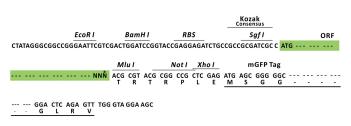


Product datasheet for RC230032L4

MEPE (NM_001184696) Human Tagged Lenti ORF Clone

Product data:

Product Type: Expression Plasmids Product Name: MEPE (NM_001184696) Human Tagged Lenti ORF Clone Tag: mGFP Symbol: MEPE OF45 Synonyms: Mammalian Cell Puromycin Selection: Vector: pLenti-C-mGFP-P2A-Puro (PS100093) E. coli Selection: Chloramphenicol (34 ug/mL) The ORF insert of this clone is exactly the same as(RC230032). **ORF** Nucleotide Sequence: **Restriction Sites:** Sgfl-Mlul **Cloning Scheme:** Cloning sites used for ORF Shuttling: ORF Sqf I Mlu I --- GCG ATC GC C ATG --- //--- NNN ACG CGT ---



* The last codon before the Stop codon of the ORF.



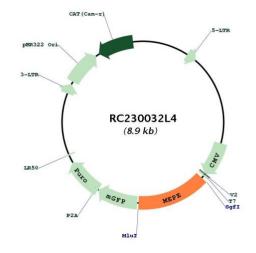
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OriGene Technologies, Inc.

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

Plasmid Map:



clone aligns with the gene accession number as a point of ual transcript sequences of the same gene can differ through g. polymorphisms), each with its own valid existence. This ent with the reference, but a complete review of all prevailing o use. <u>More info</u>
press the complete ORF with an expression tag. Expression of the gene.
olumn purified and shipped in a 2D barcoded Matrix tube ready, dried plasmid DNA (reconstitute with 100 ul of water).
n. dd 100ul of sterile water to dissolve the DNA. or 10 minutes at room temperature. en do a quick spin (less than 5000xg) to concentrate the liquid at -20°C. The DNA is stable for at least one year from date of
5.1

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	MEPE (NM_001184696) Human Tagged Lenti ORF Clone – RC230032L4
Locus ID:	56955
UniProt ID:	<u>Q9NQ76</u>
Cytogenetics:	4q22.1
Protein Families:	Secreted Protein
MW:	45.3 kDa
Gene Summary:	This gene encodes a secreted calcium-binding phosphoprotein that belongs to the small integrin-binding ligand, N-linked glycoprotein (SIBLING) family of proteins. Members of this family are components of the extracellular matrix of bone and dentin and regulate bone mineralization. Deficiency of a similar protein in mouse results in increased bone mass. Mice lacking this gene are resistant to aging-related trabecular bone loss. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Mar 2014]

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