Resource Summary Report

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QIIME 2 View

RRID:SCR_018074
Type: Tool

Proper Citation

QIIME 2 View (RRID:SCR_018074)

Resource Information

URL: https://view.qiime2.org/

Proper Citation: QIIME 2 View (RRID:SCR_018074)

Description: Web based serverless viewer of QIIME 2 artifacts and visualizations. Client side interface for viewing QIIME 2 artifacts and visualizations. Not needed working QIIME 2 installation to inspect QIIME 2 results. Supports viewing externally hosted files by automatically downloading and displaying them when links to files are provided.

Synonyms: q2view

Resource Type: data processing software, service resource, data visualization software, data acquisition software, software resource, software application

Keywords: QIIME 2, visualization, data, interface, file viewing, file download and displaying

Funding:

Availability: Free, Freely available

Resource Name: QIIME 2 View

Resource ID: SCR_018074

Alternate URLs: https://github.com/qiime2/q2view

License: BSD 3-Clause "New" or "Revised" License

Record Creation Time: 20220129T080338+0000

Record Last Update: 20250429T055941+0000

Ratings and Alerts

No rating or validation information has been found for QIIME 2 View.

No alerts have been found for QIIME 2 View.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 57 mentions in open access literature.

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

Guan W, et al. (2025) Signature of pre-pregnancy microbiome in infertile women undergoing frozen embryo transfer with gestational diabetes mellitus. NPJ biofilms and microbiomes, 11(1), 6.

Cooke DM, et al. (2024) Detection of Mycobacterium bovis in nasal swabs from communal goats (Capra hircus) in rural KwaZulu-Natal, South Africa. Frontiers in microbiology, 15, 1349163.

Dai R, et al. (2024) The influence of urban environmental effects on the orchard soil microbial community structure and function: a case study in Zhejiang, China. Frontiers in microbiology, 15, 1403443.

Schuster HJ, et al. (2024) The interrelation between microbial immunoglobulin coating, vaginal microbiota, ethnicity, and preterm birth. Microbiome, 12(1), 99.

Wei T, et al. (2024) Efficient and sustainable removal of linear alkylbenzene sulfonate in a membrane biofilm: Oxygen supply dosage impacts mineralization pathway. Water research X, 25, 100268.

Qian F, et al. (2024) Cultivation of earthworms and analysis of associated bacterial communities during earthworms' growth using two types of agricultural wastes. Bioresources and bioprocessing, 11(1), 66.

Li H, et al. (2024) Alteration of gut microbiome in goslings infected with goose astrovirus. Poultry science, 103(8), 103869.

Li A, et al. (2024) Root exudates and rhizosphere microbiota in responding to long-term continuous cropping of tobacco. Scientific reports, 14(1), 11274.

Bigeard C, et al. (2024) Molecular Xenomonitoring (MX) allows real-time surveillance of West Nile and Usutu virus in mosquito populations. PLoS neglected tropical diseases, 18(12), e0012754.

Rodríguez-García A, et al. (2024) Short-Chain Fatty Acid Production by Gut Microbiota Predicts Treatment Response in Multiple Myeloma. Clinical cancer research : an official journal of the American Association for Cancer Research, 30(4), 904.

Soh M, et al. (2024) The intestinal digesta microbiota of tropical marine fish is largely uncultured and distinct from surrounding water microbiota. NPJ biofilms and microbiomes, 10(1), 11.

García-Gamboa R, et al. (2024) Associations between bacterial and fungal communities in the human gut microbiota and their implications for nutritional status and body weigh?t. Scientific reports, 14(1), 5703.

Pan Y, et al. (2024) Drought-induced assembly of rhizosphere mycobiomes shows beneficial effects on plant growth. mSystems, 9(7), e0035424.

Abbasi AM, et al. (2024) A comparative study of the microbiomes of the ticks Rhipicephalus microplus and Hyalomma anatolicum. Parasite (Paris, France), 31, 74.

Mannochio-Russo H, et al. (2024) The microbiome diversifies N-acyl lipid pools - including short-chain fatty acid-derived compounds. bioRxiv : the preprint server for biology.

Zhang J, et al. (2024) Straw Soil Conditioner Modulates Key Soil Microbes and Nutrient Dynamics across Different Maize Developmental Stages. Microorganisms, 12(2).

Zhang RN, et al. (2024) Identification of neural alterations in patients with Crohn's disease with a novel multiparametric brain MRI-based radiomics model. Insights into imaging, 15(1), 289.

Zang X, et al. (2024) Genotype-associated heritable rumen bacteria can be a stable microbiota passed to the offspring. ISME communications, 4(1), ycad020.

Pietrasanta C, et al. (2024) Prenatal antibiotics reduce breast milk IgA and induce dysbiosis in mouse offspring, increasing neonatal susceptibility to bacterial sepsis. Cell host & microbe, 32(12), 2178.

Zhang J, et al. (2023) Rhizosphere Microbiome and Phenolic Acid Exudation of the Healthy and Diseased American Ginseng Were Modulated by the Cropping History. Plants (Basel, Switzerland), 12(16).