For Research Use Only

EXOSC2 Polyclonal antibody Catalog Number: 14805-1-AP Featured Product 7 P

Featured Product 7 Publications

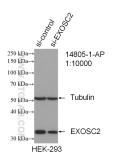


Basic Information	Catalog Number: 14805-1-AP	GenBank Accession Number: BC000747		Purification Method: Antigen affinity purification	
	Concentration:	Concentration: GeneID (NCBI): Recomme 350 μ g/ml 23404 WB 1:500- Source: UNIPROT ID: IP 0.5-4.0 gradementation Rabbit Q13868 protein lyst Isoburge: Full Name: IHC 1:20-1		Recommended Dilutions:	
	350 µg/ml			WB 1:500-1:2000 IP 0.5-4.0 ug for 1.0-3.0 mg of total protein lysate IHC 1:20-1:200 IF/ICC 1:20-1:200	
	Rabbit				
		Observed MW: 30-33 kDa			
Applications	Tested Applications:				
	WB, IHC, IF/ICC, IP, ELISA Cited Applications:			WB : HeLa cells, HepG2 cells, Jurkat cells, MCF-7 cel HEK-293 cells	
	WB, IF		IP : HeLa cells,		
	human	Species Specificity: IHC human		human skin cancer tissue,	
	Cited Species: human, mouse, yeast			CF-7 cells,	
	Note-IHC: suggested antigen retrieval with TE buffer pH 9.0; (*) Alternatively, antigen retrieval may be performed with citrate buffer pH 6.0				
	buffer pH 6.0				
Background Informatio	In the nucleus, the RNA exosome snoRNA and snRNA, in the elimina antisense RNA species and promo thereby limiting or excluding the in general mRNA turnover and spe within their 3' untranslated region [PMID:15346807]. EXOSC2 is a no	ation of RNA processing oter-upstream transcrip ir export to the cytopla ecifically degrades inh ns, and in RNA surveilla n-catalytic component	by-products an ts (PROMPTs), ar sm. In the cytopl erently unstable ince pathways, p of the RNA exos	nd of mRNAs with processing defects, asm, the RNA exosome complex is involv mRNAs containing AU-rich elements (AR preventing translation of aberrant mRNAs	
	In the nucleus, the RNA exosome snoRNA and snRNA, in the elimina antisense RNA species and promo thereby limiting or excluding the in general mRNA turnover and spe within their 3' untranslated region [PMID:15346807]. EXOSC2 is a no exoribonuclease activity and invo 17545563].	ation of RNA processing oter-upstream transcrip ir export to the cytopla- ecifically degrades inh ns, and in RNA surveilla n-catalytic component olves in a multitude of	by-products an ts (PROMPTs), ar sm. In the cytopl erently unstable ince pathways, p of the RNA exos cellular RNA pro	d non-coding 'pervasive' transcripts, such ad of mRNAs with processing defects, asm, the RNA exosome complex is involv mRNAs containing AU-rich elements (AR preventing translation of aberrant mRNAs ome complex that has 3'->5' cessing and degradation events [PMID:	
	In the nucleus, the RNA exosome snoRNA and snRNA, in the elimina antisense RNA species and promo thereby limiting or excluding the in general mRNA turnover and spi within their 3' untranslated regio [PMID:15346807]. EXOSC2 is a no exoribonuclease activity and invo 17545563].	ation of RNA processing oter-upstream transcrip in export to the cytopla- ecifically degrades inh- ns, and in RNA surveilla n-catalytic component olves in a multitude of Pubmed ID Jo	by-products an ts (PROMPTs), ar sm. In the cytopi erently unstable unce pathways, p of the RNA exos cellular RNA pro	d non-coding 'pervasive' transcripts, such ad of mRNAs with processing defects, asm, the RNA exosome complex is involv mRNAs containing AU-rich elements (AR reventing translation of aberrant mRNAs ome complex that has 3'->5' cessing and degradation events [PMID: Application	
Background Informatio	 In the nucleus, the RNA exosome snoRNA and snRNA, in the elimina antisense RNA species and promothereby limiting or excluding the in general mRNA turnover and spwithin their 3' untranslated region [PMID:15346807]. EXOSC2 is a no exoribonuclease activity and invol17545563]. Author Tobias Moll 	ation of RNA processing ter-upstream transcrip in export to the cytopla- ecifically degrades inh- ns, and in RNA surveilla n-catalytic component olves in a multitude of Pubmed ID Jo 36241425 Li	by-products an ts (PROMPTs), ar sm. In the cytopl erently unstable ince pathways, jr of the RNA exos cellular RNA pro	d non-coding 'pervasive' transcripts, such ad of mRNAs with processing defects, asm, the RNA exosome complex is involv mRNAs containing AU-rich elements (AR preventing translation of aberrant mRNAs ome complex that has 3'->5' ccessing and degradation events [PMID: Application WB	
	In the nucleus, the RNA exosome snoRNA and snRNA, in the elimina antisense RNA species and promo thereby limiting or excluding the in general mRNA turnover and spi within their 3' untranslated regio [PMID:15346807]. EXOSC2 is a no exoribonuclease activity and invo 17545563].	ation of RNA processing oter-upstream transcrip in export to the cytopla- ecifically degrades inh ns, and in RNA surveilla n-catalytic component olves in a multitude of Pubmed ID JC 36241425 Li 27259150 Cc	by-products an ts (PROMPTs), ar sm. In the cytopi erently unstable unce pathways, p of the RNA exos cellular RNA pro	d non-coding 'pervasive' transcripts, such ad of mRNAs with processing defects, asm, the RNA exosome complex is involv mRNAs containing AU-rich elements (AR reventing translation of aberrant mRNAs ome complex that has 3'->5' cessing and degradation events [PMID: Application	

For technical support and original validation data for this product please contact: T: 4006900926 E: Proteintech-CN@ptglab.com W: ptgcn.com

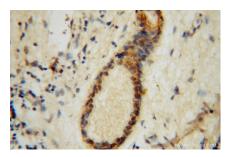
This product is exclusively available under Proteintech Group brand and is not available to purchase from any other manufacturer.

Selected Validation Data

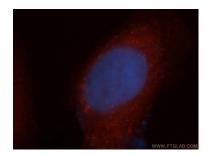


WB result of EXOSC2 antibody (14805-1-AP; 1:10000; incubated at room temperature for 1.5 hours) with sh-Control and sh-EXOSC2 transfected HEK-293 cells. $\begin{array}{c} 116 kd \rightarrow \\ 97 kd \rightarrow \\ 72 kd \rightarrow \\ 36 kd \rightarrow \\ 28 kd \rightarrow \\ 28 kd \rightarrow \end{array}$

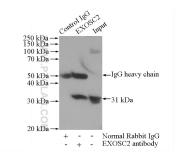
HeLa cells were subjected to SDS PAGE followed by western blot with 14805-1-AP (EXOSC2 antibody) at dilution of 1:500 incubated at room temperature for 1.5 hours.



Immunohistochemical analysis of paraffinembedded human skin cancer using 14805-1-AP (EXOSC2 antibody) at dilution of 1:100 (under 40x lens).



Immunofluorescent analysis of MCF-7 cells, using EXOSC2 antibody 14805-1-AP at 1:50 dilution and Rhodamine-labeled goat anti-rabbit IgG (red). Blue pseudocolor = DAPI (fluorescent DNA dye).



IP result of anti-EXOSC2 (IP:14805-1-AP, 4ug; Detection:14805-1-AP 1:1000) with HeLa cells lysate 1080ug.