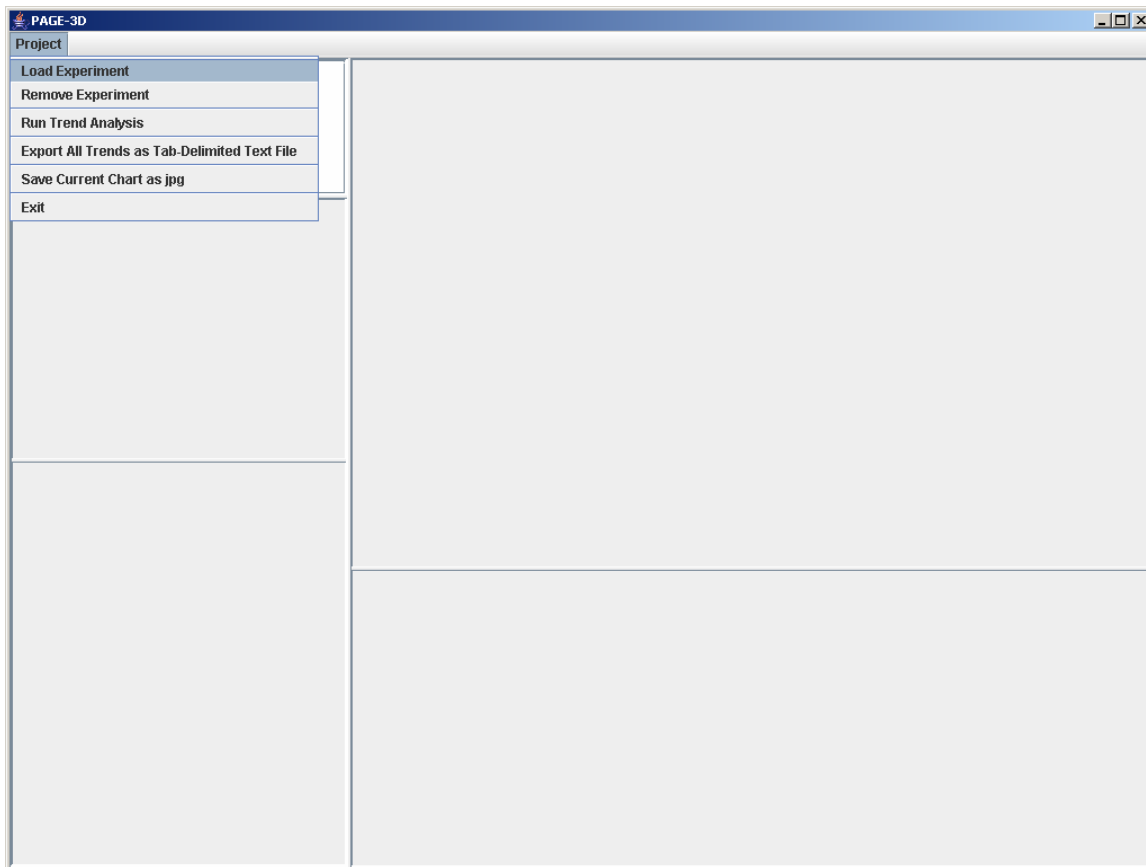
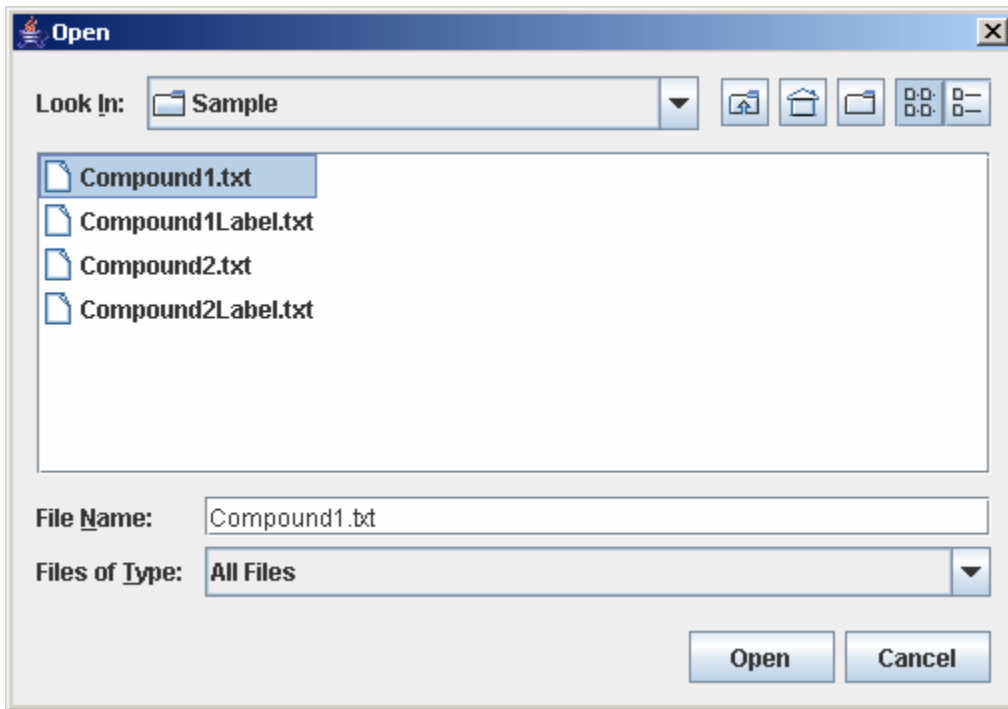


# Quick Start Guide

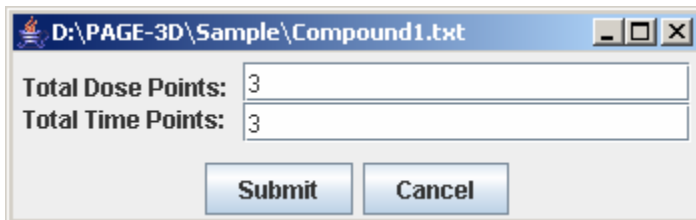
1. Double click on PAGE.bat
2. Click on Project and select Load Experiment



