

# ATM Antibody (C-term)

Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP8046b

# **Product Information**

Application	IHC-P, WB, E
Primary Accession	<u>Q13315</u>
Other Accession	<u>Q62388</u>
Reactivity	Human
Predicted	Mouse
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Clone Names	RB3113/3114
Calculated MW	350687
Antigen Region	3027-3056

## **Additional Information**

Gene ID	472
Other Names	Serine-protein kinase ATM, Ataxia telangiectasia mutated, A-T mutated, ATM
Target/Specificity	This ATM antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 3027~3056 amino acids from the C-terminal region of human ATM.
Dilution	IHC-P~~1:100~500 WB~~1:500 E~~Use at an assay dependent concentration.
Format	Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is prepared by Saturated Ammonium Sulfate (SAS) precipitation followed by dialysis against PBS.
Storage	Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.
Precautions	ATM Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

#### **Protein Information**

Name	ATM
Function	Serine/threonine protein kinase which activates checkpoint signaling upon double strand breaks (DSBs), apoptosis and genotoxic stresses such as ionizing ultraviolet A light (UVA), thereby acting as a DNA damage sensor

	(PubMed: <u>10550055</u> , PubMed: <u>10839545</u> , PubMed: <u>10910365</u> , PubMed: <u>12556884</u> , PubMed: <u>14871926</u> , PubMed: <u>15064416</u> ,
	PubMed: <u>15448695</u> , PubMed: <u>15456891</u> , PubMed: <u>15790808</u> ,
	PubMed: <u>15916964</u> , PubMed: <u>17923702</u> , PubMed: <u>21757780</u> , PubMed: <u>24534091</u> , PubMed: <u>35076389</u> , PubMed: <u>9733514</u> ). Recognizes the
	substrate consensus sequence [ST]-Q (PubMed: <u>10550055</u> , PubMed: <u>10839545</u> ,
	PubMed: <u>10910365</u> , PubMed: <u>12556884</u> , PubMed: <u>14871926</u> , PubMed: <u>15448695</u> , PubMed: <u>15456891</u> , PubMed: <u>15916964</u> ,
	PubMed: <u>13448695</u> , PubMed: <u>13436891</u> , PubMed: <u>13916964</u> , PubMed: <u>17923702</u> , PubMed: <u>24534091</u> , PubMed: <u>9733514</u> ). Phosphorylates
	'Ser-139' of histone variant H2AX at double strand breaks (DSBs), thereby
	regulating DNA damage response mechanism (By similarity). Also plays a role in pre-B cell allelic exclusion, a process leading to expression of a single
	immunoglobulin heavy chain allele to enforce clonality and monospecific
	recognition by the B-cell antigen receptor (BCR) expressed on individual
	B-lymphocytes. After the introduction of DNA breaks by the RAG complex on one immunoglobulin allele, acts by mediating a repositioning of the second
	allele to pericentromeric heterochromatin, preventing accessibility to the RAG
	complex and recombination of the second allele. Also involved in signal
	transduction and cell cycle control. May function as a tumor suppressor. Necessary for activation of ABL1 and SAPK. Phosphorylates DYRK2, CHEK2,
	p53/TP53, FBXW7, FANCD2, NFKBIA, BRCA1, CREBBP/CBP, RBBP8/CTIP,
	FBXO46, MRE11, nibrin (NBN), RAD50, RAD17, PELI1, TERF1, UFL1, RAD9,
	UBQLN4 and DCLRE1C (PubMed: <u>10550055</u> , PubMed: <u>10766245</u> , PubMed: <u>10802669</u> , PubMed: <u>10839545</u> , PubMed: <u>10910365</u> ,
	PubMed: <u>10973490</u> , PubMed: <u>11375976</u> , PubMed: <u>12086603</u> ,
	PubMed: <u>15456891</u> , PubMed: <u>19965871</u> , PubMed: <u>21757780</u> ,
	PubMed: <u>24534091</u> , PubMed: <u>26240375</u> , PubMed: <u>26774286</u> , PubMed: <u>30171069</u> , PubMed: <u>30612738</u> , PubMed: <u>30886146</u> ,
	PubMed: <u>30952868</u> , PubMed: <u>38128537</u> , PubMed: <u>9733515</u> , PubMed: <u>9843217</u> ).
	May play a role in vesicle and/or protein transport. Could play a role in T-cell
	development, gonad and neurological function. Plays a role in replication-dependent histone mRNA degradation. Binds DNA ends.
	Phosphorylation of DYRK2 in nucleus in response to genotoxic stress prevents
	its MDM2-mediated ubiquitination and subsequent proteasome degradation (PubMed: <u>19965871</u> ). Phosphorylates ATF2 which stimulates its function in
	DNA damage response (PubMed: <u>15916964</u> ). Phosphorylates ERCC6 which is
	essential for its chromatin remodeling activity at DNA double-strand breaks
	(PubMed: <u>29203878</u> ). Phosphorylates TTC5/STRAP at 'Ser-203' in the cytoplasm in response to DNA damage, which promotes TTC5/STRAP nuclear localization
	(PubMed: <u>15448695</u> ). Also involved in pexophagy by mediating
	phosphorylation of PEX5: translocated to peroxisomes in response to reactive
	oxygen species (ROS), and catalyzes phosphorylation of PEX5, promoting PEX5 ubiquitination and induction of pexophagy (PubMed: <u>26344566</u> ).
Cellular Location	Nucleus. Cytoplasmic vesicle. Cytoplasm, cytoskeleton, microtubule
	organizing center, centrosome {ECO:0000250 UniProtKB:Q62388}.
	Peroxisome matrix. Note=Primarily nuclear (PubMed:9050866, PubMed:9150358). Found also in endocytic vesicles in association with
	beta-adaptin (PubMed:9707615). Translocated to peroxisomes in response to
	reactive oxygen species (ROS) by PEX5 (PubMed:26344566)
Tissue Location	Found in pancreas, kidney, skeletal muscle, liver, lung, placenta, brain, heart,
	spleen, thymus, testis, ovary, small intestine, colon and leukocytes

