

## **Product datasheet for SC202486**

## ASPM (NM 018136) Human 3' UTR Clone

**Product data:** 

**Product Type:** 3' UTR Clones

**Product Name:** ASPM (NM\_018136) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: ASPM

Synonyms: ASP; Calmbp1; MCPH5

**ACCN:** NM\_018136

**Insert Size:** 222 bp

Insert Sequence: >SC202486 3' UTR clone of NM\_018136

The sequence shown below is from the reference sequence of NM\_018136. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site

Blue=Stop Codon

CAATTGGCAGAGCTCAGAATTCAAGCGATCGC

TGGTGATGGATACGCTTGGCATTCCTTAT**TAG**TAAATGTAAACATTTTCAGTATGTGTAAAGAAA TATTAAAGCCAATCATGAGTACGTAAAAGTGATTTTTGCTCTCCGTGTACAACTTTTAAAATCTGACTTTG TTTTAAAAAAAACATAAACTGTTCATTACATTCTTCATTTTTATCATTTATAGTTTTATGCATGTAATAAA

CTAATATGTCAT

ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCG

**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.

**RefSeg:** NM 018136.4



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## ASPM (NM\_018136) Human 3' UTR Clone - SC202486

Summary: This gene is the humar

This gene is the human ortholog of the Drosophila melanogaster 'abnormal spindle' gene (asp), which is essential for normal mitotic spindle function in embryonic neuroblasts. Studies in mouse also suggest a role of this gene in mitotic spindle regulation, with a preferential role in regulating neurogenesis. Mutations in this gene are associated with microcephaly primary type 5. Multiple transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, May 2011]

**Locus ID:** 259266