

Resource Summary Report

Generated by [dkNET](#) on Jul 30, 2026

Anti-c-Jun, phospho (Ser73) Antibody, Unconjugated

RRID:AB_330892

Type: Antibody

Proper Citation

Cell Signaling Technology Cat# 9164, RRID:AB_330892

Antibody Information

URL: http://antibodyregistry.org/AB_330892

Proper Citation: Cell Signaling Technology Cat# 9164, RRID:AB_330892

Target Antigen: c-Jun, phospho (Ser73)

Clonality: unknown

Comments: Applications: W, IF-IC. Consolidation on 10/2018: AB_10078079, AB_10078897, AB_330892, AB_330893.

Antibody Name: Anti-c-Jun, phospho (Ser73) Antibody, Unconjugated

Description: This unknown targets c-Jun, phospho (Ser73)

Target Organism: rat, mouse, human

Antibody ID: AB_330892

Vendor: Cell Signaling Technology

Catalog Number: 9164

Record Creation Time: 20250819T222809+0000

Record Last Update: 20250820T021727+0000

Ratings and Alerts

No rating or validation information has been found for Anti-c-Jun, phospho (Ser73) Antibody, Unconjugated.

No alerts have been found for Anti-c-Jun, phospho (Ser73) Antibody, Unconjugated.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 19 mentions in open access literature.

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

Joseph S, et al. (2025) MAPK14/p38 γ shapes the molecular landscape of endometrial cancer and promotes tumorigenic characteristics. Cell reports, 44(1), 115104.

Park JS, et al. (2025) L1CAM signaling through planar cell polarity generates SOX2 + metastatic progenitors in lung adenocarcinoma. bioRxiv : the preprint server for biology.

Liu F, et al. (2025) PTEN neddylation aggravates CDK4/6 inhibitor resistance in breast cancer. Oncogene, 44(33), 2997.

Zhao B, et al. (2025) PLK1 blockade enhances the anti-tumor effect of MAPK inhibition in pancreatic ductal adenocarcinoma. Cell reports, 44(4), 115541.

Lee S, et al. (2024) Ganoderma lucidum extract attenuates corticotropin-releasing hormone-induced cellular senescence in human hair follicle cells. iScience, 27(5), 109675.

Yu F, et al. (2024) Pancreatic β cell interleukin-22 receptor subunit alpha 1 deficiency impairs β cell function in type 2 diabetes via cytochrome b5 reductase 3. Cell reports, 43(12), 115057.

Capelo-Diz A, et al. (2023) Hepatic levels of S-adenosylmethionine regulate the adaptive response to fasting. Cell metabolism, 35(8), 1373.

Wei X, et al. (2022) Ablating Lgr5-expressing prostatic stromal cells activates the ERK-mediated mechanosensory signaling and disrupts prostate tissue homeostasis. Cell reports, 40(10), 111313.

Zhao K, et al. (2021) Cytidine Deaminase APOBEC3A Regulates PD-L1 Expression in Cancer Cells in a JNK/c-JUN-Dependent Manner. Molecular cancer research : MCR, 19(9), 1571.

Huang Z, et al. (2020) Histone deacetylase 6 promotes growth of glioblastoma through the MKK7/JNK/c-Jun signaling pathway. Journal of neurochemistry, 152(2), 221.

Flasse L, et al. (2020) Apical Restriction of the Planar Cell Polarity Component VANGL in Pancreatic Ducts Is Required to Maintain Epithelial Integrity. Cell reports, 31(8), 107677.

Berthenet K, et al. (2020) Failed Apoptosis Enhances Melanoma Cancer Cell Aggressiveness. Cell reports, 31(10), 107731.

Sinha NK, et al. (2020) EDF1 coordinates cellular responses to ribosome collisions. *eLife*, 9.

Crespo M, et al. (2020) Neutrophil infiltration regulates clock-gene expression to organize daily hepatic metabolism. *eLife*, 9.

Ng PK, et al. (2018) Systematic Functional Annotation of Somatic Mutations in Cancer. *Cancer cell*, 33(3), 450.

Tang C, et al. (2018) Glucose-Induced β -Cell Dysfunction In Vivo: Evidence for a Causal Role of C-jun N-terminal Kinase Pathway. *Endocrinology*, 159(11), 3643.

Yuan D, et al. (2017) Kupffer Cell-Derived Tnf Triggers Cholangiocellular Tumorigenesis through JNK due to Chronic Mitochondrial Dysfunction and ROS. *Cancer cell*, 31(6), 771.

Bajikar SS, et al. (2017) Tumor-Suppressor Inactivation of GDF11 Occurs by Precursor Sequestration in Triple-Negative Breast Cancer. *Developmental cell*, 43(4), 418.

Cheng CS, et al. (2017) Iterative Modeling Reveals Evidence of Sequential Transcriptional Control Mechanisms. *Cell systems*, 4(3), 330.