

Product datasheet for **RG213284**

FRYL (NM_015030) Human Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Tag:	TurboGFP
Symbol:	FRYL
Synonyms:	AF4p12; KIAA0826; MOR2
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)

ORF Nucleotide Sequence: >RG213284 representing NM_015030
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCCCGCATCGCC

ATGTCAAACATTACGATTGACCCAGATGTCAAACCTGGTGAATATGTCATCAAGAGCCTCTTTGCAGAAT
TTGCTGTTCAAGCTGAAAAGAAAATGAAGTTGTAATGGCCGAACCTTGGAGAAGCTATTGTCCAGATC
TCTTCAGAGGGGTGAAGATCTTCAGTTTGATCAGTTGATAAGCTCTATGAGCTCAGTAGCAGAGCACTGT
CTCCCTTCCTTACTTCGCACCTTGTGACTGGTACAGACGCCAAAATGGAACGGAAGATGAATCTTATG
AATATAGGCCTCGGTCTAGCACAAAGTCTAAGGGGGATGAACAGCAACGTGAAAAGAGATTATCTTCTTGA
AAGGAGGGACTTAGCAGTAGACTTCATTTTTTTAGTTTGAAGTTCTAAAGCAGATTCTGT
CATCCTGTACCCGATCCCTTAGTTCATGAAGTTCTAACTTAGCTTTAAGCACTTTAAACATAAGGAAG
GATATTCAGGAACCAACTGGGAATGTGCATATTATTGCTGATTTATGCAGAGGTGATAGGGTTCT
TGCCCAATCAAAGTTTCAGGCTGTAAGGAAGAAGTTTGTGACAGAATTAAGAACTGCGACAAAAGGAA
CAAAGCCCACATGTGGTACAAAGTGTCACTAGCTTAATAATGGGAATGAAATTTTTTCGAGTAAAAATGT
ATCCTGTAGAAGATTTGAAGCATCTTCAATTTATGCAGGAATGTGCTCAGTATTTCTTAGAAGTGAA
AGATAAAGATATAAACATGCACCTTGCTGGTTTATTTGTGGAGATTCTTATCCCTGTAGCTGCTGCTGTT
AAAAATGAAGTGAATGTTCCCTGTTTGAAAAATTTGTGGAGATGCTTTATCAGACTACTTTTGAATGA
GCTCGAGAAAGAAGCATTATTGGCTTTATATCCACTAATTACCTGCCTTTTATGTGTCAGTCAGAAACA
ATTTTTTTAAATAACTGGCATATTTCTACAGAAGTGTGTCACATTTAAGAATAAAGATCCGAAA
ATGTCTCGAGTTGCACTGGAATCTTTGTATAGATTATTGTGGGTTTATGAATTAGAATAAAATGTGAAA
GCAACACTGTAACCTCAAAGTCGTCTTATGAGCATAGTGTGAGCACTTTTCCAAAAGGCTCACGAAGTGT
GGTTCTCTGTGACACACCTCTCAATATATTTGTGAAGATTATTCAGTTCAATGCTCAGGAACGCTTGGAT
TTTGCAATGAAAGAAATAATTTGATCTTCTCAGTGTGGAAAACTACTAAAACCTTACCATTAAATC
CAGAGAGAATGAACATAGGTCTCAGAGTCTTCTGTAAATAGCAGACAGTTGCAGCAGAAAGATGGTGA
ACCACCATGCCAACCAAGAGTTATTCTTCCCTCAGGAAATACTCTCGTGTAAAGAAAATATTTCTC
AATAAACCTTGACAGATGAAGAAGCAAAAGTCATAGGAATGTCAGTTTACTATCCTCAAGTAAGAAAAG



