

Product datasheet for **MG215586**

Sptbn4 (NM_032610) Mouse Tagged ORF Clone

Product data:

Product Type:	Expression Plasmids
Tag:	TurboGFP
Symbol:	Sptbn4
Synonyms:	1700022P15Rik; 5830426A08Rik; dyn; lnd; nmf261; nmf379; qv; ROSA62; SpbIV; Spnb4
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)

ORF Nucleotide Sequence: >MG215586 representing NM_032610
Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTGACTGGATCCGGTACCGAGGAGATCTGCC
GCCCGCATCGCC

ATGGCACAAGTACCAGGGGAGGTGGACAACATGGAGGGCCAGCAGTCTCCAACAACAACAACCCCTTCAG
CTCGCTGGGAGAGCCAGATCGGGGCTGGGATCGGGAACCTCCCGCAGCCGCAATGCTGCGGCTTCACT
CTTTGAGTGTCTCTCGGATCAAGGCTTTAGCAGATGAGCGAGAGGCTGTGCAGAAGAAAACCTTCAACAAG
TGGGTGAATTCGCACCTTGCCCGCTGGGCTGCCACATTGGAGACCTGTATGCAGACCTTCGGGACGGGT
TTGTCCCTCACCCGGCTTCTGGAAGTGTCTCGGGGGAGCAGCTGCCGAGGCCCACTCGCGGCCGATGCG
GATCCACTCGCTGGAGAAGCTGGACAAGGCTCTGCAAGTTCCTGAAGGAGCAGCGTGTGCATCTGGAGA
GTGGGGTACATGACATCGTGGATGGGAACCCGACTGACGCTGGGGTGGTCTGGACCATCATCTGC
GCTTCCAGATCAAGTCAAGATTGAGACAGAAGACAACAGAGACCCGCTCCGCCAAGGACGCTCT
GCTTGTGGTGGCAGATGAAAACAGCCGTTACCCTGAGGTAACATCCAGAATTTCAACCACGAGCTGG
AGAGATGGCCTGGCCTTCAATGCCCTATTACCCGTCACAGGCTGATCTTGTGGACCTCAGCAAATTA
CAAAGTCCAACGCCAATAACAACCTGCAGAGAGCTTCCGCACAGCCGAGCAGCACCTGGGGCTAGCAG
TCTGCTGGACCCCAAGATGTGAACATGGAGGCTCCAGATGAGAAGTCCATTATCACCTATGTGGTCTCC
TTCTACCACTACTTCTCCAAGATGAAGGCTCTGGCGGTGGAGGAAAACGGATTGAAAAGGTCCTGGAAC
AGGTGTTGGAGGTGGACAAGATCATAGAAGCTATGAGGAGCTGGCGGCGAGCTGTTGGCCTGGATCCA
TCGCACTGTGGACCTATTAGCAACAAAAGTTTGTAACTCACTGAGTGGGGTGCAGCAGCAGCTGCAG
GCCTTACGGCTTACTGTACGCTGGAGAAGCCCGTCAAGTTTCAGGAGAAGGGAACTGGAGGTGCTAC
TCTTACGATCCAAGCAAACCTGCGCGCCACAACCGCGCCTCTTGTGCCTCGAGAAGGCTGTGGCAT
CTGGGATATTGACAAGGCGTGGGGTGGAGTGGAGAAGGAGCAGCATGAGCGAGAGGGCGGCCCTCAGAGCT
GAGCTCATCCGGCAGGAGAAGTTGGAACCTTTGGCTCAAAGATTTGACCACAAGGTGGCCATGAGGGAGA
GCTGGCTAAATGAGAACCAGCGCCTGGTCTCCAGGACAACCTCGGATATGAGCTGCCGGCAGTGGAGGC
AGCCATGAAGAAGCACGAAGCATTGAGGAGACATTGCAGCTACGAGGAGCGGGTTCAAGGTGTGGCG
GAGTTAGCCAGCGGTTGGCAGCTGAAGGCTACTATGATGCCCGCGGGTGGCTGCCAGCGTGACAGT



