

Step 1: click at the website

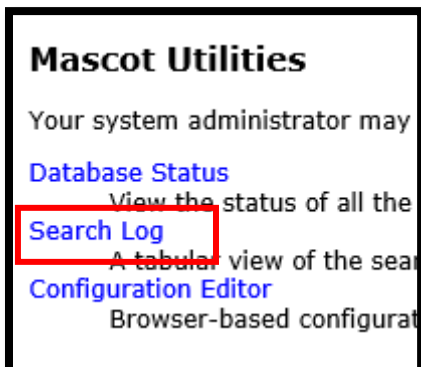
Results can be viewed by going to the web address:
<https://sysbio-mascot.wehi.edu.au/mascot>

Step 2: Login with this userID & password ... then at new page... click "HOME"

Login
Username: PI-Client337
Password: Helotries



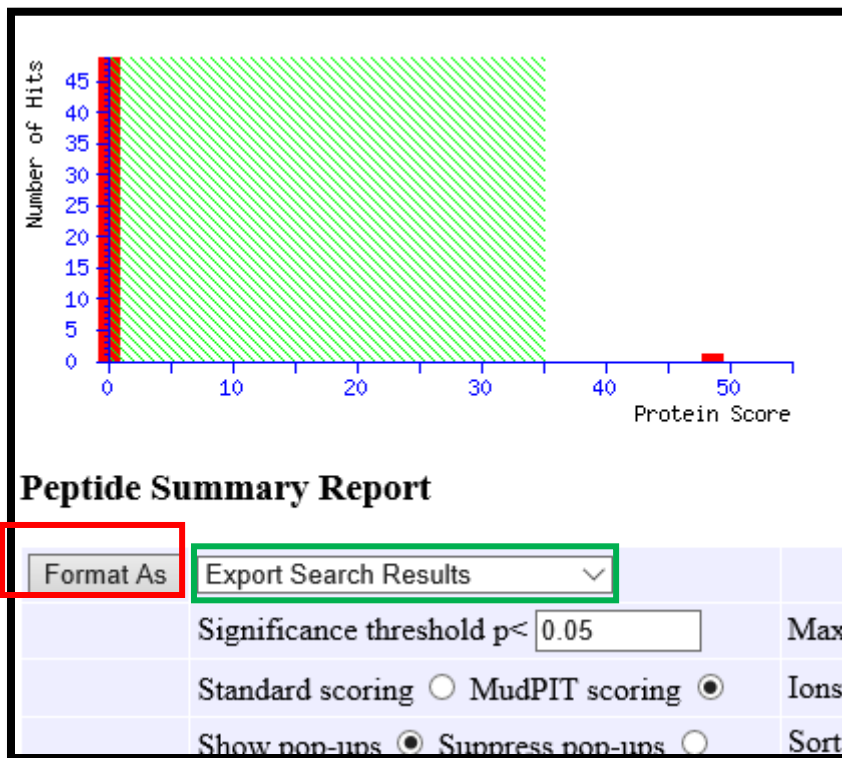
Step 3: Click Search Log



Step 4: refer to the report, click at the Job number that you want

Job#
<input type="radio"/>
<input checked="" type="checkbox"/>
<input type="text"/>
310691
300970
300969
300967
300966
300965
300964
300963
300962
300961
300960
300959
300958
300957
300956
300955
300954
300953

Step 5: At this page, at the green color square box choose “Export Search results” & click “Format As” button



Step 6: another new page pop up and choose “MGF Peak List” then website will auto run

Export search results

Export format	XML CSV pepXML mzIdentML DTASelect Mascot DAT File MGF Peak List
Significance threshold p<	
Ions score cut-off	
Threshold type	Technology <input type="radio"/>
Max. number of hits	AUTO
Protein scoring	Standard <input type="radio"/> MudPIT <input checked="" type="radio"/>

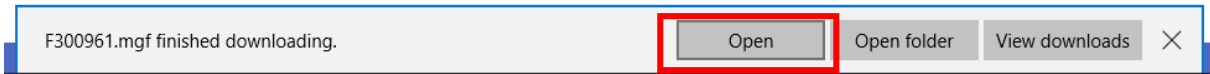
Step 7: Click at the “Export search results” button

Export search results

[Help](#)

Export format	MGF Peak List
Show command line arguments	Export search results

Step 8: if you are using windows operating system, at the end of the screen will see the raw data file been downloaded and you just click "Open".



Step 9: This is the raw data file:

