

Product datasheet for MC225362

Epg5 (NM_001195633) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Epg5 (NM_001195633) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Epg5
Synonyms:	4732475F16; 5430411K18Rik; AI661957; AW456499; mKIAA1632
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Fully Sequenced ORF:	>MC225362 representing NM_001195633 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGATCGCC**

ATGGCCGAGGCTGTGAAGCCCCCGGGCCAAGGCCAAGGCCAGCCGACCAAAGGCAAGGAAAAGAAGA
AGCATGAAGCCCTTCAGACTTGTGATGCAGGTCCCCTGCCAGAGACCTGTCGCGAGCAGGAGAGCCCGTG
TCCAGTCTCTGAAGGAGATGACCTGAAGTCGTCTGCTGACCCCAAGCTCCACAGTGTGTGT
GGATGGAATGAAAGTGAATGTTGATATACCACTCACCTCCTTGACTATAGGGGATGAAGTCCCCCGG
TACAGGACACAGAGGACCTAAAAGAAAGGGGAGAGGTCACAGCCGGTGTGGAGATGATGAAATGGAGTT
GAAGGTGGACCCTGGGGACAACGTTATAGCTAAAGGTGAACCTTGTAAAGAACTCCCGGAAGTAGAGGAC
CACACTCATCCAGTGTGGCCCTCCAGAGAGCACACTGCAGCCTGACTTCCCTGCACTCAGCAGGCAG
TGGAGGGCTCACATGCCAGAAACATCCGACTAGGAAGCAAGATGAAGCCGCCCTCGGCTGTTCAAAGGT
GTTTCAGAATGTTAGCTTGACAGTTCTTATGAAGCCAAAGAAGTTTCTCAGCCACCGAGAGTGAAGAAA
CTATATCCTGAGTTGCCAGCTGAAATTGCTGAAGTACCAGCTCTGGTGGCAGTGAAGCCATTGCTCCGTA
GTGAGCGACTCTACCCAGAATCCCGTCTCAACCAGAAGTGACACCCTTCACTAAAGAACAGCTCAAAT
CTTGGAGCCTGGCTCATGGCTGGAAAATGTTGCATCATATGTAAGAATTCGATAACATTGCTCATCAG
GACAGACAGAGTTTTATGAGCTGCTTTTGAATTACTCACGGTGCAGGAAGCAGTTGCTGTTGGCTGAAG
CTGAGCTGCTCACTCTGATGCTGACTGCCACAGTGCCAAAAGTCGACTGTGGCACTTTAAGGATGAGCA
GATGGCCGTACAGGGCATCTGTGCAGATCAAGTAAAAGTCTACGGGCATCACCACTACCAGCGAGTAGAG
ATGAATGAAAATGTCCTGGGCGAGCTGAAGAAGCTGTTTGTGATGCCAAATCTGAGCACCTCCACCAGACCC
TGACCCTCACTCGTACACTTCTGTGCTCTCGAGGTTGCAAGTGGAGTCTACATCTTACACTGCTCAA
TAGCTCAGCTGCTCTGAGATCTTAGCAGTTTATCAGGCAGACCAAGTTCCTAAACTGACAGAAAGCATC
CCCTCTGATGTGTGCTGAGCTGAAGGAGTGCATTAGTGTCTTGTTCATGTTACCAGGAGAGTGAAGTGAAG
ATGCCAGTTCATGAGGACATCCTGCTCTGGCTGCAGAAGCTGGTGTCCGTGTTACAACGAGTCGGCTG
TCCTGGGATCACTTCTTCTCCTGAACCAGTCTCCGCTGCCAGCTGGCATTGTAATGGGCTGTT



View online »

