

# Resource Summary Report

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## NCI-H2126

RRID:CVCL\_1532

Type: Cell Line

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### Proper Citation

(RRID:CVCL\_1532)

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### Cell Line Information

**URL:** [https://web.expasy.org/cellosaurus/CVCL\\_1532](https://web.expasy.org/cellosaurus/CVCL_1532)

**Proper Citation:** (RRID:CVCL\_1532)

**Sex:** Male

**Defining Citation:** [PMID:1311061](#), [PMID:8806092](#), [PMID:9559342](#), [PMID:10987304](#), [PMID:11030152](#), [PMID:11314036](#), [PMID:11416159](#), [PMID:14762065](#), [PMID:16157194](#), [PMID:19153074](#), [PMID:19472407](#), [PMID:20164919](#), [PMID:20215515](#), [PMID:20557307](#), [PMID:22460905](#), [PMID:22961666](#), [PMID:25485619](#), [PMID:25877200](#), [PMID:26361996](#), [PMID:26589293](#), [PMID:28196595](#), [PMID:29444439](#), [PMID:29681454](#), [PMID:30038707](#), [PMID:30894373](#), [PMID:31068700](#), [PMID:31803961](#), [PMID:31978347](#)

**Comments:** Omics: Transcriptome analysis by RNAseq., Omics: Transcriptome analysis by microarray., Omics: SNP array analysis., Omics: Protein expression by reverse-phase protein arrays., Omics: DNA methylation analysis., Omics: Deep quantitative proteome analysis., Omics: Deep proteome analysis., Omics: Deep exome analysis., Omics: CNV analysis., Omics: Array-based CGH., Population: Caucasian., Part of: MD Anderson Cell Lines Project., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

**Category:** Cancer cell line

**Name:** NCI-H2126

**Synonyms:** H2126, H-2126, NCIH2126

**Cross References:** BTO:BTO\_0003240, CLO:CLO\_0008056, EFO:EFO\_0002280, ArrayExpress:E-MTAB-38, ArrayExpress:E-MTAB-783, ArrayExpress:E-MTAB-2706, ArrayExpress:E-MTAB-2770, ArrayExpress:E-MTAB-3610, ATCC:CCL-256, ATCC:CRL-

5925, BioGRID\_ORCS\_Cell\_line:530, BioSample:SAMN03471320, BioSample:SAMN03471386, BioSample:SAMN05292434, BioSample:SAMN10988437, cancercellines:CVCL\_1532, CCRID:3101HUMSCSP599, Cell\_Model\_Passport:SIDM00775, ChEMBL-Cells:ChEMBL3308894, ChEMBL-Targets:ChEMBL2366160, CLS:300639, Cosmic:687814, Cosmic:877266, Cosmic:903616, Cosmic:961841, Cosmic:1006537, Cosmic:1146903, Cosmic:1219066, Cosmic:1239892, Cosmic:1995560, Cosmic:2125182, Cosmic:2629443, Cosmic-CLP:687814, DepMap:ACH-000785, EGA:EGAS00001000610, EGA:EGAS00001000978, GDSC:687814, GEO:GSM108839, GEO:GSM108840, GEO:GSM206485, GEO:GSM253405, GEO:GSM274811, GEO:GSM274812, GEO:GSM353228, GEO:GSM434271, GEO:GSM513958, GEO:GSM514343, GEO:GSM784235, GEO:GSM794354, GEO:GSM887408, GEO:GSM888487, GEO:GSM1374724, GEO:GSM1670218, IARC\_TP53:21557, IGRhCellID:NCIH2126, LiGeA:CCLE\_835, LINCS\_LDP:LCL-1627, PharmacDB:NCIH2126\_1071\_2019, PRIDE:PXD002556, Progenetix:CVCL\_1532, PubChem\_Cell\_line:CVCL\_1532, Wikidata:Q54907929

**ID:** CVCL\_1532

**Record Creation Time:** 20250131T201500+0000

**Record Last Update:** 20250131T203139+0000

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## Ratings and Alerts

No rating or validation information has been found for NCI-H2126.

**Warning:** Discontinued: ATCC; CRL-5925

Omics: Transcriptome analysis by RNAseq., Omics: Transcriptome analysis by microarray., Omics: SNP array analysis., Omics: Protein expression by reverse-phase protein arrays., Omics: DNA methylation analysis., Omics: Deep quantitative proteome analysis., Omics: Deep proteome analysis., Omics: Deep exome analysis., Omics: CNV analysis., Omics: Array-based CGH., Population: Caucasian., Part of: MD Anderson Cell Lines Project., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

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## Data and Source Information

**Source:** [Cellosaurus](#)

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## Usage and Citation Metrics

We found 10 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](#).

Horie M, et al. (2024) Exosomes secreted by ST3GAL5high cancer cells promote peritoneal

dissemination by establishing a premetastatic microenvironment. *Molecular oncology*, 18(1), 21.

Pelos G, et al. (2024) Fast proliferating and slowly migrating non-small cell lung cancer cells are vulnerable to decitabine and retinoic acid combinatorial treatment. *International journal of cancer*, 154(6), 1029.

Chen K, et al. (2023) Individualized tumor-informed circulating tumor DNA analysis for postoperative monitoring of non-small cell lung cancer. *Cancer cell*, 41(10), 1749.

Parihar K, et al. (2022) Data driven and biophysical insights into the regulation of trafficking vesicles by extracellular matrix stiffness. *iScience*, 25(8), 104721.

Limagne E, et al. (2022) MEK inhibition overcomes chemoimmunotherapy resistance by inducing CXCL10 in cancer cells. *Cancer cell*, 40(2), 136.

Koppula P, et al. (2021) KEAP1 deficiency drives glucose dependency and sensitizes lung cancer cells and tumors to GLUT inhibition. *iScience*, 24(6), 102649.

Cheng CC, et al. (2020) Sperm-specific COX6B2 enhances oxidative phosphorylation, proliferation, and survival in human lung adenocarcinoma. *eLife*, 9.

Yenerall P, et al. (2020) RUVBL1/RUVBL2 ATPase Activity Drives PAQosome Maturation, DNA Replication and Radioresistance in Lung Cancer. *Cell chemical biology*, 27(1), 105.

Ge P, et al. (2019) miR-762 activation confers acquired resistance to gefitinib in non-small cell lung cancer. *BMC cancer*, 19(1), 1203.

Shirole NH, et al. (2016) TP53 exon-6 truncating mutations produce separation of function isoforms with pro-tumorigenic functions. *eLife*, 5.