

#### 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

## Product datasheet for MR220510L3V

#### Pitrm1 (NM\_145131) Mouse Tagged ORF Clone Lentiviral Particle

### **Product data:**

| Product Type:                | Lentiviral Particles  |
|------------------------------|---|
| Product Name:                | Pitrm1 (NM_145131) Mouse Tagged ORF Clone Lentiviral Particle   |
| Symbol:                      | Pitrm1  |
| Synonyms:                    | 2310012C15Rik; AA410010; mKIAA1104; MP-1; MP1; Ntup1; PreP  |
| Mammalian Cell<br>Selection: | Puromycin   |
| Vector:                      | pLenti-C-Myc-DDK-P2A-Puro (PS100092)  |
| Tag:                         | Myc-DDK   |
| ACCN:                        | NM_145131   |
| ORF Size:                    | 3108 bp   |
| ORF Nucleotide<br>Sequence:  | The ORF insert of this clone is exactly the same as(MR220510).  |
| 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 clone is substantially in agreement with the reference, but a complete review of all prevailing variants is recommended prior to use. <u>More info</u> |
| OTI Annotation:              | This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.  |
| RefSeq:                      | <u>NM 145131.1, NP 660113.1</u>   |
| RefSeq Size:                 | 3547 bp   |
| RefSeq ORF:                  | 3111 bp   |
| Locus ID:                    | 69617   |
| UniProt ID:                  | <u>Q8K411</u>   |
| Cytogenetics:                | 13 A1   |



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# Pitrm1 (NM\_145131) Mouse Tagged ORF Clone Lentiviral Particle - MR220510L3V Gene Summary: Metalloendopeptidase of the mitochondrial matrix that functions in peptide cleavage and degradation rather than in protein processing. Has an ATP-independent activity. Specifically cleaves peptides in the range of 5 to 65 residues. Shows a preference for cleavage after small

degradation rather than in protein processing. Has an ATP-independent activity. Specifically cleaves peptides in the range of 5 to 65 residues. Shows a preference for cleavage after small polar residues and before basic residues, but without any positional preference. Degrades the transit peptides of mitochondrial proteins after their cleavage. Also degrades other unstructured peptides. It is also able to degrade amyloid-beta protein 40, one of the peptides produced by APP processing, when it accumulates in mitochondrion. It is a highly efficient protease, at least toward amyloid-beta protein 40. Cleaves that peptide at a specific position and is probably not processive, releasing digested peptides intermediates that can be further cleaved subsequently.[UniProtKB/Swiss-Prot Function]

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