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Product datasheet for SC117569

FKBP51 (FKBP5) (NM_004117) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	FKBP51 (FKBP5) (NM_004117) Human Untagged Clone
Tag:	Tag Free
Symbol:	FKBP51
Synonyms:	AIG6; FKBP51; FKBP54; P54; PPIase; Ptg-10
Mammalian Cell Selection:	None
Vector:	pCMV6-XL5
E. coli Selection:	Ampicillin (100 ug/mL)

Fully Sequenced ORF:

>OriGene ORF within SC117569 sequence for NM_004117 edited (data generated by NextGen Sequencing)

ATGACTACTGATGAAGGTGCCAAGAACAATGAAGAAAGCCCCACAGCCACTGTTGCTGAG
CAGGGAGAGGATATTACCTCCAAAAAAGACAGGGGAGTATTAAAGATTGTCAAAAGAGTG
GGGAATGGTGAGGAAACGCCGATGATTGGAGACAAAGTTTATGTCCATTACAAAGGAAAA
TTGTCAAATGGAAAGAAGTTCGATTCCAGTCATGATAGAAATGAACCATTTGTCTTTAGT
CTTGGCAAAGGCCAAGTCATCAAGGCATGGGACATTGGGGTGGCTACCATGAAGAAAGGA
GAGATATGCCATTTACTGTGCAAACCAGAATATGCATATGGCTCGGCTGGCAGTCTCCCT
AAAATTCCCTCGAATGCAACTCTCTTTTTTGAGATTGAGCTCCTTGATTTCAAAGGAGAG
GATTTATTTGAAGATGGAGGCATTATCCGGAGAACCAAACGGAAAGGAGAGGGATATTCA
AATCCAAACGAAGGAGCAACAGTAGAAATCCACCTGGAAGGCCGCTGTGGTGGAAGGATG
TTTGACTGCAGAGATGTGGCATTCACTGTGGGCGAAGGAGAAGACCACGACATTCCAATT
GGAATTGACAAAGCTCTGGAGAAAATGCAGCGGGAAGAACAATGTATTTTATATCTTGGA
CCAAGATATGGTTTTGGAGAGGCAGGGAAGCCTAAATTTGGCATTGAACCTAATGCTGAG
CTTATATATGAAGTTACACTTAAGAGCTTCGAAAAGGCCAAAGAATCCTGGGAGATGGAT
ACCAAAGAAAAATTGGAGCAGGCTGCCATTGTCAAAGAGAAGGGAACCGTATACTTCAAG
GGAGGCAAATACATGCAGGCGGTGATTCAGTATGGGAAGATAGTGTCCTGGTTAGAGATG
GAATATGGTTTATCAGAAAAGGAATCGAAAGCTTCTGAATCATTTCTCCTTGCTGCCTTT
CTGAACCTGGCCATGTGCTACCTGAAGCTTAGAGAATACACCAAAGCTGTTGAATGCTGT
GACAAGGCCCTTGGACTGGACAGTGCCAATGAGAAAGGCTTGTATAGGAGGGGTGAAGCC
CAGCTGCTCATGAACGAGTTTGAGTCAGCCAAGGGTGACTTTGAGAAAGTGCTGGAAGTA
AACCCCCAGAATAAGGCTGCAAGACTGCAGATCTCCATGTGCCAGAAAAAGGCCAAGGAG
CACAACGAGCGGGACCGCAGGATATACGCCAACATGTTCAAGAAGTTTGCAGAGCAGGAT
GCCAAGGAAGAGGCCAATAAAGCAATGGGCAAGAAGACTTCAGAAGGGGTCACTAATGAA
AAAGGAACAGACAGTCAAGCAATGGAAGAAGAGAAACCTGAGGGCCACGTATGA

Clone variation with respect to NM_004117.3
201 t=>c

5' Read Nucleotide Sequence:

