

Product datasheet for **MC229289**

Palb2 (NM_001289842) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Palb2 (NM_001289842) Mouse Untagged Clone
Tag: Tag Free
Symbol: Palb2
Synonyms: BC066140
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC229289 representing NM_001289842
Red=Cloning site **Blue**=ORF **Orange**=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**GCGATCGCC**

ATGGAAGAGCTTCCGGGAAGCCCCTCAGCTATGCGGAGAAGGAAAAGTTGAAGGAAAAGCTAGCATT
TGAAAAAGGAATACAGCAGGACACTTGCTCGACTTCAGCGTGCCAAAAGAGCTGAGAAGGCTAAGAACT
TAAGAAAGCCATAGAAGATGGTGTGCCAGCCAGAAAGCTTCCACAGCTAAGCCACTCGAATCTATA
AATAAAGGCTTCTTGTGACACATTGCAAAGCAACCATCTGGATGAGGAGACTGGAGAAAACATCTCTC
AGATACTTGATGTTGAACCTCAATCCTTTAACTGTAACAAGGCAAGAAGTATTACATACACCAAGAGC
AGGTGACATTC AAGGACAACCTTTGCAATAGCACCAGCAGCCCTGATGGCAAGAAAGAGCAGAACACGCTT
CCGGGGACAACAAAGACGCCGTGGGAGAAGTCACTGTTTACAGGAGAAGGAAGATTATTTTGACACTA
ATTCTCTGGCGCTCCTTGGTAAGCATCGAAAAGGGCAAGAATCAATCAGTAGGAAGAATTCTAGGACTCC
CGTGAGTGAAAAGACTCACCTTTAAGTCTCAGGTCTCAGATCCCTGACCCTCCAGCACTGTTACAGGA
ATTGGAGAGGGTATATTAATCCGCCATCTGGCAAATCAGAAAGGGGAATTGATACACTTGTGAGAGGAA
ATACTGTCTCCGGGAGGCTGCAGTTCCTCATGTACTGCGTCAAACAGCAATCACAGTCAAGCACTTGA
GCATACACTCCTAAAAGTGGCTGCAAAATTAATACTACTCAGGGCCCGGCTTCATCCAAAACCTGGTGGA
CAAGACCAAAAAATGACTATATTTACAGTAAACTCAGTAGTATATAAAGCTGTGCGTGCCCATGGTCAGC
TGCCAGGAAGTCCCAATCTTGTCTGTAATGATCTCACGCATAGTAACTTGCCAGCAAAATGATACCCC
AAACTCTAAATCTTTAAATCTCCAGTAACTGTTGATGAGAGAAAATGAACCTCTTCAGGAAGATGAA
ATTCTTGGTCCATCTAAGAACTTCAACCTGGCAGCAGTCTCTCCTCCTCCACAGAAAGTCAAATACATT
CTTGTACTATGCTGAAGCCTTCTGTTTCTGCAGAATACTATGTTAGAACAACCCGTCGTATGTCAGA
TTGTCAGAGAAAAATAGCTCTGGAAGCTGTAATCAAAGTCATTTGGGTGTCAAAAAGAAAGAGCTTAAA
AAGAAAACCAAAGCTACTAAGGCGTCTCCTCTCCAGTGAAGACTGACCAGAGTGAAGTGGCATGC
TGGACACGAGCAGGGACAGTCCAGTTCAGGAAGCCTCTCTCAGAACTGCTCTACCAGCTGAGGTCAG
CTCTCCTCCAGGACCTGCTGAAAAGGCCACCACACCGCCACCTGGTAGAGGACACAGAGGGAAAACGAAAA
TCAGCCCGCACCTCCACTGGTCACTGCCAGTCTGTTTTCTCCTTGTGCCGCACTGGCTGTTAAACA



