## **Resource Summary Report**

Generated by dkNET on Apr 24, 2025

# **NormqPCR**

RRID:SCR\_003388

Type: Tool

## **Proper Citation**

NormqPCR (RRID:SCR\_003388)

#### **Resource Information**

URL: http://www.bioconductor.org/packages/release/bioc/html/NormqPCR.html

Proper Citation: NormqPCR (RRID:SCR\_003388)

**Description:** Software package providing functions for the selection of optimal reference

genes and the normalization of real-time quantitative PCR data.

**Abbreviations:** NormqPCR

Synonyms: NormqPCR - Functions for normalisation of RT-qPCR data

Resource Type: software resource

**Defining Citation:** PMID:22748112

Keywords: gene expression, microtitre plate assay, qpcr, reference gene, bio.tools

Funding:

Availability: GNU Lesser General Public License, v3

Resource Name: NormqPCR

Resource ID: SCR\_003388

Alternate IDs: OMICS\_02315, biotools:normqpcr

Alternate URLs: https://bio.tools/normqpcr

**Record Creation Time:** 20220129T080218+0000

**Record Last Update:** 20250420T014141+0000

## **Ratings and Alerts**

No rating or validation information has been found for NormqPCR.

No alerts have been found for NormqPCR.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 49 mentions in open access literature.

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

Ye J, et al. (2024) Grid1 regulates the onset of puberty in female rats. The Journal of veterinary medical science, 86(5), 497.

Sun H, et al. (2024) Identification and validation of stable reference genes for RT-qPCR analyses of Kobresia littledalei seedlings. BMC plant biology, 24(1), 389.

Escoto A, et al. (2024) Nuclear EGFR in breast cancer suppresses NK cell recruitment and cytotoxicity. Oncogene.

Morikka J, et al. (2024) Toxicogenomic assessment of in vitro macrophages exposed to profibrotic challenge reveals a sustained transcriptomic immune signature. Computational and structural biotechnology journal, 25, 194.

Hornung E, et al. (2024) Neuromodulatory co-expression in cardiac vagal motor neurons of the dorsal motor nucleus of the vagus. iScience, 27(8), 110549.

Djari A, et al. (2024) A stable combination of non-stable genes outperforms standard reference genes for RT-qPCR data normalization. Scientific reports, 14(1), 31278.

Taylor MH, et al. (2024) Phase I dose escalation study of IO-108, an anti-LILRB2 antibody, in patients with advanced solid tumors. Journal for immunotherapy of cancer, 12(11).

Flora A, et al. (2023) Human dermal fibroblast subpopulations and epithelial mesenchymal transition signals in hidradenitis suppurativa tunnels are normalized by spleen tyrosine kinase antagonism in vivo. PloS one, 18(11), e0282763.

Röttgen V, et al. (2023) Early milk-feeding regimes in calves exert long-term effects on the development of ovarian granulosa cells. BMC genomics, 24(1), 485.

Loubet I, et al. (2023) A high diversity of non-target site resistance mechanisms to acetolactate-synthase (ALS) inhibiting herbicides has evolved within and among field populations of common ragweed (Ambrosia artemisiifolia L.). BMC plant biology, 23(1), 510.

Kyle Martin W, et al. (2023) Wildfire-related smoke inhalation worsens cardiovascular risk in sleep disrupted rats. Frontiers in environmental health, 2.

Lindemann A, et al. (2023) Anesthetic?specific IncRNA and mRNA profile changes in blood during colorectal cancer resection: A prospective, matched?case pilot study. Oncology reports, 49(2).

Formosa A, et al. (2023) Validation of reference gene stability for miRNA quantification by reverse transcription quantitative PCR in the peripheral blood of patients with COVID-19 critical illness. PloS one, 18(8), e0286871.

Whittaker DS, et al. (2023) Circadian modulation by time-restricted feeding rescues brain pathology and improves memory in mouse models of Alzheimer's disease. Cell metabolism, 35(10), 1704.

Hutchins E, et al. (2021) Extracellular circular RNA profiles in plasma and urine of healthy, male college athletes. Scientific data, 8(1), 276.

Lusardi TA, et al. (2021) Cerebrospinal Fluid MicroRNA Changes in Cognitively Normal Veterans With a History of Deployment-Associated Mild Traumatic Brain Injury. Frontiers in neuroscience, 15, 720778.

Posillico CK, et al. (2021) Sex differences and similarities in the neuroimmune response to central administration of poly I:C. Journal of neuroinflammation, 18(1), 193.

Levin-Schwartz Y, et al. (2021) Exosomal miRNAs in urine associated with children's cardiorenal parameters: a cross-sectional study. Epigenomics, 13(7), 499.

Leung C, et al. (2021) 3D single cell scale anatomical map of sex-dependent variability of the rat intrinsic cardiac nervous system. iScience, 24(7), 102795.

Linardi? M, et al. (2021) Identification and selection of optimal reference genes for qPCR-based gene expression analysis in Fucus distichus under various abiotic stresses. PloS one, 16(4), e0233249.