

Product datasheet for RC211475L3V

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

SIAH1 (NM_001006610) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: SIAH1 (NM 001006610) Human Tagged ORF Clone Lentiviral Particle

Symbol: SIAH1

Synonyms: BURHAS; SIAH1A

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK

ACCN: NM_001006610

ORF Size: 939 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC211475).

Sequence:

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.

RefSeq: <u>NM 001006610.1</u>

 RefSeq Size:
 2393 bp

 RefSeq ORF:
 942 bp

 Locus ID:
 6477

 UniProt ID:
 Q8IUQ4

 Cytogenetics:
 16q12.1

Protein Families: Druggable Genome

Protein Pathways: p53 signaling pathway, Ubiquitin mediated proteolysis, Wnt signaling pathway





ORIGENE

MW: 34.6 kDa

Gene Summary: This gene encodes a protein that is a member of the seven in absentia homolog (SIAH)

family. The protein is an E3 ligase and is involved in ubiquitination and proteasome-mediated degradation of specific proteins. The activity of this ubiquitin ligase has been implicated in the development of certain forms of Parkinson's disease, the regulation of the cellular response to hypoxia and induction of apoptosis. Alternative splicing results in several additional transcript variants, some encoding different isoforms and others that have not

been fully characterized. [provided by RefSeq, Jul 2008]