

Resource Summary Report

Generated by PRECISE-TBI on Aug 31, 2026

Anti-HNF3beta/FOXA2

RRID:AB_390153

Type: Antibody

Proper Citation

Millipore Cat# 07-633, RRID:AB_390153

Antibody Information

URL: http://antibodyregistry.org/AB_390153

Proper Citation: Millipore Cat# 07-633, RRID:AB_390153

Target Antigen: HNF3beta/FOXA2

Host Organism: rabbit

Clonality: polyclonal

Comments: seller recommendations: IgG; IgG Western Blot; Immunohistochemistry; IH, WB

Antibody Name: Anti-HNF3beta/FOXA2

Description: This polyclonal targets HNF3beta/FOXA2

Target Organism: h, m, r

Antibody ID: AB_390153

Vendor: Millipore

Catalog Number: 07-633

Record Creation Time: 20250819T225710+0000

Record Last Update: 20250820T034907+0000

Ratings and Alerts

- Used for flow cytometry assay by the Human Islet Research Network community.
Contact(s): [Shuibing Chen](#), [Stephen Parker](#) - Human Islets Research Network

No alerts have been found for Anti-HNF3beta/FOXA2.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 47 mentions in open access literature.

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

Borisch C, et al. (2026) Generation of human induced pluripotent stem cell line MHHi040-A from a female Fabry disease patient carrying c.644A>G missense mutation. Stem cell research, 95, 104064.

Secula S, et al. (2026) Generation of an induced pluripotent stem cell line, LGMi002-A, from a Bardet-Biedl Syndrome patient with a BBS5 homozygous pathogenic variant. Stem cell research, 95, 104046.

Secula S, et al. (2026) Generation of an induced pluripotent stem cell line from an Alström syndrome patient with biallelic ALMS1 pathogenic variants. Stem cell research, 94, 103990.

Ghiasighorveh P, et al. (2026) Generation of iPSC and isogenic gene-corrected lines from a patient with Shwachman Diamond syndrome. Stem cell research, 95, 104045.

Chen G, et al. (2026) Single-cell multi-omic analyses highlight the essential role of NKX2.2-CLEC16A/endosomal pathway for human pancreatic differentiation and function. Cell reports, 45(2), 116990.

Suryaprakash S, et al. (2026) Generation of iPSC and isogenic gene-corrected lines from a patient with RPS7 (c.277_279delGTC)-mutated Diamond-Blackfan anemia syndrome. Stem cell research, 91, 103908.

Zhang X, et al. (2026) Human pancreatic progenitor organoids define genetic and epigenetic barriers to early PDAC transformation. Developmental cell, 61(6), 1190.

D'Oliveira Albanus R, et al. (2025) Integrative single-cell multi-omics profiling of human pancreatic islets identifies T1D-associated genes and regulatory signals. Cell reports, 44(8), 116065.

Clua Provost C, et al. (2025) Generation of two isogenic-corrected control cell lines (IRMBi001-A-1; IRMBi001-A-2) from Autosomal dominant Alzheimer's disease patient-derived iPSCs carrying a G217D mutation in presenilin 1 gene. Stem cell research, 87, 103780.

Urdaneta KE, et al. (2025) Generation of patient-derived and gene-corrected hiPSC lines from Hereditary Hemorrhagic Telangiectasia type 2 patients with ACVRL1 c.1042delG

mutation. Stem cell research, 88, 103812.

Mamaeva D, et al. (2025) Generation of a human iPSC line, INMi007-A, carrying compound heterozygous DCT variants associated with oculocutaneous albinism type 8. Stem cell research, 88, 103810.

Huang T, et al. (2025) Inhibition of PRC2 enables self-renewal of blastoid-competent naive pluripotent stem cells from chimpanzee. Cell stem cell, 32(4), 627.

Baek D, et al. (2024) Generation of an induced pluripotent stem cell line (KNIHi001-A) by reprogramming peripheral blood mononuclear cells isolated from a patient with Parkinson's disease. Stem cell research, 76, 103358.

Manibarathi K, et al. (2024) An iPSC model for POLR3A-associated spastic ataxia: Generation of three unrelated patient cell lines. Stem cell research, 76, 103363.

Korneck M, et al. (2024) Generation of homozygous and heterozygous REEP1 knockout induced pluripotent stem cell lines by CRISPR/Cas9 gene editing. Stem cell research, 77, 103378.

Matsui S, et al. (2024) Pioneer and PRDM transcription factors coordinate bivalent epigenetic states to safeguard cell fate. Molecular cell, 84(3), 476.

Jahn C, et al. (2024) Generation of human induced pluripotent stem cell line MHHi029-A from a male Fabry disease patient carrying c.959A > T mutation. Stem cell research, 77, 103404.

Swisa A, et al. (2024) The evolutionarily ancient FOXA transcription factors shape the murine gut microbiome via control of epithelial glycosylation. Developmental cell, 59(16), 2069.

Matsui S, et al. (2024) Protocol for establishing inducible CRISPRd system for blocking transcription factor-binding sites in human pluripotent stem cells. STAR protocols, 5(3), 103233.

Son N, et al. (2023) Generation of a human fibroblast-derived induced pluripotent stem cell line KRIBBi009-A from a patient with breast cancer. Stem cell research, 68, 103060.