CATTAGATAGCATCCTCAGACATTTGGACAAAGAAGTTGGGAGACCAATGTGTATGACCAGTGTGCAGAT
GTCTAATAAGGAGCCTGAAGACATGATTACGGGGGAAAGAAAACCAAGATTGATTTGTTTGAAGCTTGT
ATTGCTGCGATTCCAAGGTTGATTCCTGACGGTATGAGCAGAACTGACCTGATTGAATTGTTAGCAAGGC
TCACAATTCATATGGATGAAGAACTGCGTGCTGGCTTTCAATACTCTGCAGGCACTAATGCTTGATTT
TCCAGATTGGCGGGAGGATGTTCTTTCAGGATTTGTTATTTTATTGTTTCGTGAAGTGACTGATGCCAT
CCCACACTTCTTGATAATGCCGTAAGATGTTGGTACAATTAATAAATCAGTGGAACAAGCAGCCCAAA
TGCATAATAAAAAACAGGACACTCAGCATGGTGTAGCTAATGGAGCTTCTCATCCCCCTCTCTGAAAAG
GAGCCCATATTCCAATGTATTCCATGTGGTTGAAGGCTTTGCGCTTGTCATTCTCTGTAGCAGTCGACCT
GCCACTAGGAGACTAGCCGTCAGTGTCCTTAGAGAAAACGCGGCTTTATTTGCACTTCTGAAAATACCTA
AGGGTGATGATGAATTAGCCATAGATGTGATGGACAGGCTAAGCCCATCCATTCTTGAGAGCTTCATACA
TCTCACTGGGGCTGATCAGACTACTTTGCTCTATTGCCCTAGCTCGATAGATTTACAACTTTAGCAGAA
TGGAACTCTTCTCCTATTAGCCACCAAGTTTGATGTGATTAGTCCATCACATATATGGATATTTGCACATG
TGACCCAAGGCCAAGACCATGGATTATAAGTCTCTCCAGTTTTTTAAAGCAAGAAAATCTTCTAAACA
CTGCTCTACAGCTGTGAGCTATGCTTGGATGTTTGCATACACAAGACTTCAGTTGTTGTCCCCTCAGGTC
GATATAAATAGCCCATCAATGCTAAGAAAGTAAATACCACCACAAGCAGTGACTCATACATTGGCCTGT
GGAGAAACTATCTGATCCTTTGCTGCACTGCAGCAACATCGTCATCTCCACATCTGCAGGTTCTGTGAG
ATGTTCTCCTCCTGAGACGCTGGCGTCTACCCAGATAGCGGCTATAGCATTGATTCTAAAATTATTGGC
ATCCCATCCCCTTCATCCTTGTTTAAAGCATAGTTCCAATGATGCGTTCTGAGAGCATGGAATACAG
AATCCCTTGTTCTAGGTCTTGGCAGGACCAACCCAGGAGCTTTAGGGAATAGAGGAATTACATCC
CATAATTAAAGAAGCACTCGAAAGAAGGCCAGAGAATATGAAACGGCGCAGGCTCGAGACATTTTACGA
GTACAAGTGTACGAATATTTGAAGTCTGGCAGATGCTGGTGTATTAGTCACAGTGCAAGTGGTGGCC
TTGATAATGAAACACATTTTCTCAACAACACTTTATTGGAATATGTAGATTTAACTAGACAACCTCTGGA
AGCAGAAAAAGAAAGACTCTGACACACTGAAGGATATACGATGCCATTTTAGTGCCTTAGTGGCGAAT
ATTATTCAGAAATGTTCCAGTGCAACAGAGAAGAAATTTTTCTCAACAGAGCCTTCGTACAGCTCTAT
TTATGCTGTTCACTGAGGCGAGTCTTTTAGCATCATGTTTACGCCCTTGACAGATACAGTGATAG
AAATATGCAAATTAATAGACATCAATACTGTGCGTTAAAGGCTATGTCTGCTGACTGTGTTGTGGCCCT
GTTGCAGATAATGTAGGACTTTTCATCAGATGGCTATTTGTACAAATGGTTGGATAACATTTTGATTCTC
TGGACAAAAAGGTTCAACAGCTGGGCTGTGAAGCAGTTACGTTGTTACTGGAGCTGAACCCGTGATCAGAG
