

## Product datasheet for RC203806L4V

## OriGene Technologies, Inc.

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## HSD17B8 (NM\_014234) Human Tagged ORF Clone Lentiviral Particle

**Product data:** 

**Product Type:** Lentiviral Particles

**Product Name:** HSD17B8 (NM\_014234) Human Tagged ORF Clone Lentiviral Particle

Symbol: HSD17B8

**Synonyms:** D6S2245E; dJ1033B10.9; FABGL; H2-KE6; HKE6; KE6; RING2; SDR30C1

Mammalian Cell

Selection:

Puromycin

**Vector:** pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

**ACCN:** NM\_014234

ORF Size: 783 bp

**ORF Nucleotide** 

The ORF insert of this clone is exactly the same as(RC203806).

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.

**RefSeg:** NM 014234.3

 RefSeq Size:
 1002 bp

 RefSeq ORF:
 786 bp

 Locus ID:
 7923

 UniProt ID:
 Q92506

 Cytogenetics:
 6p21.32

**Protein Families:** Druggable Genome

**Protein Pathways:** Androgen and estrogen metabolism, Metabolic pathways





ORIGENE

**MW:** 27 kDa

**Gene Summary:** 

In mice, the Ke6 protein is a 17-beta-hydroxysteroid dehydrogenase that can regulate the concentration of biologically active estrogens and androgens. It is preferentially an oxidative enzyme and inactivates estradiol, testosterone, and dihydrotestosterone. However, the enzyme has some reductive activity and can synthesize estradiol from estrone. The protein encoded by this gene is similar to Ke6 and is a member of the short-chain dehydrogenase superfamily. An alternatively spliced transcript of this gene has been detected, but the full-length nature of this variant has not been determined. [provided by RefSeq, Jul 2008]