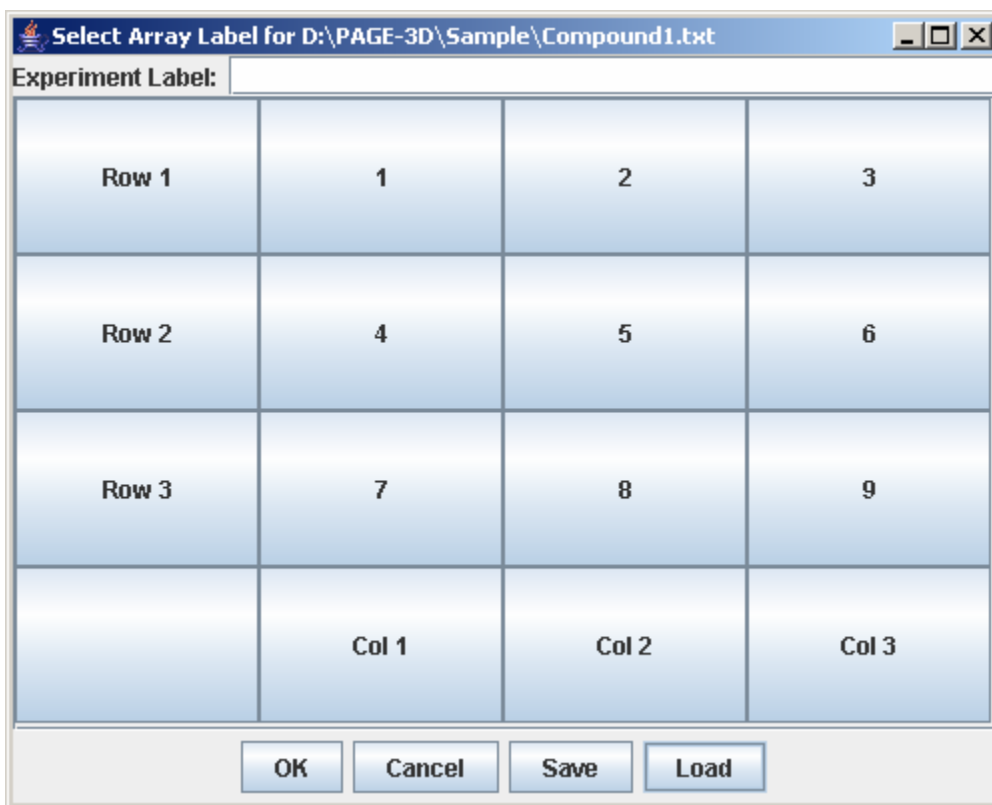
3. Select Compound1.txt and click Open



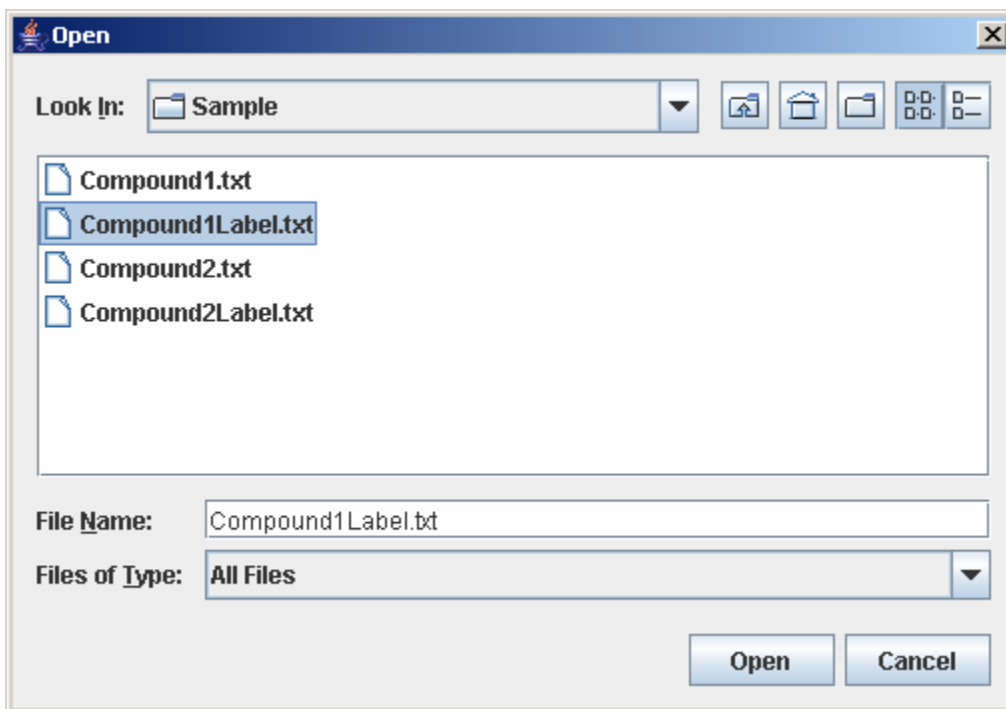
4. Enter 3 for total dose points and total time points. Then, click Submit