CCTTTTATACAGATCAAAGTGTGAATAACCCCTTCAGGGGTCTTTCATTTTATGCAGTCCCTTGGCCCTGC
 TGATGTCTCCTGTCAAAAACCGAGCGGAGTTTATGTGCCACATGAAGCCCAGTGAGTGGAAGCCATCATC
 TTCAGGGCCAGCATCTGGGAACCTGGACACTTGTAGATGAAGCGGGGAAGAGGATGAAGACCCTGAGACC
 AGTTGGATTCTCCTTAATGAAGACGACTTGGTCACCCTTTTGTCCAGTTTCCCTTTTCAGGAGCTTTTC
 AGCATCTTCTGGGTTTAAAGCCAAAGGTGATTATTTACCTGAAACAACAAGCCCCAAGAGATGATGAA
 AATTTTTGCCTTTGCCAACTCATTAGTGGAGCTTCTGGCTGTGGACTAGACACCTTTAACAGAGCCCGC
 TACAGGCAGTTTGTGAAGCGCATTGGATACCTGATCAGGATGACCCTTGGCTATGTCAGTGACCATTGGG
 CTCAGTATGTGAGCCACAGCACAGGCGCGGGCTAACACCGCAGCCCTACTCCATGGAGAAACTGCAGGT
 GGAATTTGATGAGCTGTTTCTCAGGGCTGTCTCCATGTCCTGAAAGCCAAAAGACTTGGCATTGGCTG
 TTCATGTCTGAGATGCCCTTCGGCACGCTGTGGTCCAGATGCTCTGGAAGCTCTGTACCTCATGCACC
 AGGTGGAGAGTGGGGACCTGCAGCAACTCTGTGCCTCCCTGCAGCCGGCCGAGTGCAAGAGGCGACTCCA
 AGACCCAGAACATTTTGCAGCTTTGAGAAGTGTCTTTCTCCATCAACAGCTCAGAGGAGATCTGTCTC
 CTCACGGCCTTTGCTCAGATGGCTCGAGCCAGAAGAACCAATGTGGATGAAGACTTCATAAAAATCATTG
 TCCTGGAGATCTATGAGGTATCTTATGTCACACTGTCTACCAGAGAGACTTTCTCAAAGGTTGGTCGAGA
 GCTTCTGGGAGCCATTGCAGCTGTTACCCTGAGATTATCTGTCTTTTGGACAGAGTGCAGGAGACC
 ATTGACCAAGTTGGAATGGTTTCTTATACTTGTTTAAAGAACTGCCTTTGTATCTTTGGCGACCTTCTG
 CCCCAGAGATAGCTGTGATTTCGGGACTGGTTATTGAATAATAATCTGACAGCAGTGAAGAACAAGTTGGC
 CTGTGTGATCCTAGAAGGACTGAATTGGGGATTACTGAACAGGGTACCCTTCATTTGGATCAAGCATTG
 CACACGGAAAGTGGCCTTACTGGTTCTCGAGGCTTATCAGAAATACCTGGCACAGAAACCTTATACTGGGC
 TTAATTTCTGAAAGTATGAAACAGGTTTCTATTTGGCCAGTATTGTTTCGATATGGAGAGACTCCTGAGAC
 CTCATTTAACCAGTGGCCTGGAATTTAATCTTGAGGTTAAAATTGCACAAAAATGATTTTGAAGACAG
 AACTTCCAGTCATTCCCTTCTGCAGCACAGTACCTGACATGACAGAATCGTCCATGTTTACCCTCTCT
 TGAAGGCTGTGAAATCAGGCTTGGCCATTGGCTGCTATCTCGCTTAGCTGTGACAGCTGTTGGCCACAG
 CCTTGAGAAGTTCTGTGAGAAGGCATCCCACTGCTGGCGTGCTGGTCCAGTCAGCGCATCTGAGAGCA
 GTGGTCCATGCCCTAGATAAGATCCTGCCTGTGTTCTACCCTTACCAGTGTACCTTCTGAAGAACGAGC
 AGTTTTTGTGCAACCTTCTCCTCTTCTACAATTGGACAGCGGTGTTCCCGAGGGGTCACACAGCAGGT
