

## Product datasheet for **SC205265**

### Flotillin 1 (FLOT1) (NM\_005803) Human 3' UTR Clone

#### Product data:

**Product Type:** 3' UTR Clones  
**Product Name:** Flotillin 1 (FLOT1) (NM\_005803) Human 3' UTR Clone  
**Symbol:** Flotillin 1  
**Mammalian Cell Selection:** Neomycin  
**Vector:** pMirTarget (PS100062)  
**ACCN:** NM\_005803  
**Insert Size:** 437 bp  
**Insert Sequence:** >SC205265 3'UTR clone of NM\_005803  
The sequence shown below is from the reference sequence of NM\_005803. The complete sequence of this clone may contain minor differences, such as SNPs.  
**Blue**=Stop Codon **Red**=Cloning site

```
GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC
GTGAATCACAAGCCTTTGAGAACAGCCTGAGCCTTCAGCCCTCACAGATGCCAGCCTCATAGCTGAAG
TTGCCTGAATGATCCTCCTGTTGCATGTAACCCACTGGCCTCCCTGAGCATGTCCATTGACAGTGAGGT
CCCACCCTCATCTCTCCTTGCCAAATAGTTTGTGCCTTGTCTGAAGGGGGTTGCTCCCCTTGCCAAC
CTCACACTGCTATGATTGCCAACTCCAGCGGTCCCATGTCAGCCTTCTGATGATCCCACTCCACCCAC
CTCAACTTATTTAACTTCTAATTAATCAGACTGTTTGAGCCTGTTGTCTAGAATATTTTCTGACCA
AGACTGAGGGATGGGCTGGAGGTTTTCAACTTTGCTACCCAAATAAATTGCTGTAAGTAAGTACTAATA
AAACAGAAGCAACTGGAAATTA
ACGCGTAAGCGGCCGCGGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCCTTCTATGAAAGG
```

**Restriction Sites:** SgfI-MluI

**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\\_005803.4](#)



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**Summary:** This gene encodes an protein that localizes to the caveolae, which are small domains on the inner cell membranes. This protein plays a role in vesicle trafficking and cell morphology. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jan 2016]

**Locus ID:** 10211

**MW:** 16