TCCTTCGCCAGTGGGCCCTGCTGACCGGGCTGGTAGGTGCCCGCGGACCCGGCTTGAGCAGAATCTGGC
CCTGCAGAAGTCTCCAGGAGATGGTCTACATGGTGGACTGGATGGAGGAGATGCAGACTCAGCTGCTG
TCTCGGGAATGTGGGCAGCACCTGGTGGAGGCCGATGACCTGCTGCAGAAGCACGGGCTGCTGGAGGGG
ACATCGCAGCCCAGAGTGAGCGAGTGGAGGCCCTCAATGCTGCAGCCCTGAGATTCTCCAGCTACAAGG
CTATCAGCCCTGCGACCCTCAGGTCATTTGCAACCGTGTGAACCAGTGCATGGCTGTCTGTGCGAGCTG
CAGGAGCAGGCAGCACGGCGCGCGCAGAGCTGGAAGCTTCTCGGAGCCTGTGGGCGCTGCTGCAGGAGC
TGGAGGAGGCGGAGAGTTGGGCGCGGACAAAGGAGCGCTCCTAGAGGCCACGAGCGCAGTGGTGGCGC
GGCGGGCACGGCGGGCGCGCACGACCTGTCCAGCACCCGCTCGCTGCTGGCGCAGCATAAGATCCTG
CAGGGCGAGCTGGGCGGCCGGCGGGCGCTGCTGCAGCAGGCGCTGCAGCAGCGGGGAGGAGTTGGTGGCG
CGGGCGGCTCCGTGGGCCCGGGCGCGGAGCCTTTGCAATTTGGCTGGCTGGCGGAGCGCGGGCTAGCGC
TCGGCACCAATTTGGCGCCGACCTGGACGGCTGCTGGACTGGCTTGTGACGCGTATCGTCTAGCAGCCG
CTGGAGATTTTGGTACGACGAAGCGTCCAGCCGTCGCTGGCGGCCAGCACCCGCTCTCACCGGGA
GGTGAAGCGCACCGTGGCCCTGTGGTGGCTGAGGCGCCAGCTGGCGACACTGGGAGGAGCCAGTGGC
GCAGGGCCGCTGGTGGTGGCTGTCAGGTGCGCGTGGTGAAGCCGAACAGTTGTTCCCGAGGTGACCG
AGGTGGCCGCTGCGACGACAGTGGCTTCCGGACGCGCTCGCCGTGTACCGCATGTTCCGGCAGGTACA
TGCCTGCGAGTTGGATCGGGGAGAAAGAGCAGTGGCTGCTGGCCATGCGTGTGCCGATTCCCTCGAC
GACGTTGAGGTGGTGCAGCATCGATTTGAAAGTCTTGACCAGGAAATGAACAGCCTGATGGGCCGAGTCC
TGGATGTGAACAAACAGTTCCAGGAGTTGGTAGAAGGAGGTATCCAGCTCCGATGAGGTGCGCTCTG
TCAGGACCATCTCAACAGCAGGTGGAACCGCATCGTGGAGTTGGTGAACAGCGCAAAGAGGAGATGAGC
GCGGTGCTGCTGGTGGAGAACCAGTGTGGAGGTGGCCGAGGTGCGCGCCAAAGTGCAGGAGAGGAGCAG
GGCAGTGGAGAGCGCGCCCGGGCGGGGAGCCCTGCAATGGCGCTGAGTGGCTGGAGGGCTCT
ACAGGCGCTAGAGCCGACAGGGCGCGCTGCTGGAGGAGGCGCGCTGCTGGCCGAACGCTTCCCCGCA
CAGGCCAGCGCCTGCACCAGGGCGCCGAGGAGCTAGGAGCCGAGTGGGGCGCACTGGCCGAGCGGCTC
AGGCCTGCGGCGAGGCGGTTGCGGCGCAGGTGCGCTGCAGCGTTTCTGCGGATTTGGATACTTTCT
GGACTGGCTGGTGGCGGCGCAGGAGGACAGGCGCGCTCGAGGGCGCTGCTCGCAGCCTGGAGGAG
GCGGATGGGCTGCTGGCGGCCACGCCGCTCAAGGAGGAGTGGATCAGCGTGGAGGAGACTACGCGC
GCATCGTGGCTGCCAGTGAAGCGCTGCTAGCGTCCGAGGGCGCGGAGCTGGGCCCGGCTGGCGCTAGA