CAACCTGATGACTGGGCTGTGGACCGCTGCTACACGGGCTCCGGGAGGGTGGCGGCCGGCTGCTTTAA
GCCATTGCTAATGTTTTCCAGAACAGGATTATCAATGTGACACAGTGATGCTTCTAAATCTGACTGT
TTAAAGCAGCTGATTCTCTAGAAGTATCTATGAAGTGTCTATGCAACTTTTACAGATTCTGGAACCGAA
GATGTTTCGTATGCTCACAATTGGAGGTTAGAGAACAGATGGAGTACTCAGCCAGCTGTCTCTCTA
CCACATCTCTATTCTGTTTCATATTATCAGTTGTCCGAGGAAGTACAGAGGGCGTATCCTGAGCTAACTC
TCGCCATATTCTCAGAGATAAGCCAGAGAATCCAGACAGCTCACCTGCTGGCGGCGAGGTGATGCTGCA
CTACCTGCTACCATGGATGAACAACATCGAGCTGGTGGACTTAAACCTCTCCCCACAGCAAGGCGACAT
GATGAAGATGAAGATGATTCTTAAAGACCGAGAAGTTATGGTGACTAGTAGGCGCTGGTTACGGGGAG
AAGGATGGGGATCTCCACAAGCCACTGCAATGGTTTTGAACAATCTGATGTATATGACAGCAAGTATGG
CGATGAAGTGGCTGGTGGAGGTGGAGAATGTGTGGACCACACTGCAGATGGCTGGCCCCAAAACCTG
AAAATAATTTTGCACTTTTGTATCAGCATTTGTGGGGTGAATAGCGAACCAAGCCTCTTGCTTACGTGA
AGAAGGTCATTGTATATTTAGGTAGAGATAAAACAATGCAAGTTGCTAGAAGAGCTGGTGAGTGAGCTTCA
GCTGACCGATCCTGTCACTTCAAGGGTCACTCAGATGGATAATCCCCCGTATTATCGCATCACTTCCAGC
TATAAAATCCCTTCTGTACCTCAGGAAGTACTCCAGTAGCAATACAATGGTAGCTCCACAGATGGCA
ATCCTGATAATAAGCCATTAAAGAGAATATTGAAGAGAGCTATGTGCACCTGGACATTTACAGTGGACT
AAACAGTCATTTGAATCGGCAACATCACAGACTAGAATCCCGATACAGTAGCAGCTCTGGAGGATCTTAT
GAAGAAGAAAAAGTGATTCAATGCCACTTTATTCTAATTGGCGACTGAAAGTGATGGAGCATAACCAAG
GAGAGCCACTGCCCTTCCACCAGCTGGAGGCTGCTGGTCAACCTGGTGGATTACGTGCCTGAAACGTC
ATCACCTGGATTACCTCTTACAGGTGAACATAGCAGTGATCCTTTTGACTGATCTCATCATTGATCAT
AGTGTGAAGGTGGAATGGGGAAGCTACCTCCATCTTCTTCTCATGCAATTTTATAGGGTTTGACCACT
GCCACCTGAGGTGATGAACATTGTAACGCCTGCTTCTGCACTTATTAATAGTAATGGGACCCCAATAG
TAACATCCGAAGTGTGCTTCTGCTTCTCAGGAACAAGGAGTTAATGAGCCAGGGTGTACAGTC
AAACAAGTTGCACACTTAGATTATAATTTACAGCAGGCATTAACGATTTTATACCTGATTACCAGCCCT
CCCCTATGACTGACTCAGGGCTTAGCTCAAGTTCTACCTCTTCTAGTATCAGCTTAGGAAATAACAGTGC
TGCCATTTACATCTGCACACCACTATCCTCAATGAGGTTGACATCTCAGTGGAGCAGGATGAAAAAGTC
AAAACCTCATGGAATTCATTACCTCAAGAAAAAGAGGGCCCTTTGGAACCATGAGGATGTTTCTGCCA
AGAATCCTAGCATAAAGAGTGTGAACAGTTAACTACATTTTGAACATGTGGTTTCTGTTTTAAGCA
GTCAAGCTCAGAAGGAATTCATCTGGAACATCATCTTAGTGAAGTTGCTCTGCAACAGCACTTCTCTGT