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GGTCCAAGGGCAAATTCACCAAGCATAAATGTCAGAACAGAGGCGTGGTTATTCATGACTTTGAGTTACC
TGATGAAGACTTTGGGCTTCTTAAACTTGAAAAATTGAAGTCCTGCTCGGAAAACTGATTGAGTCTCT
GACTCAAAAACTGTGGTGAGAGGCTTCTAGAGAAGGAAACCATGCTGCTCTGGAGGAAGTCAAAAGAG
ATTCGGAGACGGAGGGCTTGAAGAGGAGCTCACTGTTCCACCAGGAGAAGCGTACCGTCCAGGGCCAAC
CCTGAGAAGGCAGCCAGGGAGCAAGGACCTTCTTCATCCATAGTGCTTTTCACTCCTGCAGACTGCT
GCGCCTAACGACAGTGGCAGGCCCTCCCTCCCTGTGCTCACCTGCTTCCCCATCTTAGGCATGACTC
CAGCTCTTGGCTCCCAAGCAGCCGGTGAGACTATCTACTGAAGCTGCACAGCCTTGCTCTACATCCCA
ACCTCCTCTCTTGGGAGACACAAACAGTCTTGTAATAACAGTAAACAGTGAACAGTTCAGCCTGCTCA
CCAAAACCGGACACCAACCTGCAAGCGTCAGGTAGGCAAGGACAACCTGCCTGTGACAGTACTCTGGCC
CCCAAGCAACACCTCTACCTGTTGAGTCATTCACTTTTCAGAGAAAATCAGCTTTGTGAAAATGCATGCCT
CGAGTTACATGAACATTCCACTGAACAGACTGAACTGCAGATCGCCCTGCTTGTGACAATCTAAACCCA
GAAAACCTACAGTTGGTTTCAGAGTTAAAGAATCCTTCCAGTTCCTGCTCTGTGGATGTGAGCGCCATGT
GGTGGGAAAGAGCTGGTCTAAGGAGCCATGTATCGTAACTGCTGTGAAGATGTAGTTTCTTTGGAA
ACCCTTGAATTCTCTGCAGTGGGAGAAAGTTCATACCTGGCACTTCACAGAGTTCAGTATTACAATA
GTTCCAGTGCCTGATGTTACAATCTTATATGTGTAGCTTTGGGAAGTTTGGAAATCAGAGAAATCAGGG
CGTTGCTGTGCTTCTGGTGATGACAGTGAAAAGCAAGTCCCTCCTGAAATCTGGAGATATAAAGCAT
GCTTGGCCTGACAAAGCGGAGGCTAGTTAGTAGCACTGGGACCTTTTGAATCAACAAATACAAATCATG
ACATTTGCTGACGACGGAAGCAGCAAAGATGAACAGCTTTTGTGCTCCTGATGAGACTGTACTGACTT
TTGCTGAAGTCCAAGGGACGCAGGAAGCTCTGCTTGGTACTACCACTGTGAACAGCATTGTGATCTGGAA
TTTAAAAACCGGCCAGCTCCTGAAGAAGATGCACATTGATGACTCCTACCAGGCTTCGGTCTGTCACGGA
GCCTATTCTGAGAAGTTCCCTGTAGCCAGGCTGGCCTGCACCTGTCTATGTCTGAGTAAAGATGACCT
TGAACCTACGATTCCCTGCTTCTACCTCTCCAGTGTGGGCTTTCTGGCACAAGTCAAGCTGAGCTACAA
TCCTATTGACTGA
    
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ACGCGTACGCGGCCGCTCGAGCAGAAAACCTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites:

SgfI-MluI

ACCN:

NM_001289842

Insert Size:

3093 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).

OTI Annotation:

Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.

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.

RefSeq:

[NM_001289842.1](#), [NP_001276771.1](#)

RefSeq Size: 3347 bp

RefSeq ORF: 3093 bp

Locus ID: 233826

UniProt ID: [Q3U0P1](#)

Cytogenetics: 7 F2

Gene Summary: Plays a critical role in homologous recombination repair (HRR) through its ability to recruit BRCA2 and RAD51 to DNA breaks. Strongly stimulates the DNA strand-invasion activity of RAD51, stabilizes the nucleoprotein filament against a disruptive BRC3-BRC4 polypeptide and helps RAD51 to overcome the suppressive effect of replication protein A (RPA). Functionally cooperates with RAD51AP1 in promoting of D-loop formation by RAD51. Serves as the molecular scaffold in the formation of the BRCA1-PALB2-BRCA2 complex which is essential for homologous recombination. Via its WD repeats is proposed to scaffold a HR complex containing RAD51C and BRCA2 which is thought to play a role in HR-mediated DNA repair. Essential partner of BRCA2 that promotes the localization and stability of BRCA2. Also enables its recombinational repair and checkpoint functions of BRCA2. May act by promoting stable association of BRCA2 with nuclear structures, allowing BRCA2 to escape the effects of proteasome-mediated degradation. Binds DNA with high affinity for D loop, which comprises single-stranded, double-stranded and branched DNA structures. May play a role in the extension step after strand invasion at replication-dependent DNA double-strand breaks; together with BRCA2 is involved in both POLH localization at collapsed replication forks and DNA polymerization activity (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) contains an alternate 3' terminal exon, compared to variant 1. It encodes isoform 2, which is shorter and has a distinct C-terminus, compared to isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.