

Product datasheet for **MC218765**

Vipas39 (NM_134044) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Vipas39 (NM_134044) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Vipas39
Synonyms:	6720456H09Rik; 9330175H22Rik; AI413782; hSPE-39; SPE-39; Spe39; Vipar
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >MC218765 representing NM_134044
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGATCGC**

ATGAACCGGACAAAAGGTGATGAGGAGGAATATTGGAACAGCTCCAAGTTCAAGGCTTTCACTTTTGATG
 ATGAAGATGATGAGCTCTCACAGTTAAAGGAATCCAAACGCGCAGTGAATAGCCTTCGAGACATCGTGGA
 TGACGACGATGACGATGACCTGGAGAGAGTCAGCTGGACTGGAGAACCTGTGGGAAGTATCTCATGGTCC
 ATCAAAGAGACTGCTGGGAGTAGCGGTCAACCCCTGAGGGACGTGAACAGCTAAAGGGCCGAAACAGCT
 TCTACACACAGCTACCCAAACCTCCTTCTACTTACTCCCTCAGCAGCTTCTTCAGAGGTAGAAGTAGACC
 TGGGAGTTTCCAGTCCCTTCTGATGCTCTGTCAGACACACCTGCCAAAAGTTATGCTCCAGAGCTGGGG
 AGGCCTAAGGGAGAATACAGGTCTGTTCACTAGAGCGCTTCCGCTCCCTCCAGGATAAGCTGCAGCTCC
 TAGAGGAGGCAGTAAGCATGCATGATGGGAACGTCATCACTGCAGTTCTGATCTTCCTGAAGAGGACCT
 GAGCAAAGAGATCCTTTTTCGAGAGCTGGAGGTACGCCAGGTGCGACTGAGACATCTCATTCACTTTCTT
 AAAGAAATAGGGGACCAAAGCTGCTTTTAGATCTCTCAGGTTCTAGATAGAGCAGAAGAACTGGCGC
 TATCCCATTATCGAGAGCACTTGAACATTCAGGACCCTGAAAAACGAAAAGAATTTCTTAAGACTTGCAT
 TGGTTTGCCATTTTCAGCAGAAGACGCTGCGCATGTACAAGACCATTACACCCTGCTGGAGCGCCAGATC
 ATCATCGAGGCAAACGACCGGCACCTGGAGTCTTCAGGGCAGACTGATATCTTCCGGAAGCACCCCGTA
 AAGCTTCCATTCTCAACATGCCGTTAGTGACGACACTCTTCTACGCTGCTTCTATCACTACACGGAGTC
 TGAGGGAACCTTCAGCAGTCCCATCAACCTGAAGAAAACATTCAAGATCCCAGACAAACAGTATGTGCTG
 ACAGCCTTGGCTGCGCGTGCCAAGCTGAGAGCCTGGAATGACGTTGATGCCCTGTTCAACACAAAGAACT
 GGTGGGTTACACCAAGAAGAGAGCACCCATTGGCTTCCATCGAGTTGTGAAATTTGCACAAGAACAG
 TGCCCTGTCCAGATACTGCAGGAATATGTCAATCTGGTGGAAAGATGTGGACACCAAGTTGAACCTAGCT
 ACTAAGTTCAAGTGCCATGATGTCGTCATTGATACTTGCCGAGACCTGAAGGATCGTCAGCAGTTGCTTG
 CATACAGGAGCAAAGTAGACAAAGGATCCGCCGAGGAAGAGAAGATTGACGCCATTCTCAGCAGCTCGCA
 AATTGATGGAAGA**CTAA**

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: SgfI-MluI

ACCN: NM_134044

Insert Size: 1419 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).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. 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.

Note:	Plasmids are not sterile. For experiments where strict sterility is required, filtration with 0.22um filter is required.
RefSeq:	NM_134044.3 , NP_598805.2
RefSeq Size:	2509 bp
RefSeq ORF:	1419 bp
Locus ID:	104799
UniProt ID:	Q8BGQ1
Cytogenetics:	12 D2
Gene Summary:	Proposed to be involved in endosomal maturation implicating in part VPS33B. In epithelial cells, the VPS33B:VIPAS39 complex may play a role in the apical RAB11A-dependent recycling pathway and in the maintenance of the apical-basolateral polarity (PubMed:20190753). May play a role in lysosomal trafficking, probably via association with the core HOPS complex in a discrete population of endosomes; the functions seems to be indepenedent of VPS33B (By similarity). May play a role in vesicular trafficking during spermatogenesis (By similarity). May be involved in direct or indirect transcriptional regulation of E-cadherin.[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (3) differs in the 5' UTR and lacks an in-frame exon in the coding region, compared to variant 1. It encodes isoform b, which is shorter than isoform a. Variants 2 and 3 both encode isoform b. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.