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 TFIIH 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? 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?/? 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.