Resource Summary Report

Generated by dkNET on Sep 22, 2026

RefSeqGene

RRID:SCR_013787
Type: Tool

Proper Citation

RefSeqGene (RRID:SCR_013787)

Resource Information

URL: http://www.ncbi.nlm.nih.gov/refseq/rsg/

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Description: A data set of defined genomic sequences used as reference standards for well-characterized genes. These standard nucleotide sequences serve as foundations for locating mutations, establishing conventions for numbering exons and introns, and defining the coordinates of other variations. Sequences are aligned to reference chromosomes. RefSeqGene is a subset of NCBI RefSeq.

Synonyms: NCBI RefSeqGene, RefSeqGene Project

Resource Type: information resource

Defining Citation: PMID:18927115

Keywords: data set, genomic sequences, reference standard

Availability: Free, Public

Resource Name: RefSeqGene

Resource ID: SCR_013787

License URLs: http://www.nlm.nih.gov/privacy.html

Record Creation Time: 20220129T080318+0000

Record Last Update: 20260919T195248+0000

Ratings and Alerts

No rating or validation information has been found for RefSeqGene.

No alerts have been found for RefSeqGene.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 56 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Campbell AS, et al. (2026) Molecular insights into electroreceptor ribbon synapses from differential gene expression in sturgeon lateral line organs. Journal of anatomy, 248(5), 784.

Shostak K, et al. (2026) The lipid transfer protein STARD7 controls intestinal tumor development in a context-dependent manner. EMBO molecular medicine, 18(5), 1771.

Betti M, et al. (2026) Biopsy RNA-seq captures TROP-2-linked migration and clonal resistance to forecast aggressiveness in metastatic melanoma. Journal of experimental & clinical cancer research : CR, 45(1).

DiPietro EE, et al. (2025) Genetics of ovulatory dysfunction and infertility: a scoping review and gene ontology analysis. Frontiers in endocrinology, 16, 1458711.

Chen HH, et al. (2025) Population-specific polygenic risk scores for people of Han Chinese ancestry. Nature, 648(8092), 128.

Burykina Y, et al. (2025) A Unique Case of a Child with Two Rare Hereditary Diseases: Familial Dilated Cardiomyopathy and Arterial Calcification. International journal of molecular sciences, 26(12).

Huang X, et al. (2025) Mutation spectra and genotype?phenotype analysis of congenital hypothyroidism in a neonatal population. Biomedical reports, 22(2), 30.

Lv W, et al. (2024) Extrachromosomal circular DNA orchestrates genome heterogeneity in urothelial bladder carcinoma. Theranostics, 14(13), 5102.

Pugacheva EM, et al. (2024) BORIS/CTCFL epigenetically reprograms clustered CTCF binding sites into alternative transcriptional start sites. Genome biology, 25(1), 40.

Pranck?nien? L, et al. (2023) Microevolutionary processes analysis in the Lithuanian genome. Scientific reports, 13(1), 11941.

Chou KJ, et al. (2022) A new missense mutation of calcium sensing receptor with isoleucine replaced by serine at codon 857 leading to type V Bartter syndrome. Experimental cell research, 414(1), 113080.

Markova TV, et al. (2022) Clinical and genetic characterization of three Russian patients with pycnodysostosis due to pathogenic variants in the CTSK gene. Molecular genetics &

genomic medicine, 10(5), e1904.

Ikeda S, et al. (2022) Disruption of piRNA machinery by deletion of ASZ1/GASZ results in the expression of aberrant chimeric transcripts in gonocytes. The Journal of reproduction and development, 68(2), 125.

Ziadi W, et al. (2021) STAT3 polymorphisms in North Africa and its implication in breast cancer. Molecular genetics & genomic medicine, 9(8), e1744.

Urnikyte A, et al. (2021) Genome-Wide Landscape of North-Eastern European Populations: A View from Lithuania. Genes, 12(11).

Erdo?an M, et al. (2021) The Genetic Analysis of Cystic Fibrosis Patients With Seven Novel Mutations in the CFTR Gene in the Central Anatolian Region of Turkey. Balkan medical journal, 38(6), 357.

Markova T, et al. (2021) Clinical and genetic characterization of autosomal recessive stickler syndrome caused by novel compound heterozygous mutations in the COL9A3 gene. Molecular genetics & genomic medicine, 9(3), e1620.

Penitenti F, et al. (2021) Clinical presentation, genotype-phenotype correlations, and outcome of pancreatic neuroendocrine tumors in Von Hippel-Lindau syndrome. Endocrine, 74(1), 180.

Liu W, et al. (2021) Analysis of STAG3 variants in Chinese non-obstructive azoospermia patients with germ cell maturation arrest. Scientific reports, 11(1), 10077.

Wu CY, et al. (2021) Lipopolysaccharide stimulation test on cultured PBMCs assists the discrimination of cryopyrin-associated periodic syndrome from systemic juvenile idiopathic arthritis. Scientific reports, 11(1), 11903.