

HRX Antibody (C-term)
Purified Rabbit Polyclonal Antibody (Pab)
Catalog # AP1123b**Specification**

HRX Antibody (C-term) - Product Information

Application	IHC-P, FC,E
Primary Accession	Q03164
Other Accession	NP_005924
Reactivity	Human
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	431764
Antigen Region	3879-3908

HRX Antibody (C-term) - Additional Information**Gene ID** 4297**Other Names**

Histone-lysine N-methyltransferase 2A, Lysine N-methyltransferase 2A, ALL-1, CXXC-type zinc finger protein 7, Myeloid/lymphoid or mixed-lineage leukemia, Myeloid/lymphoid or mixed-lineage leukemia protein 1, Trithorax-like protein, Zinc finger protein HRX, MLL cleavage product N320, N-terminal cleavage product of 320 kDa, p320, MLL cleavage product C180, C-terminal cleavage product of 180 kDa, p180, KMT2A, ALL1, CXXC7, HRX, HTRX, MLL, MLL1, TRX1

Target/Specificity

This HRX antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 3879-3908 amino acids from the C-terminal region of human HRX.

Dilution

IHC-P~~1:50~100

FC~~1:10~50

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

HRX Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

HRX Antibody (C-term) - Protein Information

Name KMT2A**Synonyms** ALL1, CXXC7, HRX, HTRX, MLL, MLL1, TRX1

Function Histone methyltransferase that plays an essential role in early development and hematopoiesis (PubMed:[12453419](#), PubMed:[15960975](#), PubMed:[19187761](#), PubMed:[19556245](#), PubMed:[20677832](#), PubMed:[21220120](#), PubMed:[26886794](#)). Catalytic subunit of the MLL1/MLL complex, a multiprotein complex that mediates both methylation of 'Lys-4' of histone H3 (H3K4me) complex and acetylation of 'Lys-16' of histone H4 (H4K16ac) (PubMed:[12453419](#), PubMed:[15960975](#), PubMed:[19187761](#), PubMed:[19556245](#), PubMed:[20677832](#), PubMed:[21220120](#), PubMed:[24235145](#), PubMed:[26886794](#)). Catalyzes methyl group transfer from S-adenosyl-L- methionine to the epsilon-amino group of 'Lys-4' of histone H3 (H3K4) via a non-processive mechanism. Part of chromatin remodeling machinery predominantly forms H3K4me1 and H3K4me2 methylation marks at active chromatin sites where transcription and DNA repair take place (PubMed:[12453419](#), PubMed:[15960975](#), PubMed:[19187761](#), PubMed:[19556245](#), PubMed:[20677832](#), PubMed:[21220120](#), PubMed:[25561738](#), PubMed:[26886794](#)). Has weak methyltransferase activity by itself, and requires other component of the MLL1/MLL complex to obtain full methyltransferase activity (PubMed:[19187761](#), PubMed:[26886794](#)). Has no activity toward histone H3 phosphorylated on 'Thr-3', less activity toward H3 dimethylated on 'Arg-8' or 'Lys-9', while it has higher activity toward H3 acetylated on 'Lys-9' (PubMed:[19187761](#)). Binds to unmethylated CpG elements in the promoter of target genes and helps maintain them in the nonmethylated state (PubMed:[20010842](#)). Required for transcriptional activation of HOXA9 (PubMed:[12453419](#), PubMed:[20010842](#), PubMed:[20677832](#)). Promotes PPP1R15A-induced apoptosis (PubMed:[10490642](#)). Plays a critical role in the control of circadian gene expression and is essential for the transcriptional activation mediated by the CLOCK-BMAL1 heterodimer (By similarity). Establishes a permissive chromatin state for circadian transcription by mediating a rhythmic methylation of 'Lys-4' of histone H3 (H3K4me) and this histone modification directs the circadian acetylation at H3K9 and H3K14 allowing the recruitment of CLOCK-BMAL1 to chromatin (By similarity). Also has auto-methylation activity on Cys-3882 in absence of histone H3 substrate (PubMed:[24235145](#)).

