

Product datasheet for **MC225343**

Speg (NM_001085370) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Speg (NM_001085370) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Speg
Synonyms:	Apeg1; AW125581; BPEG; D1Bwg1450e; mKIAA1297; SPEGalpha; SPEGbeta
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
Fully Sequenced ORF:	>MC225343 representing NM_001085370 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGC**

ATGGTGGTGGCGCCAGGAGCTGACGTGCTACTTAAGTGTATCATACCGCCAACCCCCACCCCAAGTGT
CCTGGAAAAGGATGGGTCCATGTTGCACAGCGAGGGTCGTCTTCTCATCCGGGTGAAGGTGAACGGCA
CACACTGCTGCTCAGAGAGGCCAGGCTGCCGATGCTGGGAGCTACACAGCCACTGCCACCAACGAAGT
GGCCAAGCTACCTGTGCTTCTTCACTGGCTGTGAGACCTGGCGGCTCCACATCCCCTTTAGCAGCCCCA
TCACCTCTGATGAGGAGTACCTGAGCCCCCAGAGGAGTCCCAGAGCCTGGGGAGACCTGGCCCCGAAC
CCCTACCATGAAGCTCAGTCCCAGCCAGGATCATGATTCTCCGACTCTTCTTCCAAGGCACCCCCAACG
TTCAAGGTCTCACTCATGGACCAATCGGTGAGAGAAGGTCAAGATGTCATTATGAGCATCCGTGTGCAGG
GAGAGCCCAAGCCTGTGGTTTCTGGCTGAGGAATCGACAGCCCGTGCGCCAGACCAGCGGCGCTTTC
AGAGGAGGCCGAGGGTGGGCTCTGCCGCTTGGAGTCTGGCTGCTGAACGGGGCGATGCTGGTTTCTAC
ACATGCAAGGCCGTCAACGAATATGGCGCTCGGCAGTGCAGGCCCCCTGGAGTCCGAGCACACCCTG
AAAGCCGTCCCTGGCCGTGCTGGCCCCCTGCAGGACGTGGAGTGGGGCCGGGGAGATGGCGCTGTT
TGAGTGCCTGGTGGCAGGTCCAGCTGACGTGAGGTGAGTGGCTGTGCCGGGGCCGCTGCTGACGCT
GCGCTGCTCAAATGCAAGATGCATTTTATGGCCGCAAGTGAAGCTGCTGCTCACCTCTGTGCACGAGG
ACGACAGTGGTGTCTACACCTGCAAGCTGAGCAGACCCAAAGTGAAGTGAAGTGCAGTGCACCGGCTGAC
GGTTCGGCCGTGCTGGCGCTCTTTCACCCGGTGTGGAAGATGTGGAGGTGTTGGAAGGCCGTGCT
GCCCGCTTGGATTGCAAGATCAGCGGTACCCCGCTCCCTCCGTTACCTGGACTCATTTTGGCCACCCCG
TGAATGAGGGTGACAACCTGCGGCTTCAACAGGATGGAGGTCTACACTACTGCACATTGCCGGGTGGG
CAGTGAGGACGAAGACTCTATGAAGTCACTGCCACCAACTCATGGCCAGGCCACTGCTCCGCCAG
CTGTATGTGGAGGAGCCTCGGACAGCTGCCTCTGGCCCCAGCTCGAAGCTGGAGAAGATGCCATCTATCC
CTGAGGAGCCGAGCACGGAGATCTAGAGAGGCTGTCTATTCCGACTTCTACGGCCACTGCAAGATCT
GGAAGTGGGACTAGCTAAGGAGGCCATGCTGGAGTGCCAAAGTGACCGGCCTGCCCTATCCACCATCAGC
TGGTCCACAATGGCCACCGCATTCAAAGCAGTGACGACCGACGCATGACACAGTACAGGGACATACATC



[View online »](#)

