

Resource Summary Report

Generated by [dkNET](#) on Apr 28, 2025

HEK293T

RRID:CVCL_0063

Type: Cell Line

Proper Citation

(Abcam Cat# ab255449, RRID:CVCL_0063)

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_0063

Proper Citation: (Abcam Cat# ab255449, RRID:CVCL_0063)

Sex: Female

Defining Citation: [PMID:3031469](#), [PMID:15900046](#), [PMID:25182477](#), [PMID:26694163](#), [PMID:28196595](#), [PMID:28601559](#), [PMID:29468137](#), [PMID:34800366](#)

Comments: Omics: Transcriptome analysis by RNAseq., Omics: Protein expression by reverse-phase protein arrays., Omics: Mitochondrial proteome analysis., Omics: miRNA expression profiling., Omics: Genome sequenced., Omics: Deep proteome analysis., Virology: Highly susceptible to infection by Zika virus (ZIKV) (PubMed=29468137)., Part of: MD Anderson Cell Lines Project., Part of: ENCODE project common cell types; tier 3.

Category: Transformed cell line

Name: HEK293T

Synonyms: Hek293T, HEK-293T, HEK 293T, HEK-293-T, HEK 293 T, 293-T, 293 T, 293T, Human Embryonic Kidney 293T, 293tsA1609neo

Cross References: BTO:BTO_0002181, CLO:CLO_0050894, EFO:EFO_0001082, EFO:EFO_0001184, CLDB:cl7154, Abcam:ab255449, Abcam:ab255593, Abcam:ab282205, AddexBio:T0011002/445, ATCC:CRL-3216, BCRJ:0361, BioGRID_ORCS_Cell_line:267, BioSample:SAMN01821609, BioSample:SAMN03473454, BioSamples:SAMEA2168958, BioSamples:SAMEA2536418, BioSamples:SAMEA2536419, CCLV:CCLV-RIE 1018, CCRID:1101HUM-PUMC000091, CCRID:3101HUMGNHu17, CCRID:3101HUMSCSP502, CCRID:4201HUM-CCTCC00187, CCRID:5301HUM-KCB07044YJ, CCTCC:GDC0187, ChEMBL-Cells:ChEMBL3706569, ChEMBL-Targets:ChEMBL3706568, CLS:300189,

Cosmic:1326280, Cosmic:2440479, DSMZ:ACC-635, DSMZCellDive:ACC-635, ECACC:12022001, ENCODE:ENCBS276VDW, ENCODE:ENCBS319KKV, ENCODE:ENCBS323RYE, ENCODE:ENCBS582KZL, ENCODE:ENCBS634AAA, ENCODE:ENCBS986WKD, FCS-free:6-2-482-1-4-3, GEO:GSM1008573, GEO:GSM5936927, GEO:GSM5936928, GEO:GSM5936929, IBRC:C10683, ICLC:HTL04001, KCB:KCB 200744YJ, Lonza:504, MeSH:D057809, NCBI_Iran:C498, NCBI_Iran:C644, PRIDE:PXD000593, PRIDE:PXD001062, PRIDE:PXD001165, PRIDE:PXD001609, PRIDE:PXD004452, PRIDE:PXD006633, PRIDE:PXD006698, PRIDE:PXD012357, PRIDE:PXD013232, PRIDE:PXD016924, PRIDE:PXD018162, PRIDE:PXD018182, PRIDE:PXD019204, PRIDE:PXD028149, PRIDE:PXD029242, PRIDE:PXD029738, PRIDE:PXD030166, PubChem_Cell_line:CVCL_0063, RCB:RCB2202, TOKU-E:3537, Ubigen:YC-A006, Ubigen:YC-A007, Wikidata:Q27546876

ID: CVCL_0063

Vendor: Abcam

Catalog Number: ab255449

Record Creation Time: 20250131T200353+0000

Record Last Update: 20250131T201608+0000

Ratings and Alerts

No rating or validation information has been found for HEK293T.

No alerts have been found for HEK293T.

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 68026 mentions in open access literature.

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

Al Kabbani MA, et al. (2025) Effects of P301L-TAU on post-translational modifications of microtubules in human iPSC-derived cortical neurons and TAU transgenic mice. Neural regeneration research, 20(8), 2348.

Ren Z, et al. (2025) A peptide encoded by LINC00944 suppresses the growth of melanoma cells by diminishing EP400-MYC interaction. Biochemical pharmacology, 231, 116652.

Yu J, et al. (2025) Calcineurin: An essential regulator of sleep revealed by biochemical,

chemical biological, and genetic approaches. *Cell chemical biology*, 32(1), 157.

Huang H, et al. (2025) Structural insights into the biochemical mechanism of the E2/E3 hybrid enzyme UBE2O. *Structure (London, England : 1993)*, 33(2), 274.

McIntyre ABR, et al. (2025) Phosphorylation of a nuclear condensate regulates cohesion and mRNA retention. *Nature communications*, 16(1), 390.

Chakraborty A, et al. (2025) Identification of ABHD6 as a lysophosphatidylserine lipase in the mammalian liver and kidneys. *The Journal of biological chemistry*, 301(2), 108157.

Hein MY, et al. (2025) Global organelle profiling reveals subcellular localization and remodeling at proteome scale. *Cell*, 188(4), 1137.

Chu SN, et al. (2025) Dual β -globin-truncated erythropoietin receptor knockin restores hemoglobin production in β -thalassemia-derived erythroid cells. *Cell reports*, 44(1), 115141.

Lagattuta KA, et al. (2025) The T cell receptor sequence influences the likelihood of T cell memory formation. *Cell reports*, 44(1), 115098.

Zhou B, et al. (2025) IMPDH2 dephosphorylation under FGFR signaling promotes S-phase progression and tumor growth. *Cell reports*, 44(1), 115116.

Khoury Damaa M, et al. (2025) Cyclin O controls entry into the cell-cycle variant required for multiciliated cell differentiation. *Cell reports*, 44(1), 115117.

Wang J, et al. (2025) Salsolinol as an RNA m6A methylation inducer mediates dopaminergic neuronal death by regulating YAP1 and autophagy. *Neural regeneration research*, 20(3), 887.

Melo Garcia L, et al. (2025) Overcoming CD226-related immune evasion in acute myeloid leukemia with CD38 CAR-engineered NK cells. *Cell reports*, 44(1), 115122.

Vallbracht M, et al. (2025) Nucleocapsid assembly drives Ebola viral factory maturation and dispersion. *Cell*, 188(3), 704.

Abelman RO, et al. (2025) TOP1 Mutations and Cross-Resistance to Antibody-Drug Conjugates in Patients with Metastatic Breast Cancer. *Clinical cancer research : an official journal of the American Association for Cancer Research*.

Wang Q, et al. (2025) The nanoscale organization of the Nipah virus fusion protein informs new membrane fusion mechanisms. *eLife*, 13.

Xu SB, et al. (2025) KPNA3 regulates histone locus body formation by modulating condensation and nuclear import of NPAT. *The Journal of cell biology*, 224(1).

Rössler A, et al. (2025) Nonhuman primate antigenic cartography of SARS-CoV-2. *Cell reports*, 44(1), 115140.

Champagne J, et al. (2025) Adoptive T cell therapy targeting an inducible and broadly shared product of aberrant mRNA translation. *Immunity*, 58(1), 247.

Liebau RC, et al. (2024) Transcription-Coupled Repair of DNA Interstrand Crosslinks by UVSSA. *bioRxiv* : the preprint server for biology.