Cellular Location

Nucleus [MLL cleavage product C180]: Nucleus. Note=Localizes to a diffuse nuclear pattern when not associated with MLL cleavage product N320

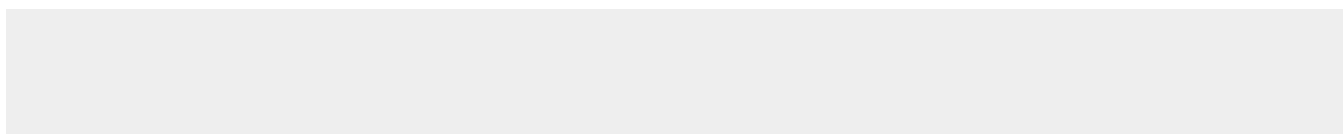
Tissue Location

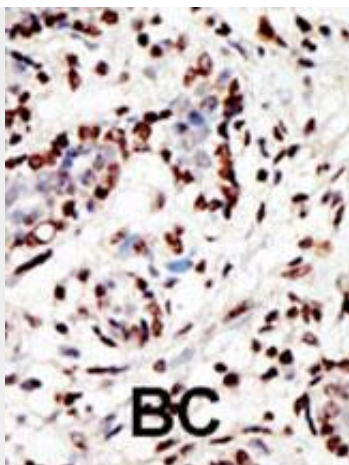
Heart, lung, brain and T- and B-lymphocytes.

HRX Antibody (C-term) - Protocols

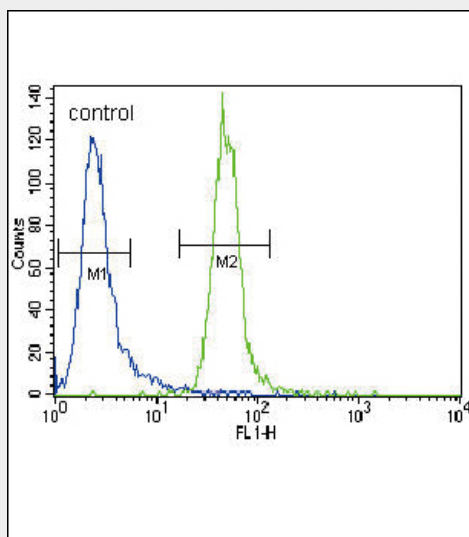
Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

HRX Antibody (C-term) - Images



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



HRX Antibody (C-term) (Cat. #AP1123b) flow cytometric analysis of CEM cells (right histogram) compared to a negative control cell (left histogram). FITC-conjugated goat-anti-rabbit secondary antibodies were used for the analysis.

HRX Antibody (C-term) - Background

The gene variously symbolized ALL1, HRX, or MLL located on 11q23 has been demonstrated to be fused with a number of translocation partners in cases of leukemia. Tse et al. (1995) characterized 2 t(1;11)(q21;q23) translocations that fused the MLL gene to a gene on chromosomal band 1q21, AF1Q, in 2 infants with acute myelomonocytic leukemia. In one of these patients, the derivative chromosome 11 represented an in-frame fusion of the N-terminal portion of the MLL gene to the complete AF1Q open reading frame, whereas the derivative chromosome 1 did not give rise to an open reading frame. This observation suggested that the N-terminal portion of the MLL gene is critical for leukemogenesis in translocations involving band 11q23.

HRX Antibody (C-term) - References

Megonigal, M.D., et al., Proc. Natl. Acad. Sci. U.S.A. 97(6):2814-2819 (2000).
Pegram, L.D., et al., Blood 96(13):4360-4362 (2000).
Sano, K., et al., Blood 95(3):1066-1068 (2000).

Cui, X., et al., Nat. Genet. 18(4):331-337 (1998).
Nilson, I., et al., Br. J. Haematol. 93(4):966-972 (1996).