

Product datasheet for **SC119779**

IGFBP3 (NM_000598) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	IGFBP3 (NM_000598) Human Untagged Clone
Tag:	Tag Free
Symbol:	IGFBP3
Synonyms:	BP-53; IBP3
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL5</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF:

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>OriGene sequence for NM_000598 edited
GAATTCGGCACGAGGTGGGCGCTGAGGATCAGCCGCTTCTGCTGGATTCCACAGCTTC
GCGCCGTGTAAGTGTGCGCCCATCCCTGCGCGCCAGCCTGCCAAGCAGCGTGCCCGGTT
GCAGGCGTCATGCAGCGGGCGGACCCACGCTCTGGGCGCTGCGCTGACTCTGCTGGTG
CTGCTCCGCGGGCCGCGGTGGCGGGGCTGGCGGAGCTCGGGGGGCTGGGTCCCCTG
GTGCGCTGCGAGCCGTGCGACGCGCGTGCAGTGGCCAGTGCGCGCTCCGCGCGCGTG
TGCGCGGAGCTGGTGGCGGAGCCGGGCTGCGGCTGCTGCTGACGTGCGCACTGAGCGAG
GGCCAGCCGTGCGGCATCTACACCGAGCGCTGTGGCTCCGGCCTTCGCTGCCAGCCGTCG
CCCGACGAGGCGCGACCGCTGCAGGCGCTGCTGGACGGCCGCGGGCTCTGCGTCAACGCT
AGTGCCGTGAGCCGCTGCGCGCTACCTGCTGCCAGCGCCGAGCTCCAGGAAATGCT
AGTGAGTCGGAGGAAGACCCGAGCGCCGCGAGTGTGGAGAGCCCGTCCGTCTCCAGCAG
CACCGGTGTCTGATCCCAAGTTCACCCCTCCATTCAAAGATAATCATCATCAAGAAA
GGGCATGCTAAAGACAGCCAGCGCTACAAAGTTGACTACGAGTCTCAGAGCACAGATAAC
CAGAACTTCTCCTCCGAGTCCAAGCGGGAGACAGAATATGGTCCCTGCCGTAGAGAAATG
GAAGACACACTGAATCACCTGAAGTTCTCAATGTGCTGAGTCCCAGGGGTGTACACATT
CCCAACTGTGACAAGAAGGGATTTATAAGAAAAAGCAGTGTGCGCCTTCCAAAGGCAGG
AAGCGGGGCTTCTGCTGGTGTGTGGATAAGTATGGGCAGCCTCTCCAGGCTACACCACC
AAGGGGAAGGAGGACGTGCACTGCTACAGCATGCAGAGCAAGTAGACGCCTGCCGCAAGG
TTAATGTGGAGCTCAAATATGCCTATTTTGCACAAAAGACTGCCAAGGACATGACCAGC
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TATTCAGAGACTCGAGCACAGCACCAGACTTCATGCGCCGTTGAATGCTCACCACATG
TTGGTCTGAAGCGCCGACCACTGACTTTGTGACTTAGGCGGCTGTGTGCCTATGTAGAG
AACACGCTTACCCCACTCCCGTACAGTGGCAGCAGGCTTTATCGAGAATAGGAAAAAC
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AAATATGTATCTAAGAATGTTCTAGGGCACTCTGGGAACCTATAAAGGCAGGATTTTCGG
GCCCTCCTTTCAGGAATCTTCTGAAGACATGGCCAGTGAAGGCCAGGATGGCTTT
TGCTGCGGCCCGTGGGGTAGGAGGGACAGAGAGACAGGGAGAGTCAAGCTCCACATTCA
GAGGCATCACAAGTAATGGCACAATTCTCGGATGACTGCAGAAAATAGTGTGTTGTAGT
TCAACAACCTAAGACGAAGCTTATTTCTGAGGATAAGCTTTTAAAGGCAAGCTTATT
TTCATCTCTCATTTTTGCTCCTTAGCACAATGTAAAAAGAATAGTAATATCAGAAC
AGGAAGGAGGAATGGCTTGTGGGAGCCCATCCAGGACTGGGAGCACATAGAGATTC
ACCCATGTTTGTGAACTTAGAGTATTCTCATGCTTTTCTTTATAATTCACACATATAT
GCAGAGAAGATATGTTCTGTAAACATTGTATACAACATAGCCCCAAATATAGTAAGATC
TACTAGATAATCCTAGATGAAATGTTAGAGATGCTATATGATACAACCTGTGGCCATGA
CTGAGGAAAGGAGCTCACGCCCAGAGACTGGGCTGCTCTCCCGGAGGCCAAACCAAGAA
GGTCTGGCAAAGTCAGGCTCAGGGAGACTCTGCCCTGCTGCAGACCTCGGTGTGGACACA
CGCTGCATAGAGCTCTCCTTGAACACAGAGGGTCTCAAGACATTCTGCCTACCTATTAG
CTTTTCTTATTTTTTAACTTTTTGGGGGAAAAGTATTTTTGAGAAGTTTGTCTTGCA
ATGTATTTATAAATAGTAATAAAGTTTTTACCATTAATAAAAAAAAAAAAAAAAAAACTCGAC
    
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5' Read Nucleotide Sequence:

>OriGene 5' read for NM_000598 unedited
 TTGTAATACGACTCACTATAGGGCGGCCGGAATTCGGCACGAGGTGGGCGCTGAGGATC
 AGCCGCTTCTGCCTGGATTCCACAGCTTCGCGCCGTGTACTGTCGCCCCATCCCTGCGC
 GCCCAGCTGCCAAGCAGCGTCCCCGGTTGCAGGCGTCATGCAGCGGGCGGACCCACG
 CTCTGGGCGCTGCGCTGACTCTGCTGGTGTCTCCGCGGGCCGCCGGTGGCGGGGCT
 GGCGCGAGCTCGGGGGCTTGGGTCCCCTGGTGCCTGCGAGCCGTGCGACGCGCGTGCA
 CTGGCCCCAGTGCAGCCCTCCGCCCGCGTGTGCGCGGAGCTGGTGCAGCGCCGGGCTGC
 GGCTGCTGCCTGACGTGCGCACTGAGCGAGGGCCAGCCGTGCGGCATCTACACCGAGCGC
 TGTGGCTCCGGCCTTCGCTGCCAGCCGTCGCCCGACGAGGCGCGACCCGCTGCAGGCGCTG
 CTGGACGGCCGCGGGCTCTGCGTCAACGCTAGTGCCGTGACCCGCTGCGCGCTACCTG
 CTGCCAGCGCCGACGCTCCAGGAAATGCTAGTGAGTCGGAGGAAGACCGCAGCGNCCGG
 CAGTGTGGAGAGCCCGTCCGTCTCCAGCACGCACCGNGTGTCTGATCCCAGTTCACC
 CCCTCCATTCAAAGATAATCATCATCAAGAAAGGGCATGCTAAAGACAGCCAGCGCTACA
 AAGTTGACTACGAGTCTCAGAGCACAGATACCCAGAACTTNCTCTNCGAGTNCAGCGGGA
 GACGGAATATGGTCCCTGGCGTAGAAAAATGGAGAACAACACTGNATCACCTGAGTNCATG
 GGCTGATCCCCAGGGGTACCATTCCCCTGACANAAAGGGATTATAC

3' Read Nucleotide Sequence:

>OriGene 3' read for NM_000598 unedited
 CGCGGCCCAATCTANAGTCGAGTTTTTTTTTTTTTTTTTTTAAATGGTAAAACTTTATTT
 ACTATTTATAAATACATTGCAAGACAACTTCTCAAAAATACTTTTCCCCCAAAAAGTT
 AAAAAATAAAGAAAAGCTAATAGGTAGGCAGAATGTCTTGAGACCCCTCTGTTTTCAAG
 GAGAGCTCTATGCAGCGTGTGTCCACACCGAGGTCTGCAGCAGGGCAGAGTCTCCCTGAA
 CCTGACTTTTGCAGACCCCTTTTTGGGTTGGGCCTCCGGGAAAACACCCCATTTTT
 GGGGGGGACCCCTTTTTCTTAATTTGGGGCCACGTGGGTTAATATAGCCCTTTTA
 ACATTCTCCCTCGAATATTCTCCACTAAAACCTTCTATTTTTTGGGGGGTTTTTTG
 GGTACCATGGGTTAAAAAAAACCTTTTTTTTTGGTTATTTTTTTGGGGGGTTTTATA
 AAAAAAGTGTGAAGTTTCTTTTTTACCACCCACGGGGGTTTCTTTTTTCCCCC
 CGCGCGGGGGGGGGCCCCACAAAAACCCCTCTCTGTTGGGGGAAAAAATTTTTTTTT
 TTTCTCGGCGGGGGGGGGCGCTTAAAAGAAAAAAAATCCGTCTCTGCTCTTACTC
 TCCCACCAAAACCTTTTTGTTGGTGGGGGAAACAACCACACAATTCTTTTTTTTCCA
 AAACACGCAACACCCCTCCCTCTCCCTTTTCAACAGACCTGAGCTTTCTTTTCTTTCTT
 TCCCCCCCCCCCCCGCTCATACTTCAATACTCTTGTGCTGCTCCTTCTTTCTTCTT
 CCGCCCTTCTCGTCTCTCCT

Restriction Sites:

NotI-NotI

ACCN:

NM_000598

Insert Size:

2440 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:	<ol style="list-style-type: none">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_000598.4 , NP_000589.2
RefSeq Size:	2620 bp
RefSeq ORF:	876 bp
Locus ID:	3486
UniProt ID:	P17936
Cytogenetics:	7p12.3
Domains:	thyroglobulin_1, IB
Protein Families:	Druggable Genome, Secreted Protein
Protein Pathways:	p53 signaling pathway
Gene Summary:	<p>This gene is a member of the insulin-like growth factor binding protein (IGFBP) family and encodes a protein with an IGFBP domain and a thyroglobulin type-I domain. The protein forms a ternary complex with insulin-like growth factor acid-labile subunit (IGFALS) and either insulin-like growth factor (IGF) I or II. In this form, it circulates in the plasma, prolonging the half-life of IGFs and altering their interaction with cell surface receptors. Alternate transcriptional splice variants, encoding different isoforms, have been characterized. [provided by RefSeq, Jul 2008]</p> <p>Transcript Variant: This variant (2) uses an alternate in-frame splice site in the 5' coding region, compared to variant 1, resulting in a shorter protein (isoform b).</p>