>OriGene 5' read for NM_004117 unedited

GGNCGGCACGCGCAATTCGGCACGAGGCCGGCGCGCGGGCGGCCTGCTGGGCGGGCTGAA
GGGTTAGCGGAGCACGGGCAAGGCGGAGAGTGACGGAGTCGGCGAGCCCCCGCGGCGACA
GGTTCTCTACTTAAAAGACAATGACTACTGATGAAGGTGCCAAGAACAATGAAGAAAGCC
CCACAGCCTCTGTTGCTGAGCAGGGAGAGGATATTACCTCCAAAAAAGACAGGGGAGTAT
TAAAGATTGTCAAAAGAGTGGGGAATGGTGAGGAAACGCCGATGATTGGAGACAAAGTTT
ATGTCCATTACAAAGGAAAATTGTCAAATGGAAAGAAGTTCGATTCCAGTCATGATAGAA
ATGAACCATTTGTCTTTAGTCTTGGCAAAGGCCAAGTCATCAAGGCATGGGACATTGGGG
TGGCTACCATGAAGAAAGGAGAGATATGCCATTTACTGTGCAAACCAGAATATGCATATG
GCTCGGCTGGCAGTCTCCCTAAAATTCCCTCGAATGCAACTCTCTTTTTTGAGATTGAGC
TCCTTGATTTCAAAGGAGAGGATTTATTTGAAGATGGAGGCATTATCCGGAGAACCAAAC
GGATAGGAGAGGGATATTCAAATCCAAACGAAGGAGCAACAGTAGAAATCCACCTGGAAT
GCCGCTGTGGTGGAAGGATGTTTGACTGCAGAGATGTGGCATTCACTGTGGGCGATAGAG
AATGACCACGACATTCCAATTTGGAATTTGACAAGCTCTGGAGAAAATGCAGNCGGGNAG
AACAATGTATTTTATATCTTGGACCAAGATATGGTTTTTNAGAAGCAGGNAAGCCTTAAT
TTGGCATGGACCCTATGCTGAGCTATTTTGAAGTACACTTAAGACTTCGAAAAGCCAAA

3' Read Nucleotide Sequence:

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>OriGene 3' read for NM_004117 unedited
CTATGGCACGCGGCACGCAATCTAGGATCGAGTTTTTTTTTTTTTTTTTTGGGAAGTACA
TTTTTTTTGGATTTATTTAAAGAACACACAAAAAAGTGCATCTAAATGATTTTTTTTTTT
AAAAAACAATGTCTTTAATTCCTCATATGCTGACTTGGCAATCCATATGCAGCCTAAAAG
GTGGTTGAAAGATGTCTGTGGACTTCTACACAAATTGTTTAAGATTATTGAATTTGATAC
AAGAATACGTGAAGTGTCTTCTTGAGTGGTAATGGGCACCCTGTAGTTATTTGCTCAAAA
CCACTCACACACCCTTCCCCTCCCCACACACACTTTTGCCAGTTCCCCCTGGTGAACCAT
AATACACACGTGGGAGACAGCTGGCATCATCTGAGTCTTGGAATAGCTTTCTTGTTTGTT
ATCTGCAAAAAGCTAGGTGGCCTTTTTAGTATCAAATTTCAATCTTGGCCTCACTGGAAC
CATGCTCTACAGAGCTTTCCCAACAGTTTAGCAAGTAAGCAAAATCAGCTTTTGCTCTTC
ATATTTCGTAGGTGAGACTGGGGCTTTTAAAAGAAATATTTCATAAAAAAGACAAAAGCC
TAAATGAGAAATTAGAGGGATTTTAGTTATCATTAAAGGAAAAAACTCACAAATTGATCC
CATTTGTTCCATGGGGAAGAAAAGTATTATAAAAGGAAATAACTTGATTTTTAGGTTTGA
AGGGGGTCCCAATCTAATAATAAAAGTCTTTTGATAAAACCATCATCAAAATAAAAACTA
ATTTAAAATATAACTGCCAGTGTTTTTTAAGATTTAAAAAGGTGAGATGTTCCCAGTTTA
AAGCAATGGNTNCTTTGGNNTGCANATGCATTCCATGAGTGTTNCTTAAGCTGAGGAATA
GAGAGCATGCTCATCTGTTTACCGCTAGACTGNANTCACTACTGTCACATTTAGACTTGC
CTATTCATGTCACTCAGTACTGAGATT
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Restriction Sites: NotI-NotI

ACCN: NM_004117

Insert Size: 3780 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

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_004117.2, NP_004108.1

RefSeq Size: 3781 bp

RefSeq ORF: 1374 bp

Locus ID: 2289

UniProt ID: Q13451

Cytogenetics: 6p21.31

Domains: FKBP, TPR

Protein Families: Druggable Genome

Gene Summary: The protein encoded by this gene is a member of the immunophilin protein family, which play a role in immunoregulation and basic cellular processes involving protein folding and trafficking. This encoded protein is a cis-trans prolyl isomerase that binds to the immunosuppressants FK506 and rapamycin. It is thought to mediate calcineurin inhibition. It also interacts functionally with mature hetero-oligomeric progesterone receptor complexes along with the 90 kDa heat shock protein and P23 protein. This gene has been found to have multiple polyadenylation sites. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2009]
Transcript Variant: This variant (1) represents the shortest transcript and encodes the longer isoform (1). Variants 1, 2 and 3 encode the same 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.