

Resource Summary Report

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QtiPlot

RRID:SCR_017311

Type: Tool

Proper Citation

QtiPlot (RRID:SCR_017311)

Resource Information

URL: <https://www.qtiplot.com/>

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Description: Software tool for data analysis and scientific visualization by IONDEV SRL.

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

Keywords: data, analysis, visualization, IONDEV SRL

Availability: Restricted

Resource Name: QtiPlot

Resource ID: SCR_017311

Record Creation Time: 20220129T080334+0000

Record Last Update: 20260919T195338+0000

Ratings and Alerts

No rating or validation information has been found for QtiPlot.

No alerts have been found for QtiPlot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 30 mentions in open access literature.

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

Fabrizio M, et al. (2026) Cryo-EM structure of TRPM1 reveals a non-canonical architecture with an inverted transmembrane domain. *Nature communications*, 17(1).

Yang JY, et al. (2026) Injectable Deep Eutectic Solvent-Based Ionic Gel With Rapid Gelation and Broad Hemostatic Functionality. *Advanced healthcare materials*, 15(3), e03038.

Zanoni JA, et al. (2025) Effects of Glycerol and Phenolics on *Myceliophthora heterothallica* Endoxylanase Expressed in *K. phaffii*. *Biotech (Basel (Switzerland))*, 14(3).

Guo C, et al. (2025) Pulmonary Toxicity of Long, Thick MWCNT and Very Long, Thin Carboxylated MWCNT Aerosols Following 28 Days Whole-Body Exposure. *Toxics*, 13(5).

Midya R, et al. (2025) Artificial transneurons emulate neuronal activity in different areas of brain cortex. *Nature communications*, 16(1), 7289.

Quirk S, et al. (2025) A multidomain hydrolase from the thermophile *Thermanaeromonas toyohensis* degrades high molecular mass forms of polyhydroxybutyrate. *Protein science : a publication of the Protein Society*, 34(8), e70223.

Zhang J, et al. (2025) Molecular basis of TRPV3 channel blockade by intracellular polyamines. *Communications biology*, 8(1), 727.

Payet RD, et al. (2024) Elucidation of *Spartina* dimethylsulfoniopropionate synthesis genes enables engineering of stress tolerant plants. *Nature communications*, 15(1), 8568.

Najer A, et al. (2024) Enhanced Antimalarial and Antisequestration Activity of Methoxybenzenesulfonate-Modified Biopolymers and Nanoparticles for Tackling Severe Malaria. *ACS infectious diseases*, 10(2), 732.

Gilbert C, et al. (2024) Species identification of ivory and bone museum objects using minimally invasive proteomics. *Science advances*, 10(4), eadi9028.

Baumgartner SA, et al. (2024) Love your wet grass! Dry season grazing reserves show highest grass regrowth in communal semi-arid rangelands of Tanzania. *PloS one*, 19(11), e0313818.

Maksaev G, et al. (2023) Blockade of TRPV channels by intracellular spermine. *The Journal of general physiology*, 155(5).

Bharadwaj AG, et al. (2023) Identification and characterization of calreticulin as a novel plasminogen receptor. *The Journal of biological chemistry*, 300(1), 105465.

Adler NS, et al. (2022) Insights into the product release mechanism of dengue virus NS3 helicase. *Nucleic acids research*, 50(12), 6968.

Yakubu HG, et al. (2022) Micro-Encapsulated Microalgae Oil Supplementation Has No Systematic Effect on the Odor of Vanilla Shake-Test of an Electronic Nose. *Foods (Basel*,

Switzerland), 11(21).

Perera RP, et al. (2021) Bucky Ball Is a Novel Zebrafish Vasa ATPase Activator. *Biomolecules*, 11(10).

Brasil S, et al. (2021) A Novel Noninvasive Technique for Intracranial Pressure Waveform Monitoring in Critical Care. *Journal of personalized medicine*, 11(12).

Shen M, et al. (2021) Structural basis for the multi-activity factor Rad5 in replication stress tolerance. *Nature communications*, 12(1), 321.

Ernst O, et al. (2021) Species-Specific Endotoxin Stimulus Determines Toll-Like Receptor 4- and Caspase 11-Mediated Pathway Activation Characteristics. *mSystems*, 6(4), e0030621.

Aslam M, et al. (2021) Putative second hit rare genetic variants in families with seemingly GBA-associated Parkinson's disease. *NPJ genomic medicine*, 6(1), 2.