCTGCTTCCGTTGAGGTTGAGTGCTCATCAGAGAACCCACAATTGTCTCCTAACCT
GGAAGATGACAGGGACAGGGGTTGAAACATGAAACAGGGACACATGAAGAAGAAAGGGCAAATCAGATT
GATGAGAAGGAACATCCTTATAAGGAAGAAGAAAGCATCTCCCCGGGCACTCCCAGGTACGCTGACTCCT
CCCCCTCCAAATCCAACCATCACTCTCCCTCCAGTCATTGGGCAATCTTGACATTAGTGTGTCAGTTC
TGAAAAATCCTTTGGGTGTTAAAGGAACAAGGCACACTGGAGAGGCTTATGATCCAAGTAACACAGAAGAA
ATTGCAAATGGGTTGCTTTTTCTTAATCAAGTCATATTTATGAAAAACAAGACAGATGTTGCCACAAGA
CAGTGCATTCCATGGCATCAAAGTTCACGGATGGTGACCTGAACAATGATGGTCCATGATGAAGGCTT
ACGCTCTAGTCAGCAAAATCCCAAAGTACAGAAATACATTAGCTTCAGCCTCCCGCTATCTGAGGCAACT
GCACACATTTACCCAGGTGACAGTGGCGTGCCCAACAACAACCCAGCCACAGCTTTCCAGTGAAGACT
CTGACAGTGACTATGAACCTTGCCAGAGATAACCCTAACCTACACCGAGGAGTTTTTCAGATGATGACCT
GGAGTATCTGGAATGTTCTGATGTTATGACGGATTACTCTAATGCAGTTTGGCAAAGGAACCTGCTGGGG
ACTGAGCATGTTTTTTTATTAGAAAAGCGATGACGAAGAGATGGAATTCGGTGAGCATTGCCTGGGTGGGT
GTGAGCATTTCTCAGTGAATGGGTTGTGGGTCTCGGTGTCGGGTGACGCTGGGCCTATGTTGCCAC
TGCTGGCTTCTGTGGTCATCACTACAACCCCAAGAAAGTTGGGGTGAGGAGCAGCAGAGTCTCCAAGCAC
GGTCCCTCATCCCACAACAGGGATGACTCTCATTTTGGGACCTCACCAGGATGGAACGCTTTCAGTGA
CAGAACAGGGGAGATATAAACTCCCCTGCTCCCGAGGCTGCTGAAAATGATTACAGGAATTCAGG
AGAAACCAGAGACAGCCCAAGCAAGAGAGGAATTTGCCAGTGACAATCTGCTCAACATGGATGAATCA



