Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 14, 2025

Human ERR beta/NR3B2 Antibody

RRID:AB_2100412
Type: Antibody

Proper Citation

(R and D Systems Cat# PP-H6705-00, RRID:AB_2100412)

Antibody Information

URL: http://antibodyregistry.org/AB_2100412

Proper Citation: (R and D Systems Cat# PP-H6705-00, RRID:AB_2100412)

Target Antigen: ERR beta/NR3B2

Host Organism: Mouse

Clonality: monoclonal

Comments: Applications: Western Blot, Immunohistochemistry, Immunoprecipitation

Antibody Name: Human ERR beta/NR3B2 Antibody

Description: This monoclonal targets ERR beta/NR3B2

Target Organism: Human

Clone ID: H6705

Antibody ID: AB_2100412

Vendor: R and D Systems

Catalog Number: PP-H6705-00

Record Creation Time: 20241016T221914+0000

Record Last Update: 20241016T223908+0000

Ratings and Alerts

No rating or validation information has been found for Human ERR beta/NR3B2 Antibody.

No alerts have been found for Human ERR beta/NR3B2 Antibody.

Data and Source Information

Source: Antibody Registry

Usage and Citation Metrics

We found 17 mentions in open access literature.

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

Ura H, et al. (2024) Establishment of a human induced pluripotent stem cell line, KMUGMCi010-A, from a patient with X-linked Ohdo syndrome bearing missense mutation in the MED12 gene. Stem cell research, 77, 103388.

Li QS, et al. (2023) ESRRB Inhibits the TGF? Signaling Pathway to Drive Cell Proliferation in Cervical Cancer. Cancer research, 83(18), 3095.

Ura H, et al. (2023) Establishment of human induced pluripotent stem cell lines, KMUGMCi006, from a patient with Tuberous sclerosis complex (TSC) bearing mosaic nonsense mutations in the Tuberous sclerosis complex 2 (TSC2) gene. Stem cell research, 70, 103129.

Ura H, et al. (2022) Establishment of a human induced pluripotent stem cell line, KMUGMCi002-A, from a patient bearing a heterozygous c.6362_6364del mutation in the NIPBL gene leading Cornelia de Lange syndrome (CdLS). Stem cell research, 63, 102860.

Ura H, et al. (2022) Establishment of a human induced pluripotent stem cell line, KMUGMCi001-A, from a patient bearing a heterozygous c.772 + 3_772 + 4dup mutation in the ACVRL1 gene leading Telangiectasia, hereditary hemorrhagic, type 2 (HHT2). Stem cell research, 61, 102743.

Ura H, et al. (2022) Establishment of a human induced pluripotent stem cell line, KMUGMCi004-A, from a patient bearing a heterozygous c.1832delG mutation in the APC gene leading familial adenomatous polyposis (FAP). Stem cell research, 63, 102867.

Kim YS, et al. (2022) Rap1 controls epiblast morphogenesis in sync with the pluripotency states transition. Developmental cell, 57(16), 1937.

Ura H, et al. (2022) Establishment of a human induced pluripotent stem cell line, KMUGMCi003-A, from a patient with trichothiodystrophy 1 (TTD1) bearing compound heterozygous missense mutations in the ERCC2 gene. Stem cell research, 64, 102885.

Ura H, et al. (2022) Establishment of a human induced pluripotent stem cell line, KMUGMCi005-A, from a patient with Epidermodysplasia verruciformis (EV) bearing homozygous splicing donor site mutation in the TMC8 gene. Stem cell research, 64, 102926.

Bayerl J, et al. (2021) Principles of signaling pathway modulation for enhancing human naive pluripotency induction. Cell stem cell, 28(9), 1549.

Hashimoto M, et al. (2019) Epiblast Formation by TEAD-YAP-Dependent Expression of Pluripotency Factors and Competitive Elimination of Unspecified Cells. Developmental cell, 50(2), 139.

Garland W, et al. (2019) A Functional Link between Nuclear RNA Decay and Transcriptional Control Mediated by the Polycomb Repressive Complex 2. Cell reports, 29(7), 1800.

McLaughlin K, et al. (2019) DNA Methylation Directs Polycomb-Dependent 3D Genome Re-organization in Naive Pluripotency. Cell reports, 29(7), 1974.

Mor N, et al. (2018) Neutralizing Gatad2a-Chd4-Mbd3/NuRD Complex Facilitates Deterministic Induction of Naive Pluripotency. Cell stem cell, 23(3), 412.

Bornelöv S, et al. (2018) The Nucleosome Remodeling and Deacetylation Complex Modulates Chromatin Structure at Sites of Active Transcription to Fine-Tune Gene Expression. Molecular cell, 71(1), 56.

Sardina JL, et al. (2018) Transcription Factors Drive Tet2-Mediated Enhancer Demethylation to Reprogram Cell Fate. Cell stem cell, 23(5), 727.

Leroy F, et al. (2014) Early intrinsic hyperexcitability does not contribute to motoneuron degeneration in amyotrophic lateral sclerosis. eLife, 3.