



Repository for
Targeted Proteomics Assays

Josh Eckels

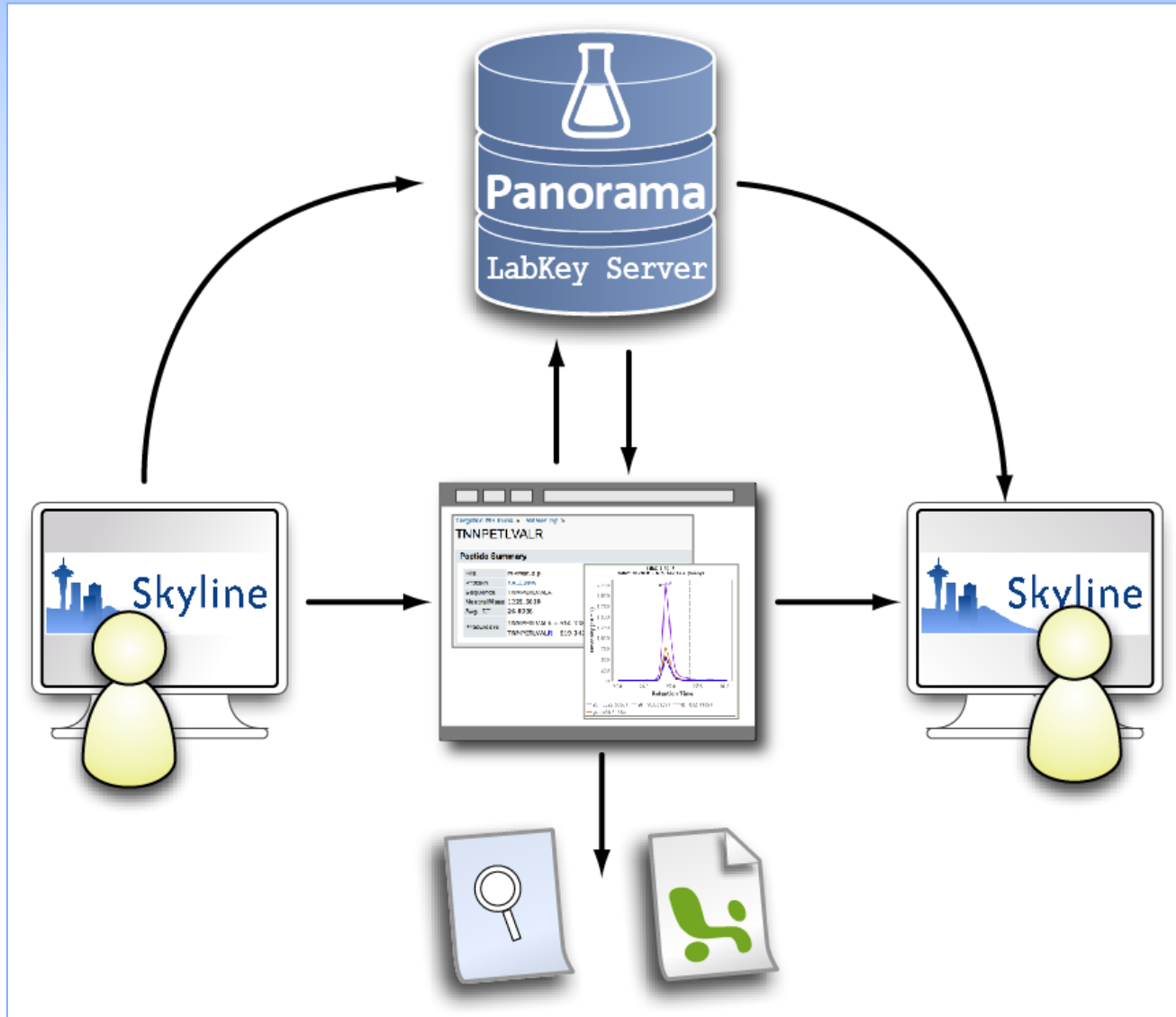
jeckels@labkey.com

Skyline Users Group - June 9, 2013

Goals

- Organize and reuse targeted assays
 - Search and review a large body of data
 - Use previous results to inform new experiments
- Facilitate data sharing
 - Private, with access for internal/external collaborators
 - Public, for published data
- Excellent integration with Skyline
 - Import and retain the complete data model
 - Support data from various workflows

