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(Mouse) Uhrf1 Antibody (C-term)

Purified Rabbit Polyclonal Antibody (Pab)
Catalog # AP21204b

Product Information

Application	WB, E
Primary Accession	Q8VDF2
Reactivity	Mouse
Predicted	Rat
Host	Rabbit
Clonality	polyclonal
Isotype	Rabbit IgG
Clone Names	RB51453
Calculated MW	88304

Additional Information

Gene ID	18140
Other Names	E3 ubiquitin-protein ligase UHRF1, 632-, Nuclear protein 95, Nuclear zinc finger protein Np95, Ubiquitin-like PHD and RING finger domain-containing protein 1, mUhrf1, Ubiquitin-like-containing PHD and RING finger domains protein 1, Uhrf1, Np95
Target/Specificity	This Mouse Uhrf1 antibody is generated from a rabbit immunized with a KLH conjugated synthetic peptide between 596-628 amino acids from the C-terminal region of Mouse Uhrf1.
Dilution	WB~~1:2000 E~~Use at an assay dependent concentration.
Format	Purified polyclonal antibody supplied in PBS with 0.05% (V/V) Proclin 300. This antibody is purified through a protein A column, followed by peptide affinity purification.
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	(Mouse) Uhrf1 Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

Protein Information

Name	Uhrf1 {ECO:0000303|PubMed:17673620}
Synonyms	Np95

Function	E3 ubiquitin-protein ligase that acts as a key epigenetic regulator by bridging DNA methylation and chromatin modification (PubMed:12058012, PubMed:12084726, PubMed:14993289, PubMed:17673620, PubMed:26065575, PubMed:30392929). Plays a key role in DNA methylation inheritance by promoting recruitment of DNMT1 to hemimethylated DNA and ensure faithful propagation of the DNA methylation patterns through DNA replication (PubMed:17673620, PubMed:21268065, PubMed:26065575). Acts both as a histone reader and writer: specifically recognizes and binds (1) hemimethylated DNA at replication forks and (2) histone H3 trimethylated at 'Lys-9' and unmethylated at 'Arg-2' (H3K9me3 and H3R2me0, respectively), thereby activating its E3 ubiquitin-protein ligase activity (PubMed:15361834, PubMed:17994007, PubMed:18772888, PubMed:18772891, PubMed:21489993, PubMed:26065575). UHRF1 then mediates histone H3 'Lys-18' monoubiquitination (H3K18ub), a docking site for DNMT1, leading to DNMT1 recruitment and replication-coupled DNA methylation maintenance (PubMed:26065575, PubMed:29053958, PubMed:30392929). Also mediates histone H3 'Lys-14' and 'Lys-23' ubiquitination (H3K14ub and H3K23ub) at lower level (PubMed:29053958). Histone ubiquitin ligase activity also stimulates the methyltransferase activity of SUV39H1 and/or SUV39H2, promoting accumulation of H3K9me3 histone mark to reinformce heterochromatin state (By similarity). Enriched in pericentric heterochromatin where it recruits different chromatin modifiers required for this chromatin replication (By similarity). Also localizes to euchromatic regions where it negatively regulates transcription possibly by impacting DNA methylation and histone modifications (By similarity). Plays a role in DNA repair by cooperating with UHRF2 to ensure recruitment of FANCD2 to interstrand cross-links (ICLs) leading to FANCD2 activation (By similarity). Also ubiquitinates non-histone proteins, such as histone H3, KIF11 and PML (By similarity). Acts as a critical player of proper spindle architecture by catalyzing the 'Lys-63'-linked ubiquitination of KIF11, thereby controlling KIF11 localization on the spindle (By similarity).
Cellular Location	Nucleus {ECO:0000255|PROSITE-ProRule:PRU00358, ECO:0000269|PubMed:10984098, ECO:0000269|PubMed:11161719, ECO:0000269|PubMed:14993289, ECO:0000269|PubMed:17994007, ECO:0000269|PubMed:21489993, ECO:0000269|PubMed:26065575, ECO:0000269|PubMed:36056023, ECO:0000269|PubMed:8634372}. Chromosome. Note=Associated with replicating DNA from early to late S phase, including at replicating pericentric heterochromatin (PubMed:36056023). Also localizes to euchromatic regions (PubMed:36056023). In non-S-phase cells, homogeneously distributed through the nucleus (PubMed:36056023)
Tissue Location	Expressed in thymus, testis, spleen and lung. Within testis, expressed in almost all cells except elongated spermatids.

Background

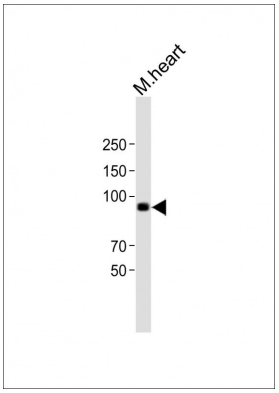
Multidomain protein that acts as a key epigenetic regulator by bridging DNA methylation and chromatin modification. Specifically recognizes and binds hemimethylated DNA at replication forks via its YDG domain and recruits DNMT1 methyltransferase to ensure faithful propagation of the DNA methylation patterns through DNA replication. In addition to its role in maintenance of DNA methylation, also plays a key role in chromatin modification: through its tudor-like regions and PHD- type zinc fingers, specifically recognizes and binds histone H3 trimethylated at 'Lys-9' (H3K9me3) and unmethylated at 'Arg-2' (H3R2me0), respectively, and recruits chromatin proteins. Enriched in pericentric heterochromatin where it recruits different chromatin modifiers required for this chromatin replication. Also localizes to euchromatic regions where it negatively regulates transcription possibly by impacting DNA methylation and histone modifications. Has E3 ubiquitin-protein ligase activity by mediating the ubiquitination of target proteins such as histone H3 and PML. It is still unclear how E3 ubiquitin-protein ligase activity is related to its role in chromatin in vivo. May

be involved in DNA repair.

References

Fujimori A.,et al.Mamm. Genome 9:1032-1035(1998).
Davenport J.W.,et al.Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
Carninci P.,et al.Science 309:1559-1563(2005).
Church D.M.,et al.PLoS Biol. 7:E1000112-E1000112(2009).
Muto M.,et al.J. Biol. Chem. 277:34549-34555(2002).

Images



All lanes: Anti-FGFR1(Y307)Antibody at 1:500 dilution + Mouse heart lysate Lysates/proteins at 20 µg per lane. Secondary: Goat Anti-Rabbit IgG, (H+L), Peroxidase conjugated (ASP1615) at 1/15000 dilution. Observed band size: 90 KDa Blocking/Dilution buffer: 5% NFDM/TBST.

Please note: All products are 'FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES'.