TCTTCTCGACACTATGCTGGGAGATCCTTTCAGATTTTCAGGGCCCTAAAGCAGCCTCTCACTGCAACTA
CACTTTCTGATGTTCTCTCCAGACTTGTAGAACTGTAGGGGATCCAGGAGAAGATGCACAGGGATTGTG
GATTGAGCTTCTTCTCACATTGGAATCTGCAATTGATACTTTGGCTGAAACCATGAAGCATTATGATCTT
CTTTCTGCCCTTTCTCAAACCTCATATCATGATCCTATAATGGGAAACAAGTATGCAGCTAACAGGAAAA
GCACTGGACAACCTCAATCTAAGCACAGTCCCATTAAATAGTAGCAGTTATTTGGGATATAACAGTAATGC
AAGAAGTAACCTTTGAGATTAAGTTTGATTGGTGACCGACGAGGTGACCGGCGGCGAGTAACACACTG
GATATAATGGATGGACGGATAAACCATAGCAGTAGTTAGCAAGGACTAGAAGCCTTCTCTCTAAGAG
AGAAAGGAATGTATGACGTGCAGTCCACTACTGAGCCTACCAACTTGATGGCCACCATTTTTTGGATAGC
AGCATCTTTATTAGAATCAGATTATGAATATGAATACCTCCTGGCTCTCAGGCTTCTCAACAACTGCTT
ATCCATTTGCCTTTGGATAAATCAGAGAGTCGAGAGAAGATTGAAAATGTACAAAGCAAATTGAAATGGA
CTAATTTTCCAGGACTTCAGCAGCTCTTCTTAAGGGTTTTACCTCAGCATCTACACAAGAAATGACCGT
GCACCTCCTCAGTAAACTCATTCTGTCTCCAAACATACATTGGTGGATCCTTCCCAATTGTCAGGCTTT
CCTCTTAACATCCTTTGCTTATTGCCTCACTTAATCCAGCATTTTGACAGCCCAACTCAGTTTTGCAAAG
AAACAGCTAGTCGAATAGCAAAGGTTTGTGCAGAAGAAAAATGCCAACACTTGTCAATCTGGCACACAT
GATGAGTTTGTACAGTACACACAGTATTCCAGAGACTGTTCTAACTGGATCAATGTCGTGTGCAGATAC
CTGCATGACTCCTTCTCAGATACAACATTTAATCTTGACTTATCTTGCAAGCTGTTAGAGAAAGGAT
TGTCCAGTATGCAGCAATCATTACTACAGATTATTTATAGTCTATTGAGTCATATTGACCTGTCTGCAGC
CCCAGCCAAGCAGTTTAACTCTGGAGATCATAAAGATTATTGGCAAAATATGTACAGAGTCCTTACTGGAAG
GAAGCCCTTAACATATTAAGCTGGTGGTGCACGCTCTGCGAGTCTTGTCTACCCAGTGATATCCCCA
AGACCTATGGAGGAGATACAGGTTCTCCTGAAATATCCTTCACTAAAATTTTAAATGTTTCTAAGGA
GTTGCCTGGGAAGACCTTAGATTTTCAATTTGATATCTGAGACACCAATTATTGGAACAAATATGTT
GATCAGCACAGTGCAGCTGGAAGAAATGGGAAACCAAAGTTATTGCTGTCACTAGAAGTACTTCTCAA
CTTCTTCTGGTTCTAATCTAATGCCTTGGTCTGTTAGTTGGAAAAGGCCACAGTTATCACAGCGAAG
AACAAGAGAAAAGCTAATGAATGTGCTTCTCTGTGGTCCAGAATCTGGCCTCCCAAAGAACCCATCA
GTTGATTTTCTTCTAATGAGGATTTGGAAGTCGGTGACCAACAGACTAGCCTAATTTCTACAACAGAAG
ACATAAATCAAGAGGAAGAAGTAGCTGTGGAAGATAATAGCAGTGAACAACAGTTTGGTGTTTTTAAGGA
TTTTGACTTTTTAGATGTTGAATTGGAAGATGCAGAGGGTGAAGTATGGACAATTTCACTGGGAGTT
CGCAGGCGCTCACTGGACAGTATTGACAAAGGGGACACTCCATCCCTCCAGGAGTACCACTGCTCTAGTA
GCACCCCCAGCCTGAACCTCACCAATCAGGAGGATACAGATGAGTCCTCGGAAGAAGAAGCGGCCTTAC
AGCAAGCCAGATACTCTCACGCACACAGATGTTAAACAGTGATTCTGCCACTGATGAAACAATACCAGAC
CATCCTGACTTACTTCTCCAGTCTGAAGATTCCACTGGCAGCATCACAACAGAGGAAGTGCTTCAAATCA
GGGATGAGACCCCAACTTTGGAGGCTTCTCTAGATAATGCTAACAGCCGGCTGCCTGAGGATACAACCTC
AGTATTAAGGAGGAACATGTTACAACCTTTGAAGTGAAGGATCCTATATAATTCAAGAACAGCAGGAA
TCTCTGTGTGTCAAGGAATCTTGATTTAGAAGAACTGAAATGCCAGAGCCTCTAGCTCCTGAAAGTT
ACCCGAGTCAGTCTGTGAAGAGGATGTTACCTTAGCTCTGAAAGAGCTAGATGAAAGATGTGAAGAAGA
AGAAGCGGATTTCTCCGACTGTCTAGTCAAGATGAAGAAGAGCAAGATGGTTTTCCAGAAGTACAGAGC
TCGCCTCTGCCGTACCATTCTTTCTGCCATCATAGCCGCTTTTCCAGCCGTGGCATATGATGATGAAG
AGGAAGCCTGGCGCTGCCAGTCAATCAGATGCTGTCTGACACCGACGGGTCTCTGCAGTGTCTTACTTT
TCATGTGTTTTCTAGGCTGTTTCTAGACAATTCAAAGAAAGTTTGGAGAAATACTAATGAGGCAGTCAGC
TTTCTTGGTGATAGTCTGCAACGCATTGGTACCAAATTTAAAGTTCTTGGAAAGTATGATGCTGTGTT
CAGAATGCCCAACAGTCTTTTGGATGCTGAAACACTGATGTCATGTGGTTTGTCTGGAACACTCAAGTT
TGGTGTTTTGGAGTTGCAAGAACACCTGGATACATACAATGTGAAAAGAGAAGCCGCTGAGCAGTGGCTA
GATGATTGTAAGAGGACATTTGGTGCCAAAGAAGACATGTATAGGATAAACACAGATGCACAACAATGG
AAATACTAGCAGAATTGGAGCTCTGCCGAAGATTATACAATTGCATTTTCAATTGCTGCTTCTGTTCCA
GGCCTACTGTAACTTATCAACCAAGTAAATACGATAAAAAATGAAGCAGAGGTATCAACATGTCAGAG
GAACTTGCCCAACTGGAAGTATCCTCAAAGAAGCTGAGTCCGCTTCCGAAAACGAAGAAATTGACATTT
CCAAAGCTGCACAACTACTATAGAACTGCCATTCTTTAATTGAACTTTGAAAAATAAAGAAATT
TATATCAGCTGTAGCACAAGTCAAAGCTTTTCAAGTCTCTCTGGCCAGTGATATCTTTGGCAGTTGTGAA
GATGACCTGTACAGACACTGTTACATATATATTTCCATCATCAGACGCTGGGCCAGACAGGAAGCTTTG
CAGTTATAGGCTCTAACCTGGACATGTGAGAAGCAACTACAACTGATGGAACCTAATCTGGAATAAG
AGAGTCTCTACGCATGGTGAATCATACCAACTTCTAGCACAGGCCAAACCAATGGGAAATATGGTGAGC
ACTGGATTCT

ACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

Protein Sequence:

>RG213284 representing NM_015030

Red=Cloning site Green=Tags(s)

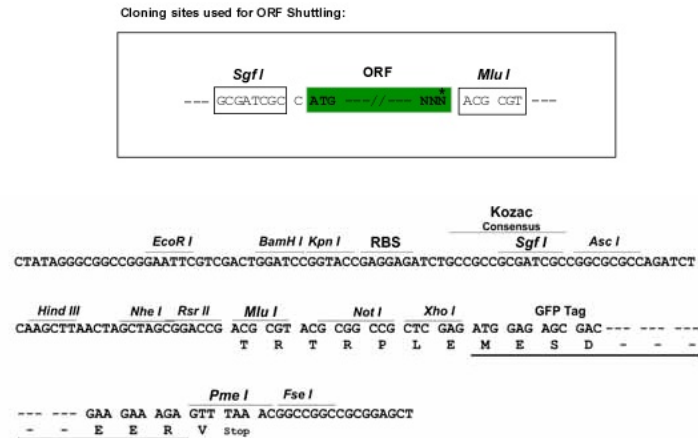
MSNITIDPDVKPGYVIKSLFAEFVQAQAEKKIEVMAEPLKLLSRSLQRGDELQFDQLISSMSSVAEHC
LPSLLRRTLFDWYRRQNGTEDESIEYRPRSSTKSKGDEQQRERDYLLERRDLAVDFIFCLVLEVLKQIPV
HPVPDPLVHEVLNLAFAKFKHKEGYSGTNTGNVHIIADLYAEVIGVLAQSKFQAVRKKFVTELKELRQKE
QSPHVQSVISLIMGKFFRVKMYPVEDFEASFQFMQCAQYFLEVKDKDIKHALAGLFVEILIPVAAAV
KNEVNPCLKNFVEMLYQTTFELSSRKKHSLALYPLITCLLCVSQKQFFLNWHIFLQNCLSHLKKNKDPK
MSRVALESYLRLWVYVIRIKCESNTVTQSRLMSIVSALFPKGSRSVVPDTPNLNIFVKIIFIAQERLD
FAMKEIIFDLLSVGKSTKFTTINPERMNIPLRVFLVIADSLQKQKDEPPMPTTGVLPSGNTLRVKKIFL
NKTLDDEEAKVIGMSVYYPQVRKALDSILRHLDEKVRPMCMSTSVQMSNKEPEDMITGERKPKIDLFRTC
IAAIPRLIPDGMSTDLIELLARLTIHMDIELRALAFNTLQALMLDFPDWREDVLSGFVYFIVREVTDVH
PTLLDNAVKMLVQLINQWKQAAQMHNKNQDQHGAVANGASHPPPLERSPYSNVFHVVEGFALVILCSSRP
ATRRLAVSVLREIRALFALLEIPKGDDELAIDVMDRLSPSILESFHILTGADQTTLLYCPSSIDLQTLAE
WNSSPISHQFDVISPISHWIFAHVTQGGDPWIIISLSSFLKQENLPKHCSTAVSYAWMFAYTRLQLLSPQV
DINSPINAKKVNTTSSDSYIGLWRNYLILCCSAATSSSSTAGSVRCSPPETLASTPDSGYSIDSKIIG
IPSPSSSLFKHIVPMRSESMEITESLVGLGRTPNGAFRELIEELHPIIKEALERRPENMKRRRRRDIR
VQLVRIFELLADAGVISHSASGGLDNETHFLNNTLLEYVDLTRQLLEAENEKSDTLKDIRCHFSALVAN
IIQNPVPHQRRSIFPQQLRHSLFMLFSHWAGPFSIMFTPLDRYSRNMQINRHQYCALKAMSAVCCGP
VADNVGLSSDGYLYKWLNDISLDDKKVHQLGCEAVTLLLELNPDQSNLMYWAVDRCYTGSGRVAAAGCFK
AIVANVFQNRDYQCDVTMLNLILFKAADSSRSIYEVAMQLQILEPKMFRYAHKLEVQRTDGVLSQLSPL
PHLYSVSYQQLSEELARAYPELTALFSEISQRIQTAHPAGRQVMLHYLLPMMNNIELVDLKLPTARRH
DEDEDDSLKDRELMTSRRWLRGEGWSPQATAMVLNNLMYMTAKYGDELAWSEVENVWTTADGWPKNL
