

Resource Summary Report

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Thermo Fisher: Qubit 2.0 Fluorometer Qubit 2.0

RRID:SCR_020553

Type: Tool

Proper Citation

Thermo Fisher: Qubit 2.0 Fluorometer Qubit 2.0 (RRID:SCR_020553)

Resource Information

URL: https://www.thermofisher.com/us/en/home/industrial/spectroscopy-elemental-isotope-analysis/molecular-spectroscopy/fluorometers/qubit/qubit-flex-fluorometer.html?gclid=CjwKCAiA6vXwBRBKEiwAYE7iS0T2xfCBHWYsu-Uhf_Y8f69WS9j4yYK1udH27vHhWXnuyA_u7Vj4xoCuOcQAvD_BwE&ef_id=CjwKCAiA6vXwBRBKEiwAYE7iS0T2xfCBHWYsu-Uhf_Y8f69WS9j4yYK1udH27vHhWXnuyA_u7Vj4xoCuOcQAvD_BwE:G:s&s_kwid=AL!3652!3!39

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Description: Invitrogen Qubit Flex Fluorometer can selectively, accurately measure the concentration of up to 8 samples of DNA, RNA or protein simultaneously. Compared to single sample microvolume fluorometers, using the Qubit Flex Fluorometer for multiple samples reduces your time to data by up to 50%, with time savings beginning with batches of as few as 8 samples. Combining the one-of-a-kind Qubit Flex Fluorometer and optimized Invitrogen Qubit reagents and tube strips offers a complete solution for DNA, RNA and protein quantification.

Resource Type: instrument resource

Keywords: Thermo Fisher, Fluorometer, Instrument Equipment, USEDit

Availability: Commercially available

Resource Name: Thermo Fisher: Qubit 2.0 Fluorometer Qubit 2.0

Resource ID: SCR_020553

Alternate IDs: Model_Number_Qubit

Record Creation Time: 20220129T080350+0000

Record Last Update: 20260801T190700+0000

Ratings and Alerts

No rating or validation information has been found for Thermo Fisher: Qubit 2.0 Fluorometer Qubit 2.0.

No alerts have been found for Thermo Fisher: Qubit 2.0 Fluorometer Qubit 2.0.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 26 mentions in open access literature.

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

Scourzic L, et al. (2026) T follicular helper cells transiently unlock a plasticity state in germinal centre B cells during the humoral immune response. *Nature cell biology*, 28(1), 35.

Barajas JM, et al. (2026) Tandem Duplications in UBTF Create XPO1-Dependent Nuclear Export Signals that Reveal a Leukemic Therapeutic Dependency. *Blood cancer discovery*, 7(1), 51.

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.

Puzio-Kuter AM, et al. (2025) Restoration of the Tumor Suppressor Function of Y220C-Mutant p53 by Rezatapopt, a Small-Molecule Reactivator. *Cancer discovery*, 15(6), 1159.

Ishikawa R, et al. (2025) Immunohistochemical and molecular evolutionary features of jejunoileal adenocarcinoma unveiled through comparative analysis with colorectal adenocarcinoma. *Neoplasia (New York, N.Y.)*, 66, 101180.

Mathios D, et al. (2025) Detection of Brain Cancer Using Genome-wide Cell-free DNA Fragmentomes. *Cancer discovery*, 15(8), 1593.

Megid RA, et al. (2025) Sotorasib resistance triggers epithelial-mesenchymal transition and activates AKT and P38-mediated signaling. *Frontiers in molecular biosciences*, 12, 1537523.

Boccardo S, et al. (2025) Dynamics of tissue repair regulatory T cells and damage in acute *Trypanosoma cruzi* infection. *PLoS pathogens*, 21(1), e1012906.

Figuerola-Bou E, et al. (2025) KDM6 Demethylases Contribute to EWSR1::FLI1-Driven Oncogenic Reprogramming in Ewing Sarcoma. *Cancer research*, 85(22), 4485.

Weidling I, et al. (2025) hiPSC-neurons recapitulate the subtype-specific cell intrinsic nature of susceptibility to neurodegenerative disease-relevant aggregation. *Acta neuropathologica communications*, 13(1), 108.

Ma D, et al. (2025) Spatial determinants of antibody-drug conjugate SHR-A1811 efficacy in neoadjuvant treatment for HER2-positive breast cancer. *Cancer cell*, 43(6), 1061.

Valvo VM, et al. (2025) Olaparib and Radiotherapy Induce Type I Interferon- and CD8+ T Cell-Dependent Sensitization to Immunotherapy in Pancreatic Cancer. *Molecular cancer therapeutics*, 24(6), 843.

Pan S, et al. (2025) A holistic genome dataset of bacteria and archaea of mangrove sediments. *GigaScience*, 14.

Campos-Iglesias D, et al. (2024) Loss of ADAM29 does not affect viability and fertility in mice but improves wound healing. *iScience*, 27(6), 110135.

Sulsenti R, et al. (2024) Intracellular Osteopontin Promotes the Release of TNF α by Mast Cells to Restrain Neuroendocrine Prostate Cancer. *Cancer immunology research*, 12(9), 1147.

Cao C, et al. (2024) CXCR4 orchestrates the TOX-programmed exhausted phenotype of CD8+ T cells via JAK2/STAT3 pathway. *Cell genomics*, 4(10), 100659.

Saggese P, et al. (2024) Glucose Deprivation Promotes Pseudohypoxia and Dedifferentiation in Lung Adenocarcinoma. *Cancer research*, 84(2), 305.

Stacchiotti S, et al. (2024) GDF-15 Predicts Epithelioid Hemangioendothelioma Aggressiveness and Is Downregulated by Sirolimus through ATF4/ATF5 Suppression. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 30(22), 5122.

Gobbini A, et al. (2023) Protocol for the detection of defined T cell clones in a heterogeneous cell population. *STAR protocols*, 5(1), 102787.

Fait A, et al. (2023) Evolutionary history of *Staphylococcus aureus* influences antibiotic resistance evolution. *Current biology : CB*, 33(16), 3389.