

Product datasheet for **SC330193**

HDLBP (NM_001243900) Human Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: HDLBP (NM_001243900) Human Untagged Clone
Tag: Tag Free
Symbol: HDLBP
Synonyms: HBP; PRO2900; VGL
Vector: pCMV6-Entry (PS100001)
Fully Sequenced ORF: >SC330193 representing NM_001243900.
 Blue=Insert sequence Red=Cloning site Green=Tag(s)

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- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001243900
- Insert Size:** 3708 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).
- 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_001243900.1](#)
- RefSeq Size:** 6291 bp
- RefSeq ORF:** 3708 bp

Locus ID: 3069

UniProt ID: [Q00341](#)

Cytogenetics: 2q37.3

MW: 138 kDa

Gene Summary: The protein encoded by this gene binds high density lipoprotein (HDL) and may function to regulate excess cholesterol levels in cells. The encoded protein also binds RNA and can induce heterochromatin formation. [provided by RefSeq, Mar 2016]
Transcript Variant: This variant (3) uses a downstream promoter compared to variant 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.