CGAGTGGCTGCCGACCTCGAGGTTGGCTGGCACAACCTGCTCGGCTTGTGGGAGGAGCGCAGGAGGCG
CTGGTCCAAGCGCACGCTATCAACTTCTGAGAGACCTGTGCCAGGCGCTTGGGTGCTGCGCAACC
AGGAGTGGCGCTGCTGGCGTGAATCCCTTGCACAGTGGAGTGGTGAAGAGGCGATGAAAAGGCA
CCGGGATTTCTCACCAGATGGAGCTGAACCAGCAAAGATGCAGGTGGCGGTTCAAGCTGCAGAGAGC
CTGCTGCGGCAAGGCAACGCTACGGGAGCAGGCGCAGGAGGAGTGGCTCGGCTGCTAGAGAAGGCC
AAGAAAATCAGCTGAGGGCCAGCAGTGGATGCAAAAATCTGACCAGCTTGTGCTGCAGCACTTTCT
TAGAGATTGTCATGAGCTGGATGGTGGATCCATGAGAAGATGCTGATGGCTCGTATGGCACCCGGGAA
GACAGGCACAAAGCTGCACAAGAGATGGCTCCGGCACCAAGCGCTTCAAGGCTGAGCTGGCTCAAGG
AGTGGCTGGAGAAGATAGAAAGGGAGGGCCAGCAACTGATGCAGGAGAAACCGAGCTGGCAGCCTCTGT
GCGCAAGAAATTAGGTGAGATCCGCGAGTGTGGGCGGAGCTGGAGAGCACACAGGCCAAGGCGCGC
CAGCTCTCGAGGCCAGCAAGGCGGACCAGCTAGTGCAGAGCTTCCGGAGCTGGACAAGCGGCTGCTT
ACATGGAAGCCAGCTGCAAGACGTAGACCCCGTGGGACCTGGCCACTGTCAACAGCCAGCTCAAGAA
GTTACAGTCCATGGAGTCTCAGGTGGAGGAGTGGTGGCGAGAGTGGGCGAGCTTCCAGGCGCAGCGCG
GCGCTTCCGCTGGAGCAGGCGAGCAAGGAGTTGGTGGGGGAGCGGAGAGTGGGTTGGGCGAGCGCTGG
TGCCCTGCTGGAGCCTTTGCAGGAGCGACGCCGCTTGTGCTGGCATCCAAGGAGCTGCATCAGGTGGC
CCACGACCTGGACGATGAACTGGCTTGGGTTCAAGAGCGGCTGCCACTGGCCATGCAGACAGAGCGAGG
ACTGGCTTGCAGGCTGTCCAGCAGCACATCAAAAAGAACCAGGGCCTGCGGCGGGAGATCCAGGCTCACG
GGCCGCGCTGGAGGAGGTGCTGGAGCGCGGGCGTGTGGCTCGCTGCGCAGCCAGAGGCCGAAGC
GGTGGCCGCTGGCCAAGAGCAGCTGCAGAGCGCTGGACCGGCTGCGGGAAGCGGCCGAACGGCGGAG
CAGACGCTGGACGCGCCTTCCAGGTGGAACAGTACTACTTTGACGTGGCTGAGGTGGAGGCAATGGCTGG
GCGAGCAGGAGCTGCTCATGATGAGTGGAGACAAGGGCAAGGATGAACAGAGCACCCCTGCAGCTGCTCAA
GAAGCATTTACAACAGGAGCGTGGAGAATATGAGGAAAGCATCGCCAGCTGTACGCCAGTGC
CGGGCGCTACTGGAATGGGACACCCAGACAGTGGAGCAATCAGCCGCGCAGTCCCAAGTGGATCGCC
TCTATGTGGCGCTCAAGGAGCTGGGCGAGGAACGACAGGCTGAGCCTGGAACAGCAGTACTGGCTACTACA
GCTCAGCCGCGAGGTGGATGAGCTGGAACACTGGATAGCCGAGAAGGAGTGGTGGTGGCTCCCCAGAG
CTGGGCGAGGACTTCAACACGTGTGGTGTACAGGAGAAATTTCTCAGAGTTGCCAGTGGAGACAGGAA
CCGAGGGCGGAGCGGCTGGCGCGGTCAACCAGATGGTGGACGAGCTGATTGAGTGGTGCACACAGC
AGCGGCCACCATGGCTGAGTGAAGGACGGCTGAACGAGGCTGGGCTGAGCTGCTGGAACATATGGG

