

## **Product datasheet for SC334464**

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## NKAIN2 (NM\_001300737) Human Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

**Product Name:** NKAIN2 (NM\_001300737) Human Untagged Clone

Tag: Tag Free Symbol: NKAIN2

Synonyms: FAM77B; NKAIP2; TCBA; TCBA1

Mammalian Cell

Selection:

Neomycin

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Fully Sequenced ORF: >NCBI ORF sequence for NM\_001300737, the custom clone sequence may differ by one or

more nucleotides

Restriction Sites: Sgfl-Mlul

**ACCN:** NM 001300737

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).





## **Reconstitution Method:**

- 1. Centrifuge at 5,000xg for 5min.
- 2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
- 3. Close the tube and incubate for 10 minutes at room temperature.
- 4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
- 5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: <u>NM 001300737.1</u>, <u>NP 001287666.1</u>

 RefSeq Size:
 3457 bp

 RefSeq ORF:
 624 bp

 Locus ID:
 154215

 UniProt ID:
 Q5VXU1

 Cytogenetics:
 6q22.31

**Protein Families:** Transmembrane

**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]

Transcript Variant: This variant (3) has an alternate exon in its 5' end compared to variant 1. This variant represents translation initiation at a downstream AUG compared to variant 1; the 5'-most initiation codon, as used in variant 1, is associated with a truncated ORF that would render the transcript a candidate for nonsense-mediated decay (NMD). Leaky scanning may allow translation initiation at the downstream AUG to encode an isoform (3) that has a shorter N-terminus, compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.