

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

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

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: 20260813T055202+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 102 mentions in open access literature.

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

Kawano A, et al. (2026) Effects of Initial Periodontal Therapy on Intraoral Bacterial Flora in Patients With Oral Contraceptive-Related Severe Gingivitis. Case reports in dentistry, 2026, 5078642.

Khodadad CLM, et al. (2026) Microbial community characterization of multi-crop growouts in the XROOTS aeroponic-hydroponic system on the International Space Station. Frontiers in microbiomes, 5, 1779816.

Du J, et al. (2026) Microbial community succession along the entrance to distant zone in karst cave ecosystem: community assembly and distance decay patterns in weathered rock and sediment habitats. Environmental microbiome, 21(1).

McConnell RD, et al. (2026) Dietary DNA Metabarcoding From Animal Fecal Samples. Current protocols, 6(1), e70226.

Yilmaz B, et al. (2026) Spatial mapping of human colonic niches reveals rapid, mucus-specific microbiota disruption after bowel cleansing. Gut microbes, 18(1), 2635866.

Obayashi N, et al. (2026) Prevalence and Ecological Role of Streptococcus toyakuensis in Saliva of Healthy Young Individuals. International dental journal, 76(1), 103987.

Ali EA, et al. (2026) A national survey of gastrointestinal nematodes in Australian dairy goats using faecal egg counts and deep amplicon sequencing. Current research in parasitology & vector-borne diseases, 9, 100390.

Ito T, et al. (2025) Comparison of the Fecal Microbiota from Long-term Captive and Newly Captured Whale Sharks (Rhincodon typus). Microbes and environments, 40(3).

Cano-Argüelles AL, et al. (2025) Protocol to study the impact of breast cancer on colonization resistance of mouse microbiota using network node manipulation. STAR protocols, 6(1), 103618.

Shen J, et al. (2025) Unveiling the Mystery of the Stimulatory Effects of Arecoline: Its Relevance to the Regulation of Neurotransmitters and the Microecosystem in Multi-Ecological Intestinal Sites. International journal of molecular sciences, 26(7).

Muzami EM, et al. (2025) Metagenomic insights to bacterial communities, functional traits, and soil health in banana smallholder agroecosystems of Kenya. *Frontiers in microbiology*, 16, 1582271.

Jeong SM, et al. (2025) Sardine Inclusion in a Food Waste-Based Substrate for Rearing Black Soldier Fly (*Hermetia illucens*) Larvae: Effects on Growth Performance, Body Composition, and Gut Microbiome. *Insects*, 16(9).

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.

Abukhiran IM, et al. (2025) Mucosal Microbiome Markers of Complete Pathologic Response to Neoadjuvant Therapy in Rectal Carcinoma. *Cancer research communications*, 5(5), 756.

Brar G, et al. (2025) High abundance of lactobacilli in the gut microbiome of honey bees during winter. *Scientific reports*, 15(1), 7409.

Guo M, et al. (2025) One of the Short-Chain Fatty Acids (SCFAs), Sodium Propionate, Can Reduce the Dosage of Sishen Pill in Regulating the Intestinal Microbiota in Diarrhea with Kidney-Yang Deficiency Syndrome. *Journal of inflammation research*, 18, 7195.

Zhang S, et al. (2025) Soil chemical factors contributing to differences in bacterial communities among tea field soils and their relationships with tea quality. *Frontiers in plant science*, 16, 1540659.

Yu D, et al. (2025) Deciphering the microbiological mechanism of Tongxie Yaofang in treating IBS-D: a multimodal mechanistic study in mice integrating network pharmacology, computational simulation, and 16S rRNA sequencing. *Experimental biology and medicine* (Maywood, N.J.), 250, 10725.

Kenneth MJ, et al. (2025) Deciphering the Post-Operative Dynamics of Opportunistic Gut Microbiota in Colorectal Cancer Patients. *Microorganisms*, 13(12).

Fu C, et al. (2025) The mitigation of spatial constraint in porous environments enhances biofilm phylogenetic and functional diversity. *Microbiome*, 13(1), 84.