

Resource Summary Report

Generated by [dkNET](#) on Aug 21, 2026

Rabbit anti-INT4 Antibody, Affinity Purified

RRID:AB_937909

Type: Antibody

Proper Citation

Bethyl Cat# A301-296A, RRID:AB_937909

Antibody Information

URL: http://antibodyregistry.org/AB_937909

Proper Citation: Bethyl Cat# A301-296A, RRID:AB_937909

Target Antigen: INT4

Host Organism: rabbit

Clonality: polyclonal

Comments: Applications: WB, IP, IHC
Original Manufacturer

Antibody Name: Rabbit anti-INT4 Antibody, Affinity Purified

Description: This polyclonal targets INT4

Target Organism: mouse, human

Antibody ID: AB_937909

Vendor: Bethyl

Catalog Number: A301-296A

Alternative Catalog Numbers: A301-296A-M, A301-296A-T

Record Creation Time: 20250819T235937+0000

Record Last Update: 20250820T070255+0000

Ratings and Alerts

No rating or validation information has been found for Rabbit anti-INT4 Antibody, Affinity Purified.

Warning: Discontinued at Thermo Fisher Scientific
Applications: WB, IP, IHC
Original Manufacturer

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Blears D, et al. (2024) Redundant pathways for removal of defective RNA polymerase II complexes at a promoter-proximal pause checkpoint. *Molecular cell*, 84(24), 4790.

Sabath K, et al. (2024) Assembly mechanism of Integrator's RNA cleavage module. *Molecular cell*, 84(15), 2882.

Sabath K, et al. (2024) Basis of gene-specific transcription regulation by the Integrator complex. *Molecular cell*, 84(13), 2525.

Bhowmick R, et al. (2023) Integrator facilitates RNAPII removal to prevent transcription-replication collisions and genome instability. *Molecular cell*, 83(13), 2357.

Stein CB, et al. (2022) Integrator endonuclease drives promoter-proximal termination at all RNA polymerase II-transcribed loci. *Molecular cell*, 82(22), 4232.

Beckedorff F, et al. (2020) The Human Integrator Complex Facilitates Transcriptional Elongation by Endonucleolytic Cleavage of Nascent Transcripts. *Cell reports*, 32(3), 107917.