

Resource Summary Report

Generated by [RRID](#) on Aug 6, 2026

Thermo Fisher: QuantStudio 3 Real Time PCR System

RRID:SCR_018712

Type: Tool

Proper Citation

Thermo Fisher: QuantStudio 3 Real Time PCR System (RRID:SCR_018712)

Resource Information

URL: <https://www.thermofisher.com/order/catalog/product/A28567#/A28567>

Proper Citation: Thermo Fisher: QuantStudio 3 Real Time PCR System (RRID:SCR_018712)

Description: System designed for users who need easy-to-use real-time PCR system. QuantStudio 3 Real Time PCR System supports 96-well (0.2ml or 0.1ml) or 384-well (QS5 only) consumables. Can detect extremely small changes in gene expression, as little as 1.5-fold. Its dynamic range of up to 10 logs allows for tracking wide range of expression changes across various sample types and conditions. When connected to Thermo Fisher Cloud, QuantStudio 3 system provides access to your data wherever and whenever you want.

Synonyms: , ABI QuantStudio3 Flex instrument, QuantStudio TM 3 Real-Time PCR System, Life Technologies QuantStudio 3 Real Time PCR System, Applied Biosystems QuantStudio 3 Real-Time PCR System, QuantStudio 3 RT PCR System

Resource Type: instrument resource

Keywords: ABRF, Applied Biosystems, Thermo Fisher Scientific, equipment, instrument, QuantStudio 3 Real-Time PCR System, 96-well, 0.2 mL, laptop, genomic data, genomic application

Specification URL: https://raw.githubusercontent.com/SciCrunch/RRID-Instruments/main/PDF/SCR_018712.pdf

Resource Name: Thermo Fisher: QuantStudio 3 Real Time PCR System

Resource ID: SCR_018712

Alternate IDs: SCR_020238

Record Creation Time: 20220129T080341+0000

Record Last Update: 20260801T190624+0000

Ratings and Alerts

No rating or validation information has been found for Thermo Fisher: QuantStudio 3 Real Time PCR System.

No alerts have been found for Thermo Fisher: QuantStudio 3 Real Time PCR System.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 76 mentions in open access literature.

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

Fernandes P, et al. (2026) Functional immune responses induced by a capsid assembly modulator in chronic hepatitis B virus-infected humanized mice. *Cell host & microbe*, 34(1), 116.

Yi E, et al. (2025) Selective Depletion of Cancer Cells with Extrachromosomal DNA via Lentiviral Infection. *Cancer research communications*, 5(8), 1458.

Seo DW, et al. (2025) Dual independent mechanisms underlying gut epithelial remodeling upon sugar substitute consumption. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 39(3), e70374.

Zampieri C, et al. (2025) The urokinase receptor/uPAR is a major effector of p53 gain-of-function mutations in gemcitabine-treated pancreatic ductal adenocarcinoma. *The Journal of biological chemistry*, 301(9), 110561.

Khan K, et al. (2025) Loss of Sorting Nexin 10 Accelerates KRAS-Induced Pancreatic Tumorigenesis. *Cancer research communications*, 5(9), 1541.

Fernandez RF, et al. (2025) Aging Effects on Lipid Metabolism in Response to Traumatic Brain Injury. *Journal of neurochemistry*, 169(9), e70227.

Chia ML, et al. (2025) Metabolic imaging distinguishes ovarian cancer subtypes and detects their early and variable responses to treatment. *Oncogene*, 44(9), 563.

Nelson KL, et al. (2025) VAX014 Activates Tumor-Intrinsic STING and RIG-I to Promote the Development of Antitumor Immunity. *Molecular cancer therapeutics*, 24(4), 587.

Yu J, et al. (2025) Resilience and vulnerabilities of tumor cells under purine shortage stress. *Clinical cancer research : an official journal of the American Association for Cancer*

Research.

Bond Newton SE, et al. (2025) ER stress tolerance is regulated by copper-dependent PERK kinase activity. *Cell reports*, 44(10), 116318.

Guerrero Ruiz VM, et al. (2025) Macrophage fatty acid metabolism under chronic hypoxia shapes ?? T cell recruitment via CCL22. *iScience*, 28(8), 113025.

Maligireddy SS, et al. (2025) Targeting a pathogenic cryptic exon that drives HLRCC to induce exon skipping. *Molecular therapy. Nucleic acids*, 36(3), 102668.

Yamasaki A, et al. (2025) Cyclin-dependent kinase inhibitor 1A mediates mouse line- and fate-dependent cellular responses in Cx3cr1-Cre genetic tools. *Cell reports*, 44(9), 116267.

Malik A, et al. (2024) Doxorubicin?induced cardiomyopathy is mitigated by empagliflozin via the modulation of endoplasmic reticulum stress pathways. *Molecular medicine reports*, 29(5).

Dhaliwal NK, et al. (2024) Protocol for the efficient and inducible generation of CRISPR-Cas9-edited human cortical neurons from the iCas9-iNgn2 hPSCs. *STAR protocols*, 5(4), 103352.

Hai L, et al. (2024) A clinically applicable connectivity signature for glioblastoma includes the tumor network driver CHI3L1. *Nature communications*, 15(1), 968.

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

Danev N, et al. (2024) Comparative transcriptomic analysis of bovine mesenchymal stromal cells reveals tissue-source and species-specific differences. *iScience*, 27(2), 108886.

Yang L, et al. (2024) Neurons enhance blood-brain barrier function via upregulating claudin-5 and VE-cadherin expression due to glial cell line-derived neurotrophic factor secretion. *eLife*, 13.

de Miranda FS, et al. (2024) MicroRNA as a promising molecular biomarker in the diagnosis of breast cancer. *Frontiers in molecular biosciences*, 11, 1337706.