

# Resource Summary Report

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## [oligo](#)

RRID:SCR\_015729

Type: Tool

### Proper Citation

oligo (RRID:SCR\_015729)

### Resource Information

**URL:** <https://bioconductor.org/packages/release/bioc/html/oligo.html>

**Proper Citation:** oligo (RRID:SCR\_015729)

**Description:** Software package to analyze oligonucleotide arrays (expression/SNP/tiling/exon) at probe-level. It currently supports Affymetrix (CEL files) and NimbleGen arrays (XYS files).

**Synonyms:** oligo package

**Resource Type:** data analysis software, data processing software, software application, software resource, source code

**Defining Citation:** [PMID:20688976](#)

**Keywords:** oligonucleotide, microarray gene expression, r, oligonucleotide array, snp, gene expression, probe-level, affymetrix array, cel file, and nimblegen array, xys file, bio.tools

**Funding:** CAPES (Coordenação de Aprimoramento Pessoal de Nível Superior) ; NCRR R01RR021967; NHGRI P41HG004059

**Availability:** Free, Available for download, Runs on Mac OS, Runs on Windows

**Resource Name:** oligo

**Resource ID:** SCR\_015729

**Alternate IDs:** biotools:oligo

**Alternate URLs:** <https://bio.tools/oligo>

**License:** LGPL

**Record Creation Time:** 20220129T080327+0000

**Record Last Update:** 20260829T182500+0000

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## Ratings and Alerts

No rating or validation information has been found for oligo.

No alerts have been found for oligo.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1808 mentions in open access literature.

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

Lin R, et al. (2026) Senescence-related gene signature predicts prostate cancer progression and identifies PCNA as a therapeutic target via multi-omics machine learning integration. British journal of cancer, 134(4), 662.

Xing H, et al. (2026) Multi-omics integration analysis reveals metabolic changes during sika deer antler growth. BMC genomics, 27(1).

Jin Y, et al. (2026) Comparative Analysis of the Transcriptome of the Chicken Breast Muscle at Different Developmental Stages. Animals : an open access journal from MDPI, 16(7).

Cicha E, et al. (2026) The Effects of ACTH and Dexamethasone on the Transcriptomic Profile of the Rat Adrenal Gland: An In Vivo Study. Current issues in molecular biology, 48(2).

Wang Y, et al. (2026) Formation Mechanisms of the Ellipsoid Egg in Silkworm (Bombyx mori): Insights from Transcriptomic Profiling. Genes, 17(2).

Ren Y, et al. (2026) Comparative Hepatic Transcriptomic Analysis Reveals Metabolic Regulatory Differences Between Qilian and Oula Sheep. Veterinary sciences, 13(5).

Li L, et al. (2026) Complete Mitochondrial Genome of Melophagus ovinus from Qinghai-Tibet Plateau Provides Evidence for D-Loop Length Polymorphism. Genes, 17(6).

Li X, et al. (2026) Differential expression of long non-coding RNA in the hypothalamus-pituitary-gonadal axis of Wanxi white geese during laying and broodiness periods. Animal bioscience, 39(3), 250348.

Li X, et al. (2026) Vitamin E acetate and its aerosol activate aryl hydrocarbon receptor signaling and exacerbate inflammation in human U937-derived macrophages. Toxicology

in vitro : an international journal published in association with BIBRA, 112, 106196.

Pan Y, et al. (2026) Integrating single-nucleus barcoding with spatial transcriptomics via Stamp-seq to reveal immunotherapy response-enhancing functional modules in NSCLC. *Cell discovery*, 12(1), 10.

Huang Z, et al. (2026) Development of an RT-ERA-CRISPR/Cas12a assay for duck Tembusu virus. *Poultry science*, 105(5), 106612.

Gao L, et al. (2026) Transcriptome Sequencing Unveils a Novel Mechanism Underlying Breed Distinctions Between Thin- and Fat-Tailed Sheep. *Genes*, 17(2).

Jiang L, et al. (2026) Establishment of duplex multi-enzyme isothermal rapid amplification detection method for bovine astrovirus and norovirus. *Frontiers in veterinary science*, 13, 1819282.

Ren Y, et al. (2026) Rapid, sensitive, and highly specific detection of *Mycoplasma pneumoniae* and its mutation using the RPA-CRISPR/Cas13a system. *Microbiology spectrum*, 14(5), e0179325.

Zhao X, et al. (2026) Integrated transcriptomic and metabolomic analyses reveal the regulatory mechanism of subcutaneous fat deposition in Baicheng Oil chickens. *Poultry science*, 105(7), 106894.

Sun Z, et al. (2026) Novel goose parvovirus as a major risk factor for red skin and bristle feather syndrome in meat ducks: Epidemiological associations with duck circovirus and reovirus. *Poultry science*, 105(9), 107281.

Xie W, et al. (2026) Exploring the Cardioprotective Spectrum-Effect Relationship of *Apocynum venetum* L. Using a Zebrafish Model. *Pharmaceuticals (Basel, Switzerland)*, 19(6).

Song Y, et al. (2026) Establishment and application of a rapid and visual detection method for *Mycoplasma pneumoniae* using enzymatic recombinase amplification combined with lateral flow dipstick. *Biosafety and health*, 8(3), 199.

Gurer Kluge EE, et al. (2026) Mucosal calprotectin is associated with severity of aGI-GVHD and poor outcomes after allogeneic stem cell transplantation. *Blood*, 147(8), 886.

Feng J, et al. (2026) Rapid growth phenotype of carbapenem-resistant *Enterobacter cloacae*: growth fitness, stability, and mechanistic insights. *BMC genomics*, 27(1).