Skyline/Panorama Workflow

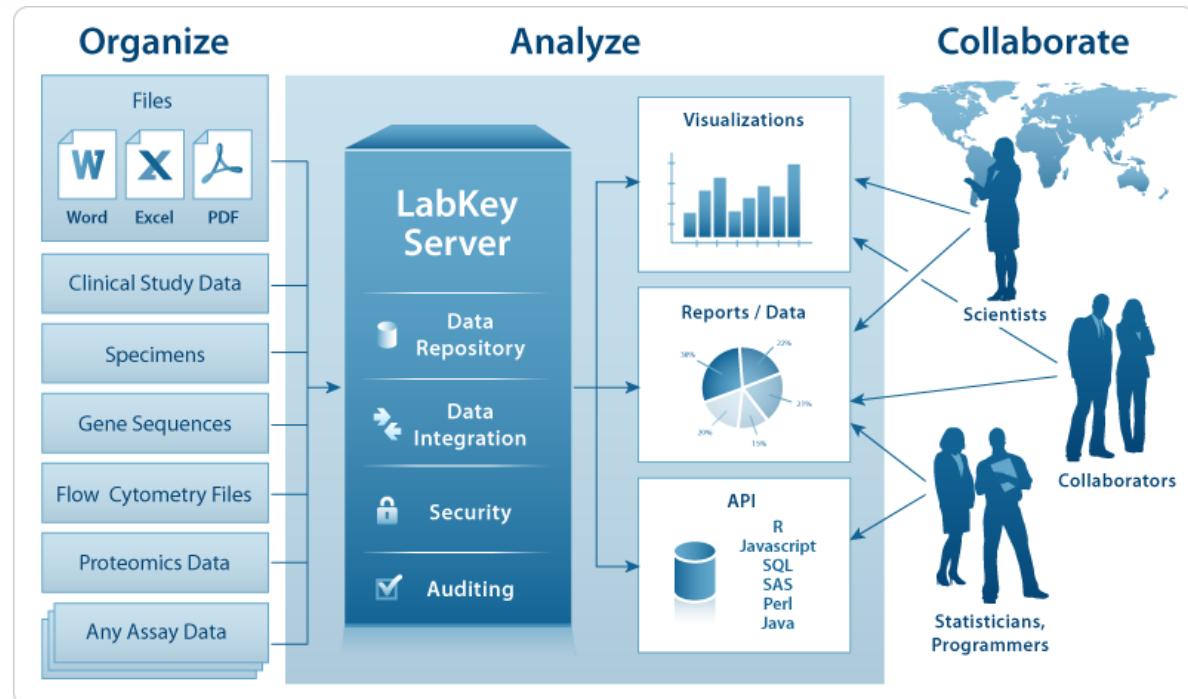


History

- Work started in late 2011
- Announced at ASMS 2012
- Publicly released in December 2012
 - Part of LabKey Server version 12.3
- Released to PanoramaWeb.org in February 2013
 - Hosted server managed by MacCoss Lab
 - Recorded webinar has full details
- Second public release in April 2013
 - Added search features, reporting enhancements, fixes
- Next release scheduled for July 2013

LabKey Server

- Open source (Apache 2.0 license)
- Web user interface
- Powerful security and permissions system
- Other assay data
 - MS/MS peptide search
 - Flow cytometry
 - Sequencing
 - Plate-based assays
- Specimen tracking
- Clinical study data
- Collaboration tools
- Extensive client APIs



LabKey Software

- Co-developers with MacCoss Lab on Panorama
- Consulting company focused on LabKey Server
 - 25 employees in Seattle and San Diego
 - 8 years history working with scientists
- Strength in computer science and data management
- Partner with researchers to develop software
- Related service offerings include:
 - Training
 - Technical support
 - Customization
 - Hosting



