For Research Use Only

SALL4 Monoclonal antibody

Catalog Number:66236-1-lg 3 Publications

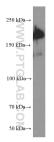


Basic Information	Catalog Number: 66236-1-lg	GenBank Accession Number: BC111714	Purification Method: Protein G purification	
	Size:	GeneID (NCBI):	CloneNo.:	
	150ul , Concentration: 1500 ug/ml by Nanodrop and 793 ug/ml by Bradford method using BSA as the standard;		2B5B2	
		UNIPROT ID: Q9UJQ4	Recommended Dilutions: WB 1:500-1:2000	
	Source: Mouse	Full Name: sal-like 4 (Drosophila)		
	lsotype: IgG1	Calculated MW: 1053 aa, 112 kDa		
	Immunogen Catalog Number: AG17480	Observed MW: 165 kDa		
Applications	Tested Applications: WB, ELISA	Positive Controls:		
	Cited Applications: WB, IHC	WB : K-562 cells,		
	Species Specificity: human			
	Cited Species: human			
Background Informatio	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i	family. SALL4 is constitutively ex ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe SALL4B, lacked the region corresp	tains 7 C2H2-type zinc fingers and belongs expressed in acute myeloid leukemia. The te symptoms and transformation to AML tha t/beta-catenin signaling pathway. Sequenc open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful pected to consist of 617 amino acids.	
	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of	family. SALL4 is constitutively ex ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe SALL4B, lacked the region corresp	xpressed in acute myeloid leukemia. The te symptoms and transformation to AML tha t/beta-catenin signaling pathway. Sequenc open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful	
	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of Author Pub	family. SALL4 is constitutively ex ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe SALL4B, lacked the region corresp encoded by SALL4B cDNA was exp	expressed in acute myeloid leukemia. The te symptoms and transformation to AML that t/beta-catenin signaling pathway. Sequence open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful pected to consist of 617 amino acids.	
	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of Author Pub Peng Wu 322	family. SALL4 is constitutively ex- ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe SALL4B, lacked the region corresp encoded by SALL4B cDNA was exp med ID Journal	kpressed in acute myeloid leukemia. The te symptoms and transformation to AML that t/beta-catenin signaling pathway. Sequence open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful pected to consist of 617 amino acids. Application IHC	
	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of Author Pub Peng Wu 322 Chiung-Chi Cheng 338	family. SALL4 is constitutively ex- ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe SALL4B, lacked the region corresp encoded by SALL4B cDNA was exp med ID Journal 22813 Pediatr Surg Int	kpressed in acute myeloid leukemia. The te symptoms and transformation to AML that t/beta-catenin signaling pathway. Sequence open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful pected to consist of 617 amino acids. Application IHC	
Notable Publications	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of Author Pub Peng Wu 322 Chiung-Chi Cheng 338	family. SALL4 is constitutively ex- ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe SALL4B, lacked the region corresp encoded by SALL4B cDNA was expe med ID Journal 22813 Pediatr Surg Int 60837 Cancer Chemoth	expressed in acute myeloid leukemia. The te symptoms and transformation to AML that t/beta-catenin signaling pathway. Sequence open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful bected to consist of 617 amino acids. Application IHC her Pharmacol WB	
Notable Publications	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of Author Pub Peng Wu 322 Chiung-Chi Cheng 338 Jun Zhang 395 Storage: Storage Storage Storage Buffer:	family. SALL4 is constitutively ex- ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe- sALL4B, lacked the region corresp encoded by SALL4B cDNA was expe- med ID Journal 22813 Pediatr Surg Int 60837 Cancer Chemoth 00780 Discov Oncol er shipment.	expressed in acute myeloid leukemia. The te symptoms and transformation to AML that t/beta-catenin signaling pathway. Sequence open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful bected to consist of 617 amino acids. Application IHC her Pharmacol WB	
Background Informatic	human SALL4, also named as Sal-like protein the sal C2H2-type zinc-finger protein constitutive expression of SALL4 in m transplantable. SALL4 is able to bind analysis of the larger cDNA fragment that started from a strong consensus i splicing variant of SALL4, designated length SALL4A. The putative protein of Author Pub Peng Wu 322 Chiung-Chi Cheng 338 Jun Zhang 395 Storage: Storage:	family. SALL4 is constitutively ex- ice is sufficient to induce MDS-lik beta-catenin and activate the Wn isolated revealed a single, large nitiation sequence and was expe- sALL4B, lacked the region corresp encoded by SALL4B cDNA was expe- med ID Journal 22813 Pediatr Surg Int 60837 Cancer Chemoth 00780 Discov Oncol er shipment.	expressed in acute myeloid leukemia. The te symptoms and transformation to AML that t/beta-catenin signaling pathway. Sequence open-reading frame, designated as SALL4A cted to encode 1053 amino acids. The other ponding to amino acids 385 to 820 of the ful bected to consist of 617 amino acids. Application IHC her Pharmacol WB	

For technical support and original validation data for this product please contact:T: 1 (888) 4PTGLAB (1-888-478-4522) (toll freeE: proteintech@ptglab.comin USA), or 1(312) 455-8498 (outside USA)W: ptglab.com

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

Selected Validation Data



K-562 cells were subjected to SDS PAGE followed by western blot with 66236-1-lg (SALL4 Antibody) at dilution of 1:1000 incubated at room temperature for 1.5 hours.