

Resource Summary Report

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BioRad QuantaSoft Analysis Software

RRID:SCR_025696

Type: Tool

Proper Citation

BioRad QuantaSoft Analysis Software (RRID:SCR_025696)

Resource Information

URL: <https://www.bio-rad.com/webroot/web/pdf/lsr/literature/10047467.pdf>

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Description: Software analysis tool that enables data analysis of multiplex experiments and advanced applications and assay strategies for Droplet Digital PC.

Synonyms: QuantaSoft Analysis software

Resource Type: software resource, software application

Keywords: Droplet Digital PCR, data files reader, QX100 Droplet Reader data, QX200 Droplet Reader data,

Funding:

Availability: Restricted

Resource Name: BioRad QuantaSoft Analysis Software

Resource ID: SCR_025696

Record Creation Time: 20240907T053256+0000

Record Last Update: 20250421T054624+0000

Ratings and Alerts

No rating or validation information has been found for BioRad QuantaSoft Analysis Software.

No alerts have been found for BioRad QuantaSoft Analysis Software.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

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

Hollander JF, et al. (2025) Serially Quantifying TERT Rearrangement Breakpoints in ctDNA Enables Minimal Residual Disease Monitoring in Patients with Neuroblastoma. Cancer research communications, 5(1), 167.

Hickerson BT, et al. (2023) Interchangeability of the Assays Used to Assess the Activity of Anti-SARS-CoV-2 Monoclonal Antibodies. Viruses, 15(8).

Tereshko L, et al. (2023) A novel method for quantitation of AAV genome integrity using duplex digital PCR. PloS one, 18(12), e0293277.

Frindt G, et al. (2022) Expression of ENaC subunits in epithelia. The Journal of general physiology, 154(10).

O'Hara BA, et al. (2021) Teriflunomide Inhibits JCPyV Infection and Spread in Glial Cells and Choroid Plexus Epithelial Cells. International journal of molecular sciences, 22(18).

Verta JP, et al. (2020) Cis-regulatory differences in isoform expression associate with life history strategy variation in Atlantic salmon. PLoS genetics, 16(9), e1009055.

Bryda EC, et al. (2019) A novel conditional ZsGreen-expressing transgenic reporter rat strain for validating Cre recombinase expression. Scientific reports, 9(1), 13330.

Usher CL, et al. (2015) Structural forms of the human amylase locus and their relationships to SNPs, haplotypes and obesity. Nature genetics, 47(8), 921.