Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 29, 2025

HNF4? (C11F12) Rabbit mAb

RRID:AB_2295208
Type: Antibody

Proper Citation

(Cell Signaling Technology Cat# 3113, RRID:AB_2295208)

Antibody Information

URL: http://antibodyregistry.org/AB_2295208

Proper Citation: (Cell Signaling Technology Cat# 3113, RRID:AB_2295208)

Target Antigen: HNF4alpha

Host Organism: rabbit

Clonality: monoclonal

Comments: Applications: WB, IHC-P, IF-IC

Antibody Name: HNF4? (C11F12) Rabbit mAb

Description: This monoclonal targets HNF4alpha

Target Organism: human

Clone ID: C11F12

Antibody ID: AB_2295208

Vendor: Cell Signaling Technology

Catalog Number: 3113

Alternative Catalog Numbers: 3113S

Record Creation Time: 20231110T075420+0000

Record Last Update: 20241115T010226+0000

Ratings and Alerts

No rating or validation information has been found for HNF4? (C11F12) Rabbit mAb.

No alerts have been found for HNF4? (C11F12) Rabbit mAb.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 37 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Trinh LT, et al. (2025) Positive autoregulation of Sox17 is necessary for gallbladder and extrahepatic bile duct formation. Development (Cambridge, England), 152(2).

Wang Z, et al. (2024) Molecular subtypes of neuroendocrine carcinomas: A cross-tissue classification framework based on five transcriptional regulators. Cancer cell, 42(6), 1106.

Namoto K, et al. (2024) NIBR-LTSi is a selective LATS kinase inhibitor activating YAP signaling and expanding tissue stem cells in vitro and in vivo. Cell stem cell, 31(4), 554.

Bi G, et al. (2024) Modeling alcohol-associated liver disease in humans using adipose stromal or stem cell-derived organoids. Cell reports methods, 4(5), 100778.

Khampang S, et al. (2023) CRISPR/Cas9 mediated approach to generate YAP-depleted human embryonic stem cell line (MUSIe002-A-1). Stem cell research, 66, 102990.

Ebisudani T, et al. (2023) Genotype-phenotype mapping of a patient-derived lung cancer organoid biobank identifies NKX2-1-defined Wnt dependency in lung adenocarcinoma. Cell reports, 42(3), 112212.

He S, et al. (2023) Spatial-temporal proliferation of hepatocytes during pregnancy revealed by genetic lineage tracing. Cell stem cell, 30(11), 1549.

Roberts MA, et al. (2023) Parallel CRISPR-Cas9 screens identify mechanisms of PLIN2 and lipid droplet regulation. Developmental cell, 58(18), 1782.

Liu X, et al. (2023) Genetic recording of in vivo cell proliferation by ProTracer. Nature protocols.

Wei T, et al. (2023) Loss of Mettl3 enhances liver tumorigenesis by inducing hepatocyte dedifferentiation and hyperproliferation. Cell reports, 42(7), 112704.

Qi S, et al. (2023) Two Hippo signaling modules orchestrate liver size and tumorigenesis. The EMBO journal, e112126.

Qi S, et al. (2022) WWC proteins mediate LATS1/2 activation by Hippo kinases and imply a tumor suppression strategy. Molecular cell, 82(10), 1850.

Ma H, et al. (2022) The nuclear receptor THRB facilitates differentiation of human PSCs into more mature hepatocytes. Cell stem cell, 29(5), 795.

Khampang S, et al. (2022) Derivation of the MUSIe002-A human embryonic stem cell line. Stem cell research, 59, 102660.

Ungricht R, et al. (2022) Genome-wide screening in human kidney organoids identifies developmental and disease-related aspects of nephrogenesis. Cell stem cell, 29(1), 160.

Huang XT, et al. (2022) Embryogenic stem cell-derived intestinal crypt fission directs de novo crypt genesis. Cell reports, 41(11), 111796.

Trinh LT, et al. (2022) Differential regulation of alternate promoter regions in Sox17 during endodermal and vascular endothelial development. iScience, 25(9), 104905.

Orstad G, et al. (2022) FoxA1 and FoxA2 control growth and cellular identity in NKX2-1-positive lung adenocarcinoma. Developmental cell, 57(15), 1866.

Johnson HJ, et al. (2022) A scalable and tunable thermoreversible polymer for 3D human pluripotent stem cell biomanufacturing. iScience, 25(10), 104971.

Han X, et al. (2021) A suite of new Dre recombinase drivers markedly expands the ability to perform intersectional genetic targeting. Cell stem cell, 28(6), 1160.