

## Product datasheet for **SC207274**

### **JAKMIP2 (NM\_014790) Human 3' UTR Clone**

#### **Product data:**

|                       |                       |
|-----------------------|-----------------------|
| <b>Product Type:</b>  | 3' UTR Clones         |
| <b>Symbol:</b>        | JAKMIP2               |
| <b>Synonyms:</b>      | JAMIP2; NECC1         |
| <b>Mammalian Cell</b> | Neomycin              |
| <b>Selection:</b>     |                       |
| <b>Vector:</b>        | pMirTarget (PSI00062) |
| <b>ACCN:</b>          | NM_014790             |
| <b>Insert Size:</b>   | 2000 bp               |



**Insert Sequence:**

>SC207274 3'UTR clone of NM\_014790

The sequence shown below is from the reference sequence of NM\_014790. The complete sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

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GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAAGGCCAAGAAGGGCGGAAAGATCGCCGTG
TAACAATTGGCAGAGCTCAGAATTCAGCGATCGCC
GAAGAAAAGAGTAACCGAAAACATGGATAGATCCCAGGAAGACAAGTGCTTCTAAACCTTCAAAGATG
GCAAAATTGTTTACACCAAGTGAGAGGGAGATCAAAAGCTAAGAACTACCTGTAGCCAGGACTACAACT
GTGTATTTTAAAGCCATTATTCAGGTTTCTTACTTGACAGTTCTACACAACCTGTTGAAATCTAC
AATATATGCTGCATTTAATGAAACATGTATATGTCAAATCAGAAGAGAAGAACTATAACATATATTGT
GTAAAGAAAAGTTTCAGCAATGGAAGTCTGTCAGATCAAGCAAGATGTGCTTTGGGCATGGAAC
CAAAGTTACATGAAATATTCAACCCCTGCTGTGACGGGGGTCATTTTAAATGTAACACCACACCCCAT
GGAAACACTAGTCTGATAATAAACATCATTTTAAAGATCAAAACAAACAAAAACAAAGGT
GGGTGGGAGTGAAACACGAGGAATACCTATGAAGAGCTATTTACAATAAAATGTTTCATTTGAAAAGT
CTGGTTTCTTACTACAGGATTTGCTTTTCTGTCTTTATGATTGTTTTAACTCAGGAACAGAGACATC
CGCATCGTAAATGGGAAGAAAAACACTCTCTGTTTTGTTTTCTGCAAAAGGTGGAGCTTGGAATTC
ATTCCACAGACGACAGATGAGAAGTGGATCACAGTGACCACAACAAACTGGTAACGAAGTCAAAGGAC
CCGAGCGAATGTGACATGTGAAATAAAGCAGATTCGATGCTTCTTGTCTACTCTCACATCTGCATT
CCTCTATCAACAATGGACTGATCTCATTCATATTGAAAACCTAAGAAGATGATACTGTAATCACATGCT
CCCTGACTTAGTTCTGTTTAGAACGCCAACATCTATAGGGAACATGGGCCAGAGAGCATCTCTGAATCT
TTTATTTCCAAAATACTGTTTTTCAAATGGCTGCTTAAATTCCTGCAATGAATATGCTATATGATT
TGGGCAAAACACAGCAGAATCAATGAGTAAATTTCTACCAAGAGTCGGGTTTCCACAGAGATGTGAA
TATTAACCTAACTATTATTTTTTAAAGAACTCCTCACATGCCTAGAGCATTCTGACCTCTTCTGTTTA
GAGTGATAACATTTTTTCCAGACAGATGACAGCAACAAGTAGCTCTACTTTTTATATGGAAATAAC
ATTATGTTACTGACAAGTCATTTTCTACCAGAGACAATTAGAGGACTGGTACTTTTCTAAGTTGTGT
ATTACTGAGTCTTGAGTCATCTCTGAGAGCAATGTATGTTCTGAAAAGCTAATTTTACACCTCATT
TACTGTAGATAAACAGAACAACTCAATTTTTCAAATTTTTGAAGTTTTATTTATGGGGCTAGAATCCC
ATGCATAGAGAATATGTGTAGGGAGCCAGAACTCTTCTATGAAATGCTATACAAATCTTTCCATATG
TGGGTTCACTATCATCTTCAGTTTGTAGCACTGGCATTGCAAGGCCAGAACAACTGAAGAATGTA
CAATAAACTAGAAATGCTGGGGTCCCACTTTTGTCTTACATTTACTGCTGACCCAGGTATCTTTGG
CCCAGTCACTTAACCTCTCTGTGCCTCAGTTTCTCATCTGCAAAATGGGGTAAAAATGCTTTTTCATA
AAGTTTTATTTGCAATTTTAGAACAAGAGGTGCTTAAAGTAGCACAATGTGCTTTGTAATTATTACAAG
TGCTGTATTATTTGACTTTCTGTGTTGTGTGACAAAAGATTCCTCATGTAATAAAATAGTGCTTCAG
TTTGGTTAAATTTAAGCACTATAGCACTGAAGCTTGCTTATCAAGCTACTTGGGACATTATTAGCAATA
AAAATTGAGAGATTCAAAACCAGGATCTTTAATAATCTATAGGAAATAGATCATCTCATGCTATTGAA
ACGCGTAAGCGGCCGCGGCATCTAGATTGCAAGAAAATGACCGACCAAGCGACGCCAACCTGCCATCA
CGAGATTTGATTCCACCGCCGCTTCTATGAAAGG
```

**Restriction Sites:**

Sgfl-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.

**Note:**

Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.

**RefSeq:** NM\_014790.5

**Summary:** The protein encoded by this gene is reported to be a component of the Golgi matrix. It may act as a golgin protein by negatively regulating transit of secretory cargo and by acting as a structural scaffold of the Golgi. Alternative splicing results in multiple transcript variants.  
[provided by RefSeq, Aug 2012]

**Locus ID:** 9832

**MW:** 78.1