GTAAGAGAGACAGAGATGAAGCTCTTGTCTGGTGTGAGTCAGAAAACCTCAGGGATGAGCCAGTGTGGGAGA
 CCGCAGCTGACAAGAGAGTGGGGGAAAGGACTTATGGAGCAAGAGGGGTTCAAGGAAATCTGCCAGGGT
 GAGGCAGCCGGGAATGAAGGGAAATCCCAAGAAGCCGAATGCCAACCTGAGAGAAAGTACAACAGAAGGT
 ACCCTTCATCTCTGCTCTGCCAAAGAATCTGCTGAGCCCCACTAACCCAGAGTGATAAAAGAGAGACTT
 CTCACACCACAGCAGCAGCGACTGGTCGGAGTTCCCATGCTGATGCAAGAGAATGTGCTATTTCAACCCA
 GGCAGAGCAAGAAGCAAAAACCCCTCAAACCTCAACAGACTCAGTCTCAAAGAAGGCAACACAAATTGC
 AAGGGAGAAGGCATGCAAGTTAATACTCTATTTGAAACAAGCCAGGTTCCAGACTGGAGTGATCCTCCTC
 AGGTACAAGTTCAGGAAACAGTCAGAGAGACAATCTCTTGACGCCAGATGCCAGCTTTCTCAGAGCCTGC
 TGGGGAGGAGTCCCCATTCACTGGGACCAACAATTTCTTCTCAAACCTAGGAGGGGTCCACAAGGAA
 AATGCATCATTAGCTCAACACTCGGAGGTCAAACCCTGTACCTGTGGTCCACAGCATGAAGAAAAACAAG
 ACAGAGATGGCAACATACCTGACAATTTAGGGGAAAGACCTAAAATATGAGCAGAGCATCTCAGAAGCCAA
 TGATGAGACTATGTCCCAGGTGTGTTCTCAAGGCATCTCCCAAGGATGCTCGTGTGACTTCAGGGAG
 CCTGTGGTGTCTCTGTTGCTTCCCTGAACCCACAGATACTGCCCTCACCTGGAAAATGTGTGTGATG
 AGCCAAGGGACAGAGAAGCAGTGTGTGCAATGGAGTGTGTTGAGGCTGGTGACCAAGGAACGTGTTTTGA
 TACCATAGATTCTTGTGGGAGACCAGTTGATAAATATTCGCCTCAAGAAATTTGCTCTGTAGATACG
 GAACTGGCAGAAGGTCAAACAAAGTATCTGATTTATGTTCTTCTAATGACAAGACTGGAAAGTCTTTT
 TTCAGACACAAGTGTCTGAGACTTCAGTGTCTACGTGCAAAAAGCAGCAAGGACGGCAACTCAGTCATGTC
 CCCTCTTTTTACCACTTTTACCTTGAACATTTACACACAGCTAGTGAAGGTGCCACAGGAGAAAAAT
 CTAGCCAAGGTGGAGAATTCACCTACCCACTGGCCTCCACAGTACATGCTGGCCAGGAGCAGCCAAGCC
 CCAGCAACTCAGGAGGGCTTGATGAAACACAGCTCCTTTCTTCTGAGAACAATCCTTTAGTGCAATTTAA
 AGAAGGAGGTGACAAGAGCCCCAGTCTAGTGCCGACAGACACCACAGCCACACCAGCCAGTTATAGTTCA
 ATTGTGAGTTTTCTTGGGAGAAGCCAAACAACATTAACGTCTAATAATGAGTGTCTTCAAGCGACCAGGG
 AGACTGAGGACACATCAACTGTTACCATTTGCCACCGAAGTCCACCCAGCCAAATACCTTGTGTGCAAT
 TCCTGAGGACAAGCATGCAGGTGGCACTGAGGAGAGGTTCCCTCGTGATCCCATGAAAAGGTTTCCCAA
 TTTCTTCCCAAGTGCAGTTGGATCATATTTAAGTGGTGTACCATCAAATCTACAAAAGACTACTTT
 GCAGGGCACCCAGTGTGCCAGGAGTCCCACACCATGTCCTGCAGCTCCCAGAGGGAGAGGGTTTTCTGCAG
