

Product datasheet for **SC336893**

ASB15 (NM_001290258) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: ASB15 (NM_001290258) Human Untagged Clone
Tag: Tag Free
Symbol: ASB15
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC336893 representing NM_001290258.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

ATGGATACTAATGATGACCCTGATGAAGACCATCTTACAAGTTATGATATTCAGCTAAGTATTCAAGAA
 TCCATTGAAGCCAGCAAGACTGCACTTTGTCTGAAAGATTTGTACCCCTAAGTGCTCAAAACAGAAAA
 CTTGTGGAGGCCATAAAACAAGTCCATTCTGAGCTCCAGGAGTATGTAAATATAAATATGCAATG
 GATGAAGCTGATGAAAAAGGATGGTTTCCATTGCATGAAGCTGTTGTTCAACCCATTCAACAACTCTT
 GAGATTGTTCTGGATGCATCCTATAAGACACTCTGGGAATTCAAGACCTGTGATGGAGAAACACCCTTG
 ACTTTGGCAGTCAAAGCTGGTCTGGTGGAAAATGTAAGAACTTTATTAGAAAAGGGAGTGTGGCCCAAC
 ACAAAAAATGATAAAGGAGAGACCCCTTCTGATTGCTGTGAAAAAGGGCTCCTATGACATGGTGTCTG
 ACTCTGATCAAACATAACACTAGCCTAGACCAGCCCTGTGTCAAGCGATGGTCAGCAATGCATGAAGCA
 GCCAAGCAAGGCCGAAAAGATATCGTAGCTCTGCTGCTGAAACATGGAGGCAATGTCCACCTGAGAGAT
 GGATTTGGAGTCACACCACTAGGCGTCGCTGCCGAGTATGGTCACTGTGACGTGTTAGAACATCTAATC
 CACAAAGTGGTGTGCTTGTCTTGGCGGATGATGGGCGTCGGTGTGTTGAGGCAGCAGGAGGT
 GGCAATCCCGACTGCATTTCCCTCCTGCTGGAATATGGAGGAAGCGAAATGTACCTAACCGAGCAGGA
 CATCTTCTATACACCGAGCTGCCTATGAGGGGCATTATCTTGCACTGAAATATCTTATCCAGTAACA
 TCTAAAAATGCAATTCGAAAAAGTGGGCTAACACCAATCACTCAGCAGCAGATGGACAAAATGCACAG
 TGTCTAGAACTGCTCATTGAAAATGGTTTTGATGTCAACACTCTACTTGCTGACCACATTTCCAGAGC
 TATGACGATGAGAGGAAGACTGCGCTGTATTTGGCGTTTCTAATAATGACGTTCAATTGCACAGAAGTC
 CTTCTGGCTGCAGGTGCAGACCCAACTTAGATCCCTCAACTGTCTACTTGTTGCAGTGAGGGCCAAT
 AATTATGAAATGTCAGGCTGCTTCTCTCCCATGGAGCTAATGTCAATTGTTATTTATGCATGTGAAT
 GACACTCGTTTCCCAGTGTCATTCAATATGCTCTAAACGACGAGGTAATGCTGAGGCTATTGCTGAAT
 AATGGCTATCAAGTGGAGATGTGCTTTGACTGCATGCATGGTGACATCTTTGGAAATTCATTGTGTGG
 TCAGAGATACAGGAAGAGGTGCTGCCAGGATGGACATCTTGTTAATAAAAGATAACCCGTTCTGTGAG
 TTTATTACAGTTCCTTGGATGAAGCACTTGGTAGCAGAGTACTCGTGTACTAATAGATTACATGGAT
 TATGTTTCTCTGTGTCTAAACTGAAGTCTGCACTAGAAGTACAGAGAGAATGGCCAGAAATCCGCCAA
 ATACTAGAGAATCCTTGTTTCATTGAAGCATTGTGTGCGTTAAAAATTCGAAGGCTTATGGGTCTCCAG
 AAATCTGCCAGCCAGCCTCAGTGGAGAAGCTTCTCTACCACCAGCTATTCAAAGATACATATTATTT
 AAAGAGTATGATCTCTATGGACAAGAGCTAAAATTGACATAA

Restriction Sites: SgfI-MluI
ACCN: NM_001290258



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Insert Size:	1767 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:	<u>NM_001290258.1</u>
RefSeq Size:	2975 bp
RefSeq ORF:	1767 bp
Locus ID:	142685
UniProt ID:	<u>Q8W XK1</u>
Cytogenetics:	7q31.32
Protein Families:	Druggable Genome
MW:	65.8 kDa
Gene Summary:	This gene encodes a member of the suppressor of cytokine signaling box superfamily. The proteins in this superfamily participate in the ubiquitin-proteasome system for the degradation of proteins in the cell cycle and signal transduction pathways. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2014] Transcript Variant: This variant (2) differs in the 5' UTR compared to variant 1. Variants 1 and 2 encode the same protein. 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.