# Background

ATM is involved in signal transduction, cell cycle control and DNA repair, and may function as a tumor suppressor. It is necessary for activation of ABL1 and SAPK, and phosphorylates p53, NFKBIA, BRCA1, CTIP, NIBRIN (NBS1), TERF1, and RAD9. This protein has potential roles in vesicle and/or protein transport, T-cell

development, gonad and neurological function. ATM is also part of the BRCA1-associated genome surveillance complex. ATM is induced by ionizing radiation. Defects in ATM are the cause of ataxia talangiectasia (AT), also known as Louis-Bar syndrome, a rare recessive disorder characterized by progressive cerebellar ataxia, dilation of the blood vessels in the conjunctiva and eyeballs, immunodeficiency, growth retardation and sexual immaturity. About 30% of AT patients develop lymphomas and leukemias. Defects in ATM also contribute to T-cell acute lymphoblastic leukemia (TALL) and T-prolymphocytic leukemia (TPLL). TPLL is characterized by a high white blood cell count, with a predominance of prolymphocytes, marked splenomegaly, lymphadenopathy, skin lesions and serous effusion. Defects in ATM also contribute to B-cell non-Hodgkin's lymphomas, and to B-cell chronic lymphocytic leukemia, a disease characterized by accumulation of mature CD5+ B lymphocytes, lymphadenopathy, immunodeficiency and bone marrow failure.

# References

Suzuki, A., et al., J. Biol. Chem. 278(1):48-53 (2003). Kishi, S., et al., J. Biol. Chem. 276(31):29282-29291 (2001). Schaffner, C., et al., Proc. Natl. Acad. Sci. U.S.A. 97(6):2773-2778 (2000). Gatei, M., et al., Nat. Genet. 25(1):115-119 (2000). Becker-Catania, S.G., et al., Mol. Genet. Metab. 70(2):122-133 (2000).

### Images



Formalin-fixed and paraffin-embedded human cancer tissue reacted with the primary antibody, which was peroxidase-conjugated to the secondary antibody, followed by DAB staining. This data demonstrates the use of this antibody for immunohistochemistry; clinical relevance has not been evaluated. BC = breast carcinoma; HC = hepatocarcinoma.

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