

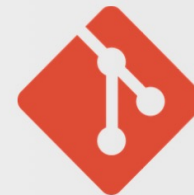
MAY 9-12, 2022

FIFTH ANNUAL

# TODOS SANTOS GENOMICS DATA ANALYSIS WORKSHOP

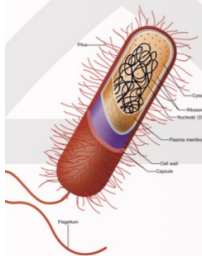
RNA-sequencing  
and Gene Expression

Command Line  
Computing



Genome Assembly and  
Comparative Genomics

Workflow Management  
and Reproducibility





## Learning Objectives for this Workshop

- Understand the principles of genomics and bioinformatics.
- Become familiar with bioinformatics tools and workflows.
- Use genomics resources and computational tools to address biological questions.
- Utilize the command line to interact with a computer.
- Incorporate best practices into your research.

The background of the slide is a dark blue, textured image featuring a large, central virus particle with numerous spikes, resembling a coronavirus. Several other smaller, similar virus particles are scattered in the background, creating a sense of depth and biological context.

# What is bioinformatics?

**Bioinformatics:** Development of computational tools (computer algorithms) and techniques for the analysis of biological data.

**Computational biology:** Applying computational tools to the analysis of biological data to address questions and hypotheses about living systems.



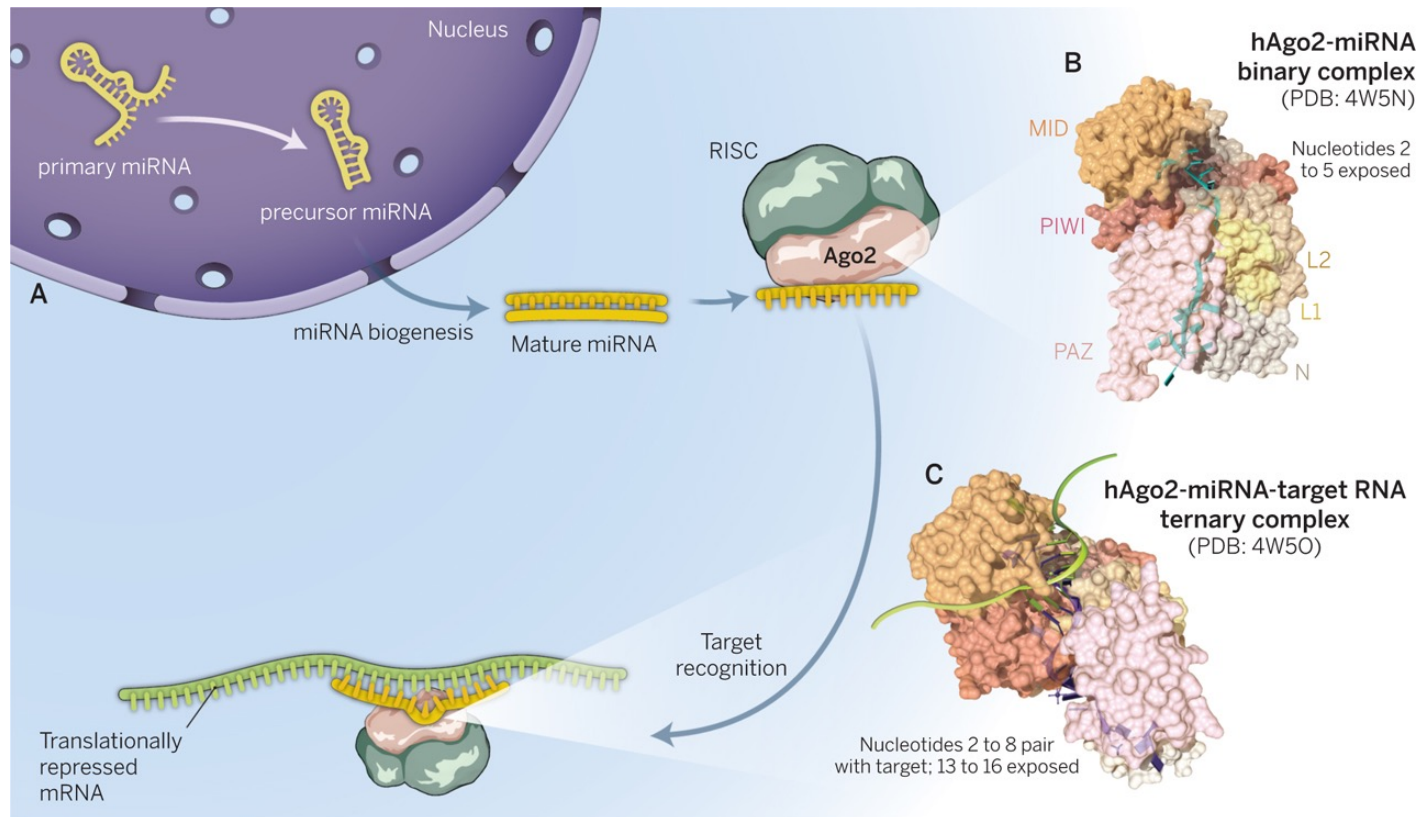


## What is genomics?

Genomics is the application of high-throughput techniques and technologies to examine the sequence, structure, and function of genomes and to analyze gene expression on a genome-wide scale.

