MetaPhlAn
RRID:SCR_004915
Type: Tool

Proper Citation

MetaPhlAn (RRID:SCR_004915)

Resource Information

URL: http://huttenhower.sph.harvard.edu/metaphlan2

Description: Computational tool for profiling the composition of microbial communities from metagenomic shotgun sequencing data. It relies on unique clade-specific marker genes identified from reference genomes.

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Resource Type: Resource, data analysis software, data processing software, software application, sequence analysis software, software resource, data analysis resource

Keywords: microbial community, microbial composition, microbial genome, microbial sequence

Resource ID: SCR_004915

Parent Organization: Bitbucket, Harvard T.H. Chan School of Public Health

References: PMID:22688413

Availability: Free, Public, Acknowledgement requested

Website Status: Last checked up

Alternate IDs: OMICS_02286
Ratings and Alerts

No rating or validation information has been found for MetaPhlAn.

No alerts have been found for MetaPhlAn.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 158 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch Infrastructure.


Shi X, et al. (2019) Microfluidics-Based Enrichment and Whole-Genome Amplification Enable
Strain-Level Resolution for Airway Metagenomics. mSystems, 4(4).


