

Product datasheet for SC328571

IL1RAP (NM_001167930) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: IL1RAP (NM_001167930) Human Untagged Clone

Tag: Tag Free Symbol: IL1RAP

Synonyms: C3orf13; IL-1RAcP; IL1R3

Mammalian Cell

Selection:

Neomycin

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

Fully Sequenced ORF: >SC328571 representing NM_001167930.

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

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGACACTTCTGTGGTGTGTGGTGAGTCTCTACTTTTATGGAATCCTGCAAAGTGATGCCTCAGAACGC TGCGATGACTGGGGACTAGACACCATGAGGCAAATCCAAGTGTTTGAAGATGAGCCAGCTCGCATCAAG TGCCCACTCTTTGAACACTTCTTGAAATTCAACTACAGCACAGCCCATTCAGCTGGCCTTACTCTGATC TGGTATTGGACTAGGCAGGACCGGGACCTTGAGGAGCCAATTAACTTCCGCCTCCCCGAGAACCGCATT AGTAAGGAGAAAGATGTGCTGTGGTTCCGGCCCACTCTCCTCAATGACACTGGCAACTATACCTGCATG TTAAGGAACACTACATATTGCAGCAAAGTTGCATTTCCCTTGGAAGTTGTTCAAAAAAGACAGCTGTTTC AATTCCCCCATGAAACTCCCAGTGCATAAACTGTATATAGAATATGGCATTCAGAGGATCACTTGTCCA AATGTAGATGGATATTTTCCTTCCAGTGTCAAACCGACTATCACTTGGTATATGGGCTGTTATAAAATA CAGAATTTTAATAATGTAATACCCGAAGGTATGAACTTGAGTTTCCTCATTGCCTTAATTTCAAATAAT GTAAAGGTAGTAGGCTCTCCAAAAAATGCAGTGCCCCCTGTGATCCATTCACCTAATGATCATGTGGTC TATGAGAAAGAACCAGGAGAGGAGCTACTCATTCCCTGTACGGTCTATTTTAGTTTTCTGATGGATTCT CGCAATGAGGTTTGGTGGACCATTGATGGAAAAAAACCTGATGACATCACTATTGATGTCACCATTAAC GAAAGTATAAGTCATAGTAGAACAGAAGATGAAACAAGAACTCAGATTTTGAGCATCAAGAAAGTTACC TCTGAGGATCTCAAGCGCAGCTATGTCTGTCATGCTAGAAGTGCCAAAGGCGAAGTTGCCAAAGCAGCC AAGGTGAAGCAGAAAGGTAATAGATGCGGTCAG<mark>TGA</mark>

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul



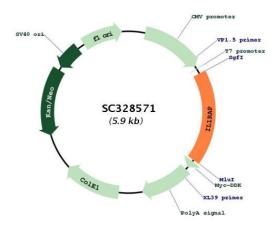
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Plasmid Map:



ACCN: NM 001167930

Insert Size: 1071 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: NM 001167930.1

RefSeq Size: 2027 bp
RefSeq ORF: 1071 bp
Locus ID: 3556



IL1RAP (NM_001167930) Human Untagged Clone - SC328571

UniProt ID: Q9NPH3
Cytogenetics: 3q28

Protein Families: Druggable Genome, Secreted Protein, Transmembrane

Protein Pathways: Apoptosis, Cytokine-cytokine receptor interaction

MW: 41 kDa

Gene Summary: This gene encodes a component of the interleukin 1 receptor complex, which initiates

signalling events that result in the activation of interleukin 1-responsive genes. Alternative splicing of this gene results in membrane-bound and soluble isoforms differing in their C-terminus. The ratio of soluble to membrane-bound forms increases during acute-phase

induction or stress. [provided by RefSeq, Jul 2018]

Transcript Variant: This variant (5) has multiple differences, compared to variant 1. It encodes the soluble isoform (2). 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.