

Resource Summary Report

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NGSCheckMate

RRID:SCR_022994

Type: Tool

Proper Citation

NGSCheckMate (RRID:SCR_022994)

Resource Information

URL: <https://github.com/parklab/NGSCheckMate>

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Description: Software package for validating sample identity in next generation sequencing studies within and across data types. Used for identifying next generation sequencing data files from the same individual. Used for checking sample matching for NGS data.

Resource Type: software resource, software toolkit

Defining Citation: [PMID:28369524](https://pubmed.ncbi.nlm.nih.gov/28369524/)

Keywords: Next Generation sequencing data file, identifying next generation sequencing data files, next generation sequencing, same individual data files, checking sample matching, NGS data.

Funding: Harvard Medical School Eleanor and Miles Shore Fellowship ;
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NIA K01AG051791;
NIMH 1P50MH106933;
NIMH 1U01MH106883;
NEI R01EY024230;
Korean Health Technology

Availability: Free, Available for download, Freely available

Resource Name: NGSCheckMate

Resource ID: SCR_022994

License: MIT license

Record Creation Time: 20221124T050201+0000

Record Last Update: 20250429T060219+0000

Ratings and Alerts

No rating or validation information has been found for NGSCheckMate.

No alerts have been found for NGSCheckMate.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Alcala N, et al. (2024) Multi-omic dataset of patient-derived tumor organoids of neuroendocrine neoplasms. GigaScience, 13.

Di Genova A, et al. (2022) A molecular phenotypic map of malignant pleural mesothelioma. GigaScience, 12.

Grant RA, et al. (2021) Circuits between infected macrophages and T cells in SARS-CoV-2 pneumonia. Nature, 590(7847), 635.

Amin SB, et al. (2020) Comparative Molecular Life History of Spontaneous Canine and Human Gliomas. Cancer cell, 37(2), 243.