 TAATTCCTCTTTCAGGTTGATAACCTGTCTGGAGATAAGAGCCAGACTGTGGACAGAGCAGACTTTAGG
 AGCTATGAAGAGAATTTCCAAGAAAGAGGAAGTAAACAAGCAGGGGTCCAGCAGCAGAGCCTGTCCC
 AGCAGGGTTCTCTTCTGCACCTGATTTCAACAAAGTTGCCTACGACATCTGCTGCACAAGAGGAAAG
 AAACCTGGTGCCACGGCCACTCACCCGAAGCTCTAGGGAAGGAGCAGGGCAGCGCTCAGGTTGGGGG
 ACGAGGGTCTCCGTGGTGTGAACTGCTGGGAAGAAGACAGTCAAGGCTCTGAGCAACGTTCCATCTC
 TCTCTGATATCCTTTTGAAGAGTCTAAAGAATATAGACCTGAAAATGGGAGGCAGGCAACAAGCTGAA
 GATTATAACTCTAGAGGCTTCCGCTTCTGAAATCTGGCCACCACGACAACCTGACAATTTCTGAGAGCAAG
 GCATCAGACGGTGGTCTCATAATTCCTGACAAGGTCTGGGCTGTACCTGATAGTCTAAAGGCAGATGCTG
 TTGTGCCGTAATGGCCCCCTCTGAAATAGCAGCATTGGCTCACAGTCCAGAGGATGCTGAGTCAGCCCT
 TGCTGATAGCAGAGAAAGCCATAAAGGGCAAGAGCCACCATCAGTGTACATTGGAGAAGTCTTTCTTCC
 CGGGGTTTCAGCAACCCAGACTCCTGGAGTCACTCCGTGGACCCTGTAGATGAAAAGGAGTTATCTGTCA
 CAGATTCAGTGTGAGCGGCTTCTGAACTGGAGGGAAGGAAAATGTTAACAATGTGAGTCAAGACCAGGA
 GGAAAAACAACCTCAAGATGGATCACACTGCCTTCTTAAAAAGTTCTGACCTGCCCTAAAATCCTAGAG
 TCCTCTGTAGATCCCATTTGATGAGATAAGTGTGATAGAGTACACCAGGGCTGAAAAACCAGAGCCCTCTG
 AAACCACACCACAGGGCCAGAGAAGGGGTCAATCAAATGACGAAAACATGGGCCACGAAGCGGAAAT
 CCAGCCGGCCATTTTGAAGTTCATGTCTCCAGGGAACCATTTCTGAGTAAAAATAGAATCAGCAGAAGC
 CAAGAAGGCAGTATGAAGCAGGAGGCTGAACAAATTAACCTGAGGAGGCAAAAACCTGCCATTTGGCAAG
 TCCTGCAACCCAGCGAAGGCGGTGAAAGAATCCAAGTGGATGTAGCATAGGCCAAATACAAGAAAGCAG
 TGATGGGAGCTTAGGGGAGGCTGAGCAAGCAAAAAGGACAAAGCAGAATTGATTTCCCCACTTCACT
 CTTTCTAGTTGTCTTCAATAATGACTCACGCTTCTCTTGGGGTTGACACGCACAACCTCCACAGGCCAAA
 TTCATGACGTCCCTGAAAATGACATAGTTGAGCCAGAAAAGCGTCAAGTATGTGTTTCTGTTTACAGAA
 AAGGGGAACTATTGAGAATGAGCGTGGGAAACCTTTGCCCTCTTCTCCTGATCTTACCAGGTTCCCTTGT
 ACTTCATCTCCTGAAGGAAATGTACAGACTTTTTGATAAGCCACAAAATGGAGGAACTAAAATAGAGG
 TGCTTCAAATTTGGGAAACCAACCCCAAGCTCATCTAGCTCCTCAGCGAAGACCTTGGCATTATTTTC
 AGGAGAACGTGAGTTAGAGAAAGCCCTAAATTAAGTGCAGGATCCATGTCAAAGGGCACCTGGGCTGT
 GCGAAAAGTCCAGGGAGAGAGAGAAGTCCCTGGAAGCCCGAGCAGGCAATCGCCAGGGACCTCACAG

