Nucleotide database

RRID:SCR_004630
Type: Tool

Proper Citation

Nucleotide database (RRID:SCR_004630)

Resource Information


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Description: Nucleotide database as collection of sequences from several sources, including GenBank, RefSeq, TPA and PDB. Genome, gene and transcript sequence data provide the foundation for biomedical research and discovery.

Abbreviations: nucest

Resource Type: storage service resource, database, service resource, data or information resource, data repository

Defining Citation: PMID:8401577

Keywords: Genome, gene, transcript sequence data, GenBank, RefSeq, TPA, PDB, gold standard

Availability: Free, Freely available

Resource Name: Nucleotide database

Resource ID: SCR_004630

Alternate IDs: nlx_62971, SCR_016578, SCR_016578

Ratings and Alerts
No rating or validation information has been found for Nucleotide database.

No alerts have been found for Nucleotide database.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 265 mentions in open access literature.

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


Zhao P, et al. (2021) Mining Unknown Porcine Protein Isoforms by Tissue-Based Map of Proteome Enhances the Pig Genome Annotation. Genomics, proteomics & bioinformatics.


Shen WW, et al. (2020) EHD2 is a Predictive Biomarker of Chemotherapy Efficacy in Triple Negative Breast Carcinoma. Scientific reports, 10(1), 7998.


Mathew R, et al. (2019) The genome of the migratory nematode, Radopholus similis, reveals
signatures of close association to the sedentary cyst nematodes. PloS one, 14(10), e0224391.


Tran HTM, et al. (2018) Use of a draft genome of coffee (Coffea arabica) to identify SNPs associated with caffeine content. Plant biotechnology journal, 16(10), 1756-1766.

