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Datasheet for ABIN6140380 anti-EXOSC7 antibody (AA 1-291)

Image



Overview

Quantity:	100 μL
Target:	EXOSC7
Binding Specificity:	AA 1-291
Reactivity:	Human
Host:	Rabbit
Clonality:	Polyclonal
Conjugate:	This EXOSC7 antibody is un-conjugated
Application:	Western Blotting (WB)

Product Details

Immunogen:	Recombinant fusion protein containing a sequence corresponding to amino acids 1-291 of human EXOSC7 (NP_055819.2).
Sequence:	MASVTLSEAE KVYIVHGVQE DLRVDGRGCE DYRCVEVETD VVSNTSGSAR VKLGHTDILV
	GVKAEMGTPK LEKPNEGYLE FFVDCSASAT PEFEGRGGDD LGTEIANTLY RIFNNKSSVD LKTLCISPRE HCWVLYVDVL LLECGGNLFD AISIAVKAAL FNTRIPRVRV LEDEEGSKDI
	ELSDDPYDCI RLSVENVPCI VTLCKIGYRH VVDATLQEEA CSLASLLVSV TSKGVVTCMR KVGKGSLDPE SIFEMMETGK RVGKVLHASL QSVVHKEESL GPKRQKVGFL G
lsotype:	IgG
Cross-Reactivity:	Human, Mouse, Rat
Characteristics:	Polyclonal Antibodies
Purification:	Affinity purification

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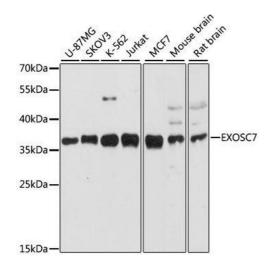
Target:	EXOSC7		
Alternative Name:	EXOSC7 (EXOSC7 Products)		
Background:	Non-catalytic component of the RNA exosome complex which has 3'->5' exoribonuclease		
	activity and participates in a multitude of cellular RNA processing and degradation events. In		
	the nucleus, the RNA exosome complex is involved in proper maturation of stable RNA species		
	such as rRNA, snRNA and snoRNA, in the elimination of RNA processing by-products and non-		
	coding 'pervasive' transcripts, such as antisense RNA species and promoter-upstream		
	transcripts (PROMPTs, and of mRNAs with processing defects, thereby limiting or excluding		
	their export to the cytoplasm. The RNA exosome may be involved in Ig class switch		
	recombination (CSR and/or Ig variable region somatic hypermutation (SHM by targeting AICDA		
	deamination activity to transcribed dsDNA substrates. In the cytoplasm, the RNA exosome		
	complex is involved in general mRNA turnover and specifically degrades inherently unstable		
	mRNAs containing AU-rich elements (AREs within their 3' untranslated regions, and in RNA		
	surveillance pathways, preventing translation of aberrant mRNAs. It seems to be involved in		
	degradation of histone mRNA. The catalytic inactive RNA exosome core complex of 9 subunits		
	(Exo-9 is proposed to play a pivotal role in the binding and presentation of RNA for		
	ribonucleolysis, and to serve as a scaffold for the association with catalytic subunits and		
	accessory proteins or complexes.,EXOSC7,EAP1,RRP42,Rrp42p,hRrp42p,p8,Epigenetics &		
	Nuclear Signaling,EXOSC7		
Molecular Weight:	31 kDa		
Gene ID:	23016		
JniProt:	Q15024		
Application Details			
Application Notes:	WB,1:200 - 1:2000		
Comment:	HIGH QUALITY		
Restrictions:	For Research Use only		
Handling			
	Liquid		
Format:	Liquid		
Format: Buffer:	PBS with 0.02 % sodium azide,50 % glycerol, pH 7.3.		

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Precaution of Use:	This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only.	
Storage:	-20 °C	
Storage Comment:	Store at -20°C. Avoid freeze / thaw cycles.	

Images



Western Blotting

Image 1. Western blot analysis of extracts of various cell lines, using EXOSC7 antibody.