Panorama - Setup

The screenshot shows a web browser window displaying the 'Configure Targeted MS Folder' setup page on PanoramaWeb (beta). The browser's address bar shows the URL: <https://panoramaweb.org:9443/labkey/targetedms/MacCoss/brendanx/Josh%27s%20TargetedMS/setup.view?wizard=true>. The page features a navigation sidebar on the left with three steps: 1. Create Folder, 2. Users / Permissions, and 3. Configure Targeted MS Folder (which is currently selected and highlighted). The main content area is titled 'Configure Targeted MS Folder' and offers two radio button options: 'Experimental data' (a collection of published Skyline documents for various experimental designs) and 'Chromatogram library' (curated precursor and product ion expression data for use in designing and validating future experiments). A checked checkbox option is present: 'Rank peptides within proteins by peak area'. A 'FINISH' button is located in the bottom right corner of the main content area. The browser window also shows other tabs like 'Create Folder' and 'Users / Permissions'.

1. Create Folder

2. Users / Permissions

3. **Configure Targeted MS Folder**

Configure Targeted MS Folder

Experimental data - a collection of published Skyline documents for various experimental designs

Chromatogram library - curated precursor and product ion expression data for use in designing and validating future experiments

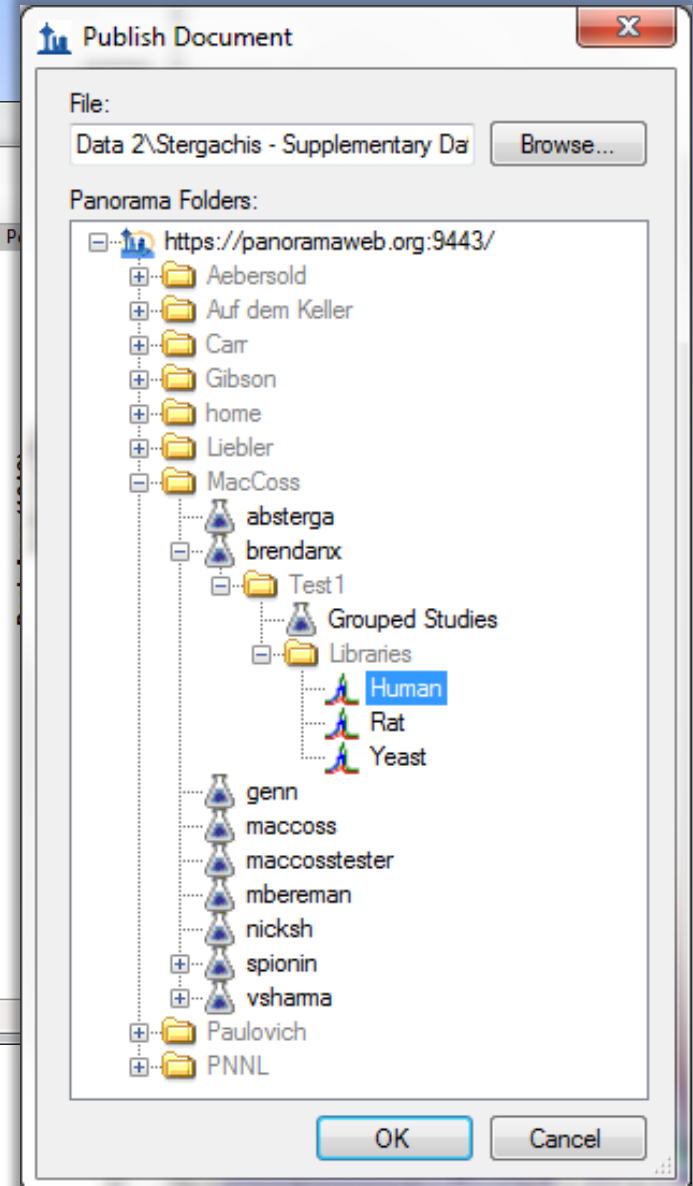
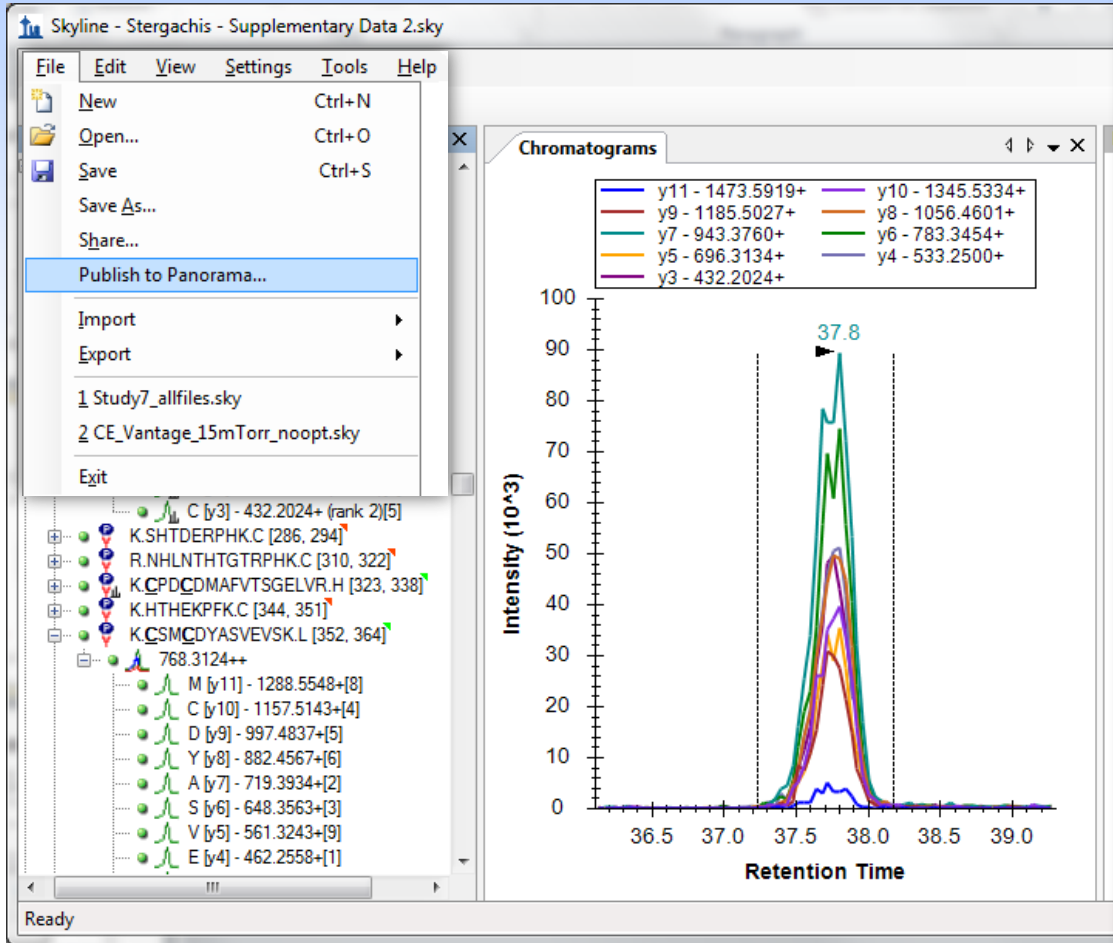
Rank peptides within proteins by peak area

[FINISH](#)

Publishing to Panorama - Setup

The screenshot displays the Skyline software interface for 'Supplementary Data 2.sky'. The 'Edit Servers' dialog box is open, showing a list of servers with 'https://panoramaweb.org:9443/' selected. The 'Edit Server' sub-dialog is also visible, showing the URL, username 'jeckels@labkey.com', and a masked password. The background shows a chromatogram with 'Intensity (10^{^3})' on the y-axis and 'Retention Time' on the x-axis. A peptide list is visible at the bottom right, including peptides like TFQ, YAL, CSM, YHD, WVG, YCD, YPCD, FHC, CDQ, FTQ, QLL, SDLG, HAD, DPDY, QHSY, THTGKPYA, HTH, THSG, NHLN, SHTD, MEG, SHTG, and HTE.