GCCTGGTGTCCAGCTGTGGGCGCTCAGCATGCTGGTGTCTATAAGAGTGAATCGCCAACAAGCTGGG
CAAAGCGGCTGCTACGCCATCTCTATGTACAGATGTGGTTCCAGGCCCTCCAGATGGTGTCCGGAA
GTAGTAGCTGTGACCGGAAGGATGGTCACACTATCGTGGAAACCCAGGAGTCTGGACATGGCTATTG
ACCCAGATTCCCTAACATACTGTGCAGCACCAGGTGTTGGGCTCGGACCAGTGGACGGCACTGGTCAC
AGGCTGCGGGAGCCTGCGTGGGAGCCACAGGGCTGAAGAAGGGGATCCAGCACATCTCCGGTCCCTC
AGCTCCAGTGGCAAGAGCAGCAGCAAGCCCTCAGCACCCTCGAAACCCGTGCAGTGTAGACATGGCC
CACCCCTGGAAGAGCGCCTGCTGTGCTGGACAAGCAGGACATTGTGTATGTGGTAGAGGCCAGCCTGC
CTGTGTACCCGCACTTTTAACCATGTGGAGGCCAGGTTGTCTGGAGGAGCTGCCGGGAGCCCTCCTC
GAGGCGGCACAGGTGTGTATGAACTGAGTCAGCCAGATGACGACCAGTATTGTCTGCGGATCTGCCGGG
TGAGCCGACGAGACCTGGGACCCTTACTTGCAGTGCCCGAAACCGCCACGGCACAAAGGCTGTCTCGT
CACCTGGAGCTGGCAGAGGCACCTCGTGTGAGTCTATCATGGAAGATGTGGAGGTGGGCGCTGGAGAA
ACCGCTCGCTTGTGTGTAGTTGAGGGAAACCACTGCCAGACATCATGTGTACAAGGATGAGGTGC
TGCTGGCCGAGAGCAACCATGTGAGCTTGTGTATGAGGAGAACGAGTGTCCCTAGTGTCTCAGCGC
AGGAAGCCAGGATGGAGGCGTCTATACCTGTACCGCCCGAACCTGGCAGGCGAAGTCTCTGCAAGGCG
GAGCTGTCTGTGCTTTCAGCTCAGACAGCCATGGAGGTGAGGGGGTTCGAGAGGATGAGGAACCCGAG
GAAGAAGACTTAGCGACTACTAGACATCCACCAAGAGATTGGCAGGGGTGCCTTCTCTACCTGCGGCG
TGATGTGGAGAGGAGCTCTGGCCTGGAGTTGACAGCAAGTTCATCCCTAGCCAGGCTAAGCCTAAAGCA
TCAGCAAGACGCGAGGCGCGCTTGTGGCCCGCTCCAGCATGGCTGTGTCTACTTCCACGAAGCCT
TTGAGAGGCGCAGGGGACTTGTGCATAGTCACGGAGCTCTGCACAGAAGAGCTTCTAGAGCGCATGGCCAG
GAAGCCACCCTGTGTGAGTCTGAGACTCGGACCTATATGCGTCAAGTGTAGAGGGGATATGCTATCTG
CATCAGAGCCATGTGCTGCACCTGGATGTAAGCCGGAGAACCTGCTTGTGTGGATGGTGCAGGGGGT
AAGACAGGTGCGCATCTGTGACTTCGGGAACGCCAGGAACTCACTCCGGGAGAGCCCCAGTACTGTCA
GTATGGCAGCCAGAGTTTGTAGCACCTGAGATTGTCAACCAGAGCCCTGTGTCTGGAGACTGACAT
TGGCCCTGGGGTGTGCTTTCCTGTGTTGACGGGATCTCCCATTTGTTGGGAGAAATGACCGGA
CAACTCATGAATATCCGAACTACAACGTGGCATTGAGGAGACTACGTTCCCTGAGTCTGAGCAGGGA
GGCCCGGGGCTTCCCTCATCAAAGTGTGGTGCAGGATCGACTGAGACCTACTGCAGAAGAGACATTGGAA
