

# Resource Summary Report

Generated by [RRID](#) on Aug 16, 2026

## Knut and Alice Wallenberg Foundation

RRID:SCR\_004778

Type: Tool

### Proper Citation

Knut and Alice Wallenberg Foundation (RRID:SCR\_004778)

### Resource Information

**URL:** <http://www.wallenberg.com/kaw/en>

**Proper Citation:** Knut and Alice Wallenberg Foundation (RRID:SCR\_004778)

**Description:** One of the largest financiers of research in Sweden that promotes scientific research, teaching and/or education beneficial to the Kingdom of Sweden. 20-30 percent of the Foundation's grants relate to high-value research equipment for use within the technical, natural sciences and biomedical fields, and about 50 percent relate to major research projects and substantial scientific networks. Universities, academies and similar research and educational institutions are eligible for grants within the following fields: \* Expensive scientific equipment \* Fellowship programs initiated by the Foundation \* Selected research projects of significant potential \* Larger educational research projects Applications may be submitted by individual scientists, or research groups affiliated to a Swedish university or equivalent institution. An application may also be submitted by the management of a university, university college, scientific academy, or national research associations. Applications from private individuals without connection to a scientific institution are not considered.

**Abbreviations:** Wallenberg Foundation

**Synonyms:** Knut och Alice Wallenbergs Stiftelse

**Resource Type:** institution

**Keywords:** technical, natural science, biomedical, scientific network, grant, research equipment, education

**Resource Name:** Knut and Alice Wallenberg Foundation

**Resource ID:** SCR\_004778

**Alternate IDs:** grid.453057.0, Crossref funder ID: 501100004063, nlx\_77359

**Alternate URLs:** <https://ror.org/004hzzk67>

**Old URLs:** [http://www.wallenberg.com/kaw/in\\_english/](http://www.wallenberg.com/kaw/in_english/)

**Record Creation Time:** 20220129T080226+0000

**Record Last Update:** 20260815T182304+0000

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## Ratings and Alerts

No rating or validation information has been found for Knut and Alice Wallenberg Foundation.

No alerts have been found for Knut and Alice Wallenberg Foundation.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 12 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](#) .

P??verlein MC, et al. (2026) Protein-induced membrane strain drives supercomplex formation. eLife, 13.

Wang JC, et al. (2026) Nociceptor neurons control pollution-mediated neutrophilic asthma. eLife, 13.

Domingo J, et al. (2026) Nonlinear transcriptional responses to gradual modulation of transcription factor dosage. eLife, 13.

Lenz T, et al. (2026) Epigenetics and chromatin structure regulate var2csa expression and the placental-binding phenotype in Plasmodium falciparum. eLife, 13.

Balodis M, et al. (2026) 1H-1H Interatomic Distances in Paracetamol-Based Structures Unveiled by Double-Quantum NMR and DFT Calculations. Molecules (Basel, Switzerland), 31(10).

Zhao Q, et al. (2026) Global molecular landscape of early MASLD progression in human obesity. eLife, 14.

Puno A, et al. (2026) Adrenomedullin restores the human cortical interneurons migration defects induced by hypoxia. eLife, 14.

Lallemend F, et al. (2025) Functional and molecular insights into muscle proprioceptors. eLife, 14.

Habuka M, et al. (2015) The Urinary Bladder Transcriptome and Proteome Defined by Transcriptomics and Antibody-Based Profiling. PloS one, 10(12), e0145301.

Habuka M, et al. (2014) The kidney transcriptome and proteome defined by transcriptomics and antibody-based profiling. PloS one, 9(12), e116125.

Song J, et al. (2014) Essential genetic interactors of SIR2 required for spatial sequestration and asymmetrical inheritance of protein aggregates. PLoS genetics, 10(7), e1004539.

Lampa S, et al. (2013) Lessons learned from implementing a national infrastructure in Sweden for storage and analysis of next-generation sequencing data. GigaScience, 2(1), 9.