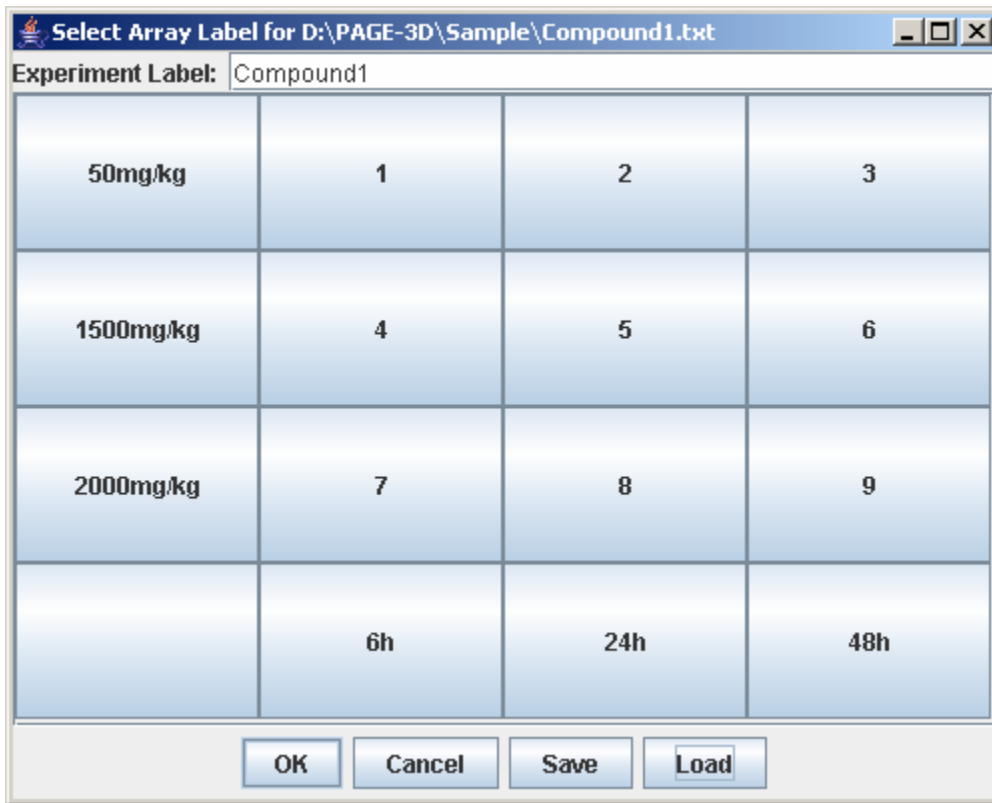
5. Click on Load



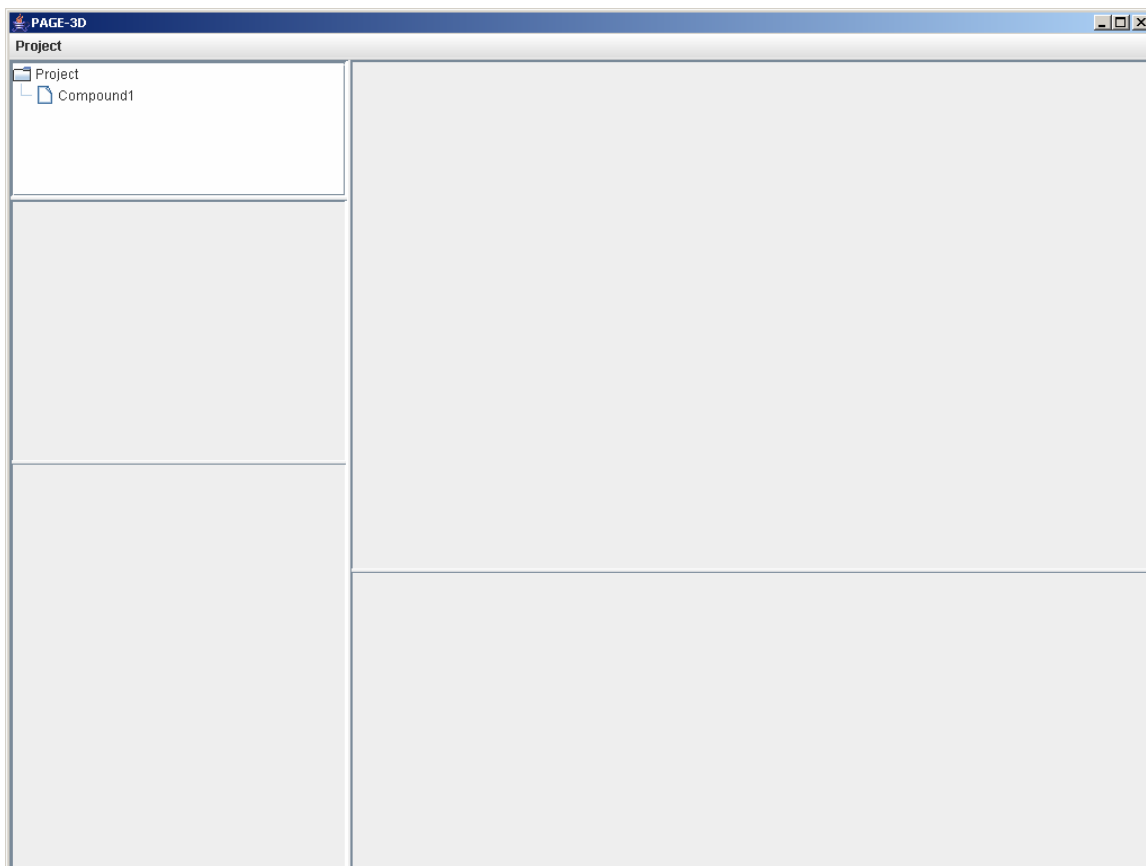
6. Select Compound1Label.txt and click Open



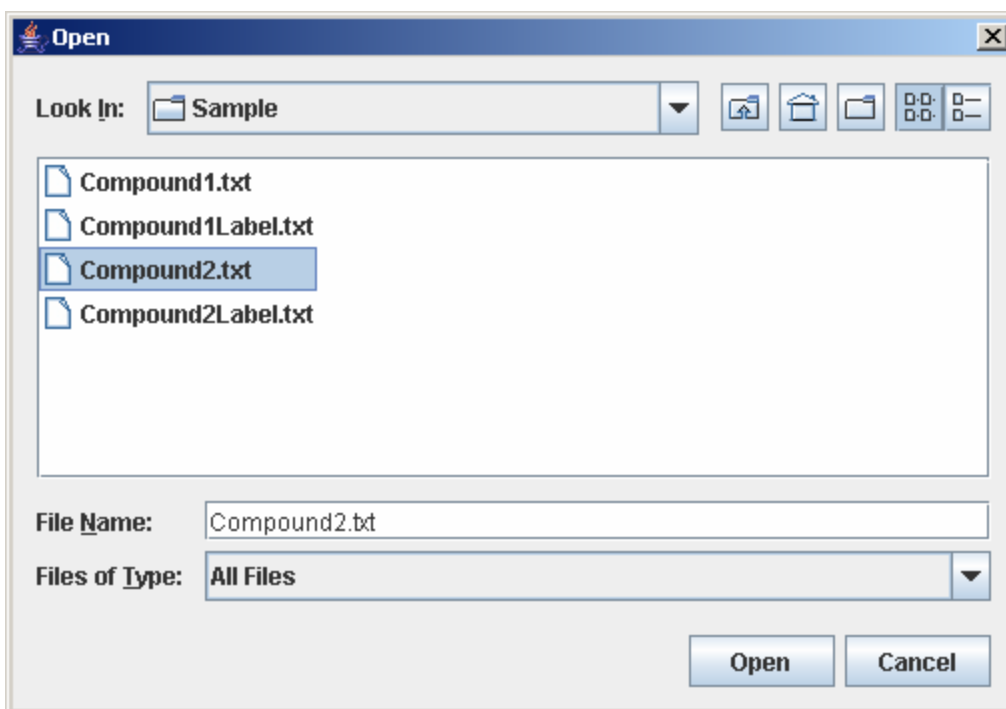
7. Click OK



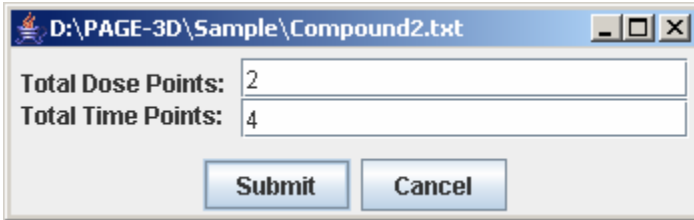
8. Click on Project and select Load Experiment again