CAGTGACGGGGTCAGAGGAGGTCAAGAGGAAGCCAGAAGCCCCAGGCAGTGGACATTTAGCTGAGGGAGT
AAAGAAGAAAATTTTGTCCAGGGTGGCAGCACTGAGGCTGAAACTGGAAGAAAAGGAAAATATCAGAAAAG
AACTCAGCCTTTCTTAAAAAGATGCCAAACTCGAAACATCATTATCACACACAGAAGAGAAAACAAGACC
CAAAAAGCCATCTTGCAAAAGAGAAGGAAGAGCTCCAGTATTACTGAAAAAGATCCAAGCTGAGATGTT
CCCTGAACACTCTGGAATGTAAATTAAGCTGCCAATTTGCAGAAATTCATGAAGATTCTACTATCTGC
TGGACAAAAGATTCAAAGTCCATAGCCCAAGTGCAGAGAAGTGCAGGGGACAACCTCCACTGTTTTCTTTG
GACTGCTGAATTTAACCTCACAGCTGAAGTTCTCAAACAGCTGTCAAGTCGCCAGGATACTAAAGGATGT
GAAGAGATTGAATTCAGCCAACCTCATCTTCAAAGAAGACTTCCTCCATGACAGCTACTTTGGGGGCCGCC
TGCGTGGTCAGATCGCCACGGAGGAGCTGCACTTTGGAGAAGGGGTTACCCGCAAAGCCTTCCGCAGCAC
AGTGATGCACGGCCTCATGCCTGTCTTCAAACCTGGCCATGCCTGTGTGCTTAAGGTGCACAATGCCATT
GCCTATGGGACCAGAAAATATGATGAGCTCATCCAAGGAACACAACTCGCTGCCAGGAATGCTATG
TTCAAAATACTGCCAGGTATTATGCCAAGATCTACGCTGCTGAAGCACAGCCTCTGGAAGGCTTTGGAGA
AGTACCTGAGATCATTCTATTTTTCTTATCCATCGGCCTGAGAACAATATCCCGTATGCTACAGTGGAG
GAGGAGCTGATTGGAGAATTTGTGAAGTATTCCATCAGGGATGGGAAAGAAAATAAAGTTCTTGAGAAGAG
AATCAGAAGCTGGTCAGAAATGTTGCACCTTCCAGCACTGGGTGTACCAGAAAACAAGTGGCTGCCTCCT
GGTGACGGACATGCAAGGTGTAGGAATGAAGCTAACTGACGTTGGCATAGCAACGCTGGTAAAGGGTAC
AAGGGATTTAAAGGCAACTGTTCCATGACCTTCAATGATCAGTTTAAAGCACTACACCAGTGAACAAGT
ATTGCAAAATGCTGGGACTGAAATCCCTTCAAACAACAACCAGAAAACAGAAGCAGCCGAGCATTGGGAA
AAGCAAAGTTCAAACAACCTCTATGACAATAAAGAAGGCAGGGCCTGAGACCCCAGGCGAAAAGAAAACC

