

Resource Summary Report

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

TFIIH p89 (S-19)

RRID:AB_2262177

Type: Antibody

Proper Citation

Santa Cruz Biotechnology Cat# sc-293, RRID:AB_2262177

Antibody Information

URL: http://antibodyregistry.org/AB_2262177

Proper Citation: Santa Cruz Biotechnology Cat# sc-293, RRID:AB_2262177

Target Antigen: ERCC3

Host Organism: rabbit

Clonality: polyclonal

Comments: Discontinued: 2016; validation status unknown check with seller; recommendations: ELISA; Immunofluorescence; Immunoprecipitation; Western Blot; Western Blotting, Immunoprecipitation, Immunofluorescence, ELISA

Antibody Name: TFIIH p89 (S-19)

Description: This polyclonal targets ERCC3

Target Organism: rat, mouse, human

Clone ID: S-19

Antibody ID: AB_2262177

Vendor: Santa Cruz Biotechnology

Catalog Number: sc-293

Record Creation Time: 20250820T043609+0000

Record Last Update: 20250820T192536+0000

Ratings and Alerts

No rating or validation information has been found for TFIH p89 (S-19).

Warning: Discontinued: 2016

Discontinued: 2016; validation status unknown check with seller; recommendations: ELISA; Immunofluorescence; Immunoprecipitation; Western Blot; Western Blotting, Immunoprecipitation, Immunofluorescence, ELISA

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Sakellariou D, et al. (2022) MutS γ regulates G4-associated telomeric R-loops to maintain telomere integrity in ALT cancer cells. *Cell reports*, 39(1), 110602.

Chappidi N, et al. (2020) Fork Cleavage-Religation Cycle and Active Transcription Mediate Replication Restart after Fork Stalling at Co-transcriptional R-Loops. *Molecular cell*, 77(3), 528.

Gillespie MA, et al. (2020) Absolute Quantification of Transcription Factors Reveals Principles of Gene Regulation in Erythropoiesis. *Molecular cell*, 78(5), 960.

Legrand N, et al. (2019) PPAR γ recruits NCOR and regulates transcription reinitiation of ANGPTL4. *Nucleic acids research*, 47(18), 9573.

Sun F, et al. (2019) Promoter-Enhancer Communication Occurs Primarily within Insulated Neighborhoods. *Molecular cell*, 73(2), 250.

Piquet S, et al. (2018) The Histone Chaperone FACT Coordinates H2A.X-Dependent Signaling and Repair of DNA Damage. *Molecular cell*, 72(5), 888.

Przetocka S, et al. (2018) CtIP-Mediated Fork Protection Synergizes with BRCA1 to Suppress Genomic Instability upon DNA Replication Stress. *Molecular cell*, 72(3), 568.

Vujanovic M, et al. (2017) Replication Fork Slowing and Reversal upon DNA Damage Require PCNA Polyubiquitination and ZRANB3 DNA Translocase Activity. *Molecular cell*, 67(5), 882.

Alekseev S, et al. (2017) Transcription without XPB Establishes a Unified Helicase-Independent Mechanism of Promoter Opening in Eukaryotic Gene Expression. *Molecular cell*, 65(3), 504.

Di Marco S, et al. (2017) RECQ5 Helicase Cooperates with MUS81 Endonuclease in Processing Stalled Replication Forks at Common Fragile Sites during Mitosis. *Molecular cell*, 66(5), 658.

Kuschal C, et al. (2016) GTF2E2 Mutations Destabilize the General Transcription Factor Complex TFIIE in Individuals with DNA Repair-Proficient Trichothiodystrophy. American journal of human genetics, 98(4), 627.