

# Resource Summary Report

Generated by [dkNET](#) on Sep 3, 2026

## HCC461

RRID:CVCL\_5135

Type: Cell Line

### Proper Citation

RRID:CVCL\_5135

### Cell Line Information

**URL:** [https://web.expasy.org/cellosaurus/CVCL\\_5135](https://web.expasy.org/cellosaurus/CVCL_5135)

**Proper Citation:** RRID:CVCL\_5135

**Description:** Cell line HCC461 is a Cancer cell line with a species of origin Homo sapiens (Human)

**Sex:** Male

**Disease:** Lung adenocarcinoma

**Defining Citation:** [PMID:10353731](#), [PMID:16187286](#), [PMID:20679594](#), [PMID:22961666](#), [PMID:25485619](#), [PMID:25877200](#), [PMID:26589293](#), [PMID:29681454](#), [PMID:31068700](#), [PMID:31803961](#)

**Comments:** Miscellaneous: Direct author submission of STR profile from Girard, Luc & Gazdar, Adi F., Population: Caucasian., Part of: Cancer Dependency Map project (DepMap) (includes Cancer Cell Line Encyclopedia - CCLE).

**Category:** Cancer cell line

**Organism:** Homo sapiens (Human)

**Name:** HCC461

**Synonyms:** HCC0461, Hamon Cancer Center 461

**Cross References:** EFO:EFO\_0003134, ArrayExpress:E-MTAB-2706, BioSample:SAMN03471006, BioSample:SAMN10989603, cancercellines:CVCL\_5135, Cell\_Model\_Passport:SIDM01595, Cosmic:903632, Cosmic:1146938, Cosmic:2699648, DepMap:ACH-000339, EGA:EGAS00001000610, GEO:GSM62336, GEO:GSM108877, GEO:GSM108878, GEO:GSM253392, GEO:GSM434262, GEO:GSM794380, GEO:GSM817115, IARC\_TP53:11446, IGRhCellID:HCC461GEO, PharmacDB:HCC461\_505\_2019, Progenetix:CVCL\_5135, Wikidata:Q54881757

**ID:** CVCL\_5135

**Record Creation Time:** 20220427T220319+0000

**Record Last Update:** 20260904T013921+0000

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## Ratings and Alerts

No rating or validation information has been found for HCC461.

No alerts have been found for HCC461.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Watanabe H, et al. (2013) Integrated cistronic and expression analysis of amplified NKX2-1 in lung adenocarcinoma identifies LMO3 as a functional transcriptional target. Genes & development, 27(2), 197.