9. Select Compound2.txt and click Open



10. Enter 2 for total dose points and 4 for total time points. Then, click Submit.



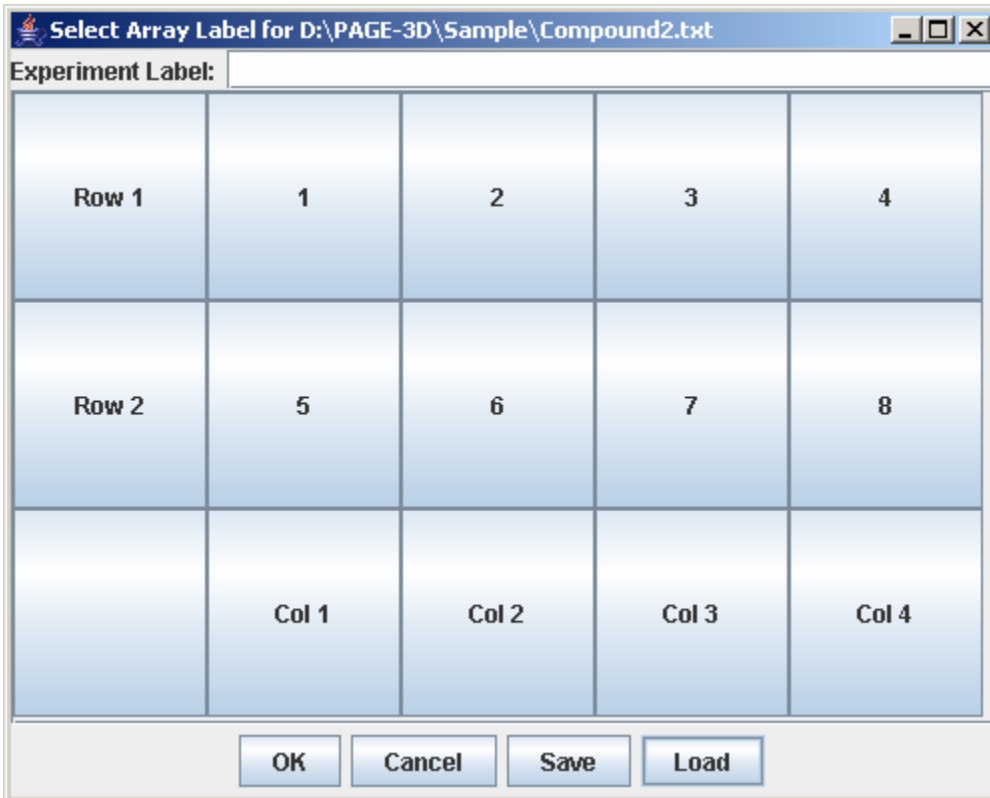
D:\PAGE-3D\Sample\Compound2.txt

Total Dose Points: 2

Total Time Points: 4

Submit Cancel

11. Click on Load



Select Array Label for D:\PAGE-3D\Sample\Compound2.txt

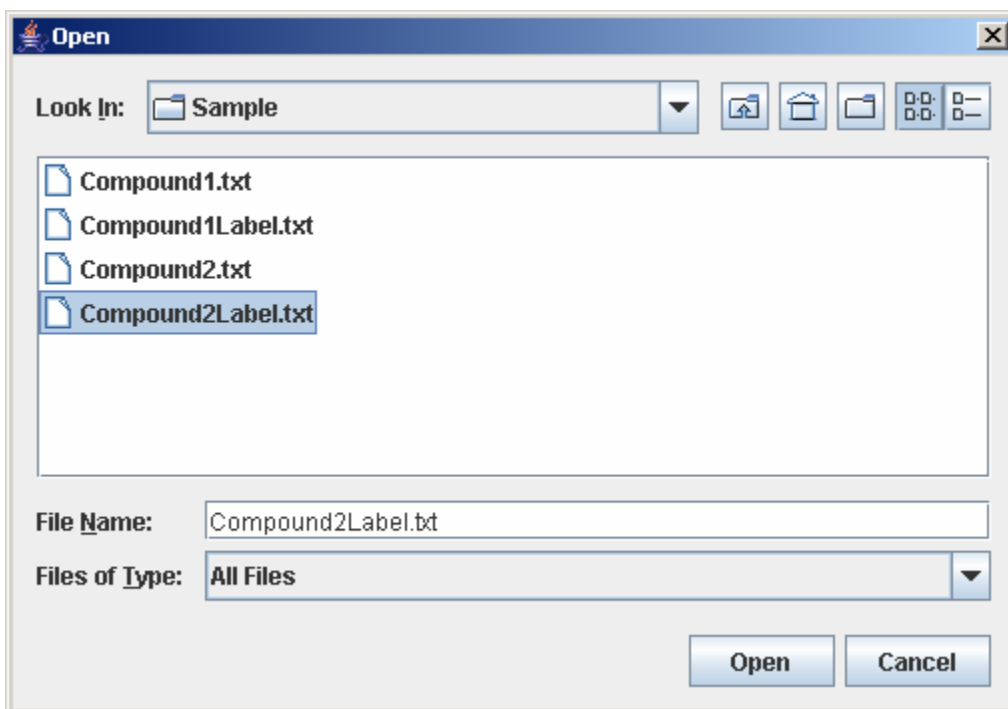
Experiment Label:

Row 1	1	2	3	4
Row 2	5	6	7	8

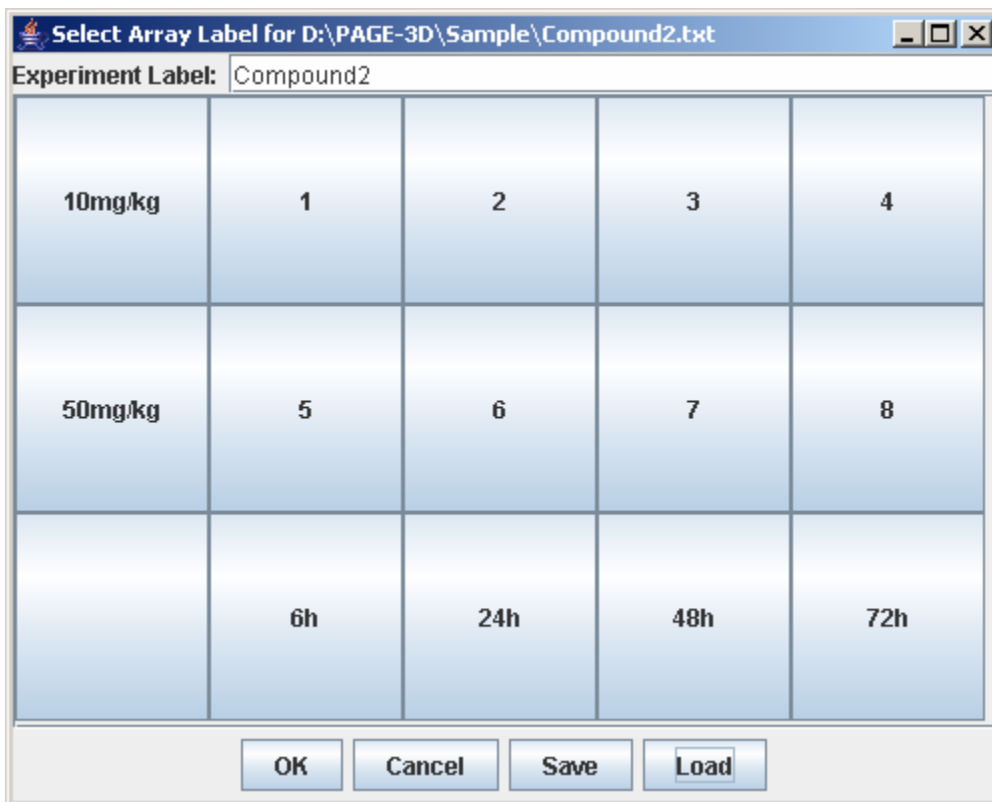
Col 1 Col 2 Col 3 Col 4

OK Cancel Save Load

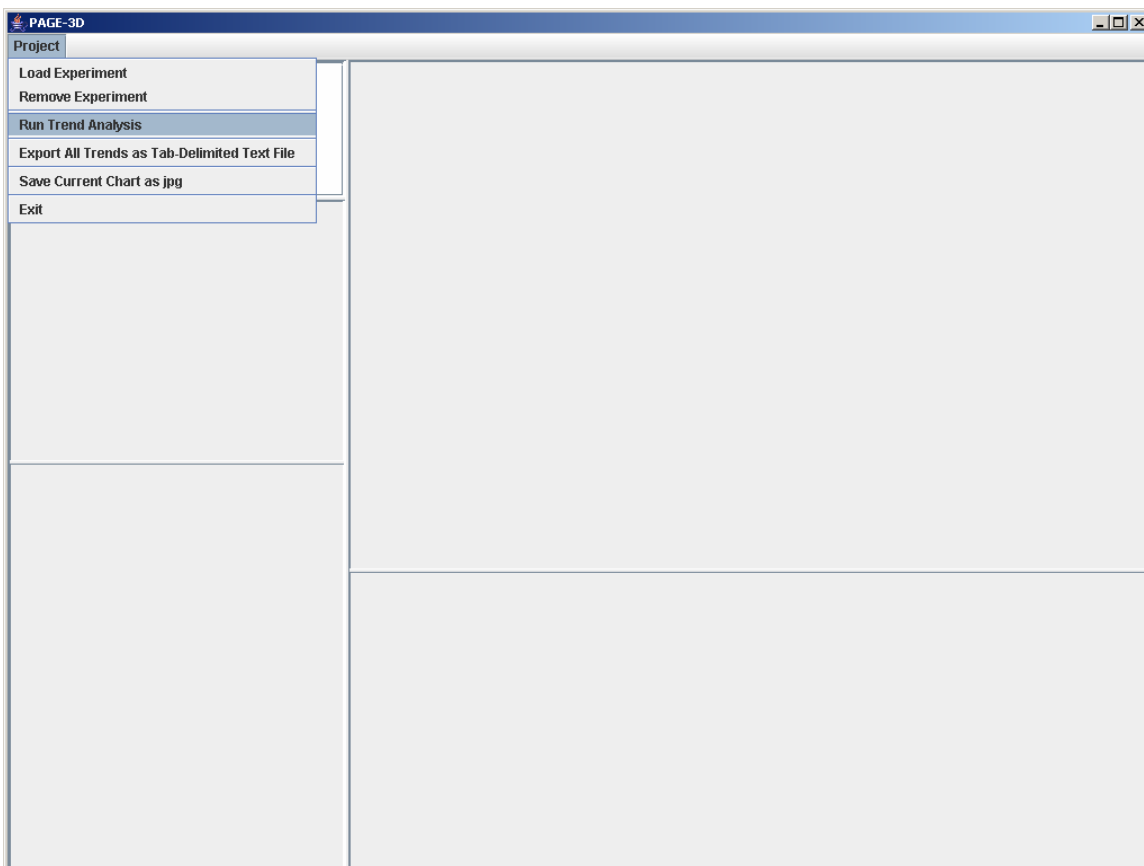
12. Select Compound2Label.txt and click Open