Publishing to Panorama



Viewing Runs - Precursors

Stergachis - Supplemental x

https://panoramaweb.org:9443/labkey/targetedms/MacCoss/brendanx/Test1/Libraries/Human/showPrecursorList.view?id=229

Human Targeted MS Dashboard Data Pipeline +

Targeted MS Runs >

Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip

Document Summary

Name: Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip [DOWNLOAD](#)

Protein Count: 96 Peptide Count: 1475

Precursor Count: 1475 Transition Count: 10806

[RENAME](#)

Precursor List

VIEWS CHARTS EXPORT PRINT PAGE SIZE 1 - 10 of 96 Next Last

Protein / Label	Description	Protein Annotations									
<input type="checkbox"/> TAF11	HsCD00301672 (1-A03)										
Peptide	Peptide Annotations	Missed Cleavages	Peptide Neutral Mass	Rank	Precursor	Precursor Annotations	Label	Precursor Neutral Mass	Q1 m/z	Q1 Z	Transition Count
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK		light	1230.4823	616.2484	2+	7
EAAAEEGELESQDVSDLTTVER	Quality Score: 4	0	2377.077		EAAAEEGELESQDVSDLTTVER		light	2377.0769	1189.5457	2+	11
EDSLLNPAAK	Quality Score: 1	0	1143.5771		EDSLLNPAAK		light	1143.5771	572.7958	2+	7
VDEDEIQK	Quality Score: 2	0	974.4556		VDEDEIQK		light	974.4556	488.2351	2+	4
MQILVSSFEEQLNR	Quality Score: 1	0	1779.8826		MQILVSSFEEQLNR		light	1779.8826	890.9485	2+	10
LIQSITGTSVSNVVIAMSGISK	Quality Score: 1	0	2332.267		LIQSITGTSVSNVVIAMSGISK		light	2332.2671	1167.1409	2+	12
VFVGEVVEEALDVCEK	Quality Score: 1	0	1820.8866		VFVGEVVEEALDVCEK		light	1820.8866	911.4506	2+	11
WGEMPLLQPK	Quality Score: 1	0	1181.5903		WGEMPLLQPK		light	1181.5903	591.8024	2+	6
GQIPNSK	Quality Score: 4	0	742.39734		GQIPNSK		light	742.3973	372.2060	2+	3
<input type="checkbox"/> POLR2F	HsCD00301192 (1-A06)										
Peptide	Peptide	Missed	Peptide Neutral	Rank	Precursor	Precursor	Label	Precursor Neutral	Q1 m/z	Q1	Transition Count

Viewing Runs - Transitions

Stergachis - Supplemental x

https://panoramaweb.org:9443/labkey/targetedms/MacCoss/brendanx/Test1/Libraries/Hum...

PanoramaWeb (beta)

Admin Help jeckels

Human

Targeted MS Runs >

Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip

Document Summary

Name	Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip	DOWNLOAD
Protein Count	96	Peptide Count
Precursor Count	1475	Transition Count

RENAME

Transition List

VIEWS CHARTS EXPORT PRINT PAGE SIZE

Protein / Label	Description
TAF11	HsCD00301672 (1-A03)

Peptide	Peptide Annotations	Missed Cleavages	Peptide Neutral Mass	Rank	Precursor
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
MDDAHESPSDK	Quality Score: 4	0	1230.4823		MDDAHESPSDK
EAAAEEGELESQDVSLTTVER	Quality Score: 4	0	2377.077		EAAAEEGELESQDVSLTTVER

Opening Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip

You have chosen to open:

Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip
which is a: WinZip File
from: https://panoramaweb.org:9443

What should Firefox do with this file?

Open with WinZip (default)

Save File

Do this automatically for files like this from now on.