ACTCGGGCCAGCTGCTCGCTGCCTCTCGGGAGCTGCATAAGTTCTTCAGCGATGCCCGGGAGCTTCAAG
GGCAGATTGAGGAGAAGCGGAGGAGGCTGCCCCGCTGACGGCACCACCTGAACCCAGACCCAGTGCCAG
CTCCATGCAGCGGACCCGCGAGCCTTTGAGCAGCAGCTGCAGCTCCTCGTGTCCCAGGTACGGCAGCTG
CAGGAAGGGGCGGCCAGCTGCGGACGGTGTACGCTGGCGAGCACGCCGAGGCCATCGCCAGCCGCGAGC
AGGAGGTGTTGCAGGGCTGGAAGGAGCTGTAGCAGCCTGTGAAGATGCTCGTGTGCACGTAGCTCCAC
GGCCGACGCCCTGCGCTTCCACAGCCAGGCCCGGACCTGCTCTCCTGGATGGACGGCATCGCGGGCCAG
ATCGGGGACAGTGACAAACCCAGGGATGTGTATCGGTGGAGGTGCTTATGAACTACCACAGGGCCTGA
AGACAGAGTTGGAGGCGCGCTGCCTGAGCTGGCAACCTGCCAGGAGCTGGGGCGGTCTCTATTACTCAA
CAAAAGCGCCATGGCTGATGAGATCCAGGCCAGCTGGACAAGCTGGGAAGCCGGAAGGAGGAGGTGTCG
GAGAAGTGGACCGTCACTGGGAGTGGCTGCAGCAGATGTGGAGGTACACCAGTTTGTCCAGGAGGCCG
TGAGGTGGAGCAGCTTATCCGGCGACACGAGGCCCTCCGAAAAGCAGCTGCGGCCTGGGAGGAGAGGTTT
AGCTCTCTGCGGCGCTTACCACGATCGAGAACTCAAGGCAGAACAGAGTAAGCAGCCACCCACCCCT
TACTGGGGCGCAAGTTCTTTGGGGACCCACAGAGCTGGCGCCAAGGCAGCGCCTTTGCTCGCTCCAGG
GGGCTATGACAGGGGCCGAGCCTTTGGCCCGAGGGCCTCGGACACGCTGTAGCGGAGGTGCGGACC
CGGGTTGGGTATGTGCGCCAGGAGCTCAAGCCCGAGCGTCTCCAGCCGCGCATTGACCGGCTGCCAGAGA
CCTCGGGGAAGGTAGAACC CGCGGCCCGACAGCCGACGCTTGGACACGACGACACCCCTGGGACTCC
GGCGGCGACGGAGTTGGTCCGGCCCGGTGCGAGCGCCAGGAGTTAGCGGATCGTGGGAGGAGCTGCCA
CGGAGACGAAGATCAGAACCAGGAGTCCGTGGATCAACCAGAGGAGACAGCGCGGAGGCCGGCCCTG
AGCGCCAGGAGTCCGCGGACACGAGGGGCCGACAGCCTTACTCTAGGTCGCTACGAGCAGATGGAGAG
ACGGCGGGAGCGGAGGGAGCGGCGGATAGAACGACAGGAGTCTAGCGAACAGGAGACTCCCACAGGGGA
GAGCTGGTCAAAGGGAAGGCCACTCTGGCTGATATTGTGGAGCACTTCAGGAGAAAGAGGCAGGCCAG
GGATCCCTGCTGGGTGCCGTGCTGCCTCAGCCTCGCGAGCTTCCCCCTGGTGCCTGCCAACGGGCT
TGAGCCGCCCGAGCGGACACCTCGGCCCGATCGGCCGCGGGCACGGGACCGGCCAAGCCGCGCCGGCGG
CCACGGCCTCGAGAGGGTGGTGGGGCGGGGAAGCCGGCGCTCGCGCTCCGCCCTGCCACGGGAGGCT
CCGCCCCCGCGCCACCTCCGCCACTCACACAGTGCAGCACGAGGGCTTCTGTGTCGCAAGCGCGA
GCTCGACGCTAACCGAAGTCGTCCAACCGGTATGGGTGAGCCTGTACTGTGTGCTCAGCAAGGGGGAG
CTGGGCTTCTACAAGGACTCCAAAGGCCAGCATCAGGGGGCACGATGGTGGGGAACCACTGCTCAGCC
TGCACAAAGCCACAGCGAGGTGGTAGTACTACAAGAAAAAGCATGTCTTCAAGCTCCAGACCCA
GGATGGCAGTGAGTTCTTGTCCAGGCTAAAGATGAGGAGGAGATGAACGGCTGGCTGGAGGCTGTGGCT
AACTCCGTGGCAGAACACGCCGAGATCGCCGATGGGGTCAGACTACCTACCACATCATCCACAGATG
AGGGCAACCCCAAGCGTGAAGGCGGGGAGCGTAGGGCCAGCGGGCGCCGGAAG

