

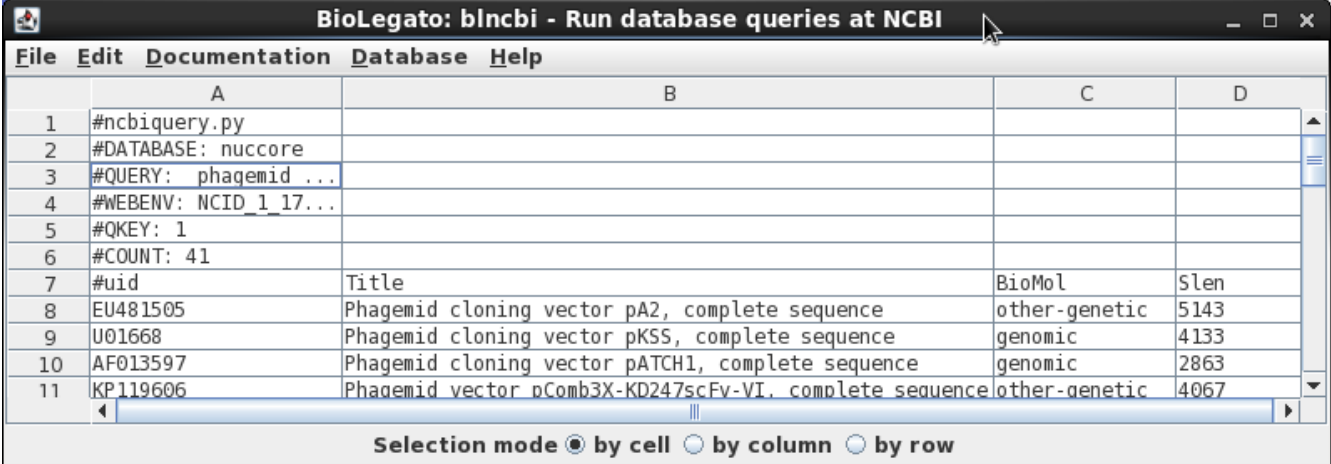
Assignment 2

How to display blncbi results

Because the QUERY term in column A might be very long, copy the term from the blncbi results to your report as text, and then for the screenshot, display the results window such that most of the space is taken up by column B. (Copy the QUERY by clicking on the QUERY cell in column A, typing CTRL-c, and then pasting the text into your report using CTRL-v.)

Using the bluescript example from the tutorial, if query term 1 was set to Title: phagemid and Division: syn, you would display the results as follows:

```
#QUERY: phagemid [TITL] AND syn [Division] AND 1:500000[SLEN]
```



The screenshot shows a window titled "BioLegato: blncbi - Run database queries at NCBI". The window contains a table with the following data:

	A	B	C	D
1	#ncbiquery.py			
2	#DATABASE: nuccore			
3	#QUERY: phagemid ...			
4	#WEBENV: NCID_1_17...			
5	#QKEY: 1			
6	#COUNT: 41			
7	#uid	Title	BioMol	Slen
8	EU481505	Phagemid cloning vector pA2, complete sequence	other-genetic	5143
9	U01668	Phagemid cloning vector pKSS, complete sequence	genomic	4133
10	AF013597	Phagemid cloning vector pATCH1, complete sequence	genomic	2863
11	KP119606	Phagemid vector pComb3X-KD247scFv-VI, complete sequence	other-genetic	4067

At the bottom of the window, there is a selection mode control: "Selection mode" with three radio buttons: "by cell" (selected), "by column", and "by row".

In this example, only a few results are shown. In your report, there should be a small enough number of hits that you can display all hits in the window by adjusting the height of the window.