

Resource Summary Report

Generated by [dkNET](#) on Sep 18, 2026

Vireo

RRID:SCR_027463

Type: Tool

Proper Citation

Vireo (RRID:SCR_027463)

Resource Information

URL: <https://github.com/single-cell-genetics/vireo>

Proper Citation: Vireo (RRID:SCR_027463)

Description: Software tool for Bayesian demultiplexing of pooled single-cell RNA-seq data without genotype reference.

Resource Type: software application, software resource, source code

Defining Citation: [PMID:31836005](#)

Keywords: Bayesian demultiplexing, pooled single-cell RNA-seq data, without genotype reference,

Funding: EMBL-EBI ;
Medical Research Council ;
National Health and Medical Research Council of Australia ;
University of Cambridge ;
Wellcome Trust

Availability: Free, Available for download, Freely available

Resource Name: Vireo

Resource ID: SCR_027463

License: Apache-2.0 license

Record Creation Time: 20250926T061848+0000

Record Last Update: 20260912T200527+0000

Ratings and Alerts

No rating or validation information has been found for Vireo.

No alerts have been found for Vireo.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Apostolov E, et al. (2026) Single-Cell and Spatial Transcriptomic Profiling Reveals Epithelial Functional States and Fibroblast Phenotypes in Hormone Therapy-Na??ve Localized Prostate Cancer. Cancer research, 86(8), 1836.

Yan R, et al. (2025) Convergent flow-mediated mesenchymal force drives embryonic foregut constriction and splitting. Nature communications, 16(1), 10643.