KIIILHFLISICGVNSEPSLLPYVKKVIVYLGRDKTMQLLEELVSELQLTDPVSSGVTHMDNPPYYRITSS
YKIPSVTSGTTSSSNTMVAPTGDGPNKPIKENIEESYVHLDIYSLNSHLNRQHRLRESYSSSSGGSY
EEEKSDSMPLYSNWRLKVMENQGEPLFPFPAGGCWSPVDYVPETSSGPLPHRCNIAVILLTDLIIDH
SVKVEWGSYLHLLHAIIFIGFDHCHPEVYEHCKRLLHLLIVMGPNSNIRTVASVLLRNKEFNEPRVLT
KQVAHLDDYNFTAGINDFIPDYQPSMTDSGLSSSSTSSISLGNNSAAISHLHTTILNEVDISVEQDGKV
KTLMEFITSRKRGLWNHEDVSAKNPSIKSAEQLTTFLKHVSVFVKQSSSEGIHLEHHLSEVALQTALSC
SSRHVAGRSFQIFRALQPLTATLSDVLSRLVETVGDGPGEDAQGFVIELLLTLESAIDTLAETMKHYDL
LSALSQTSYHDPIMGNYAANKSTGQLNLSTSPINSSSYLGYNSNARSNSRLSLIGDRRGDRRRSNTL
DIMDGRINHSSSLARTSLSSLREKGMVDVQSTTEPTNLMATIFWIAASLLESDYIEYLLALRLNKL
IHLPLDKSESREKIENVQSKLWNTNFPGLQQLFLKGFSTASTQEMTVHLLSKLISVSKHTLVDPSSLSGF
PLNILLCLPHLIQHFDSTPQFCKETASRIAKVCAEEKPTLVNLAHMMSLYSTHTYSRDCSNWINVVCY
LHDSFSDTTFNLTLYLAELLEKGLSSMQQLLQIIYSLLSHIDLAAAPAKQFNLEIIKIIGKYVQSPYWK
EALNILLVVSRSASLVPSDIPKTYGGDTGSPEISFTKIFNNVSKELPGKTLDFHFDISETPIIGNKY
DQHSAAGRNGKPKVIAVTRSTSSSSGNSNALVPVSWKRPQLSQRRTREKLMNVLSLCPESGLPKNPS
VVFSSNEDLEVGDQQTSLISTEDINQEEVAVEDNSSEQQFGVFKDFDLDVELEDAEGESMDNFWGV
RRRSLDSIDKGDTPSLQEYQCSSTPSNLNTNQEDTDESSEEEAALASQILSRTQMLNSDSATDETIPD
HPDLLQSEDSTGSITTEEVLQIRDETPTLEASLDNANSRLPEDTTSVLKEEHVTTFEDEGSYIIQEQQE
SLVCQGILDLEETEMPEPLAPESYPESVCEEDVTALKEKELDERCEEEEDFSGLSSQDEEEQDGFPEVQT
SPLPSPFLSAIIAFAQPVAYDDEEEAWRCHVNQMLSDTDGSSAVFTFHVFSRLFQTIQRKFGEITNEAVS
FLGDSLQRIGTKFKSSLEVMMLCSECPVTFVDAETLMSCGLLETLFKGVLELQEHLDYTNVKREAAEQWL
DDCKRTFGAKEDMYRINTDAQMEILAELELCRRLYKLFQLLLLFQAYCKLINQVNTIKNEAEVINMSE
ELAQLESILKEASASENEEIDISKAQTIIETAIHSLIETLKNKEFISAVAQVKAFRSLWPSDIFGSC
DDPVQTLHLHIYFHHQTLGQTGSFAVIGSNLDMSEANYKLMELNLEIRESLRMVQSYQLLAQAKPMGMNVS
TGF

