

Product datasheet for **RC229577L3V**

NKAIN2 (NM_153355) Human Tagged ORF Clone Lentiviral Particle

Product data:

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|---------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Product Type: | Lentiviral Particles |
| Product Name: | NKAIN2 (NM_153355) Human Tagged ORF Clone Lentiviral Particle |
| Symbol: | NKAIN2 |
| Synonyms: | FAM77B; NKAIP2; TCBA; TCBA1 |
| Mammalian Cell Selection: | Puromycin |
| Vector: | pLenti-C-Myc-DDK-P2A-Puro (PS100092) |
| Tag: | Myc-DDK |
| ACCN: | NM_153355 |
| ORF Size: | 423 bp |
| ORF Nucleotide Sequence: | The ORF insert of this clone is exactly the same as(RC229577). |
| 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_153355.3 , NP_699186.2 |
| RefSeq ORF: | 426 bp |
| Locus ID: | 154215 |
| UniProt ID: | Q5VXU1 |
| Cytogenetics: | 6q22.31 |
| Protein Families: | Transmembrane |
| MW: | 16.7 kDa |



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Gene Summary:

This gene encodes a transmembrane protein that interacts with the beta subunit of a sodium/potassium-transporting ATPase. A chromosomal translocation involving this gene is a cause of lymphoma. Alternative splicing results in multiple transcript variants encoding distinct isoforms. [provided by RefSeq, Jul 2014]