

Product datasheet for **SC206623**

SIRPG (NM_018556) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones
Product Name: SIRPG (NM_018556) Human 3' UTR Clone
Vector: pMirTarget (PS100062)
Symbol: SIRPG
Synonyms: bA77C3.1; CD172g; SIRP-B2; SIRPB2; SIRPgamma
ACCN: NM_018556
Insert Size: 517 bp
Insert Sequence: >SC206623 3'UTR clone of NM_018556
The sequence shown below is from the reference sequence of NM_018556. 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
ATCTACGTCCCCTGGAAGCAGAAGACCTGACTCTCCTTCTCCCTGCCACGTGGGACCCCTCATC
TCTGCTGCCTCCTTCTTCTGAGAGGCTCAGCTTGAGAGAATGAGCCAGTGAGAAGCTTCTCTAGAC
TTGGCTCCAAACATCTCCCTCCCAAGACATCTGCCTGCCACAGGCTCCTGTTGCCTTACACACAGA
CCTGGATGCCCCAGAGCAAGGTCTTCATTCATGGTCTGAGCAGGTGCCATGGGATTGGGCTCTGGGCA
CTGACTTAACGGCACCTCCCTAGAAGGCGAGAAACATGCCAAATCTAAACACACCAGGACTCCCATCCA
TCGCCTTGAGACTGACCGTAAACCACAGACGCTCTCCAGGTTCTCAAGAGTTATCCTGCCTTCCAGATT
CCTGCCTATCCCAACTCCCAGCCTTGTGAGGTTCTCTATTGCCTTTGAATACAAATGCACTCCCAA
AGTGGTTTTAAGAAAATAAAAAGATTATCCTTCC
ACGCGTAAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA
CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG
```

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_018556.4](#)



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Summary: The protein encoded by this gene is a member of the signal-regulatory protein (SIRP) family, and also belongs to the immunoglobulin superfamily. SIRP family members are receptor-type transmembrane glycoproteins known to be involved in the negative regulation of receptor tyrosine kinase-coupled signaling processes. Alternatively spliced transcript variants encoding different isoforms have been described. [provided by RefSeq, Jul 2008]

Locus ID: 55423

MW: 19.1