

# **Product datasheet for SC205996**

### OriGene Technologies, Inc.

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## BAF53A (ACTL6A) (NM\_178042) Human 3' UTR Clone

### **Product data:**

**Product Type:** 3' UTR Clones

Product Name: BAF53A (ACTL6A) (NM\_178042) Human 3' UTR Clone

**Vector:** pMirTarget (PS100062)

Symbol: ACTL6A

Synonyms: ACTL6; Arp4; ARPN-BETA; BAF53A; INO80K

**ACCN:** NM\_178042

**Insert Size:** 445 bp

Insert Sequence: >SC205996 3' UTR clone of NM\_178042

The sequence shown below is from the reference sequence of NM\_178042. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site

Blue=Stop Codon

CAATTGGCAGAGCTCAGAATTCAAGCGATCGC

AGCCATTCCAGAAATAAAAAAAAA

**ACGCGT**AAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCG

**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).

**Components:** The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.

**RefSeq:** <u>NM 178042.2</u>





### BAF53A (ACTL6A) (NM\_178042) Human 3' UTR Clone - SC205996

**Summary:** 

This gene encodes a family member of actin-related proteins (ARPs), which share significant amino acid sequence identity to conventional actins. Both actins and ARPs have an actin fold, which is an ATP-binding cleft, as a common feature. The ARPs are involved in diverse cellular processes, including vesicular transport, spindle orientation, nuclear migration and chromatin remodeling. This gene encodes a 53 kDa subunit protein of the BAF (BRG1/brm-associated factor) complex in mammals, which is functionally related to SWI/SNF complex in S. cerevisiae and Drosophila; the latter is thought to facilitate transcriptional activation of specific genes by antagonizing chromatin-mediated transcriptional repression. Together with beta-actin, it is required for maximal ATPase activity of BRG1, and for the association of the BAF complex with chromatin/matrix. Three transcript variants that encode two different protein isoforms have been described. [provided by RefSeq, Jul 2008]

Locus ID:

86