

Package: GeoMxWorkflows (via r-universe)

October 11, 2026

Title GeoMx Digital Spatial Profiler (DSP) data analysis workflows

Description Workflows for use with NanoString Technologies GeoMx Technology. Package provides bioconductor focused workflows for leveraging existing packages (e.g. GeomxTools) to process, QC, and analyze the data.

Version 1.18.0

Encoding UTF-8

Depends R (>= 4.0), NanoStringNCTools, GeomxTools

Imports Biobase, S4Vectors, rjson, readxl, EnvStats, dplyr, reshape2, methods, utils, stats, data.table, outliers, BiocGenerics, ggplot2, ggrepel, ggforce, cowplot, scales, umap, Rtsne, pheatmap, BiocStyle, networkD3

Suggests rmarkdown, knitr

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BiocType Workflow

biocViews GeneExpressionWorkflow, ImmunoOncologyWorkflow, SpatialWorkflow

VignetteEngine knitr::rmarkdown

VignetteBuilder knitr

RoxygenNote 7.2.3

Config/testthat/edition 3

Config/pak/sysreqs libcairo2-dev cmake libfontconfig1-dev libfreetype6-dev libglpk-dev make libicu-dev libpng-dev libuv1-dev libxml2-dev libssl-dev python3 zlib1g-dev

Repository Bioconductor 3.23

Date/Publication 2026-04-28 12:44:52 UTC

RemoteUrl <https://github.com/bioc/GeoMxWorkflows>

RemoteRef RELEASE_3_23

RemoteSha 3d45faa4e65ee8da035d1d844b8d45205bae3fc1