

Product datasheet for MR229636

Palb2 (NM_001289845) Mouse Tagged ORF Clone

Product data:

Product Type: Expression Plasmids
 Product Name: Palb2 (NM_001289845) Mouse Tagged ORF Clone
 Tag: Myc-DDK
 Symbol: Palb2
 Synonyms: BC066140
 Vector: pCMV6-Entry (PS100001)
 E. coli Selection: Kanamycin (25 ug/mL)
 Cell Selection: Neomycin
 ORF Nucleotide Sequence: >MR229636 representing NM_001289845
 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**GCGATCGCC**

ATGGAAGAGCTTCCGGGAAGCCCCTCAGCTATGCGGAGAAGGAAAAGTTGAAGGAAAAGCTAGCATTTT
 TGAAAAAGGAATACAGCAGGACACTTGCTCGACTTCAGACTGAACTGCAGATCGCCCTGCTTGTGACAA
 TCTAAACCCAGGAAACCTACAGTTGGTTTCAGAGTTAAAGAATCCTTCCAGTTCTGCTCTGTGGATGTG
 AGCGCCATGTGGTGGAAAGAGCTGGTGTCTAAGGAGCCATGTATCGTAACCTGCTTGTGAAGATGTAGTTT
 CTCTTTGGAAACCCTTGAATTCTCTGCAGTGGGAGAAAGTTCATACCTGGCACTTACAGAGGTTCCAGT
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ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
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Protein Sequence: >MR229636 representing NM_001289845
 Red=Cloning site Green=Tags(s)

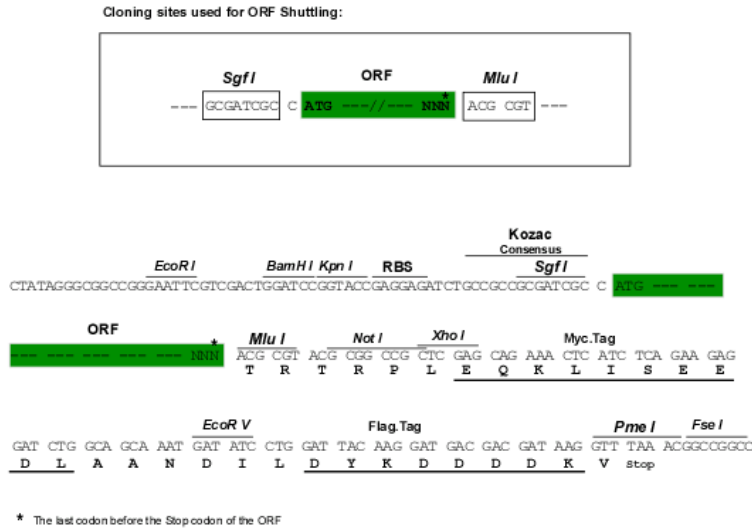
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 VLTFAEVQGTQEALLGTTTVNSIVIWNLKTGQLLKKMHIDDSYQASVCHGAYSEKGLLFVVVSQPCAKES
 QALGSPVFQLLVINPKTAQSVGVLLCSLPQOQAGRFLEGDVYKDHVAAAVLTSGTIAIWDLLLGHCTALLP
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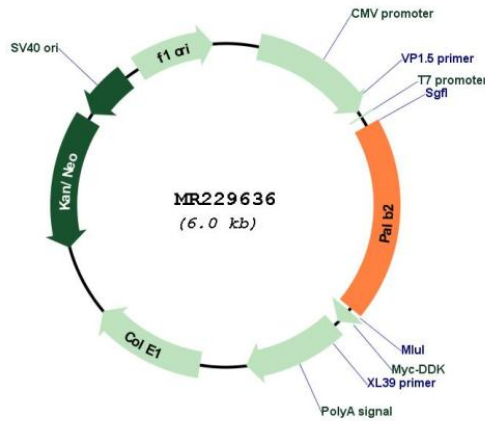
Restriction Sites:

Sgfl-MluI

Cloning Scheme:



Plasmid Map:



ACCN: NM_001289845

ORF Size:	1152 bp
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. More info
OTI Annotation:	This clone was engineered to express the complete ORF with an expression tag. Expression varies depending on the nature of the gene.
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:	NM_001289845.1 , NP_001276774.1
RefSeq Size:	1590 bp
RefSeq ORF:	1155 bp
Locus ID:	233826
UniProt ID:	Q3U0P1
Cytogenetics:	7 F2
MW:	42.6 kDa

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]