

Resource Summary Report

Generated by [PRECISE-TBI](#) on Jul 29, 2026

Find My Understudied Genes

RRID:SCR_025047

Type: Tool

Proper Citation

Find My Understudied Genes (RRID:SCR_025047)

Resource Information

URL: <https://fmug.amaral.northwestern.edu/>

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Description: Software data-driven tool to identify understudied genes and characterize their tractability. Users submit list of human genes and can filter these genes down based on list of factors. Code to generate Find My Understudied Genes app for Windows, iOS and macOS platforms.

Abbreviations: FMUG

Resource Type: software application, source code, software resource

Defining Citation: [DOI:10.7554/eLife.93429](https://doi.org/10.7554/eLife.93429)

Funding: NIGMS T32GM008449;
Northwestern University ;
Moderna Inc ;
NSF ;
NAIAD U19AI135964;
Simons Foundation ;
NIA K99AG068544

Availability: Free, Available for download, Freely available

Resource Name: Find My Understudied Genes

Resource ID: SCR_025047

Alternate URLs: <https://github.com/amarallab/fmug>

Record Creation Time: 20240305T200904+0000

Record Last Update: 20260729T044614+0000

Ratings and Alerts

No rating or validation information has been found for Find My Understudied Genes .

No alerts have been found for Find My Understudied Genes .

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Rodemann MM, et al. (2025) Identification of a Growth-Promoting Gene Cluster in the Region 2q24 as a Driver of Tumorigenesis in Childhood Hepatoblastoma. The American journal of pathology, 195(8), 1553.

Richardson R, et al. (2024) Meta-Research: Understudied genes are lost in a leaky pipeline between genome-wide assays and reporting of results. eLife, 12.