

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 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: database, data or information resource

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: 20260725T191158+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 [nidm-terms](#) .

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.