

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

NAPPA

RRID:SCR_003419

Type: Tool

Proper Citation

NAPPA (RRID:SCR_003419)

Resource Information

URL: <http://cran.r-project.org/web/packages/NAPPA/>

Proper Citation: NAPPA (RRID:SCR_003419)

Description: Software that enables the processing and normalization of the standard mRNA data output from the Nanostring nCounter software.

Abbreviations: NAPPA

Synonyms: NAPPA: Performs the processing and normalisation of Nanostring miRNA and mRNA data

Resource Type: software resource

Keywords: normalization, processing, nanostring, mirna, mrna, os x, windows

Availability: GNU General Public License, v3

Resource Name: NAPPA

Resource ID: SCR_003419

Alternate IDs: OMICS_02310

Record Creation Time: 20220129T080218+0000

Record Last Update: 20260725T190541+0000

Ratings and Alerts

No rating or validation information has been found for NAPPA.

No alerts have been found for NAPPA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Liang T, et al. (2024) Antibody Profiling of Pan-Cancer Viral Proteome Reveals Biomarkers for Nasopharyngeal Carcinoma Diagnosis and Prognosis. *Molecular & cellular proteomics* : MCP, 23(3), 100729.

O'Kell AL, et al. (2022) Exploration of autoantibody responses in canine diabetes using protein arrays. *Scientific reports*, 12(1), 2490.

Mathilakathu A, et al. (2021) Mitogen signal-associated pathways, energy metabolism regulation, and mediation of tumor immunogenicity play essential roles in the cellular response of malignant pleural mesotheliomas to platinum-based treatment: a retrospective study. *Translational lung cancer research*, 10(7), 3030.

Brcic L, et al. (2021) Digital Gene Expression Analysis of Epithelioid and Sarcomatoid Mesothelioma Reveals Differences in Immunogenicity. *Cancers*, 13(8).

Cho S, et al. (2021) Automated analysis of lexical features in frontotemporal degeneration. *Cortex; a journal devoted to the study of the nervous system and behavior*, 137, 215.

Cho S, et al. (2020) Automated analysis of lexical features in Frontotemporal Degeneration. *medRxiv : the preprint server for health sciences*.

Nevler N, et al. (2019) Validated automatic speech biomarkers in primary progressive aphasia. *Annals of clinical and translational neurology*, 6(1), 4.

Gillen J, et al. (2019) Experimental Analysis of Viral-Host Interactions. *Frontiers in physiology*, 10, 425.

Wu C, et al. (2019) Prognostic Immune Cell Profiling of Malignant Pleural Effusion Patients by Computerized Immunohistochemical and Transcriptional Analysis. *Cancers*, 11(12).

Arévalo-Pinzón G, et al. (2018) Self-assembling functional programmable protein array for studying protein-protein interactions in malaria parasites. *Malaria journal*, 17(1), 270.

Yu X, et al. (2017) Multiplexed Nucleic Acid Programmable Protein Arrays. *Theranostics*, 7(16), 4057.

Gervits F, et al. (2016) Transcranial direct current stimulation for the treatment of primary progressive aphasia: An open-label pilot study. *Brain and language*, 162, 35.