AGCGGACCGACGCGTACGCGGCCGCTCGAG - GFP Tag - GTTTAA

Protein Sequence:

>MG215586 representing NM_032610

Red=Cloning site Green=Tags(s)

```

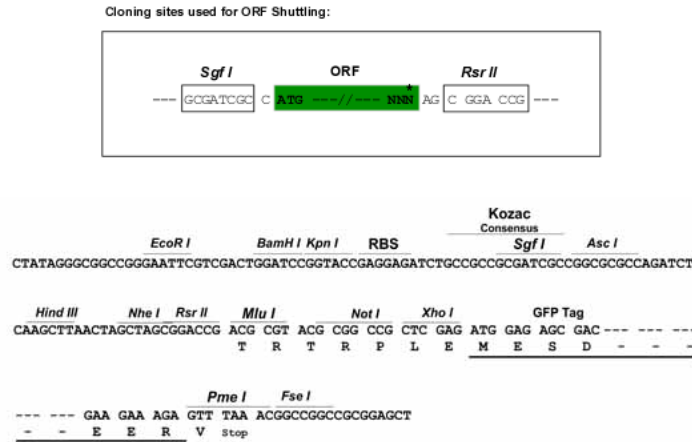
MAQVPGEVDNMEGPAVSNNNNPSARWESPDGRGWDREPPAAAANAASLFECSRIKALADEREAVQKKFTFK
WYNHSLARVVGCHIGDLYADLRDGFVLRLLLEVLSGEQLPRPTRGRMRIHSHLENDKALQFLKEQRVHLEN
VGSHDIVDGNHRLTLGLVWTIILRFQIQVIKIEIETEDNRETRSAKDALLWCQMKTAGYPEVNIQNFITTSW
RDGLAFNALIHRHRPDLVDLSKLTNSNANYNLQRAFRTAEQHLGLARLLDPEDVNMEAPDEKSIITYVVS
FYHYFSKMKALAVEGKRIGKVLQVLEVDKIIERYEELAAELLAWIHRTVDLISNQKFANSLSGVQQQLQ
AFTAYCTLEKPVKQFEKGNLEVLLFSIQSKLRAHNRRLFVPREGCGIWDIDKAWGELEKAEHEREAALRA
ELIRQEKLELLAQRFQDHKVMRESWLNENQRLVSDNFGYELPAVEAAMKKHEAIEADIAAYEERVQGV
ELAQALAAEGYDARRVAAQRDSVLRQWALLTGLVGARTRLEQNLALQKVFQEMVYVMDWMEEMQTQLL
SRECGQHLVEADDLLQKHGLLEGDIAAQSERVEALNAAALRFSQLQGYQPCDPQVICNRVNHVHGCLSEL
QEQAARRRAEAEASRSLWALLQELEEASWARDKERLLEATSGSGGAAGTAGGAHDLSTARLLAQHKIL
QGELGGRRALLQALRRGEELAAAGGSVGPGEPLHLAGLAERAASARRRWRLEEEAARRRRLQEARA
LHQFGADLGLLDWLRDAYRLAAAGDFGHDEASSRRLARQHRALTGEVEAHRGPPVGLRRLQATLGGASG
AGPLVVALQVRVVEAEQLFAEVTEVAALRRQWLRDALAVYRMFGEVHACELWIGEKEQWLLAMRVPSLD
DVEVQHRFESLDQEMNSLMGRVLDVNQTVQELVEGGHPSSDEVRSQDHLNSRWNRIVELVEQRKEEMS
AVLLVENHVLEVAEVRAQVREKRRAVE SAPRAGGALQWRLSGLEAALQALEPRQAALLEEAALLAERFPA
QATRLHQGAEEELGAEWALAGAAQACGEAVAAAGRLQRFLRDLDTFLDWLVRAQEAAGAVEGPLPRSLEE
ADGLLARHAALKEEVDQREEDYARIVAASEALLASEGAELGPGALDEWLPHEVGVHKKLLGLWEERREA
LVQAHVYQLFLRDLQALAVLRNQEVALSGAELPCTVESVEEAMKRHRDFLTTMELNQQKMQVAVQAAES
LLRQGNAYGEQAQEAVALRLEKSQENQLRAQQWMQKLLDQLVLQHFRLDCHLDGWIHEKMLMARDGTRE
DSHKLHKRWLRHQAFMAELAQNKWELEKIEREGQQLMQEKPELAASVRKKLGEIRQCWAELESTTQAKAR
QLFEASKADQLVQSF AELDKRLLHMESQLQDVPDGGDLATVNSQLKKLQSMESQVEEWCREVGEQQAQTA
