CPTC-PALB2-1 (CAB080238)

Uniprot ID: Q86YC2

Protein name: PALB2 HUMAN Full name: Partner and localizer of BRCA2

Function: Plays a critical role in homologous recombination repair (HRR) through its ability to recruit BRCA2 and RAD51 to DNA breaks (PubMed:16793542, PubMed:19423707, PubMed:19369211, PubMed:22941656, PubMed:24141787, PubMed:28319063). 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) (PubMed:20871615). Functionally cooperates with RAD51AP1 in promoting of D-loop formation by RAD51 (PubMed:20871616). Serves as the molecular scaffold in the formation of the BRCA1-PALB2-BRCA2 complex which is essential for homologous recombination (PubMed:19369211). 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 (PubMed:24141787). Essential partner of BRCA2 that promotes the localization and stability of BRCA2 (PubMed:16793542). Also enables its recombinational repair and checkpoint functions of BRCA2 (PubMed:16793542). May act by promoting stable association of BRCA2 with nuclear structures, allowing BRCA2 to escape the effects of proteasome-mediated degradation (PubMed:16793542). Binds DNA with high affinity for D loop, which comprises single-stranded, double-stranded and branched DNA structures (PubMed:20871616). 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 (PubMed:24485656).

Subcellular location:

Nucleus (experimental evidence)

NOTE: Colocalizes with BRCA2 and BRCA1 in nuclear foci. Protein existence: Experimental evidence at protein level

IHC protocol:

IHC test staining:

RNA similarity:

Literature conformance: Literature significance:

RNA tissue specificity:

RNA tissue distribution:

Comment:

Immunohistochemistry



Immunofluorescence

A Star S	IHC Sibling similarity:	Other antibody sho	ows dissimilar IHC staining pattern
	IHC fail comment:	ANTIBODY FAILED: Not consistent with RNA	
100			
	IF Overlay:		antibody (green), anti-tubulin (red) and DAPI (blue)
	main location:		
	IF additional location:		
	IF ensured for sublication on LIDA.		NI-

HIER pH6, Dilution 1:100

Not consistent with gene/protein characterization data

Very low consistency between antibody staining and RNA expression data

Negative in all tissues.

Low tissue specificity Detected in all



IF Overlay:	antibody (green), anti-tubulin (red) and DAPI (blue)
IF main location:	
IF additional location:	
IF approved for publication on HPA:	No
IF in THP-1:	Negative
IF in U-2 OS:	Negative

Western blot

•	WB Size markers (kDa):	250, 130, 100, 70, 55, 35, 25, 15, 10
	WB Lanes:	Marker (1), RT4 (2), U-251 MG (3), Plasma (4), Liver (5), Tonsil (6)
	WB Target weight (kDa):	5, 10, 98, 131
	WB Validation:	Uncertain (No bands detected.)
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