CATCCTTGGTTCAAACAGAAGCAAAGGTGCAGAGGTGAGCACAGATCACCTGAAGCTTCTCTGTAC
GGAGGAGGTGGCAGCGCTCTCAAATCAGCTACAAGTGCCACCTGGTGTGCGCCCATCCGGAAGTGT
ACGGGCTCCCCGGAGCGGTGTGGTGGCCATGCCTAGAAGACAACCCCGAGTGGGGACTCTCATCC
TCTTCACTCAGAAGAGGAAGAGTTGGAGGAACTTCCCTCAGTACCCCGCCCCCTGCAGCTGAGTTCT
CAGGTTCTCGGATATCCCTCACGACATCCCCACGAAGATGAGGCTCTGGGACCCCGAAGCTGGGGC
TGCCACCCCATGGACTGGCAAGAGCAGGAAAGGACTCCCTCTAAGGACCAGGAGGCCCCAGCCCTGAG
GCACTCCCTCTCCAGGCCAGGAGTCCCTGATGGGCCTAGCCCCAGGCGGCAGAGCTCCGAGGGGTA
GCTCTGCTGAGAGTGCCCTGCCCGAGTCCGGTGCAGGGAACCGGGCCGGAGCTTGCACAAGGCAGCATC
CGTGGAGCTGCCGAGCGCAGGAGCCCCAGCCGGGTGCCACCCGCTGACTCGGGGAGGCGCTGGGTGAG
GGCAGATGCCAGAGGCTGCAGGCCCTGCGTCAGCGGCTGTGAGGGGAGGCCCGAGGATGGCAAGG
TCAGCGGCTCAGGGGACCCCTGCTGGAGAGCCTGGGGGGCGTGCACGGGATCCAGGATGGCAAGGGC
TGCTCCAGCGAGGCTGCCCCACCACAGCCCCCTCCGAGTCCCGGGTCTGCAAAAGAGCAGCAGC
TTCTCTCAGGGCAGGCAGAGCCCCGGGTGCGCACCCGCGGGGGCACCCCTCGAAATCCCGTGG
CCAGACTTGGGGCTCGCAGGCTGCAGGAGTCTCCCTCTTATCTGCCCTTAGCGAGACCCAGCCACCGAG
CCCTGCGAGGCCAGCGTCCCCAACTCAGTATCACCAAGTCCCGGAACCTTCCGCTGTACATCCAGA
GACTCTCCCCAGCCCCGGAACCTCAGCCTGTCCAGAAAAGTCCCTGAGCCAAAACAGAACAGTCC
GAGCTGCAAAACAGCACAACCCCTTTGGCTGTGCAAAATGCCAACGCAGCCTCTCACCCCTATGCCCA
GATCATGCAGTCGCTCCAGTGTCCAGTCCCACCCTGTCCACAGGACCCCGTGTGCTCCATCAGAG
CCCAAGCCCCACGCGGTGTGTTGCCAGGGTGGCTTCCCCACCACCCGAGTCTCTGAGAAGCGCTGC
CTTCGGCCAGAACTCCCCAGTGTAGCCGAGAAAGCCGAGTCCCACGGTCCCCCAGGCCAGGCAG
CAGCCTCAGCGCAGCATCGAAAACCTGAATCAGAGGCTGTGTTGAGGCTAAGTTCAAACGCAGCCGC
GAATCGCACTGTGCGGGGCTGAGGCTACTGAGCCGCTGCGCTCGGAGGAGCGCGTCCCTCCGCTG
GGGCGGAGGACGACGGCATATACCGGCCAGTCCGGCAGGAACGCGCTAGAGCTGGTGCAGCGCCGA
GCGCTGCGCTCCGTGCAGGACCTCAGGGTAGCCGGAGAGCCGGGCTTGTGCGTGCCTTTCGCTGTCC
CTGTGCGAGAAGTGGCGGGACACCCGGGACAGCGACACCCTGCCTGGGAGTCCCGCAGCGGAGACG
GGGAGAGTCCGAGGGCGGAGCTCAGCTCGGGATCCCAGTGTGGCGGTACGAGGAGACTCAGTCT

