

Product datasheet for RC216075L4V

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

BLM (NM_000057) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: BLM (NM_000057) Human Tagged ORF Clone Lentiviral Particle

Symbol: BLM

Synonyms: BS; MGRISCE1; RECQ2; RECQL2; RECQL3

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_000057 **ORF Size:** 4251 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC216075).

Sequence:

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This

naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeg: NM 000057.1

RefSeq Size: 4437 bp
RefSeq ORF: 4254 bp

Locus ID: 641

 UniProt ID:
 P54132

 Cytogenetics:
 15q26.1

Protein Families: Druggable Genome, Stem cell - Pluripotency

Protein Pathways: Homologous recombination





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MW: 158.8 kDa

Gene Summary: The Bloom syndrome is an autosomal recessive disorder characterized by growth deficiency,

microcephaly and immunodeficiency among others. It is caused by homozygous or compound heterozygous mutation in the gene encoding DNA helicase RecQ protein on chromosome 15q26. This Bloom-associated helicase unwinds a variety of DNA substrates including Holliday junction, and is involved in several pathways contributing to the

maintenance of genome stability. Identification of pathogenic Bloom variants is required for

heterozygote testing in at-risk families. [provided by RefSeq, May 2020]