## **Resource Summary Report**

Generated by <u>dkNET</u> on May 20, 2025

# NAT/NCS2 Hound

RRID:SCR\_016473 Type: Tool

### **Proper Citation**

NAT/NCS2 Hound (RRID:SCR\_016473)

## **Resource Information**

URL: http://bioinf.bio.uth.gr/nat-ncs2

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**Description:** Web server for the detection and evolutionary classification of prokaryotic and eukaryotic nucleobase-cation symporters of the NAT/NCS2 family. Used to scan, identify and evolutionary classify NAT/NCS2 nucleobase transporter proteins.

Abbreviations: NAT, NCS2

Synonyms: Nucleobase Ascorbate Transporter, NCS2:Nucleobase Cation Symporter 2

**Resource Type:** software application, web application, data analysis software, data processing software, sequence analysis software, software resource

Defining Citation: DOI:10.1101/332452

**Keywords:** protein, sequence, scan, identify, evolutionary, classify, prokaryotic, nucleobase, transporter, protein, amino acid, conserved

#### Funding:

Availability: Free, Available to download, Freely available

Resource Name: NAT/NCS2 Hound

Resource ID: SCR\_016473

**Record Creation Time:** 20220129T080330+0000

## **Ratings and Alerts**

No rating or validation information has been found for NAT/NCS2 Hound.

No alerts have been found for NAT/NCS2 Hound.

## Data and Source Information

Source: <u>SciCrunch Registry</u>

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

Chaliotis A, et al. (2018) NAT/NCS2-hound: a webserver for the detection and evolutionary classification of prokaryotic and eukaryotic nucleobase-cation symporters of the NAT/NCS2 family. GigaScience, 7(12).