

# Resource Summary Report

Generated by [dkNET](#) on Aug 16, 2026

## Rabbit anti-ASH2 Antibody, Affinity Purified

RRID:AB\_451024

Type: Antibody

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### Proper Citation

Bethyl Cat# A300-489A, RRID:AB\_451024

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### Antibody Information

**URL:** [http://antibodyregistry.org/AB\\_451024](http://antibodyregistry.org/AB_451024)

**Proper Citation:** Bethyl Cat# A300-489A, RRID:AB\_451024

**Target Antigen:** ASH2

**Host Organism:** rabbit

**Clonality:** polyclonal

**Comments:** Applications: WB, IP, IHC, ChIP, ChIP-chip

**Antibody Name:** Rabbit anti-ASH2 Antibody, Affinity Purified

**Description:** This polyclonal targets ASH2

**Target Organism:** human

**Antibody ID:** AB\_451024

**Vendor:** Bethyl

**Catalog Number:** A300-489A

**Alternative Catalog Numbers:** ENCAB947YBH

**Record Creation Time:** 20250820T030223+0000

**Record Last Update:** 20250820T151110+0000

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### Ratings and Alerts

- ENCODE PROJECT External validation for lot: 2 is available under ENCODE ID: ENCAB947YBH - ENCODE

<https://www.encodeproject.org/antibodies/ENCAB947YBH>

**Warning:** Discontinued at Thermo Fisher Scientific  
Applications: WB, IP, IHC, ChIP, ChIP-chip

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## Data and Source Information

**Source:** [Antibody Registry](#)

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Nguyen N, et al. (2022) Recruitment of MLL1 complex is essential for SETBP1 to induce myeloid transformation. *iScience*, 25(1), 103679.

Nayak A, et al. (2019) Regulation of SETD7 Methyltransferase by SENP3 Is Crucial for Sarcomere Organization and Cachexia. *Cell reports*, 27(9), 2725.

Li L, et al. (2019) The COMPASS Family Protein ASH2L Mediates Corticogenesis via Transcriptional Regulation of Wnt Signaling. *Cell reports*, 28(3), 698.

Hu D, et al. (2017) Not All H3K4 Methylations Are Created Equal: Mll2/COMPASS Dependency in Primordial Germ Cell Specification. *Molecular cell*, 65(3), 460.

Wang SP, et al. (2017) A UTX-MLL4-p300 Transcriptional Regulatory Network Coordinately Shapes Active Enhancer Landscapes for Eliciting Transcription. *Molecular cell*, 67(2), 308.