

Resource Summary Report

Generated by [dkNET](#) on Sep 1, 2026

GM06990

RRID:CVCL_9587

Type: Cell Line

Proper Citation

RRID:CVCL_9587

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_9587

Proper Citation: RRID:CVCL_9587

Description: Cell line GM06990 is a Transformed cell line with a species of origin Homo sapiens (Human)

Sex: Female

Defining Citation: [PMID:11416159](#), [PMID:20215515](#), [PMID:20856902](#), [PMID:23861383](#)

Comments: Part of: International Histocompatibility Workshop cell lines., Part of: ENCODE project common cell types; tier 3., Part of: CEPH/Utah pedigree cell line collection.

Category: Transformed cell line

Organism: Homo sapiens (Human)

Name: GM06990

Synonyms: GM06990A, CEPH-1331-NA06990, 1331-8233

Cross References: CLO:CLO_0036556, EFO:EFO_0002783, BioSample:SAMN00004982, Coriell:GM06990, dbMHC:48538, ENCODE:ENCBS057ENC, ENCODE:ENCBS058ENC, ENCODE:ENCBS059ENC, ENCODE:ENCBS060ENC, ENCODE:ENCBS061ENC, ENCODE:ENCBS062ENC, ENCODE:ENCBS063ENC, ENCODE:ENCBS064ENC, ENCODE:ENCBS146QHV, ENCODE:ENCBS177AAA, ENCODE:ENCBS185XLT, ENCODE:ENCBS414HDG, ENCODE:ENCBS875KTK, ENCODE:ENCBS987SKT, GEO:GSE38809, GEO:GSM207754, GEO:GSM420468, GEO:GSM472903, GEO:GSM472904, GEO:GSM660029, GEO:GSM660236, GEO:GSM660440, GEO:GSM736558, GEO:GSM736635, GEO:GSM749705, GEO:GSM749708, GEO:GSM749731,

GEO:GSM827333, GEO:GSM905817, GEO:GSM905912, GEO:GSM906007, GEO:GSM923443, GEO:GSM931323, GEO:GSM931324, GEO:GSM945179, GEO:GSM945205, GEO:GSM945213, GEO:GSM945237, IHW:IHW01003, IPD-IMGT/HLA:25804, Wikidata:Q54842431

ID: CVCL_9587

Record Creation Time: 20220427T215951+0000

Record Last Update: 20260829T232857+0000

Ratings and Alerts

No rating or validation information has been found for GM06990.

No alerts have been found for GM06990.

Data and Source Information

Source: [CelloSaurus](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Wu X, et al. (2023) Genome-wide measurement of DNA replication fork directionality and quantification of DNA replication initiation and termination with Okazaki fragment sequencing. Nature protocols, 18(4), 1260.

Joshi I, et al. (2021) Genome-wide mapping of DNA double-strand breaks from eukaryotic cell cultures using Break-seq. STAR protocols, 2(2), 100554.

Chakraborty A, et al. (2020) Replication Stress Induces Global Chromosome Breakage in the Fragile X Genome. Cell reports, 32(12), 108179.