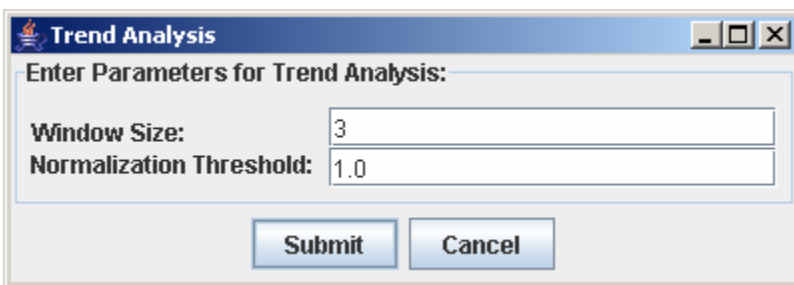
13. Click OK



14. Click on Project and select Run Trend Analysis.

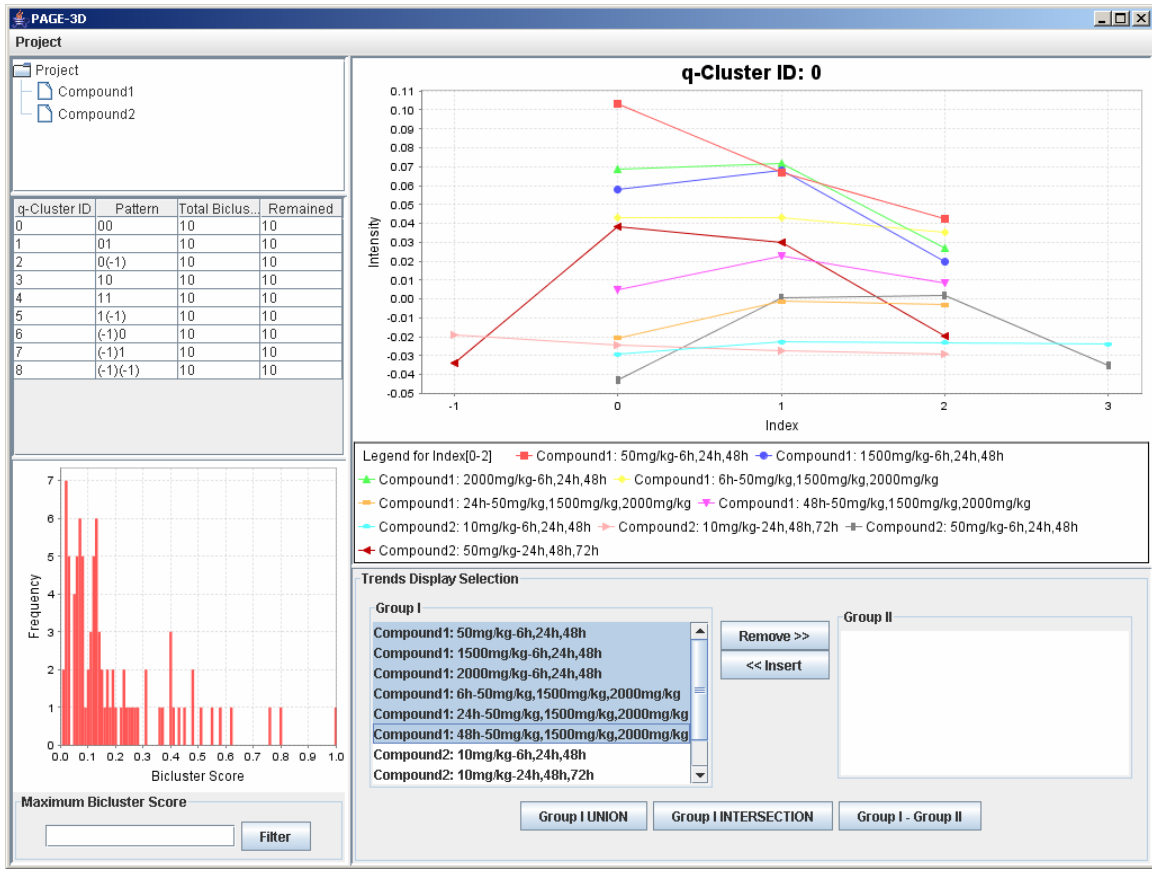


15. Enter 3 for window size. Then, click Submit.

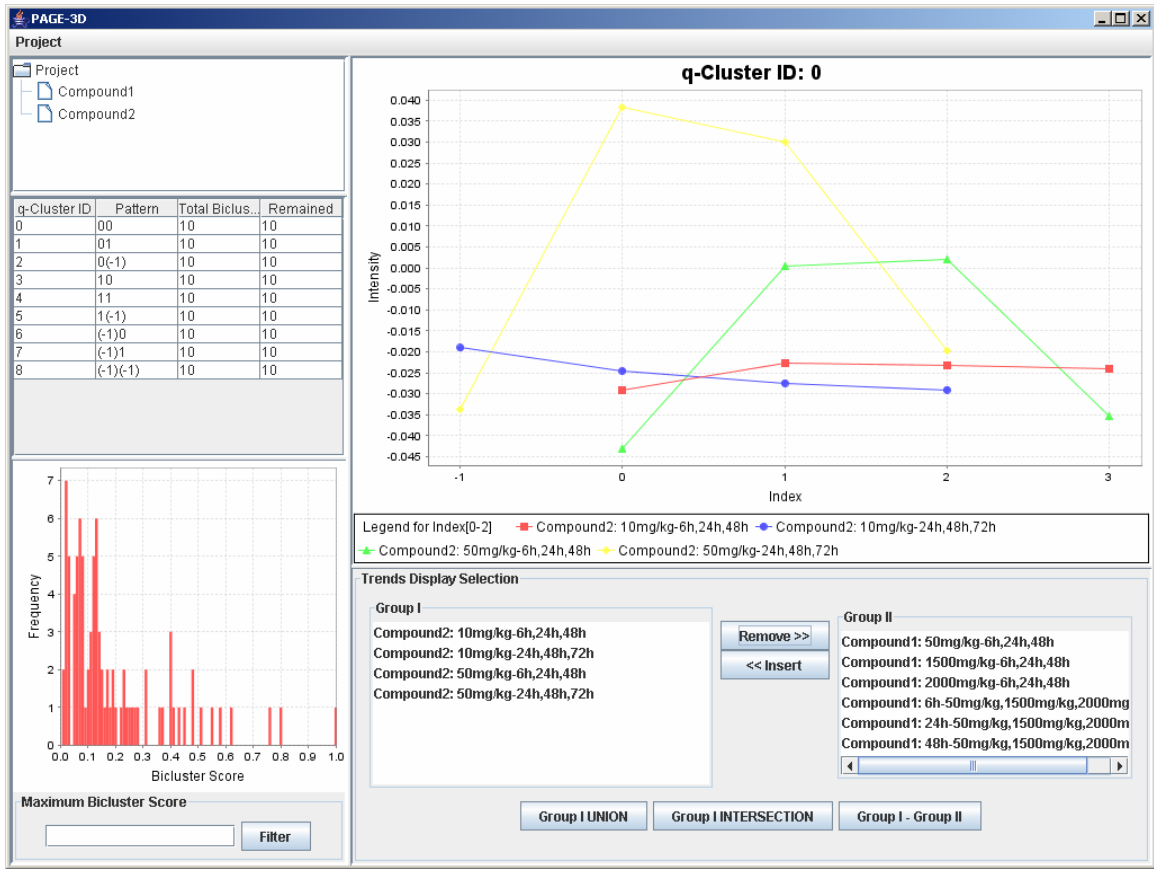


16. Select all the trends from Compound1 in the Trend Display Selection Panel





17. Click the “Remove >>” button.



18. Click on the “Group I Union” button

Group I Union

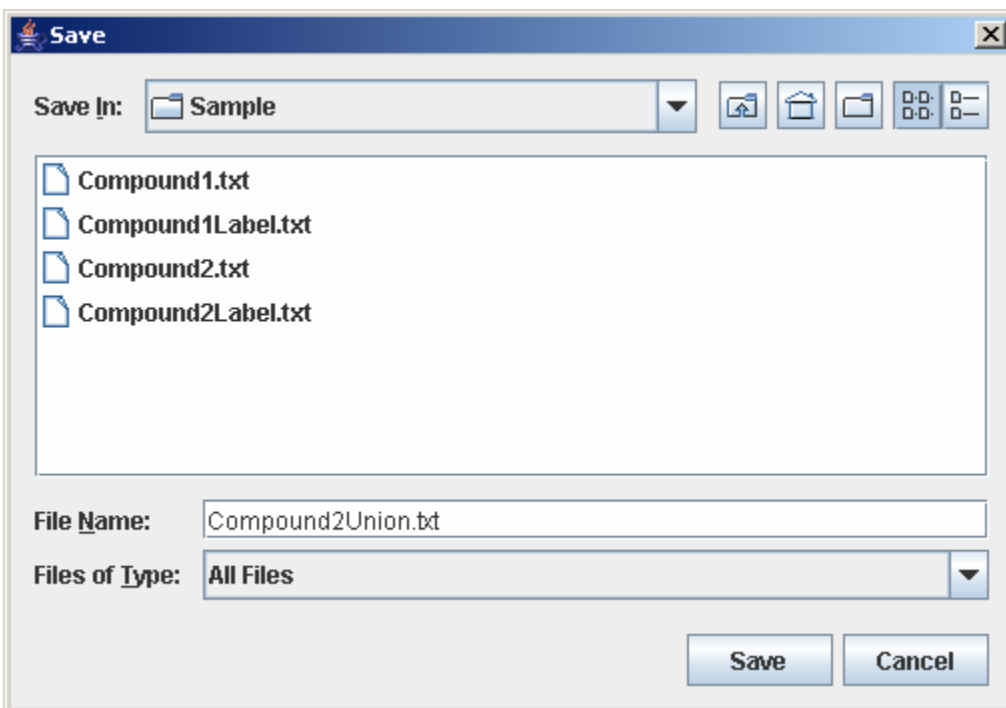
File

Row ID	Gene Accession
3	Gene 3
4	Gene 4
10	Gene 10
13	Gene 13
14	Gene 14
18	Gene 18
20	Gene 20
21	Gene 21
27	Gene 27
32	Gene 32
33	Gene 33
37	Gene 37
38	Gene 38
39	Gene 39
46	Gene 46

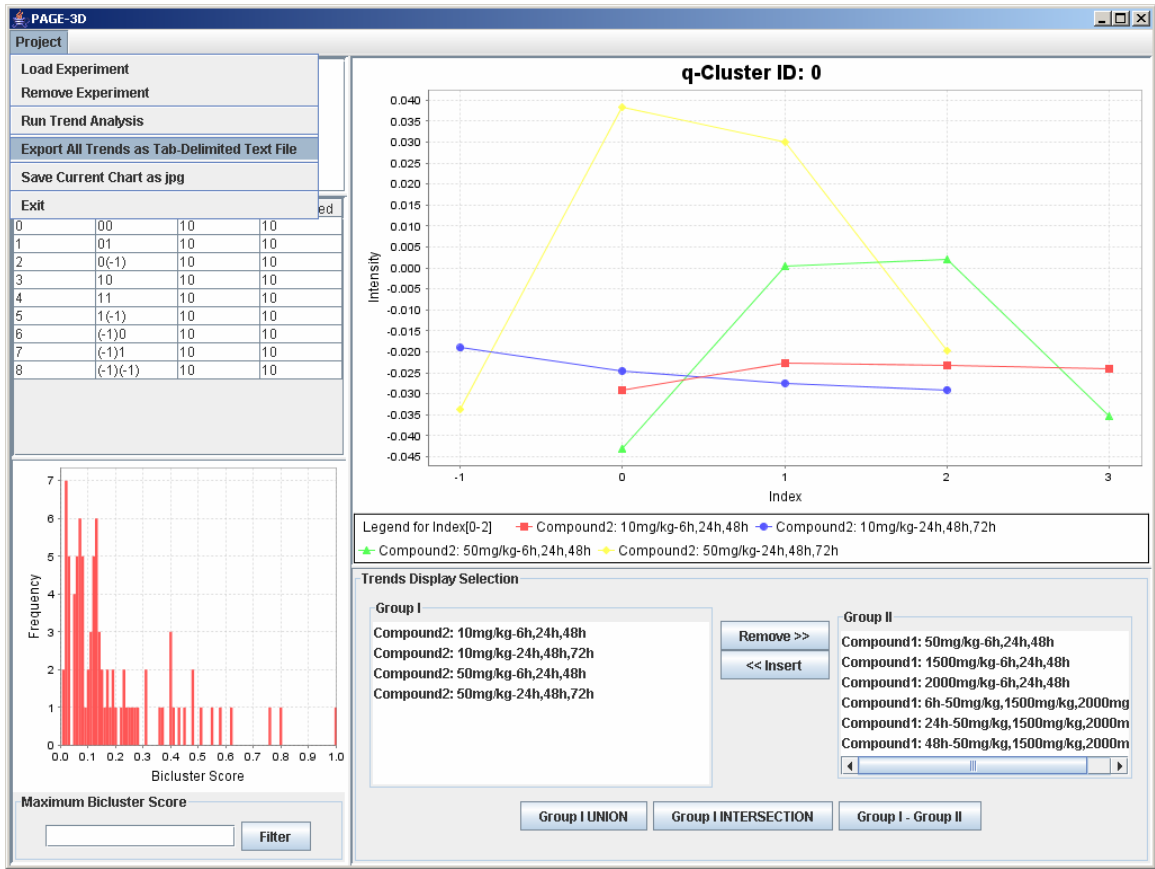
19. Click on File and select Save As Tab-Delimited

