

Product datasheet for SC115667

Unrip (STRAP) (NM_007178) Human Untagged Clone

Product data:

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|---------------------------|---|
| Product Type: | Expression Plasmids |
| Product Name: | Unrip (STRAP) (NM_007178) Human Untagged Clone |
| Tag: | Tag Free |
| Symbol: | Unrip |
| Synonyms: | MAWD; PT-WD; UNRIP |
| Mammalian Cell Selection: | None |
| Vector: | <u>pCMV6-XL5</u> |
| E. coli Selection: | Ampicillin (100 ug/mL) |
| Fully Sequenced ORF: | >OriGene ORF within SC115667 sequence for NM_007178 edited (data generated by NextGen Sequencing) |

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ATGGCAATGAGACAGACGCCGCTCACCTGCTCTGGCCACACGCGACCCGTGGTTGATTTG
GCCTTCAGTGGCATCACGCCTTATGGGTATTTCTTAATCAGCGCTTGCAAAGATGGTAAA
CCTATGCTACGCCAGGGAGATACAGGAGACTGGATTGGAACATTTTTGGGTCATAAAGGT
GCTGTTTGGGGTGCAACTGAATAAGGATGCCACCAAAGCAGCTACAGCAGCTGCAGAT
TTCACAGCCAAAGTGTGGGATGCTGTCTCAGGAGATGAATTGATGACCCTGGCTCATAAA
CACATTGTCAAGACTGTGGATTTACGCAGGATAGTAATTATTTGTTAACCGGGGACAG
GATAAACTGTTACGCATATATGACTTGAACAACTGAAGCAGAACCTAAGGAAATTAGT
GGTCATACTTCTGGTATAAAAAAAGCTCTGTGGTGCAGTGAGGATAAACAGATTCCTTCT
GCTGATGACAAACTGTTGACTTTGGGATCATGCTACTATGACAGAAGTAAAATCTCTA
AATTTTAAATATGTCTGTTAGTAGTATGGAATATATTCCTGAGGGAGAGATTTTGGTTATA
ACTTATGGACGATCTATTGCTTTTCATAGTGACAGTAAGTTTGGACCCAATTAATCCTTT
GAAGCTCCTGCAACCATCAATTCTGCATCTCTTCATCCTGAGAAAAGAATTTCTTGTGCA
GGCGGTGAAGATTTTAACTTTATAAGTATGATTATAATAGTGGAGAAGAATTAGAATCC
TACAAGGGACACTTTGGTCCTATCACTGTGTGAGATTTAGTCCTGATGGAGAATCTAT
GCCAGTGGTTCAGAAGATGGAACATTGAGACTATGGCAAATGTGGTAGGAAAAACGTAT
GGCCTTTGGAAATGTGTGCTTCTGAAGAAGATAGTGGTGAAGCTGGCAAAGCCAAAGATT
GGTTTTCCAGAGACAACAGAAGAGGAGCTAGAAGAAATTGCTTCAGAGAATTCAGATTGC
ATCTTTCTTCAGCTCCTGATGTTAAGGCCTGA

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Clone variation with respect to NM_007178.3



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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_007178.2](#), [NP_009109.2](#)

RefSeq Size: 1867 bp

RefSeq ORF: 1053 bp

Locus ID: 11171

UniProt ID: [Q9Y3F4](#)

Cytogenetics: 12p12.3

Domains: WD40

Protein Families: Druggable Genome

Gene Summary: The SMN complex plays a catalyst role in the assembly of small nuclear ribonucleoproteins (snRNPs), the building blocks of the spliceosome. Thereby, plays an important role in the splicing of cellular pre-mRNAs. Most spliceosomal snRNPs contain a common set of Sm proteins SNRPB, SNRPD1, SNRPD2, SNRPD3, SNRPE, SNRPF and SNRPG that assemble in a heptameric protein ring on the Sm site of the small nuclear RNA to form the core snRNP. In the cytosol, the Sm proteins SNRPD1, SNRPD2, SNRPE, SNRPF and SNRPG are trapped in an inactive 6S pICln-Sm complex by the chaperone CLNS1A that controls the assembly of the core snRNP. Dissociation by the SMN complex of CLNS1A from the trapped Sm proteins and their transfer to an SMN-Sm complex triggers the assembly of core snRNPs and their transport to the nucleus. STRAP plays a role in the cellular distribution of the SMN complex. Negatively regulates TGF-beta signaling but positively regulates the PDPK1 kinase activity by enhancing its autophosphorylation and by significantly reducing the association of PDPK1 with 14-3-3 protein.[UniProtKB/Swiss-Prot Function]