P.E.T. v2.0 Website User/Admin Guide



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^{*}Super Admin Only

Contact Support

Email: ulmpet@gmail.com

Requirements

You must have one of the following to access the <PhageEnzymeTool_URL> website.

- Google Chrome-version 27 or higher (recommended)
 - Mozilla Firefox-version 22 or higher
 - Microsoft Internet Explorer- version 10 or higher
 - Apple Safari-version 6 or higher
 - Opera-version 12 or higher

Website performance will vary with the speed of your Internet connection.

Overview

This website focused on improving on an existing P.E.T. tool maintained by Paul D. Wiedemeier. You can read more about the research and contributors of the original in the link found below.

<u>A web-based restriction endonuclease tool for mycobacteriophage cluster prediction</u>.

Contributors to the this <PhageEnzymeTool_URL> website are as follows:

Samuel Butler Michael Bruscato Micah Gautreaux Heather Terrell

Accessing the Website

Website URL

Enter www.websitename.com/



Create User Account

1. Click Sign up for an account at the bottom of the login page



2. Captcha

To make sure you are a human, we will need to confirm using a captcha system. Check the "*I'm not a robot*" box. Then type in the characters seen in the image, and submit using the **Verify** button.



3. Enter all of the information required on the Sign Up form and press Submit



Login

To login to your Phage Enzyme Tool account:

- 1. Enter your email address.
- 2. Enter your password.
- 3. Press Log in.



Log Out

There are two ways to log out:

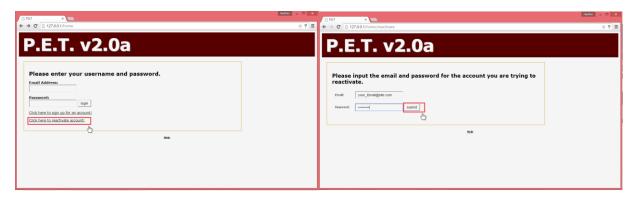
- Click Log out found at the bottom of the left-hand menu.
 -OR-
- Close the browser

Forgot Password

Contact your super admin.

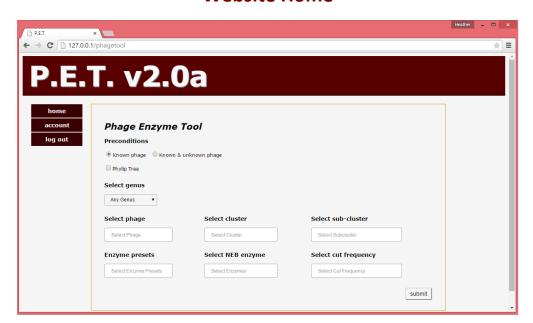
Reactivate Account

If you have previously deactivated an account, go to **Reactivate account** at the login page to do so.



Enter your previous account email and password and click submit.

Website Home



Phage Enzyme Tool

1. Preconditions

Select radio button **Known phage** if you would like the search the database for information about a known phage and its enzyme interactions.

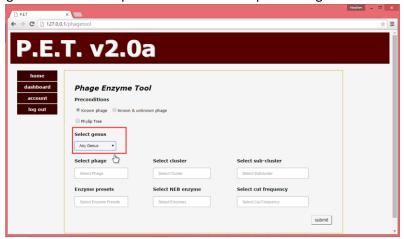
Select radio button **Known & unknown phage** if you would like the search the database for information about known phage and their enzyme interactions and have the phage tool calculate a similarity score (or sim score) based on cut count comparisons.

2. Phylip Tree

Select the **Phylip Tree** option if you would like to have a phylip tree generated from the comparisons of the cuts of the selected phage.

3. Select Genus

Select a genus from the drop down menu where up-to-date genus names can be found:



4. Select Phage

Select one, many, or all phages from the **Select phage** list which will become visible after clicking inside of the **Select phage** text box.

Phages may be unselected by clicking the **x** to the left of the selected phage's name box.

5. Select Cluster

Select one, many, or all cluster options from the **Select cluster** list which will become visible after clicking inside of the **Select cluster** text box.