	Gene Accession
3	Gene 3
4	Gene 4
10	Gene 10
13	Gene 13
14	Gene 14
18	Gene 18
20	Gene 20
21	Gene 21
27	Gene 27
32	Gene 32
33	Gene 33
37	Gene 37
38	Gene 38
39	Gene 39
46	Gene 46

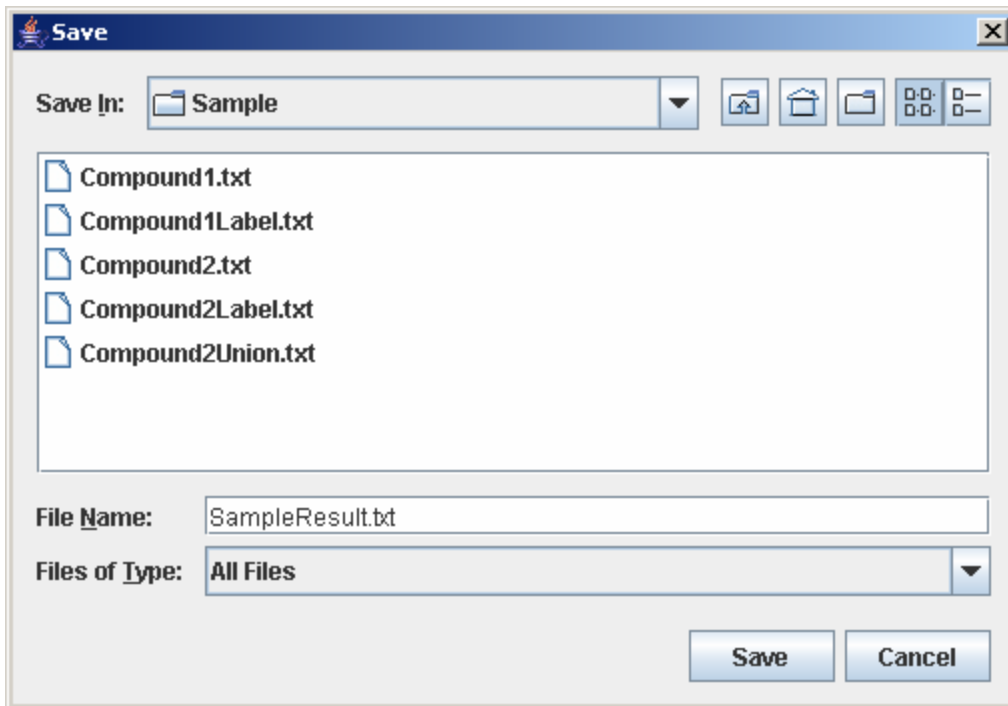
20. Enter Compound2Union.txt and click Save.



21. Click on Project and select Export All Trends as Tab-Delimited Text File



22. Enter SampleResult.txt and click Save



23. Click on Project and select Save Current Chart as jpg

PAGE-3D

Project

Load Experiment  
Remove Experiment  
Run Trend Analysis  
Export All Trends as Tab-Delimited Text File  
Save Current Chart as jpg  
Exit

0	00	10	10
1	01	10	10
2	0(-1)	10	10
3	10	10	10
4	11	10	10
5	1(-1)	10	10
6	(-1)0	10	10
7	(-1)1	10	10
8	(-1)(-1)	10	10

Frequency

Bicluster Score

Maximum Bicluster Score

Filter

**q-Cluster ID: 0**

Intensity

Index

Legend for Index[0-2]

- Compound2: 10mg/kg-6h,24h,48h
- Compound2: 10mg/kg-24h,48h,72h
- Compound2: 50mg/kg-6h,24h,48h
- Compound2: 50mg/kg-24h,48h,72h

Trends Display Selection

Group I

- Compound2: 10mg/kg-6h,24h,48h
- Compound2: 10mg/kg-24h,48h,72h
- Compound2: 50mg/kg-6h,24h,48h
- Compound2: 50mg/kg-24h,48h,72h

