## **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 1, 2025

# NCI-H513

RRID:CVCL\_A570 Type: Cell Line

#### **Proper Citation**

(KCLB Cat# 90513, RRID:CVCL\_A570)

#### **Cell Line Information**

URL: https://web.expasy.org/cellosaurus/CVCL\_A570

Proper Citation: (KCLB Cat# 90513, RRID:CVCL\_A570)

Sex: Male

Defining Citation: PMID:8806092, PMID:11030152, PMID:16630136, PMID:23830731, PMID:27397505, PMID:30894373, PMID:31803961, PMID:35839778

**Comments:** Omics: Transcriptome analysis by microarray., Omics: DNA methylation analysis., Omics: Deep quantitative proteome analysis., Omics: Deep exome analysis., Omics: Array-based CGH., Population: African American., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE)., Problematic cell line: Contaminated. Shown to be a NCI-H125 derivative. Originally thought to originate with the pleural effusion of a mesothelioma in a 64 year old male patient..

Category: Cancer cell line

Name: NCI-H513

Synonyms: H513, H-513, NCIH513, HUT513

**Cross References:** ArrayExpress:E-MTAB-3610, ATCC:CRL-5830, BioSample:SAMN03151854, Cell\_Model\_Passport:SIDM00114, Cosmic:877271, Cosmic:877408, Cosmic:980997, Cosmic:1032388, Cosmic:1152503, Cosmic:1995420, Cosmic-CLP:1240141, DepMap:ACH-002138, DepMap:ACH-002341, EGA:EGAS00001000978, GDSC:1240141, GEO:GSM1669835, KCLB:90513, PharmacoDB:H513\_438\_2019, PRIDE:PXD030304, Wikidata:Q54908072 ID: CVCL\_A570

Vendor: KCLB

Catalog Number: 90513

Record Creation Time: 20250131T201503+0000

Record Last Update: 20250131T203143+0000

#### **Ratings and Alerts**

No rating or validation information has been found for NCI-H513.

Warning: Problematic cell line: Contaminated. Shown to be a NCI-H125 derivative. Originally thought to originate with the pleural effusion of a mesothelioma in a 64 year old male patient.

Registration: International Cell Line Authentication Committee, Register of Misidentified Cell Lines; ICLAC-00414.

Omics: Transcriptome analysis by microarray., Omics: DNA methylation analysis., Omics: Deep quantitative proteome analysis., Omics: Deep exome analysis., Omics: Array-based CGH., Population: African American., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE)., Problematic cell line: Contaminated. Shown to be a NCI-H125 derivative. Originally thought to originate with the pleural effusion of a mesothelioma in a 64 year old male patient.. Warning: Discontinued: ATCC; CRL-5830

Omics: Transcriptome analysis by microarray., Omics: DNA methylation analysis., Omics: Deep quantitative proteome analysis., Omics: Deep exome analysis., Omics: Array-based CGH., Population: African American., Part of: COSMIC cell lines project., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE)., Problematic cell line: Contaminated. Shown to be a NCI-H125 derivative. Originally thought to originate with the pleural effusion of a mesothelioma in a 64 year old male patient..

### Data and Source Information

Source: Cellosaurus

#### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

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

Kolluri KK, et al. (2018) Loss of functional BAP1 augments sensitivity to TRAIL in cancer cells. eLife, 7.