TRTRPLE - GFP Tag - V

Restriction Sites:

SgfI-MluI

Cloning Scheme:



ACCN: NM_015030

ORF Size: 9039 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.

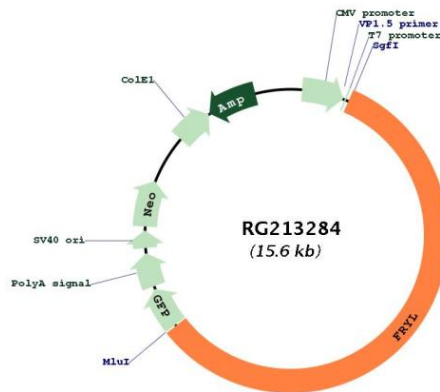
Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

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.

Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	<u>NM_015030.2</u>
RefSeq Size:	11706 bp
RefSeq ORF:	9042 bp
Locus ID:	285527
UniProt ID:	<u>O94915</u>
Cytogenetics:	4p11
Gene Summary:	Plays a key role in maintaining the integrity of polarized cell extensions during morphogenesis, regulates the actin cytoskeleton and plays a key role in patterning sensory neuron dendritic fields by promoting avoidance between homologous dendrites as well as by limiting dendritic branching (By similarity). May function as a transcriptional activator.[UniProtKB/Swiss-Prot Function]

Product images:



Circular map for RG213284