Resource Summary Report

Generated by RRID on Aug 6, 2026

Bio-Rad QuantaSoft Analysis Pro

RRID:SCR_025321
Type: Tool

Proper Citation

Bio-Rad QuantaSoft Analysis Pro (RRID:SCR_025321)

Resource Information

URL: https://www.bio-rad.com/webroot/web/pdf/lsr/literature/QuantaSoft-Analysis-Pro-v1.0-Manual.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 PCR (ddPCR). For use with data files generated on QX100 or QX200 Droplet Reader using QuantSoft Standard Edition Software versions 1.4 or later.

Synonyms: QuantaSoft AP

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

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

Resource Name: Bio-Rad QuantaSoft Analysis Pro

Resource ID: SCR_025321

Alternate URLs: https://www.bio-rad.com

Record Creation Time: 20240511T053259+0000

Record Last Update: 20260806T054815+0000

Ratings and Alerts

No rating or validation information has been found for Bio-Rad QuantaSoft Analysis Pro.

No alerts have been found for Bio-Rad QuantaSoft Analysis Pro.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1221 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

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.

Wagner CB, et al. (2025) Rad53 regulates RNase H1, which promotes DNA replication through sites of transcription-replication conflict. Cell reports, 44(11), 116565.

Marchand V, et al. (2025) Monocytes generated by interleukin-6-treated human hematopoietic stem and progenitor cells secrete calprotectin that inhibits erythropoiesis. iScience, 28(1), 111522.

Alhadidy MM, et al. (2025) O-GlcNAc modification differentially regulates microtubule binding and pathological conformations of tau isoforms in vitro. The Journal of biological chemistry, 301(3), 108263.

De Elías-Escribano A, et al. (2025) Schistosoma mansoni x S. haematobium hybrids frequently infecting sub-Saharan migrants in southeastern Europe: Egg DNA genotyping assessed by RD-PCR, sequencing and cloning. PLoS neglected tropical diseases, 19(3), e0012942.

Atas E, et al. (2025) The anti-diabetic PPAR? agonist Pioglitazone inhibits cell proliferation and induces metabolic reprogramming in prostate cancer. Molecular cancer, 24(1), 134.

Velde CD, et al. (2025) Clinical and Genetic Characteristics of Congenital Hyperinsulinism in Norway: A Nationwide Cohort Study. The Journal of clinical endocrinology and metabolism, 110(2), 554.

Das NK, et al. (2025) Structural basis for 3C and 3CD recruitment by enteroviral genomes during negative-strand RNA synthesis. Nature communications, 16(1), 9293.

Zhang S, et al. (2025) Immunomodulation of behavior impairment via spleen-specific targeting lipid nanoparticles in a MeCP2 transgenic mouse model. Scientific reports, 15(1), 38135.

Das A, et al. (2025) Adenylate cyclase 10 promotes brown adipose tissue thermogenesis. iScience, 28(2), 111833.

Santos-Peral A, et al. (2025) Basal T cell activation predicts yellow fever vaccine response independently of cytomegalovirus infection and sex-related immune variations. Cell reports. Medicine, 6(2), 101946.

Delma FZ, et al. (2025) Emergence of Flucytosine-Resistant Candida tropicalis Clade, the Netherlands. Emerging infectious diseases, 31(7), 1354.

Parenteau J, et al. (2025) Cells resist starvation through a nutrient stress splice switch. Nucleic acids research, 53(12).

Lawford HLS, et al. (2025) Integrated serological surveillance for neglected tropical diseases, vaccine-preventable diseases, and arboviruses in Samoa, 2018. Scientific reports, 15(1), 12667.

Boehm AB, et al. (2025) Detection of Monkeypox Virus Clade Ib DNA in Wastewater Solids at Wastewater Treatment Plants, United States. Emerging infectious diseases, 31(10), 1984.

Deng J, et al. (2025) Monkeypox Virus Partial-Genome Amplicon Sequencing for Improvement of Genomic Surveillance during Mpox Outbreaks. Emerging infectious diseases, 31(11), 2128.

Sun X, et al. (2025) Effect of Prior Influenza A(H1N1)pdm09 Virus Infection on Pathogenesis and Transmission of Human Influenza A(H5N1) Clade 2.3.4.4b Virus in Ferret Model. Emerging infectious diseases, 31(3), 458.

Alonso Domínguez J, et al. (2025) SARS-CoV-2 Viral Load and Cytokine Dynamics Profile as Early Signatures of Long COVID Condition in Hospitalized Individuals. Influenza and other respiratory viruses, 19(1), e70068.

Halurkar MS, et al. (2025) The widely used Ucp1-Cre transgene elicits complex developmental and metabolic phenotypes. Nature communications, 16(1), 770.

Matsuu A, et al. (2025) Oz Virus Infection in 6 Animal Species, Including Macaques, Bears, and Companion Animals, Japan. Emerging infectious diseases, 31(4), 720.