

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

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

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: 20260729T044437+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 71 mentions in open access literature.

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

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

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.

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

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.

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.

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.

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

Soh M, et al. (2025) Spatial and temporal changes in gut microbiota composition of farmed Asian seabass (Lates calcarifer) in different aquaculture settings. Microbiology spectrum, 13(5), e0198924.

Abdelrahman O, et al. (2025) Mining Actinomycetes' Metabolomes and Genomes for Anti-Phytophthora infestans Compounds. Microbial biotechnology, 18(12), e70269.

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

Zhang R, et al. (2025) Altered gut microbiome-metabolite interactions link intestinal inflammation severity and MR enterography abnormalities in Crohn's disease. *iScience*, 28(5), 112310.

Nakanishi E, et al. (2025) Microbiome Associated with *Polypedilum* sp. (Diptera; Chironomidae), a Midge Adapted to an Extremely Acidic Environment. *Microbes and environments*, 40(2).

McMullen BN, et al. (2025) Metatranscriptomic analysis of colonic mucosal samples exploring the functional role of active microbial consortia in complicated diverticulitis. *Microbiology spectrum*, 13(7), e0243124.

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.

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.

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

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 weight. *Scientific reports*, 14(1), 5703.

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.