

# **Biological Databases**

# Purpose

In this activity, students will learn to search the same online databases used by scientists to collect information about a set of genes and present them. This will be an opportunity to engage in inquiry-based learning and apply the concepts in molecular biology and genetics from this course.

# Learning Objectives

1. Use online databases to look up information about a gene.

# Activity 1 - Databases

Estimated time: 15 min

### **Instructions**

- 1. Follow these directions to launch your first LearnR tutorial: Biological Databases Tutorial.
- 2. To move through the activities click "Continue" at the bottom of the screen. When you are done with a topic, click "Next Topic" to move on.
- 3. As you complete the tutorial, fill in the table below. This will help you know which database to go back to later on.

### **Questions**

Table 1. Databases

Database	Description
GenBank	
ОМІМ	
Human Protein Atlas	
PDB	

# Activity 2 - FlyBase

Estimated time: 15 min

### **Instructions**

1. Work through the FlyBase Tutorial in SciServer.

# C-MOOR

### miniCURE-RNA-seq

2. To move through the activities click "Continue" at the bottom of the screen. When you are done with a topic, click "Next Topic" to move on.

### **Questions**

2A. What is one question you have about using FlyBase?		
2B. What is something that surprised you or that you found interesting about using FlyBase?		

# Activity 3 - Human Protein Atlas

Estimated time: 15 min

#### **Instructions**

- 1. Work through the Human Protein Atlas in SciServer.
- 2. To move through the activities click "Continue" at the bottom of the screen. When you are done with a topic, click "Next Topic" to move on.

### Questions

3A. What is one question you have about using HPA?		
3B. What is something that surprised you or that you found interesting about using HPA?		

# Activity 4 - Research a Gene!

Estimated time: 45 min

#### **Instructions**

1. Before getting started on this activity, your instructor will assign your group a letter that corresponds to a group of 4 genes.





### miniCURE-RNA-seq

## **Group Assigned Letter**

- 2. <u>Look up your letter here</u> and write the names of the four genes your group is assigned at the top of each column in the table below.
- 3. In your group, assign each student one of the four genes to research.

Individual Assigned Gene

- 4. Use FlyBase to look up the information in Table 2 below.
- 5. Use HPA to look up the information in Table 3 below.

Table 2. FlyBase Information

Category	Information
General Information	
Full Gene Name	
FlyBase ID	
Sequence Location	
Function	
Biological Process	
Cellular Component	
Expression Data	
Anatomical Expression	
Developmental Stage	
Orthologs	
Orthologs in other species	
Human Orthologs	





#### Table 3. Human Protein Atlas

Category	Information
Function	
Is the gene tissue specific? Which tissue?	
Where is it localized in cells?	

# Activity 5 - Present to your Group

Estimated time: 15 min

### **Instructions**

- 1. Present your gene to your group.
- 2. Take turns presenting your genes amongst your group and decide on one gene that you think is the most interesting.

# Activity 6 - Class Presentation

Estimated time:30 mins

### **Instructions**

- 1. With your group, create a short presentation to present your chosen gene to the class.
- 2. Your presentation should have about four slides and be thorough:
  - 1. Slide 1: The GENE you picked to share with your group, your name and date
  - 2. Slide 2 4: Present the information you collected about the gene. For full credit, include relevant images/ diagrams on your slides.
- 3. One student in the group should post your slides on Canvas to the discussion board. Make sure you mention everyone in your group by name so they also get credit for the presentation.

### **Footnotes**

#### Resources

Google Doc







## **Contributions and Affiliations**

- Rosa Alcazar, Ph.D., Clovis Community College
- Katherine Cox, Ph.D., John Hopkins
- Stephanie R. Coffman, Ph.D., Clovis Community College

Last Revised: February 2023

