

Product datasheet for SC313211

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GLEPP1 (PTPRO) (NM_030668) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: GLEPP1 (PTPRO) (NM_030668) Human Untagged Clone

Tag: Tag Free Symbol: GLEPP1

Synonyms: GLEPP1; NPHS6; PTP-OC; PTP-U2; PTPROT; PTPU2; R-PTP-O

Mammalian Cell

Selection:

Neomycin

Vector: pCMV6-Entry (PS100001)

E. coli Selection: Kanamycin (25 ug/mL)

Fully Sequenced ORF: >SC313211 representing NM_030668.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGGTTACAGAGATGAATCCCAATGTGGTAGTGATCTCCGTGCTGGCCATCCTTAGCACACTTTTAATT GGACTGTTGCTTGTTACCCTCATTATTCTTAGGAAAAAGCATCTGCAGATGGCTAGGGAGTGTGGAGCT GGTACATTTGTCAATTTTGCATCCTTAGAGAGGGATGGAAAGCTTCCATACAACTGGAGTAAAAATGGT TTAAAGAAGAGGAAACTGACAAACCCGGTTCAACTGGATGACTTTGATGCCTATATTAAGGATATGGCC TTTGCTGCAGATCTTCCACTGAATCGATGTAAAAAACCGTTACACAAACATCCTACCATATGACTTCAGC CGTGTGAGATTAGTCTCCATGAATGAAGAGGAAGGTGCAGACTACATCAATGCCAACTATATTCCTGGA TACAACTCACCCAGGAGTATATTGCCACCCAGGGGCCACTGCCTGAAACCAGAAATGACTTCTGGAAG TGTGACCATTACTGGCCATTCACGGAAGAACCTATAGCCTATGGAGACATCACTGTGGAGATGATTTCA GAGGAAGAGCAGGACGACTGGGCCTGTAGACACTTCCGGATCAACTATGCTGACGAGATGCAGGATGTG ATGCATTTTAACTACACTGCATGGCCTGATCATGGTGTGCCCACAGCAAATGCTGCAGAAAGTATCCTG CAGTTTGTACACATGGTCCGACAGCAAGCTACCAAGAGCAAAGGTCCCATGATCATTCACTGCAGTGCT GGCGTGGGACGGACAGGACATTCATTGCCCTGGACAGGCTCTTGCAGCACATTCGGGATCATGAGTTT TACATTTTTATCCATCAGTGTGCAACTGATGTGGATGAAGAAGAAGCAGCAGTTCTGCATCAGTGAT

GTCATATACGAGAATGTTAGCAAGTCCTAG

 ${\color{blue} \textbf{ACGCGTACGCGGCCGCTC} \textbf{GAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT} }$

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul



GLEPP1 (PTPRO) (NM_030668) Human Untagged Clone - SC313211

ACCN: NM_030668 **Insert Size:** 1134 bp

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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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 030668.2</u>

 RefSeq Size:
 3886 bp

 RefSeq ORF:
 1134 bp

 Locus ID:
 5800

 UniProt ID:
 Q16827

Cytogenetics: 12p13-p12

Domains: Y_phosphatase, PTPc_motif

Protein Families: Phosphatase, Transmembrane

MW: 43.8 kDa



Gene Summary:

This gene encodes a member of the R3 subtype family of receptor-type protein tyrosine phosphatases. These proteins are localized to the apical surface of polarized cells and may have tissue-specific functions through activation of Src family kinases. This gene contains two distinct promoters, and alternatively spliced transcript variants encoding multiple isoforms have been observed. The encoded proteins may have multiple isoform-specific and tissue-specific functions, including the regulation of osteoclast production and activity, inhibition of cell proliferation and facilitation of apoptosis. This gene is a candidate tumor suppressor, and decreased expression of this gene has been observed in several types of cancer. [provided by RefSeq, May 2011]

Transcript Variant: This variant (4) differs in the 5' UTR and has multiple differences in the coding region but maintains the reading frame, compared to variant 1. Variants 4 and 6 encode the same isoform (d), which is shorter than isoform a. 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.