

Resource Summary Report

Generated by [RRID](#) on Sep 6, 2026

GM00639

RRID:CVCL_7299

Type: Cell Line

Proper Citation

RRID:CVCL_7299

Cell Line Information

URL: https://web.expasy.org/cellosaurus/CVCL_7299

Proper Citation: RRID:CVCL_7299

Description: Cell line GM00639 is a Transformed cell line with a species of origin Homo sapiens (Human)

Sex: Male

Disease: Galactosemia

Defining Citation: [PMID:191830](#), [PMID:1766867](#), [PMID:2011574](#), [PMID:2762303](#), [PMID:3413074](#), [PMID:9175740](#), [PMID:10606667](#), [PMID:11416159](#)

Comments: Population: African American.

Category: Transformed cell line

Organism: Homo sapiens (Human)

Name: GM00639

Synonyms: GM-639, GM0639, GM 639, GM639, GM52VA

Cross References: CLO:CLO_0003516, CLDB:cl1483, Coriell:GM00639, Wikidata:Q54836366

ID: CVCL_7299

Hierarchy: cvcl_2779

Record Creation Time: 20220427T215908+0000

Record Last Update: 20260905T175409+0000

Ratings and Alerts

No rating or validation information has been found for GM00639.

Warning: Discontinued: Coriell; GM00639
Population: African American.

Data and Source Information

Source: [Cellosaurus](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Lazarchuk P, et al. (2024) Werner syndrome RECQ helicase participates in and directs maintenance of the protein complexes of constitutive heterochromatin in proliferating human cells. *Aging*, 16(20), 12977.

Kovacs MT, et al. (2023) DNA damage induces nuclear envelope rupture through ATR-mediated phosphorylation of lamin A/C. *Molecular cell*, 83(20), 3659.

Shukla S, et al. (2022) Genome-wide survey of D/E repeats in human proteins uncovers their instability and aids in identifying their role in the chromatin regulator ATAD2. *iScience*, 25(12), 105464.

Lazarchuk P, et al. (2022) Innate immunity mediator STING modulates nascent DNA metabolism at stalled forks in human cells. *Frontiers in molecular biosciences*, 9, 1048726.

Nitiss JL, et al. (2021) Topoisomerase Assays. *Current protocols*, 1(10), e250.