

RN213 rabbit pAb

Catalog_no:	AT7705
Applications :	IHC-p
Reactivity :	Human, Mouse
Category :	抗原抗体
Size :	100µg/50µg/20µg
Gene_name :	RNF213 ALO17 C17orf27 KIAA1554 KIAA1618 MYSTR
Protein_name :	RN213
Humangene_id :	<u>57674</u>
Humanswisspro _no:	t <u>Q63HN8</u>
Mouseswissprot _no :	<u>E9Q555</u>
Immunogen :	Synthesized peptide derived from human RN213
	Synthesized peptide derived from human RN213 This antibody detects endogenous levels of RN213 at Human/Mouse
Immunogen :	
Immunogen : Specificity :	This antibody detects endogenous levels of RN213 at Human/Mouse
Immunogen : Specificity : Formulation :	This antibody detects endogenous levels of RN213 at Human/Mouse Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.
Immunogen : Specificity : Formulation : Source :	This antibody detects endogenous levels of RN213 at Human/Mouse Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide. Rabbit
Immunogen : Specificity : Formulation : Source : Dilution :	This antibody detects endogenous levels of RN213 at Human/Mouse Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide. Rabbit IHC-p 1 : 50-200 The antibody was affinity-purified from rabbit serum by affinity-chromatography using specific immunogen.

Background : This gene encodes a protein containing a C3HC4-type RING finger domain, which is a specialized type of Zn-finger that binds two atoms of zinc and is thought to be involved in mediating protein-protein interactions. The protein also contains an AAA domain, which is associated with ATPase activity. This gene is a susceptibility gene for Moyamoya disease, a vascular disorder of intracranial arteries. This gene is also a translocation partner in anaplastic large cell lymphoma and inflammatory myofibroblastic tumor cases, where a t(2;17)(p23;q25) translocation has been identified with the anaplastic lymphoma kinase (ALK) gene on chromosome 2, and a t(8;17)(q24;q25) translocation has been identified with the MYC gene on chromosome 8. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Dec 2011],

