

## Product datasheet for **RC219331L3V**

### **CSPS (SULT1A3) (NM\_001017387) Human Tagged ORF Clone Lentiviral Particle**

#### **Product data:**

<b>Product Type:</b>	Lentiviral Particles
<b>Product Name:</b>	CSPS (SULT1A3) (NM_001017387) Human Tagged ORF Clone Lentiviral Particle
<b>Symbol:</b>	CSPS
<b>Synonyms:</b>	STM, HAST, HAST3, M-PST, ST1A5, TL-PST, SULT1A4, MGC117469
<b>Mammalian Cell Selection:</b>	Puromycin
<b>Vector:</b>	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
<b>Tag:</b>	Myc-DDK
<b>ACCN:</b>	NM_001017387
<b>ORF Size:</b>	885 bp
<b>ORF Nucleotide Sequence:</b>	The ORF insert of this clone is exactly the same as(RC219331).
<b>OTI Disclaimer:</b>	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. <a href="#">More info</a>
<b>OTI Annotation:</b>	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
<b>RefSeq:</b>	<a href="#">NM_001017387.1</a> , <a href="#">NP_001017387.1</a>
<b>RefSeq Size:</b>	1441 bp
<b>RefSeq ORF:</b>	887 bp
<b>Locus ID:</b>	6818
<b>Cytogenetics:</b>	16p11.2
<b>Protein Pathways:</b>	Sulfur metabolism
<b>MW:</b>	34.6 kDa



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**Gene Summary:**

Sulfotransferase enzymes catalyze the sulfate conjugation of many hormones, neurotransmitters, drugs, and xenobiotic compounds. These cytosolic enzymes are different in their tissue distributions and substrate specificities. The gene structure (number and length of exons) is similar among family members. This gene encodes a phenol sulfotransferase with thermolabile enzyme activity. Four sulfotransferase genes are located on the p arm of chromosome 16; this gene and SULT1A4 arose from a segmental duplication. This gene is the most centromeric of the four sulfotransferase genes. Read-through transcription exists between this gene and the upstream SLX1A (SLX1 structure-specific endonuclease subunit homolog A) gene that encodes a protein containing GIY-YIG domains. [provided by RefSeq, Nov 2010]