

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

NetAge Database

RRID:SCR_010224

Type: Tool

Proper Citation

NetAge Database (RRID:SCR_010224)

Resource Information

URL: <http://netage-project.org/>

Proper Citation: NetAge Database (RRID:SCR_010224)

Description: Database that contains gene sets and microRNA-regulated protein-protein interaction networks for longevity, age-related diseases and aging-associated processes.

Abbreviations: NetAge

Resource Type: data or information resource, database

Defining Citation: [PMID:20186480](#)

Keywords: longevity, protein-protein interaction, mirna-regulated protein-protein interaction network, gene, protein, mirna, biogerontology

Related Condition: Age-related disease, Aging

Availability: Acknowledgement requested, Public

Resource Name: NetAge Database

Resource ID: SCR_010224

Alternate IDs: nlx_156769

Record Creation Time: 20220129T080257+0000

Record Last Update: 20260905T133156+0000

Ratings and Alerts

No rating or validation information has been found for NetAge Database.

No alerts have been found for NetAge Database.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Talwar P, et al. (2019) Viral Induced Oxidative and Inflammatory Response in Alzheimer's Disease Pathogenesis with Identification of Potential Drug Candidates: A Systematic Review using Systems Biology Approach. Current neuropharmacology, 17(4), 352.

Tacutu R, et al. (2018) Human Ageing Genomic Resources: new and updated databases. Nucleic acids research, 46(D1), D1083.

O'Connor JE, et al. (2014) Systems Biology and immune aging. Immunology letters, 162(1 Pt B), 334.

Ayyadevara S, et al. (2014) Rec-8 dimorphism affects longevity, stress resistance and X-chromosome nondisjunction in C. elegans, and replicative lifespan in S. cerevisiae. Frontiers in genetics, 5, 211.

Zhu W, et al. (2014) Computational developments in microRNA-regulated protein-protein interactions. BMC systems biology, 8, 14.

Kubrycht J, et al. (2012) Virtual interactomics of proteins from biochemical standpoint. Molecular biology international, 2012, 976385.