 CACCCACAGGGTGGCCAGCATCTGACAGGAGCCGTCCATGGAGACAATGTGAACTTCTCAGCAGCATG
 ATCCAGGCACACATATGTGTAAGCACTCAGCCGGATGGAGTGGTCTGTGGCTGTGTTGGAGTTCTGGG
 TCCAGGCTCTCATAAGCCAGCATCTGTGGTACCGAGAACAGCCATTCTCTTCTCATGGACCATTGTG
 TAAGACAGCGTTTACCCTGATGCAGGAAGATTGTGTGAAAAATTGCTCTACCAACAGCACAGAATGCT
 TTGGGTTACCCTGTGACCGGAGTCTGCTCTTCTTGGTGAAGTGGATTGTGGCAGGGAACATCACGC
 CTTCTTTCGTGGAGGGCTTGTCCACGTCCACACAGGTTTGGTTTGCATGGACTGTGTTGAACATGGAATC
 CATCTTTGAAGAGGACTCTCAGCTCCGAAGAGTTGTTGAAAGGGAATTGGTTATAAATGCGTTTAGCCCT
 GACCAAGCTCTAAAGAAAGCCAGGTCCAGCTGAAGCTGCCCATCGTCCCGTCCCTGCAGCGGCTGTTGA
 TTTATCGTGGGCCACCAGGCTCTGGTACACCTTCTGATCATCCCCTTTTGGCACTCATTGGCAGAA
 GTTCTTCTCCTGTATCTTCATCGCCCAGGACCACAATATGGGTTACCTGTGGATGGTTGTATTGGAAGA
 AGGTTTTTCAAAGTCCCTCGCACGTTAATTTGTTGAAAGACATGAAGAGACGCCTGACGGAGGTGGCTG
 ACTTCCACTATGCTGCCAGCAAGGCCCTCCGTGTTCCAGCAGAGGGCAGTGAAGGGACACCAGAAGGGCA
 GGCTGGCACCCCTGGCTTCTGACTTCCCAGAACTGCACAGGGAAGTGTGAGGCTCTTTAATGTATAT
 GTATTATGGCTAGAAGATGAAAAATTTTCAGAAAGGAGATACCTATATTCCTTCTTTACCAAAGCACTACG
 ATGTTACAGGCTAGCAAAAGTATGCAGAATCAGCAGGACCTGTGGATGGAGTATGTGAATATGGAGCG
 CATAACAGCAGGATCCAGGAGACAGTTGCTCTGTGGACACAGGCCAAGCTTGTGATCCCATGCCGCACCC
 TGCAGCTCTTACGACAGCTGGATTTACGGACCCCTTGTGGCTAAAGCAAGGGTTCTGAGTAACCTTAG
 AGAAGCACGAGGCTCCACATCCTCCTTCTTCTGCACCCCGTGAGGCCTCCCGTCCACTATTCCCTC
 TGCTGCGCTGTTGACACAGAAGGACAGCACCCAGCTCATGTGCACAGACCTGAATCTGCTGCAGCAGCAG
 GCCAGGAGTGCCACTCTGCGGGAGTCGACGAGGTTGCCCTGGATGGCGAGCTGTTGGAGACCATGCCCA
 AACAGTACGTTAACCGGGAAGATCAGGCCACCCTGCATCTGGAGTGCAGAGGCAGCAGTGGCAAGAAGTG
 CCAAGGAGCAGCAGTTGTAACGGTCCAGTTTGAAGGCATGAACAAGAATGAAGCCGTAAGCCAGCAGATC
 CATGTTCTGCAGAAAGAAGTGAAGCAGCTCCAAGCGGAAGCTGCTCAGCCCCGGCACTGAATGTTGTGG
 AGGCCGCTGTGCATGCAGAAAACCTGATCACGGCCTTAGTGAACACTTATAAGTTACAGCCTACACCTGG
 AGTTTCAGAAACTTGGCATTAGCCTTTTCTTACTGTTGTGGACCATGTACAGCGATGAGACCCAGCGCCAT
 CCTCTACAAGGCAGTTCTTCACTTCATGCATCGAGATCTTGGGACAGGTGTTTGTAGCGGCACTAAGT

