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Product datasheet for RC223112L3V

SERPING1 (NM_001032295) Human Tagged ORF Clone Lentiviral Particle

Product data:

| Product Type: | Lentiviral Particles |
|------------------------------|---|
| Product Name: | SERPING1 (NM_001032295) Human Tagged ORF Clone Lentiviral Particle |
| Symbol: | SERPING1 |
| Synonyms: | C1IN; C1INH; C1NH; HAE1; HAE2 |
| Mammalian Cell Selection: | Puromycin |
| Vector: | pLenti-C-Myc-DDK-P2A-Puro (PS100092) |
| Tag: | Myc-DDK |
| ACCN: | NM_001032295 |
| ORF Size: | 1500 bp |
| ORF Nucleotide Sequence: | The ORF insert of this clone is exactly the same as(RC223112). |
| 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. <u>More info</u> |
| OTI Annotation: | This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene. |
| RefSeq: | <u>NM 001032295.1, NP 001027466.1</u> |
| RefSeq Size: | 1832 bp |
| RefSeq ORF: | 1503 bp |
| Locus ID: | 710 |
| UniProt ID: | <u>P05155</u> |
| Cytogenetics: | 11q12.1 |
| Protein Families: | Druggable Genome, Secreted Protein |
| Protein Pathways: | Complement and coagulation cascades |



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| | SERPING1 (NM_001032295) Human Tagged ORF Clone Lentiviral Particle – RC223112L3V |
|---------------|---|
| MW: | 55.2 kDa |
| Gene Summary: | This gene encodes a highly glycosylated plasma protein involved in the regulation of the complement cascade. Its encoded protein, C1 inhibitor, inhibits activated C1r and C1s of the first complement component and thus regulates complement activation. It is synthesized in the liver, and its deficiency is associated with hereditary angioneurotic oedema (HANE). Alternative splicing results in multiple transcript variants encoding the same isoform. [provided by RefSeq, May 2020] |

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