

Product datasheet for SC204540

OriGene Technologies, Inc.

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26S proteasome non ATPase regulatory subunit 12 (PSMD12) (NM_174871) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: 26S proteasome non ATPase regulatory subunit 12 (PSMD12) (NM_174871) Human 3' UTR

Clone

Vector: pMirTarget (PS100062)

Symbol: PSMD12

Synonyms: p55; Rpn5; STISS

ACCN: NM_174871

Insert Size: 364 bp

Insert Sequence: >SC204540 3' UTR clone of NM_174871

The sequence shown below is from the reference sequence of NM_174871. The complete sequence of this clone may contain minor differences, such as SNPs. Red=Cloning site

Blue=Stop Codon

CAATTGGCAGAGCTCAGAATTCAAGCGATCGC

AACATATTCCAATA

ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCG

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.





26S proteasome non ATPase regulatory subunit 12 (PSMD12) (NM_174871) Human 3' UTR Clone – SC204540

RefSeq: <u>NM 174871.2</u>

Summary: The 26S proteasome is a multicatalytic proteinase complex with a highly ordered structure

composed of 2 complexes, a 20S core and a 19S regulator. The 20S core is composed of 4 rings of 28 non-identical subunits; 2 rings are composed of 7 alpha subunits and 2 rings are composed of 7 beta subunits. The 19S regulator is composed of a base, which contains 6 ATPase subunits and 2 non-ATPase subunits, and a lid, which contains up to 10 non-ATPase subunits. Proteasomes are distributed throughout eukaryotic cells at a high concentration and cleave peptides in an ATP/ubiquitin-dependent process in a non-lysosomal pathway. An essential function of a modified proteasome, the immunoproteasome, is the processing of class I MHC peptides. This gene encodes a non-ATPase subunit of the 19S regulator. A pseudogene has been identified on chromosome 3. Alternatively spliced transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Oct 2015]

Locus ID: 5718