

## Research Leads

## Non-NIH Funding Options

# Lab Notebooks

- [global\\_ph\\_prevalence](#)
- [Ella](#)

# Open Source Resources

## Datasets

- [NCBI Developer Tools](#)
- [NCBI Github](#)
- [Single Cell Atlas](#)
- [PhysioNet](#)
- [UMMC Datasets](#)

## Tools

- [STRING, Protein-Protein Interaction Networks \(PPINs\)](#)
- [Jeremy Warner, HemOnc.org](#)
- [DeepPhe](#)
- [cTAKES](#)
- [Computational Health Informatics Program](#)
- [Galaxy Cloud Computing](#)

# Research Types

## Systematic Reviews

- [Covidence](#)
- [PRISMA](#)
  - [PRISMA Diagram R package](#)

# Coding

## R

- [UMMS CDABS, R Workshop](#)
- [rOpenSci](#)

## python

- **Tutorials**
  - [SciPy](#)
  - [NumPy](#)

## MAKE files

- [Minimal MAKE files](#)
- [MAKE files for Latex](#)
- [MAKE Reproducible Research](#)

## Statistics

- <https://hbiostat.org>
  - [R workflow](#)
  - [Reproducible Research](#)
- [Free R-based Biostats Book](#)
- [Epidemiologist R Handbook](#)
- [Nature: Statistics for Biologists](#)

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