

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Pseudogene

RRID:SCR_010453

Type: Tool

Proper Citation

Pseudogene (RRID:SCR_010453)

Resource Information

URL: <http://purl.bioontology.org/ontology/pseudo>

Proper Citation: Pseudogene (RRID:SCR_010453)

Description: Ontology for pseudogenes including biological feature, classified type, Evidence code, and subcellular origin.

Abbreviations: pseudo

Resource Type: controlled vocabulary, data or information resource, ontology

Keywords: obo

Resource Name: Pseudogene

Resource ID: SCR_010453

Alternate IDs: nlx_157640

Record Creation Time: 20220129T080258+0000

Record Last Update: 20260905T132642+0000

Ratings and Alerts

No rating or validation information has been found for Pseudogene.

No alerts have been found for Pseudogene.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 141 mentions in open access literature.

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

Selem SI, et al. (2025) SOFCs integrated with SMES under dynamic power control using Chernobyl disaster optimizer. *Scientific reports*, 15(1), 3120.

Svaricek R, et al. (2025) INSIGHT: Combining Fixation Visualisations and Residual Neural Networks for Dyslexia Classification From Eye-Tracking Data. *Dyslexia* (Chichester, England), 31(1), e1801.

Husnoo N, et al. (2025) The patient and clinician perspective on 'early' bowel resection for terminal ileal Crohn's disease (EBRIC): protocol for a multicentre mixed-methods study. *Colorectal disease : the official journal of the Association of Coloproctology of Great Britain and Ireland*, 27(3), e70042.

Ghulam A, et al. (2025) StackAHTPs: An explainable antihypertensive peptides identifier based on heterogeneous features and stacked learning approach. *IET systems biology*, 19(1), e70002.

Usai G, et al. (2025) Haplotype-resolved genome assembly of *Ficus carica* L. reveals allele-specific expression in the fruit. *The Plant journal : for cell and molecular biology*, 121(4), e70012.

Komosny D, et al. (2025) Synthetic Browsing Histories for 50 Countries Worldwide: Datasets for Research, Development, and Education. *Scientific data*, 12(1), 128.

Ruiz-Ruano FJ, et al. (2025) Programmed DNA elimination drives rapid genomic innovation in two thirds of all bird species. *bioRxiv : the preprint server for biology*.

Schwandner IA, et al. (2025) Predicting the impact of targeted fence removal on connectivity in a migratory ecosystem. *Ecological applications : a publication of the Ecological Society of America*, 35(1), e3094.

Zhang L, et al. (2025) Nicotinamide Phosphoribosyltransferase Acetylation Mediating Muscle Dysfunction Contributes to Sleep Apnoea in Obesity. *Journal of cachexia, sarcopenia and muscle*, 16(1), e13693.

Galeote-Checa G, et al. (2025) Time series segmentation for recognition of epileptiform patterns recorded via microelectrode arrays in vitro. *PloS one*, 20(1), e0309550.

Nath R, et al. (2025) Lineage-Specific Class-A GPCR Dynamics Reflect Diverse Chemosensory Adaptations in Lophotrochozoa. *Molecular biology and evolution*, 42(3).

Rostamian R, et al. (2025) A deep learning-based multi-view approach to automatic 3D landmarking and deformity assessment of lower limb. *Scientific reports*, 15(1), 534.

Chatterjee A, et al. (2024) Exploring online public survey lifestyle datasets with statistical analysis, machine learning and semantic ontology. *Scientific reports*, 14(1), 24190.

Vorstenbosch T, et al. (2024) Venturing Into the Unknown: The Importance of Variable Selection When Modelling Alien Species Under Non-Analogue Climatic Conditions. *Ecology and evolution*, 14(10), e70490.

Jayaprasad S, et al. (2024) Orthopteran Neo-Sex Chromosomes Reveal Dynamics of Recombination Suppression and Evolution of Supergenes. *Molecular ecology*, 33(23), e17567.

Azzam SM, et al. (2024) An improved Differential evolution with Sailfish optimizer (DESFO) for handling feature selection problem. *Scientific reports*, 14(1), 13517.

Chen G, et al. (2024) A lightweight Color-changing melon ripeness detection algorithm based on model pruning and knowledge distillation: leveraging dilated residual and multi-screening path aggregation. *Frontiers in plant science*, 15, 1406593.

Zhao J, et al. (2024) Metabolite-based genome-wide association studies enable the dissection of the genetic bases of flavonoids, betaine and spermidine in wolfberry (*Lycium*). *Plant biotechnology journal*, 22(6), 1435.

Chen R, et al. (2024) SpiDe-Sr: blind super-resolution network for precise cell segmentation and clustering in spatial proteomics imaging. *Nature communications*, 15(1), 2708.

Huang X, et al. (2024) A neuromorphic dataset for tabletop object segmentation in indoor cluttered environment. *Scientific data*, 11(1), 127.