Resource Summary Report

Generated by NIF on Sep 16, 2026

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_Y8f69WS9j4yYKm1udH27vHhWXnuyA_u7Vj4xoCuOcQAvD_BwE&ef_id=CjwKCAiA6vXwBRBKEiwAYE7iS0T2xfCBHWYsu-Uhf_Y8f69WS9j4yYKm1udH27vHhWXnuyA_u7Vj4xoCuOcQAvD_BwE:G:s&s_kwcid=AL!3652!3!393

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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: 20260912T195939+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 39 mentions in open access literature.

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

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.

Valentín López JC, et al. (2026) Evolution of AR and Non-AR Alterations in ctDNA of Patients with Metastatic Prostate Cancer Treated in the Phase III Alliance A031201 Trial. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(12), 2413.

Nasr MM, et al. (2026) PRDM16 Regulates Prostate Cancer Cell Dormancy and Prevents Bone Metastatic Outgrowth. Cancer research, 86(3), 604.

Foster M, et al. (2026) A Personalized Therapeutic Approach for Liver Cancers Expressing the African-Centric P47S Variant of TP53. Molecular cancer research : MCR, 24(7), 618.

Popov TM, et al. (2026) Two Molecular Subgroups Predict Most Recurrences in Advanced Laryngeal Squamous Cell Carcinoma. Cancer research communications, 6(1), 60.

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.

Fu ZH, et al. (2026) Synthetic rewriting of the IGH locus. Cell genomics, 6(6), 101252.

Monteagudo-García Ó, et al. (2026) ZC3H13 Loss Drives Cancer Metastatic Progression by Disrupting m6A RNA Methylation. Cancer research, 86(3), 622.

Adhikary U, et al. (2026) Cancer Susceptibility to Stapled Oncolytic Peptides Is Dictated by Membrane Cholesterol and Inflammatory Signaling. Cancer research, 86(11), 2810.

Pastrana CC, et al. (2026) Nonsense-Mediated RNA Decay Is a Targetable Vulnerability in Splicing Factor Mutant Myeloid Neoplasms by Enhancing R-Loop Accumulation and DNA Damage. Cancer research, 86(14), 3537.

Shah UA, et al. (2026) A High-Fiber, Plant-Based Diet in Myeloma Precursor Disorders: Results from the NUTRIVENTION Clinical Trial and Preclinical Vk*MYC Model. Cancer discovery, 16(4), 697.

Belanger K, et al. (2026) Cancer-Associated Fibroblasts Promote Epithelial-Mesenchymal Transition and Classical to Basal Subtype Shift in Pancreatic Cancer. Cancer research, 86(14), 3604.

Chang S, et al. (2026) Nascent chain folding status modulates ribosome dynamics and mRNA stability through the ribosome-associated chaperone Zuo1. Molecular cell.

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.

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

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.

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

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

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

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