# **Manuscript Title**

This manuscript (<u>permalink</u>) was automatically generated from <u>BiocWrites/bioc bib chapter 2023@f693a9b</u> on November 2, 2023.

#### **Authors**

- Vincent J. Carey
- Sean Davis <sup>™</sup>

University of Colorado Anschutz School of Medicine

☑ — Correspondence possible via <u>GitHub Issues</u> or email to Sean Davis <seandavi@gmail.com>.

#### **Abstract**

#### Introduction

### **Results**

How Bioconductor enhances cancer data reuse and value

Multiassay cancer data integration using Bioconductor (MAE + viz, iSEE)

Copy number variation analysis (Workshop, DNAcopy, PureCN, CNVRanger, Sehyun, Ludwig)

Approaches to providing reproducible, reusable, robust software for cancer data science (VC)

Bioconductor approaches to single-cell and spatially resolved cancer data and analysis (Workshops, Ludwig, Davide)

Pharmacogenomics resources in Bioconductor (Ben HK)

bibliometrics as code (MM) - maybe

## References