

Resource Summary Report

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HTS Mappers

RRID:SCR_011991

Type: Tool

Proper Citation

HTS Mappers (RRID:SCR_011991)

Resource Information

URL: http://wwwdev.ebi.ac.uk/fg/hts_mappers/

Proper Citation: HTS Mappers (RRID:SCR_011991)

Description: Data set providing an up-to-date compendium of HTS mappers initially provided in the article Tools for mapping high-throughput sequencing data. Please let fill out the provided form if you are a developer and your mapper is not listed or you want to update the data for your mapper.

Abbreviations: HTS Mappers

Resource Type: software repository, software resource, data or information resource, data set

Defining Citation: [PMID:23060614](#)

Keywords: mapping, high-throughput sequencing, dna, rna, mirna, bisulfite, mapper

Resource Name: HTS Mappers

Resource ID: SCR_011991

Alternate IDs: OMICS_01733

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260728T043400+0000

Ratings and Alerts

No rating or validation information has been found for HTS Mappers.

No alerts have been found for HTS Mappers.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Oppenheim SJ, et al. (2015) We can't all be supermodels: the value of comparative transcriptomics to the study of non-model insects. *Insect molecular biology*, 24(2), 139.

Li JW, et al. (2013) The NGS WikiBook: a dynamic collaborative online training effort with long-term sustainability. *Briefings in bioinformatics*, 14(5), 548.