

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

Gen-Probe

RRID:SCR_003202

Type: Tool

Proper Citation

Gen-Probe (RRID:SCR_003202)

Resource Information

URL: <http://www.hologic.com/>

Proper Citation: Gen-Probe (RRID:SCR_003202)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented July 7, 2017. Formerly a commercial organization for clinical diagnostics, blood screening, transplant products and research products, it had been acquired by Hologic in 2012.

Resource Type: commercial organization

Keywords: clinical services, clinical products, transplant products, research services

Availability: Free, Freely available

Resource Name: Gen-Probe

Resource ID: SCR_003202

Alternate IDs: nlx_152369, ISNI: 0000 0004 0641 3850, grid.481838.e, Wikidata: Q30343851

Alternate URLs: <https://ror.org/013x7n115>

Old URLs: <http://www.gen-probe.com>

Record Creation Time: 20220129T080217+0000

Record Last Update: 20260801T190205+0000

Ratings and Alerts

No rating or validation information has been found for Gen-Probe.

No alerts have been found for Gen-Probe.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 88 mentions in open access literature.

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

Silva-Klug AC, et al. (2025) Using Anal Cytology and Human Papillomavirus DNA and E6/E7 mRNA Detection to Optimize High-Resolution Anoscopy Referrals in Men Who Have Sex With Men With HIV. *Open forum infectious diseases*, 12(1), ofae735.

Yan Z, et al. (2025) From cervix to multisite: Detection of lower genital tract lesions in a 10-year cross-sectional colposcopy clinic study. *PloS one*, 20(12), e0338489.

Moreland ZG, et al. (2025) Myosin-based nucleation of actin filaments contributes to stereocilia development critical for hearing. *Nature communications*, 16(1), 947.

Gueuning M, et al. (2024) Resolving Genotype-Phenotype Discrepancies of the Kidd Blood Group System Using Long-Read Nanopore Sequencing. *Biomedicines*, 12(1).

Dang X, et al. (2024) Exploring the potential prompting role of cervical human papilloma virus detection in vulvar lesions: a cross-sectional study in China. *Frontiers in oncology*, 14, 1353580.

Heysell SK, et al. (2023) Pharmacokinetic-Pharmacodynamic Determinants of Clinical Outcomes for Rifampin-Resistant Tuberculosis: A Multisite Prospective Cohort Study. *Clinical infectious diseases : an official publication of the Infectious Diseases Society of America*, 76(3), 497.

Sanogo F, et al. (2023) Hematological profiles of patients with tuberculosis and nontuberculous mycobacteria infections in Bamako, Mali. *International journal of mycobacteriology*, 12(3), 235.

Di Germanio C, et al. (2023) A Stable Dried Tube Specimen for Quality Assurance and Training Programs for HIV Rapid Test for Recent Infection. *Microbiology spectrum*, 11(1), e0339822.

Portero V, et al. (2022) Chronically elevated branched chain amino acid levels are pro-arrhythmic. *Cardiovascular research*, 118(7), 1742.

Cicuéndez M, et al. (2021) Effects of Human and Porcine Adipose Extracellular Matrices Decellularized by Enzymatic or Chemical Methods on Macrophage Polarization and Immunocompetence. *International journal of molecular sciences*, 22(8).

Cicuéndez M, et al. (2021) Benefits in the Macrophage Response Due to Graphene Oxide Reduction by Thermal Treatment. *International journal of molecular sciences*, 22(13).

Cicuéndez M, et al. (2021) Candida albicans/Macrophage Biointerface on Human and Porcine Decellularized Adipose Matrices. Journal of fungi (Basel, Switzerland), 7(5).

Mwau M, et al. (2021) Prospective evaluation of accuracy of HIV viral load monitoring using the Aptima HIV Quant Dx assay with fingerstick and venous dried blood spots prepared under field conditions in Kenya. PloS one, 16(4), e0249376.

Banks GT, et al. (2020) Forward genetics identifies a novel sleep mutant with sleep state inertia and REM sleep deficits. Science advances, 6(33), eabb3567.

Muhsin NIA, et al. (2020) A novel mutation in the mouse Pcsk1 gene showing obesity and diabetes. Mammalian genome : official journal of the International Mammalian Genome Society, 31(1-2), 17.

Falcone S, et al. (2019) Modification of an aggressive model of Alport Syndrome reveals early differences in disease pathogenesis due to genetic background. Scientific reports, 9(1), 20398.

Ahmed RM, et al. (2019) Neural networks associated with body composition in frontotemporal dementia. Annals of clinical and translational neurology, 6(9), 1707.

Gannon AL, et al. (2019) Ablation of glucocorticoid receptor in the hindbrain of the mouse provides a novel model to investigate stress disorders. Scientific reports, 9(1), 3250.

Najgebauer H, et al. (2019) Integrated omics profiling reveals novel patterns of epigenetic programming in cancer-associated myofibroblasts. Carcinogenesis, 40(4), 500.

Agnew T, et al. (2018) A Wars2 Mutant Mouse Model Displays OXPHOS Deficiencies and Activation of Tissue-Specific Stress Response Pathways. Cell reports, 25(12), 3315.