ACGCGTACGCGGGCCGCTCGAG - GFP Tag - GTTTAA

Protein Sequence: >RG215838 representing NM_052947
 Red=Cloning site Green=Tags(s)

MKDSEGPQRPLCFLSTLLSQVPEKSDAVLRCCIISGQPKPEVTWYKNGQAIDGSGIISNYEFFENQYIH
 VLHLSCCTKNDAAVYQISAKNSFGMICCSASVEVECCSENPLSPNLEDDRRDRGWKHETGTHEEERANQI
 DEKEHPYKEEESI SPGTPRSADSSPKSNHLSLQSLGNLDISVSSSENPLGVKGRHTGEAYDPSNTEE
 IANGLLFLNSSHIYEKQDRCCCHKTVHSMASKFTDGDNLNDGPHDEGLRSSQQNPKVQKYISFSLPLSEAT
 AHYYPGDSAVANKQPSQLSSESDSDYELCPEITLTYTEEFSDDDLEYLECSDVMTDYSNAVWQRNLLG
 TEHVFLLESDDDEMEFGEHCLGGCEHFLSGMCGSRVSGDAGPMVATAGFCGHHSQPQEVGRSSRVSKH
 GPSSPQTGMTLILGPHQDGTSSVTEQGRYKLPATAPEAAENDYPGIQGETRDSHQAREEFASDNLNMDDES
 VRETEMKLLSGESENSGMSQCWETAADKRVGGKDLWSKRGSRKSARVRQPGMKGNPKPNANLRESTTEG
 TLHLCSAKESAEPPLTQSDKRETSHTTAAATGRSSHADARECAISTQAEQEAKTQTSTDSVSKEGNTNC
 KEGEMQVNTL FETSQVPDWSPPQVQVQETVRETI SCSQMPAFSEPAGEESPFTGTTTTISFNLGGVHKE
 NASLAQHSEVKPCTCGPQHEEKQDRDGNIPDNFREDLKYEQSISEANDETMSPGVF SRHLPKDARADFRE
 PVAVSVASPEPTDALTLENVCDEPRDREAVCAMECFEAGDQGTCFDTIDSLVGRPVDKYSPQEICSVDT
 ELAEGQNKVSDLCSSNDKTLEVFFQTQVSETSVSTCKSSKDGNSVMSPLFTSTFTLNI SHTASEGATGEN
 LAKVENSTYPLASTVHAGQE QSPSNSGGLDETQLLSENPLVQFKEGGDKSPSPSAADTTATPASYSS
 IVSFPWEKPTTLTANNECFQATRETEDTSTVTIATEVHPAKYLAVSIPEDKHAGGTEERFPRASHEKVSQ
 FPSQVQLDHILSGATIKSTKELLCRAPSVPGVPHHLVQLPEGEGFCSNSPLQVDNLSGDKSQTVDRAFR
 SYEENFQERGSETKQGVQQQLSQQGSLSAPDFQQSLPTTSAAQEERNLVPTAHPASSREGAGQRSGWG
 TRVSVVAETAGEEDSQALSNVPSLSDILLEESKEYRPGNWEAGNKKIITLEASAEIWPFRQLTNSESK
 ASDGGLIIPDKVWAVPDSLKADAVPELAPSEIAALAHSPEDAESALADSRESHKGEETISVHWRSLSS
 RGFSPRLLLESSVDPVDEKELSVTDSL SAASETGGKENVNNVSQDQEEKQLKMDHTAFFKKFLTCPKILE
 SSVDPIDEISVIEYTRAGKPEPSETTPQGAREGGQSDGNMGHEAEIQPAILQVPCLQGTILSENIRSRS
 QEGSMKQEAQIQPEEAKTAIWQVLQPSSEGERIPSGCSIGQIQESSDGS LGAEQSKKDKAELISPTSP
 LSSCLPIMTHASLGVDTNHTGQIHDVPENDIVEPRKRQYVFPVSQKRGTIENERGKPLPSSPDLTRFPC
 TSSPEGNVDFLISHKMEEPKIEVLQIGETKPPSSSSSSAKTLAFISGERELEKAPKLLQDPCQKGTLC
 AKKSREREKSLEARAGKSPGTLTAVTGSEEVKRKPEAPGSGHLAEGVKKKILSRVAALRLKLEEKENIRK
 NSAFLLKMPKLETSLSHTEEKQDPKPKSCKREGRAPVLLKKIQAEMFPEHSGNVKLSQCF AEIHEDSTIC
 WTKDSKSI AQVRSAGDNSTVSFAIVQASPKDQGLYYCCIKNSYGVTAEFNLTAEV LKQLSSRQDTKGC
 EEIEFSQLIFKEDFLHDSYFGGRLRGQIATEELHFGEVHRKAFRSTVMHGLMPVFKPGHACVLKVHNAI
 AYGTRNDEL IQRNYKLAQECYVQNTARYYAKIYAAEAQPLEGFGEVPEIIPIFLIHRPENNIPIYATVE
 EELIGEFVKYSIRDGKEINFLRRESEAGQKCTFQHWVYQKTSGLLVTD MQGVGMKLT DVGIATLAKGY
 KGFKGNCSTMFIDQFKALHQCNKYCKMLGLKSLQNNNQKQKQPSIGKSKVQTNMSTIKKAGPETGPEKKT

TRTRPLE - GFP Tag - V

Restriction Sites: Sgfl-MluI

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_052947.4](#)

RefSeq Size: 7303 bp

RefSeq ORF: 6513 bp

Locus ID: 115701

UniProt ID: [Q86TB3](#)

Cytogenetics: 18q21.31-q21.32

Protein Families: Druggable Genome, Protein Kinase

Gene Summary: Kinase that recognizes phosphorylation sites in which the surrounding peptides have an alpha-helical conformation.[UniProtKB/Swiss-Prot Function]