

#### OriGene Technologies, Inc.

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# Product datasheet for SC331112

#### C1orf86 (FAAP20) (NM\_001256945) Human Untagged Clone

### **Product data:**

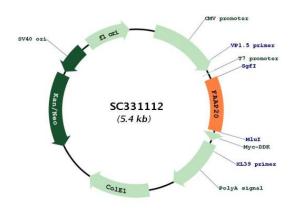
Product Type:	Expression Plasmids
Product Name:	C1orf86 (FAAP20) (NM_001256945) Human Untagged Clone
Tag:	Tag Free
Symbol:	FAAP20
Synonyms:	C1orf86; FP7162
Vector:	pCMV6-Entry (PS100001)
Fully Sequenced ORF:	>SC331112 representing NM_001256945. Blue=Insert sequence <mark>Red=</mark> Cloning site Green=Tag(s)
	ATGGAGGCGGCGCGGAGGCCGCGGCTGGGGTTGAGCCGCCGGAGGCCGCCCCGGCGGGGGGGG
<b>Restriction Sites:</b>	SgfI-Mlul



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#### Plasmid Map:



ACCN: Insert Size:	NM_001256945 570 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).

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## **C1orf86 (FAAP20) (NM\_001256945) Human Untagged Clone – SC331112**

Reconstitution Method:	<ol> <li>Centrifuge at 5,000xg for 5min.</li> <li>Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li> <li>Close the tube and incubate for 10 minutes at room temperature.</li> <li>Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li> <li>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.</li> </ol>
RefSeq:	<u>NM 001256945.1</u>
RefSeq Size:	1431 bp
RefSeq ORF:	570 bp
Locus ID:	199990
UniProt ID:	<u>Q6NZ36</u>
Cytogenetics:	1p36.33
MW:	20.5 kDa
Gene Summary:	Component of the Fanconi anemia (FA) complex required to recruit the FA complex to DNA interstrand cross-links (ICLs) and promote ICLs repair. Following DNA damage recognizes and binds 'Lys-63'-linked ubiquitin generated by RNF8 at ICLs and recruits other components of the FA complex. Promotes translesion synthesis via interaction with REV1.[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (3) lacks an alternate exon that results in a frameshift in the 5'

Transcript Variant: This variant (3) lacks an alternate exon that results in a frameshift in the 5' and subsequent coding region, and it uses an alternate 3' exon and thus differs in the 3' UTR, compared to variant 2. The encoded isoform (3) has a distinct C-terminus and is longer than isoform 2.

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