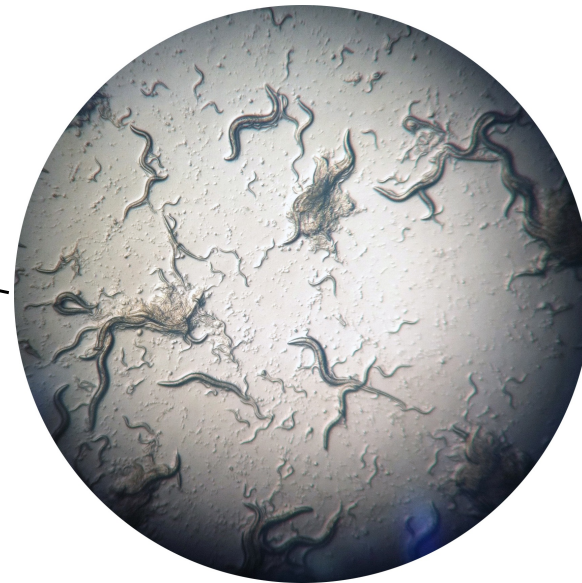
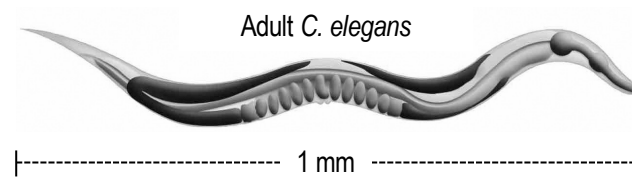
What does my lab study?

## microRNA pathway

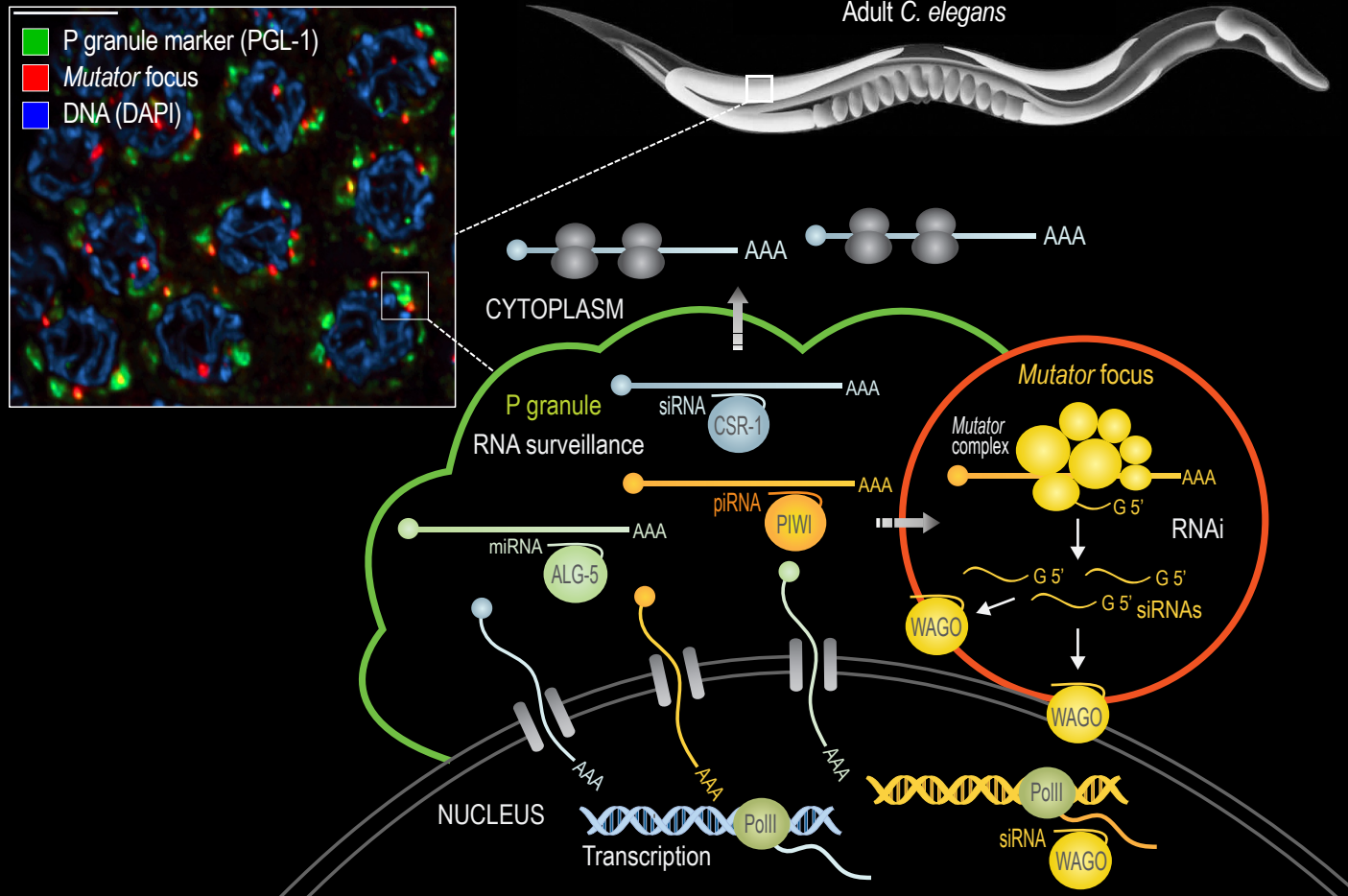


Dinshaw Patel, 2014

## *C. elegans* as a pioneer for small RNA discovery



# RNA surveillance in *C. elegans*





MontgomeryLab.org