CACGCTGGAGCGTCTGTCCAGCCGTTTGCAGCGCAGCGGCAGCAGCGAGGACTCCGGGGGCGCCTCGGGT
 CGCAGCACGCCGCTGTTTGGACGGCTGCGCAGGGCTACTTCGGAAGGGGAGAGCCTGCGACGCCCTCGGTG
 TTCCGCACAATCAGCTGGGCTCTCAGACTGGTGCCACTACACCTTCCGCTGAGTCTCTGGGCTCAGAGGC
 CAGTGGCACATCGGGTCTTTCAGCTCCTGGAGAAAGCCGAAGCCGGCACCCGGTGGGGCCTTTCCAGGCTG
 CGCAAGGACAAGGATTATCTCAGCCAAACCTCTCTCCAGTGTCCAAGAGGACTTGGGTACCAGTATG
 TGCCCAGTGAAGTCAAGTTTCCCCCAGTCTTCCACATCAAACCAAGGACCAGGTGCTGTTGGAGGGGA
 AGCAGCACCCCTGCTCTGCCTACCCGCAGCCTGCCAGCACCCCGCATCTCCTGGATGAAAGATAAGCAG
 TCCTTGCCTTTCAGAGCCCTCAGTGGTCAATTGTGCTGCAAAAGATGGACGGCAGCTGCTGAGCATACCCC
 GGGCCGGCAAGCGGCATGCTGGGCTCTATGAGTGTCTAGCCACCAATGTCTGGGCGAGCATTACCAGCTC
 CTGACTGTGGCCGTGGCCCGCATTCCAGGGAAGCTAGCTCCTCCAGAGGTGCCCCAGACCTACCATGAC
 ACAGCACTAGTAGTGTGGAAGCCAGGAGACGGTCCGGCACCTTGACATAACTCTGGAGCGGGGGTGG
 ATGGGGAATCTGTCTGGCACCCTGTGAGCTCAGGCATCCCCGACTGTTATTACAATGTGACTCAGCTGCC
 TGTCCGCGTGACCGTGCCTTCCGTGTGGCCTGTTCCAATCGTGTGGCAGGGGCCCTTTCAGCAACCCCT
 TCTGAGAAGGTCTTCATCCGAGGCACTCCAGATTCTCCTGCTCAGCCAGCTGCTGCTCCCCGAGATGCC
 CTGTTACCTCAGGGCCAACCAGGGCCCCACCGCTGACTCTCCTACCTACTGGCCCCCACCCTGCTCT
 TGCGCCCCCAGCCTCCCAGGCATCCACTCTCAGCCCCCAACTTCTCCTCATGTCTGCCAACCAGGCTTG
 TCCTCACTCAAGGCTGTGGGTCCACCACCCGCAACCCCTCCACGAAAGCACAGGGGCTTCTGGCCACCC
 AGCAAGCAGAGCCATCCCCACCCAGTATCGTGGTCAACCCAAAGTGAACCCAGGTCTTTTGTCCCTGACAC
 TGGAACCTGACCCCAACCTCTAGTCTCAAGGGGTTAAACCAGCACCTTCTCAACTTCTTGTATATG
 GTGACTTCTTTCGTGTCTGCACCTCCAGCCCCCAGGGCCCAGCCCCGAGCCCCCTCTGAGCCACCA
 AGGTGACTGTGCGGAGCCTCAGCCCAGCAAGGAGGTGGTCACTCCCAACTCTGAGAGCACCCTCT
 TCGGCAGGGTCCCCCTCAGAAACCTACACCTTCTGGAGGAGAAGGCCAGGGGCCGCTTGGCGTTGTG
 CGGTCAATGCCGGGAGAATGCTACGGGCCGAACGTTTGTGCGCAAGATCGTGCCGTACGCCGCGGAGGGCA
 AGCGACCGGTGCTGCAGGAATACGAGGTGCTTCGGACGCTGCACCACGAGCGGCTCATGTCCCTGCACGA
 GGCCTACATCACCCCTCGGTACCTCGTCTTATCGCTGAGAGCTGTGGCAACCGGGAACCTCTGTGGG
 CTCAGCGACAGATTCCGGTATTCAGAAGATGACGTGGCCACCTATGTGGTGCAGCTGCTACAAGGCTGG
 ACTACCTCCACGGCCACCAGTGTCTACACCTGGACATCAAGCCGGACAATCTACTGTTGGCCGCTGACAA
 TGCCCTTAAAATTGTGGACTTCGGCAGTGTCTCAGCCCTATAACCCCTCAGGCCCTGAAGCCACTTGGCCAC
 CGCACAGGCACTCTGGAGTTCATGGCTCCTGAGATGGTGAAGGGAGACCCCATGGCTCTGCCACGGACA
 TCTGGGAGCGGGTGTGCTCACCTACATCATGCTTAGTGGTACTCCCATTCTATGAGCCAGACCCCCA
 GGAAACAGAGGCTCGGATTGTTGGGGTGCCTTTGATGCCTTCCAGTTGTATCCTAACACATCCCAGAGT
 GCCACCCTCTTCTGAGAAAGTCTCTCAGTACATCCCTGGAGCCGGCCCTCTCTGCAGGACTGCTTGG
 CCCACCATGGCTGCAAGATGCCTACCTGATGAAGCTGCGCCGCCAGACACTCACCTTACCACCAACCG
 GCTCAAGGAATTCCTGGGCGAGCAGCGGCGACGTGGGCTGAGGCTGCTACCCGTCAAGGTGCTGCTC
 CGCTCCTACCCTGGCAGCCCCTAG

AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC
 TGGATTACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** Sgfl-RsrII
- ACCN:** NM_001085370
- Insert Size:** 7584 bp
- OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
- Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001085370.1](#), [NP_001078839.1](#)

RefSeq Size: 9885 bp

RefSeq ORF: 7584 bp

Locus ID: 11790

UniProt ID: [Q62407](#)

Cytogenetics: 1 38.88 cM

Gene Summary: This gene encodes a protein with similarity to members of the myosin light chain kinase family. This protein family is required for myocyte cytoskeletal development. Studies have determined that a lack of this protein affected myocardial development. Multiple alternatively spliced transcript variants that encode different protein isoforms have been defined. [provided by RefSeq, Mar 2010]

Transcript Variant: This variant (2) differs in the 5' UTR and 5' coding region, compared to variant 1. These differences produce a unique 5' UTR and cause translation initiation at a downstream start codon, compared to variant 1. The encoded protein (isoform 2) is shorter than isoform 1. This variant is similar, but not identical, to variant SPEGalpha described in PubMed ID: 10973969.