

# Product datasheet for SC201924

## CD41 (ITGA2B) (NM\_000419) Human 3' UTR Clone

### **Product data:**

#### **Product Type:** 3' UTR Clones **Product Name:** CD41 (ITGA2B) (NM 000419) Human 3' UTR Clone Vector: pMirTarget (PS100062) ITGA2B Symbol: Synonyms: BDPLT2; BDPLT16; CD41; CD41B; GP2B; GPIIb; GT; GT1; GTA; HPA3; PPP1R93 ACCN: NM 000419 Insert Size: 213 bp >SC201924 3'UTR clone of NM\_000419 **Insert Sequence:** The sequence shown below is from the reference sequence of NM\_000419. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GCTACCTGCACCGCCCCTTCTCCAACAAGTTGCCTCCAAGCTTTGGGTTGGAGCTGTTCCATTGGGTCC CCCTGA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG **Restriction Sites:** Sgfl-Mlul **OTI Disclaimer:** Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences, e.g., single nucleotide polymorphisms (SNPs). The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The **Components:** package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials. **RefSeq:** NM 000419.5



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#### OriGene Technologies, Inc.

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CD41 (ITGA2B) (NM_000419) Human 3' UTR Clone – SC201924
This gene encodes a member of the integrin alpha chain family of proteins. The encoded preproprotein is proteolytically processed to generate light and heavy chains that associate through disulfide linkages to form a subunit of the alpha-IIb/beta-3 integrin cell adhesion receptor. This receptor plays a crucial role in the blood coagulation system, by mediating platelet aggregation. Mutations in this gene are associated with platelet-type bleeding disorders, which are characterized by a failure of platelet aggregation, including Glanzmann thrombasthenia. [provided by RefSeq, Jan 2016]
3674
7.7

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