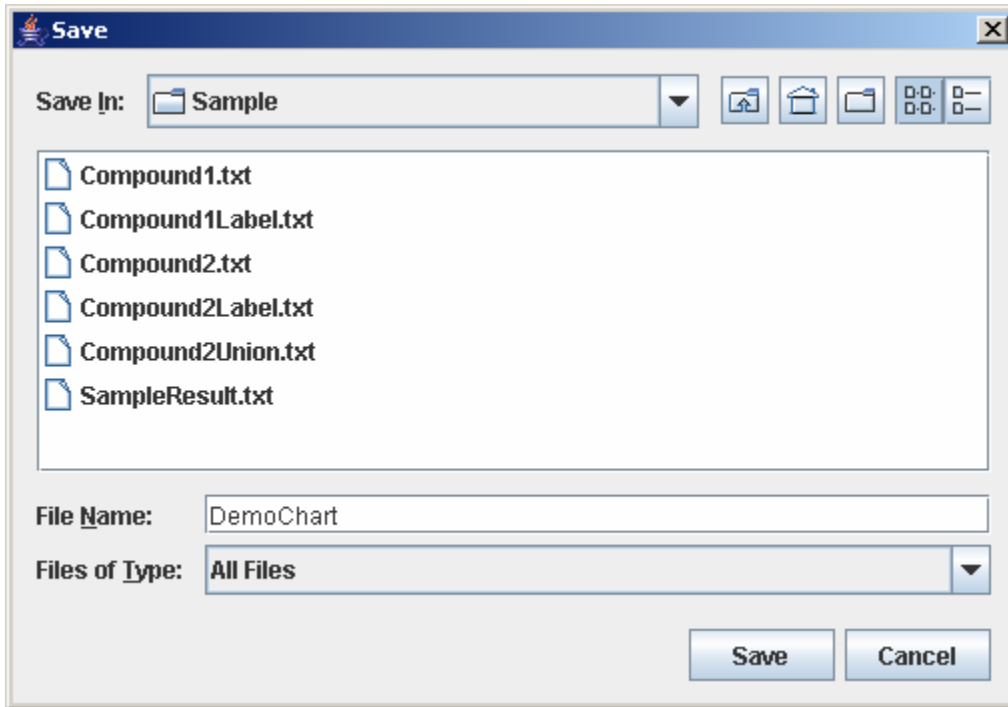
Remove >>  
<< Insert

Group II

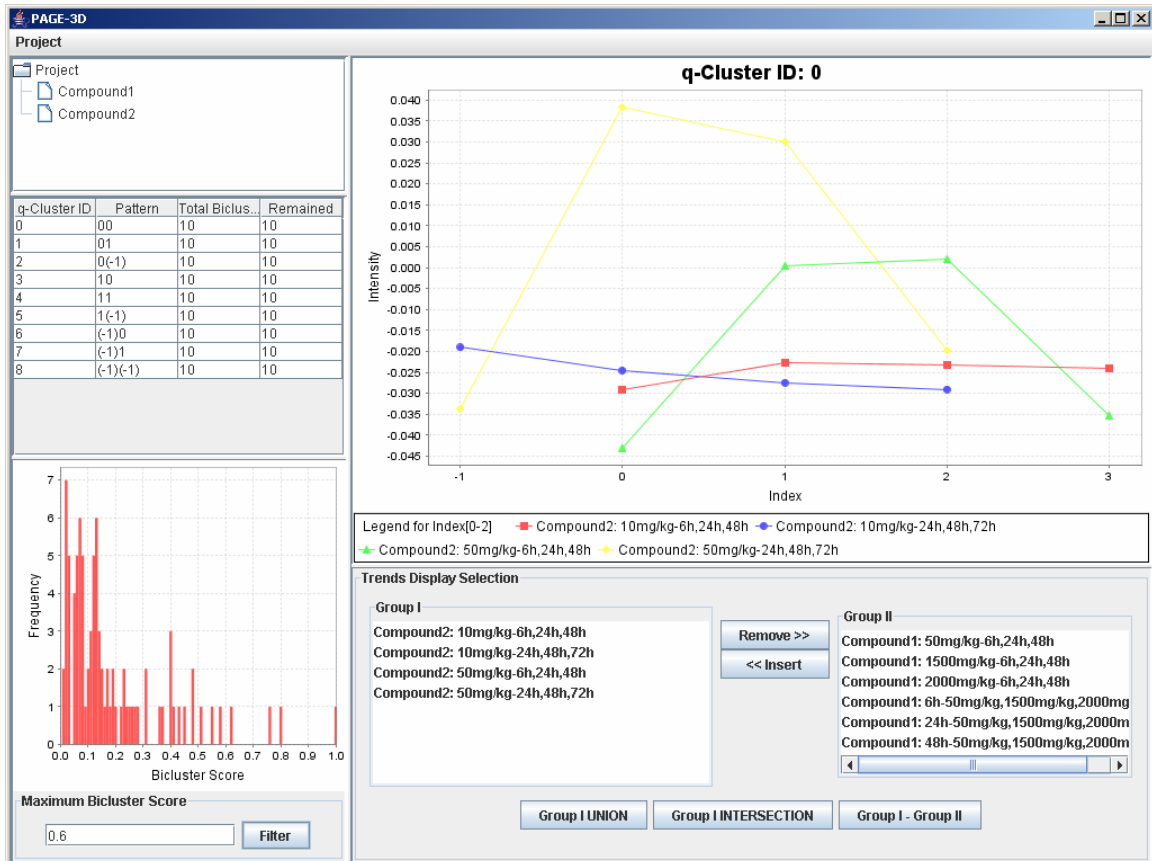
- Compound1: 50mg/kg-6h,24h,48h
- Compound1: 1500mg/kg-6h,24h,48h
- Compound1: 2000mg/kg-6h,24h,48h
- Compound1: 6h-50mg/kg,1500mg/kg,2000mg
- Compound1: 24h-50mg/kg,1500mg/kg,2000m
- Compound1: 48h-50mg/kg,1500mg/kg,2000m

Group I UNION    Group I INTERSECTION    Group I - Group II

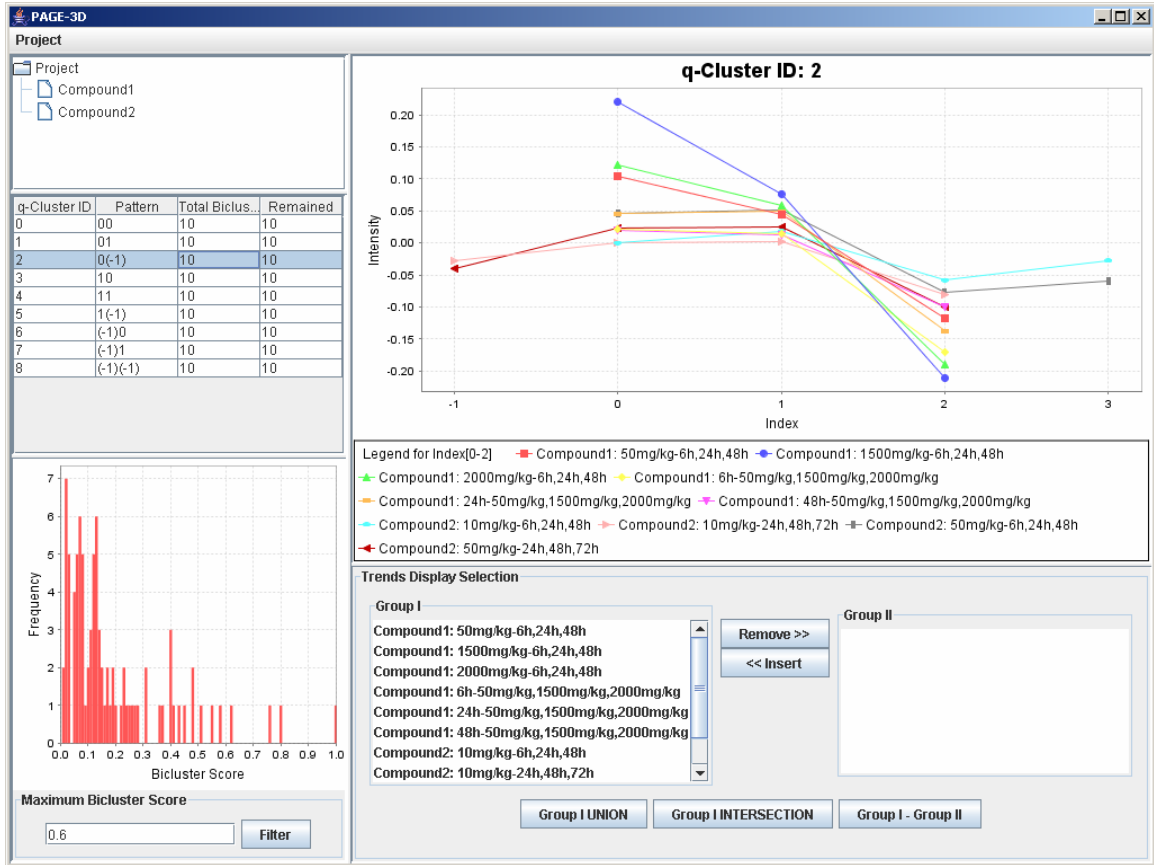
24. Enter DemoChart and click Save



25. Enter 0.6 for Maximum Bicluster Score in the lower left panel and click Filter



26. Click on q-ClusterID 2 in the Table listing the q-Cluster Ids.



27. Click on Project and select Exit.