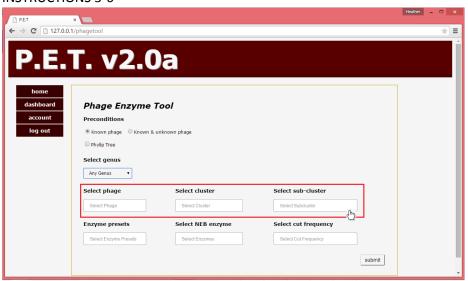
Clusters may be unselected by clicking the ${\bf x}$ to the left of the selected cluster's name box.

6. Select Sub-cluster

Select one, many, or all Sub-cluster options from the **Select cluster** list which will become visible after clicking inside of the **Select cluster** text box.

Sub-clusters may be unselected by clicking the ${\bf x}$ to the left of the selected cluster's name box.

INSTRUCTIONS 3-6



7. Select NEB Enzyme

Select one, many, or all NEB enzyme options from the **Select NEB enzyme** list which will become visible after clicking inside of the **Select cluster** text box.

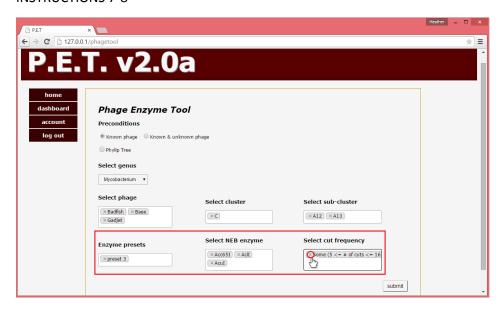
NEB enzymes may be unselected by clicking the \mathbf{x} to the left of the selected enzyme's name box.

8. Select Cut Frequency

Select one, many, or all of the cut frequency options from the **Select cut frequency** list which will become visible after clicking inside of the **Select cluster** text box.

Cut frequencies may be unselected by clicking the \mathbf{x} to the left of the selected frequency name box.

INSTRUCTIONS 7-8



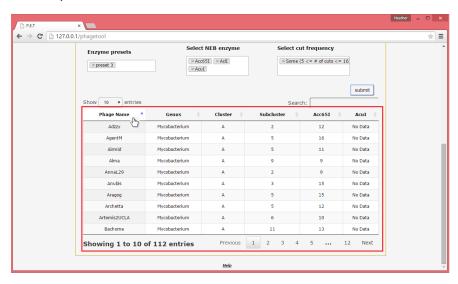
Once you are satisfied with all of your selections, click the **Submit** button to obtain results. The load time of your results may vary.

Viewing Results

Your results will be in a table format. Each row in the table holds the information from one phage and its cut data. The column names indicate what data is being shown or which enzyme created the number of shown cuts.

If you have selected the unknown and **known** information, the **unknown** phage data will always be the first row in the chart and its cuts will be shown as selected ranges. There will also be a similarity score associated with each phage row comparing the percentage of enzymes that cut in the same range as the unknown phage the user entered.

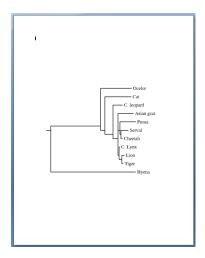
The table is <u>sortable</u> by any column. You can click on any column's header to sort by that criteria alphanumerically.



You can also <u>choose</u> the number of <u>entries</u> shown at a time, as well as <u>search</u> for specific phages.

Your **Phylip tree** results will be output to a pdf file. This tree is generated by the PHYLIP executables who's documentation can be found at http://evolution.genetics.washington.edu/phylip.html. The pdf is generated by ghostScript. An open source PostScript to PDF converter for linux. You can read more about it at http://ghostscript.com/doc/current/Ps2pdf.htm.

Example of a Phylip Tree layout:



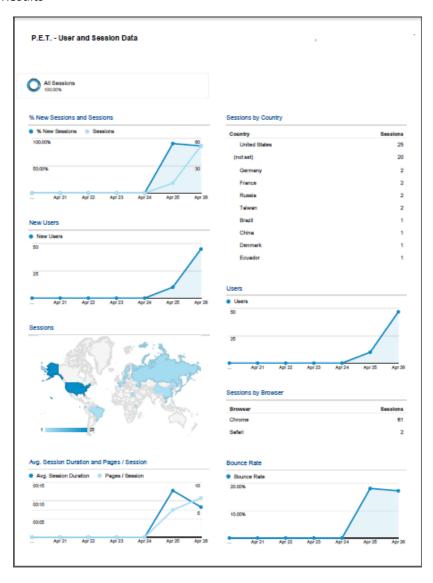
Administrative Dashboard

The Dashboard is available and viewable by administrative users only.

Demographics

The first tab on the Dashboard, **Demographics**, contains a hyperlink to the Google Analytics dashboard for the ulmpet Google account. Where demographic information on the P.E.T. users can be found.

Ex. of Results



Upload to Update

An administrator can update the tool with the latest and most up-to-date data on phage-enzyme findings.

To use the uploader you must first select the type of file you wish to upload.

The **nebCutter** file type must have the correct phage that corresponds with the file selected. This can be accomplished by either selecting the phage from the pre-populated list, or you may type in the name of the phage and it will search for it in the list. This makes the selection process much quicker.

The short **CSV** files require that you select the corresponding Genus that you will be loading. These selections are few and should be quickly found.

The last file type is uploaded by selecting the **FASTA** file that you wish to upload to the database. This process is simply browsing for the file, selecting, and sending it. Once the file is sent, the database will handle the file appropriately and you will be taken back to a fresh page in the dashboard.

"Where do I get the files to upload?"

Fasta Files can be found on phagesDB.org and associated websites.

Short CSV Files are also found on phagesDB.org and accociated websites.

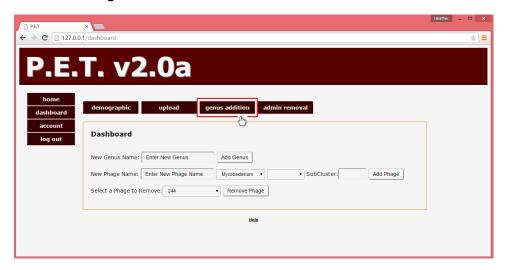
From the Navigation bar click *Data->Download Data* and select an option under "Simple (just phage names and clusters) Tab-Delimited Text Files"

<u>Neb Cutter Data Files</u> are obtained from uploading a *FASTA* file to the nebcutter website. (http://nc2.neb.com/NEBcutter2/). After cutting your phage with the tool click "save all sights" in the leftmost menu at the bottom of the page. This will download a .txt file upload to PET2.



Genus/Phage Addition or Removal

To add a new genus name, click the text box in the Dashboard **Genus Addition** tab, type in new genus name and click the **add genus** button to send it in.



This same step can be repeated for addition or removal of phages.

*Grant/Removal of Administrative Privileges

To grant or remove administrative privilege from any administrator, find their name in the drop down menu of a list of users located under the **Dashboard** \rightarrow **admin removal** tab, and click the **remove admin** or **grant admin** button to finalize your decision.

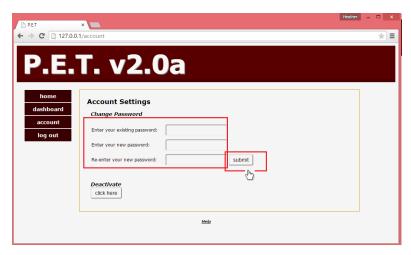


Account Settings

Account settings are found on the page linked to the **Account** menu button.

Change Password

To change your password, go to the menu and select **Account**.

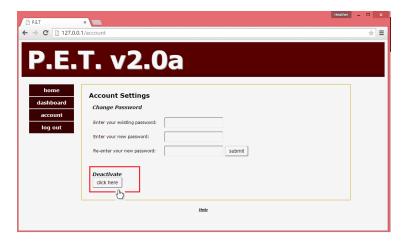


Here you will enter your current password, the new password you wish to change it to, and the new password once again for confirmation. Once you have filled in all three text boxes click the **submit** button to make the change final.

When logging in the next time, remember to enter this new password.

Deactivate Account

To deactivate your existing account, go to the menu and select Account.



At the bottom of the account setting page you will see the words "**Deactivate account**", and a **click here** button following. Once you click, your account will be deactivated.

If you wish to reactivate your account, see the **Reactivate account** option at the login page of the website.

*The super admin cannot remove his/her self