ALPLEQASKELVGERQSAVGERLVRLLLEPLQERRRLLASKELHQVAHDLDELAWVQERLPLAMQTERG
TGLQAVQQHIKKNQGLRREIQAHGPRLEEVLERAGVLSASRSPEAEAVRRGQEQLQSAWTGLREAAERRQ
QTLDAAFQVEQYFDVAEVEAWLGEQELMMSEDKGKDEQSTLQLLKKHLQLEQGVENYEEIAQLSRQC
RALLEMGHPDSEQISRRQSVDRLYVALKELGEERRVSLQYQYWLQLSRQVDELEHWIAEKEVVAAGSPE
LGQDFEHVSVLQEKSEFASETGTAGRERLAAVNQMDELIECGHTAAATMAEWDGLNEAWAELELMG
TRAQLLAASRELHKFFSDARELQGGIEEKRRRLPRLTAPPEPRPSASSMQRTRLRAFEHDLQLLVSQVRQL
QEGAAQLRTVYAGEHAETIASREQEVLQGWKELLAACEDARLHVSSSTADALRFHSQARDLLSWMDGIAGQ
IGAADKPRDVSSVEVLMNYHQGLKTELEARVPELATCQELGRSLLLNKSAMADEIQAQLDKLGSRKEEVS
EKWDRHWELQQLMEVHQFAQEAADVADAWLTAQEPLQSRRELSSVDEVEQLIRRHEAFRKAANAWEERF
SSLRRLTTIEKLKAEQSKQPPPTLLGRKFFGDPTELAAKAAPLLRPGGYDRGLEPLARRASDTLSAEVRT
RVGYVRQELKPERLQPRIDRLPETSQKVEPAAPTAAALDTPGTPAATELVRPRSERQELADRAEELP
RRRRSERQESVDQPEETARRRRPERQESADHEGPHSLTLGRYEQMERRRERRRRIERQESSEQETPTRG
ELVKGKATLADIVEQLQEKEAGPGIPAGVPSLPQPRELPPGRLPNGLEPPERTPRPDRPRARDRPPRRR
PRPREGGEGGSRRSAPAQGGSAPAPPPPTHTVQHEGFLLRKRELDANRKSNSRSWVSLYCVLSKGE
LGFYKDSKGPASGGTHGGEPLL SLHKATSEVASDYKKKKHVFKLQTDGSEFLLQAKDEEEMNGWLEAVA
NSVAEHAETIARWGQTLPTTSSSTDEGNPKREGGERRASGRRK
    
```

SGPTRRRLE - GFP Tag - V

Restriction Sites:

Sgfl-RsrII

Cloning Scheme:



ACCN: NM_032610

ORF Size: 7683 bp

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. [More info](#)

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.

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.

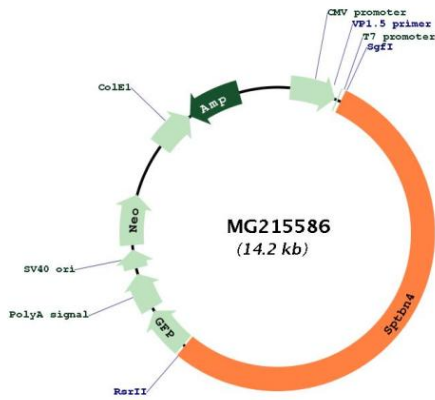
Note: Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.

RefSeq: [NM_032610.2](#), [NP_115999.2](#)

RefSeq Size: 8737 bp

RefSeq ORF: 7686 bp
 Locus ID: 80297
 Cytogenetics: 7 15.88 cM

Product images:



Circular map for MG215586