

Resource Summary Report

Generated by [RRID](#) on Sep 3, 2026

HEXIM1 antibody - ChIP Grade

RRID:AB_2233058

Type: Antibody

Proper Citation

Abcam Cat# ab25388, RRID:AB_2233058

Antibody Information

URL: http://antibodyregistry.org/AB_2233058

Proper Citation: Abcam Cat# ab25388, RRID:AB_2233058

Target Antigen: HEXIM1

Host Organism: rabbit

Clonality: polyclonal

Comments: validation status unknown, seller recommendations provided in 2012:western blot, immunoprecipitation, immunohistochemistry

Antibody Name: HEXIM1 antibody - ChIP Grade

Description: This polyclonal targets HEXIM1

Target Organism: mouse, human

Antibody ID: AB_2233058

Vendor: Abcam

Catalog Number: ab25388

Record Creation Time: 20231110T045614+0000

Record Last Update: 20260830T032202+0000

Ratings and Alerts

No rating or validation information has been found for HEXIM1 antibody - ChIP Grade.

No alerts have been found for HEXIM1 antibody - ChIP Grade.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Uhl L, et al. (2026) MYC binding to nascent RNA suppresses innate immune signaling by R-loop-derived RNA-DNA hybrids. *Cell*, 189(5), 1371.

Chen SM, et al. (2025) SLFN14 functions as a P-TEFb inhibitor to modulate the transcription of HIV-1 and cellular genes. *Science advances*, 11(51), eadt7982.

Studniarek C, et al. (2021) The 7SK/P-TEFb snRNP controls ultraviolet radiation-induced transcriptional reprogramming. *Cell reports*, 35(2), 108965.

Mota de S?? P, et al. (2020) Bromodomain and Extraterminal Inhibition by JQ1 Produces Divergent Transcriptional Regulation of Suppressors of Cytokine Signaling Genes in Adipocytes. *Endocrinology*, 161(2).

Factor DC, et al. (2020) Cell Type-Specific Intralocus Interactions Reveal Oligodendrocyte Mechanisms in MS. *Cell*, 181(2), 382.

Edwards DS, et al. (2020) BRD4 Prevents R-Loop Formation and Transcription-Replication Conflicts by Ensuring Efficient Transcription Elongation. *Cell reports*, 32(12), 108166.

Faust TB, et al. (2018) The HIV-1 Tat protein recruits a ubiquitin ligase to reorganize the 7SK snRNP for transcriptional activation. *eLife*, 7.

Bowry A, et al. (2018) BET Inhibition Induces HEXIM1- and RAD51-Dependent Conflicts between Transcription and Replication. *Cell reports*, 25(8), 2061.

Winter GE, et al. (2017) BET Bromodomain Proteins Function as Master Transcription Elongation Factors Independent of CDK9 Recruitment. *Molecular cell*, 67(1), 5.