

## Product datasheet for MR208975L3V

### OriGene Technologies, Inc.

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# Cybb (NM 007807) Mouse Tagged ORF Clone Lentiviral Particle

#### **Product data:**

**Product Type:** Lentiviral Particles

**Product Name:** Cybb (NM\_007807) Mouse Tagged ORF Clone Lentiviral Particle

Symbol:

C88302; Cgd; CGD91-phox; Cyd; gp91-1; gp91 Synonyms:

**Mammalian Cell** 

Selection:

Puromycin

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

Tag: Myc-DDK NM 007807 ACCN: **ORF Size:** 1710 bp

**ORF Nucleotide** 

Sequence:

The ORF insert of this clone is exactly the same as(MR208975).

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.

NM 007807.5, NP 031833.3 RefSeq:

RefSeq Size: 4750 bp RefSeq ORF: 1713 bp Locus ID: 13058 **UniProt ID:** Q61093 Cytogenetics: X A1.1







#### **Gene Summary:**

This gene encodes the heavy chain component of a heterodimeric transmembrane ion transporter composed of both a heavy and a light chain. This transporter mediates the transfer of electrons from nicotinamide adenine dinucleotide phosphate (NADPH) to oxygen to generate superoxide. This reaction is important in the innate immune response to pathogens. However, increased activity of the encoded protein also leads to the generation of reactive oxygen species that result in oxidative stress and can cause tissue damage. Conversely, loss of function of the related gene in human causes chronic granulomatous disease. Alternative splicing results in multiple transcript variants, although the full-length nature of some of these variants has not been determined. [provided by RefSeq, May 2013]