CCGAGTGCAGGAAGCTGCTCCAGACCATCCTGAAGAACAGGCGGCTCTGCTCGCTGCTGGCTCCTTTCTT
CACGCCAATGCTGCGCCCGCCGAGTTCATCCAGCTGTATGAGAGAGTGGTACTTGTCTGCGCAGGAC
AACAGCGACGTGATCTTCATGCTGCTCACCAAGTTTGATATTCAGCAGTGGTTAAACAGCACTAAGCCCC
CTCTGTCTGACCCGACCAGGCTCCTGGAGTCCATTCACTTGGCACTCACTGCCTGGGGCCTTGAACCAGA
GGAAGATATTCTGATGCCATTTAATCTTTCTGCAAGCACTGGACTCACCTCCTACTCTACCAGTCCCT
GACCAGTACAGTGATGTCCTCAGGCTGCTGGTGCAGAGCTCTGCCGAGCAGCTGAGTCCCGAGTGCT
GGAAAGCCACTTTGAGAGCCCTGGGCTGTTATGCCCAAGCAGCAGAGGGGGCGGCTGTGGAGAG
CTCTGGGCTTACAGTGCCTCCCGTGTCTCTGTGTCAGACAAGCAGGTAATGGAGACGGTACAGTGGCTT
TCTGACTTCTTTTATAAGTTGCGGCTCTCCAAGTTGGACTTTAAAAGCTTTGGATTATTCTCAAAGTGGA
GTCCTTACATGGCTGATGTGAAGACATTTCTGGGATACCTTGTGAAAAGGCTGACTGACTTAGAAATAGC
CTCCTTGTCTCAGGACCAACCGCCAGCAGCAAAGAAGTGTGAGGTCCCTGCACGCGCAGATCATCCAG
CTCTTTAAGCCCTGGATCCTGGTTTTAGAGGACGCGAAAGTAGCCACCAGCGACATTACCCGTGGCTGG
AGAGTGACACTGCGGTGGCTCCAGCATCGTGCAGCTTCTCTGACTGTGTCGGCTCATTGCACACGAG
TTTTAAAGACAGGCTGTTGCCAGGCGATGAGGGAGCCCTGCGGTTGCACCTGCTGCACTACTGTAAACA
TGCACAGCACCAAGATGCCGGAGTTCATTCTCTATGCCTTCCACAGTGCATACCAGAGACTGGAGTGGA
AGGACTTGCACCCCTGACCAGAGGCTCATGGAGGCTTCTTTAAGGTGGAACGTGGAAGCCCAAGAGTTG
TTTCTTGTTTTTGGGGTCTGTACTTTGCAGAGTCAACTGGGTTAGTGTGCTGTCTGACGCCTGGAATCCC
AGCCCCCTTCCGAAACCCAGAGCATGGCTGTCTGCCTCCTTTTCATGATGGTTTTACTAGCAAAGGAGG
CTCAGCTCGTTGACGAGCCAGATTCACCTTTACTGAGTCTCCTTGGACAGACAAGCTCACTCTCATGGCA
TCTTGTGGACCTGTGTACATACCAGAGTGTGTTGGGTTATTTGAGCAGTCACTACCCACCATCCGTCGTC
CTGGCAAATGACTGTTCCCGAACTGATAGTGAAGCTGTTAAAAGTGTCTGCAGGGCTTTCTGCTCATG
CTGACGGCCGGAAGCACGTGGACATAGTTCCAAAATGCCAGGCTTTCATCATCAGATGGTTCAGTTTCT
CAGTGCCTTGGAAACAGACTGAAAAATCACCTTCCCAGCACTGGAGCGGAAATATCTAAGCTGTTGGAT
GACATCATTATCTTTAACCCACCGACATGGACAGCCAGACCCGCCACATGGCCCTCAGCAGCTTCTTTG
TGAAGTCTGATGATGATGAACAATGCGGCTGTTCCAACAGCAGAGTTCCTTGTGTCAGTATTCGGAC
CTGGATTGGCCAAAGAGTGATGGCTTGATAGTGTGCCCTCTTAACAGCAGCCTGCCAAAGCCTGGCG
TCTGTCCGCCACATGGCGGAGATCACCGAAGCCTGCATCATGGCCTACTTCAAGGAAAGCTCCCTCGATC
AGAACTTAGGATGGGGCCCGTCTGGTGTCCCTTCAAGTTCAGCTCACCCGCGGAGACTTTCTGGA
GGAGTGTAGCCCTCGGTAGTTGTTGACTCTCTATGTCTACCTGCTTCAAGTGTCTGAACAGTGAAGCAG
ACGGTGAAGAACGACATGAAAATGCTCCTCGTCTGTCAGCGGCTGGCTGGAGCAGGTGATCCCGAGCTCTG
CACAGGAAGAGGCAAAGCTGTTTCTGTGGTGGCATCAAATCCTGCAGCTGTCTCTGATTAGTTGGAGCA
GAACGACTCCGTCTGACAGAGTCCGTATCCGGATCCTGCTCATGCTGCAGAGCAGGCAGAGCCTTATG
GCCGAGGAGAGGCTCAGCTCTGGGATTCTGGGGCAATCGGGTGGGCGGAGATCACCTTGTCCAACA
GGTTCCGAGTGGCTGCCGAAGCATGGCTGCCTTCTGTTGGTTGAGTCCCTGCAGAAGACCAGATTTCG
TCTGAAACCCAGCTCTGAGTTACATCTGGCTCCAAAAGCTCAGCAGGTGCTGACAGCTCTGGAGTCCATG
ACGCTCAGTAAGCAGTATGTGGAGTACCAGGACCAGATACTGCATGCTCTTCAAGTTTATAAGGCATCCTG
GCCACTGCCTTCAGAATGGGAAAAGCTTCTGGCCCTTCTGTCAACCGCCTCTATCCAGAAGTGCATTA
CCTGGACAACATCCGGTAG

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

SgfI-MluI

ACCN:

NM_001195633

Insert Size:

7719 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:	<ol style="list-style-type: none">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.
RefSeq:	<u>NM_001195633.1</u> , <u>NP_001182562.1</u>
RefSeq Size:	9672 bp
RefSeq ORF:	7719 bp
Locus ID:	100502841
UniProt ID:	<u>Q80TA9</u>
Cytogenetics:	18 E3
Gene Summary:	Involved in autophagy. May play a role in a late step of autophagy, such as clearance of autophagosomal cargo. Plays a key role in innate and adaptive immune response triggered by unmethylated cytidine-phosphate-guanosine (CpG) dinucleotides from pathogens, and mediated by the nucleotide-sensing receptor TLR9. It is necessary for the translocation of CpG dinucleotides from early endosomes to late endosomes and lysosomes, where TLR9 is located.[UniProtKB/Swiss-Prot Function]