

Product datasheet for **RC214831L3V**

VKORC1 (NM_206824) Human Tagged ORF Clone Lentiviral Particle

Product data:

| | |
|---------------------------|--|
| Product Type: | Lentiviral Particles |
| Product Name: | VKORC1 (NM_206824) Human Tagged ORF Clone Lentiviral Particle |
| Symbol: | VKORC1 |
| Synonyms: | EDTP308; MST134; MST576; VKCFD2; VKOR |
| Mammalian Cell Selection: | Puromycin |
| Vector: | pLenti-C-Myc-DDK-P2A-Puro (PS100092) |
| Tag: | Myc-DDK |
| ACCN: | NM_206824 |
| ORF Size: | 276 bp |
| ORF Nucleotide Sequence: | The ORF insert of this clone is exactly the same as(RC214831). |
| 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: | NM_206824.1 |
| RefSeq Size: | 907 bp |
| RefSeq ORF: | 279 bp |
| Locus ID: | 79001 |
| UniProt ID: | Q9BQB6 |
| Cytogenetics: | 16p11.2 |
| Protein Families: | Transmembrane |
| MW: | 9.7 kDa |



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Gene Summary:

This gene encodes the catalytic subunit of the vitamin K epoxide reductase complex, which is responsible for the reduction of inactive vitamin K 2,3-epoxide to active vitamin K in the endoplasmic reticulum membrane. Vitamin K is a required co-factor for carboxylation of glutamic acid residues by vitamin K-dependent gamma-carboxylase in blood-clotting enzymes. Allelic variation in this gene is associated with vitamin k-dependent clotting factors combined deficiency of 2, and increased resistance or sensitivity to warfarin, an inhibitor of vitamin K epoxide reductase. Pseudogenes of this gene are located on chromosomes 1 and X. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Aug 2015]