

## Product datasheet for **SC217332**

### ASXL1 (NM\_015338) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	ASXL1 (NM_015338) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	ASXL1
Synonyms:	BOPS; MDS
ACCN:	NM_015338
Insert Size:	2000 bp



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**Insert Sequence:**

>SC217332 3'UTR clone of NM\_015338

The sequence shown below is from the reference sequence of NM\_015338. The complete sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
CTCTGTGTATTGTGCCTTGTGGTGAGATAATAAATTATGGCCATGGGAAACATTGTATATTTAGTGTGT
GTATTTTGATAATGATTGATCTTAAATCTGTATACAGAATATCATTGATATAATACTCTTTAGGCAGGA
GCACTCTTGCCTTCCCCAAAATTTACTGCTAAAGCCCTCTGTCACTTGGCGACCCTTCTGGTCTTG
CTGGAGGGGTTTCTGGGTATAACCCATTGGGCTGCCAAGGCCAGCCAGCCTGAGCTCTCTGCAAGA
CAGAGCCTGATGTGCACGGAGTGGGGTTCGGGGGGTGGGGGGACTGCCTGACTCCCAGAGGGACTTG
AAACTGAAGCAAGAAGTTGCATTCTCCACCAAGGGAGTAACTACCTGAACAAAGTAGAAATGCCAG
TCTTCCACTACCCCTCCCTGCCATCTTTCTTCTGCTACTTTGGGGAGTTGATGGCCAGGAAAGAAGC
CAGCACAGGGTTAAAGTAACTCCTGGCATTGCCACCAGGGGGCTGGTGCACCTGCTGACCTCAGGGTC
ACAGTTGAGTCATTTGCCAGTTGACGGAGCAAGTTTACCTTGGTTCTGTTGCTGAAGCAAAATTTGGAA
CTTTTCTGTCTCAGTGTGATCCACTAACCCACAGGATCATTGGAACTTGAATAGCTCTGCTTGGACA
ATGGGGTTGGGAATAGGGTTGTCTTTCCTATGAAAATGCCATCTGTAGACCTTGTGAGTCAGCCGTCC
AGATGTTTGCAGGTGAATTCCTCTGCTTGCATCTCCCTGTCACCTTTGGACCCTATGGGAGTGGGCAT
CTCCACGCACCTGTGATGTGAAAGTCATTTACATTTCAAAGCAGTGTGTGTTTCTATTTTTATATT
TTAACTCTTTATTCTTGGATGTATAAAGTGAACTTTTGGCTTCTGTAAGTATGCTCTATGCACCTCT
AATGTTTTATCATGTATTTATATGTTGTACACAGTACTGGCTGATTCTGTAATGGATGTATTGTACAG
AGAACGAACGTCTCTTCCCTAATTTTACATCTTCAGCATCATTGCATTAAGTGGTGAATCTCCCTTC
TCTACATCTGTTGTCAGAGCCACTGAGTGTCTGTGCTGCTCGACGTGAGGGTGAATGATTGACTTGTGA
CCTGCCAGGTTGCCCGATGCCCTGTTGGGTACCGGCTGGACCTGCTGCAGCCTGCAGAGCCACAGTCA
GCCTGCCACATGCCACCGAGCAAACGCATCTTCTTTTACATCTCTCCTCCTACAGCCTTAATGGCT
GCTTGTGCCATATGTGACAAATCACCACCACAGTGTAAAGTCTTCTGGATTGATGGGTGAGTCCC
TGGGCAGCCCCAGGAAGGCTTCCAGATCTGGCTCCAGGGTACCACCTGTACAGCAATACCTGGGA
CCATGCTCTCCTGGGACTGTGAGGCTCCTTTTACGTACTTTTGCATCAGGCAGTTTGGGAAGAAAC
AAAGCCATGCCTGCTCCTGCCTCTCTCCAACATGTTTCCAGCAAGTAGATGCCCTGTGTGTGTTTC
CCTTGCCTTGTTCCTGCCTTATATCTGTATTTTCGACTTATTACAGAGTTGAGGGTCTTGTCTTAATT
TAGATCAAGTATAAAATTTGTATGACTTCAAGTCTCATTTTATCTGAAAGTTTTTTTCTCATTTAATC
TGATGTGGCATTTCGTGATCTGAAGCATGAGTGACAAGTTGGGAATGATGTGGTGAATTAAGTGCAG
TATTGGCCAAGTCCAAGTTGTCACTTAAAGCGTCTGTTTACCAAAGACCGGGAACAGGGGCCAAACAT
GTCCAGTCTCTTCTTCCCTCTGCTGGAACCTTTGGGACACTCAAGGGTACAGTTTGCAGTCTGCTG
GTCCATGAGGCTGCCAGAGAAAGCACTGCTTCTGTATGTCTCTTGTGGTATTGGAACAATAAACCAG
ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
    
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**Restriction Sites:**

Sgfl-Mlul

**OTI Disclaimer:**

Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms (SNPs).

**Components:**

The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.

**RefSeq:**

[NM\\_015338.6](#)

**Summary:**

This gene is similar to the *Drosophila* additional sex combs gene, which encodes a chromatin-binding protein required for normal determination of segment identity in the developing embryo. The protein is a member of the Polycomb group of proteins, which are necessary for the maintenance of stable repression of homeotic and other loci. The protein is thought to disrupt chromatin in localized areas, enhancing transcription of certain genes while repressing the transcription of other genes. The protein encoded by this gene functions as a ligand-dependent co-activator for retinoic acid receptor in cooperation with nuclear receptor coactivator 1. Mutations in this gene are associated with myelodysplastic syndromes and chronic myelomonocytic leukemia. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Sep 2009]

**Locus ID:**

171023

**MW:**

73.5