OK Cancel

Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip

Organize Extract all files

human_consensus_final_true_lib.blib

Stergachis - Supplementary Data 2.sky

Stergachis - Supplementary Data 2.sky.view

Stergachis - Supplementary Data 2.skyd

Computer

OSDisk (C:)

Recovery (D:)

2TBDrive (F:)

Network

BRN001BA9EE6DAF

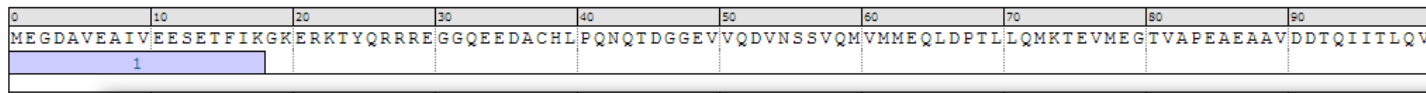
4 items

Viewing Proteins

Peptide Group

Protein / Label
 Description
 Decoy false

Sequence Coverage



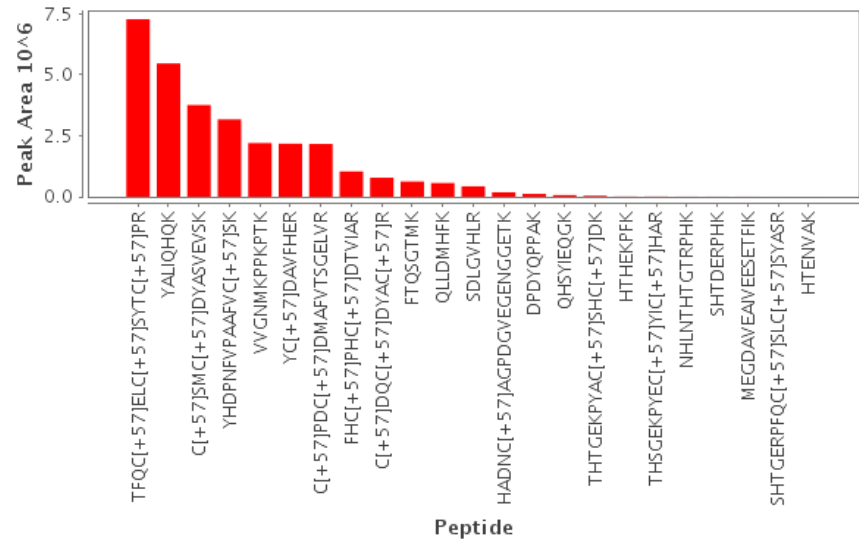
Peptides

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VIEWES	CHARTS	EXPORT	PRINT	PAGE SIZE	
Peptide	Peptide Neutral Mass	Missed Cleavages	Rank	Avg. Measured RT	Predicted RT
DETAILS ▶ MEGDAVEAIVEESETFIK	1995.9347	0		24.9907	
DETAILS ▶ DPDYQPPAK	1029.4767				
DETAILS ▶ VVGNMKPPKPTK	1294.7432				
DETAILS ▶ TFQCELCSYTCPR	1720.7008				
DETAILS ▶ SHTDERPHK	1105.5265				
DETAILS ▶ NHLNHTGTGRPHK	1511.7705				
DETAILS ▶ CPDCDMAFVTSGLVR	1855.7903				
DETAILS ▶ HTHEKPFK	1022.5298				
DETAILS ▶ CSMCDYASVEVSK	1534.6102				
DETAILS ▶ SHTGERPFQCSLCSYASR	2141.937				
DETAILS ▶ THSGEKPYECYICAR	2006.8727				
DETAILS ▶ FTQSGTMK	898.4219				
DETAILS ▶ HTENVAK	797.40314				
DETAILS ▶ FHCPCDVTIAR	1511.6761				
DETAILS ▶ SDLGVHLR	895.48755				
DETAILS ▶ QHSYIEQ GK	1088.525				
DETAILS ▶ YCDAVFHER	1195.508				
DETAILS ▶ YALIQHQK	999.5502				
DETAILS ▶ CDQCDYACR	1246.4165				
DETAILS ▶ THTGEKPYACSHCDK	1789.7512				
DETAILS ▶ QLLDMHFK	1030.527				
DETAILS ▶ YHDPNFVPAAFVCSK	1750.8137				
DETAILS ▶ HADNCAGPDGVEGENGGETK	2012.813				

Peak Areas

CTCF

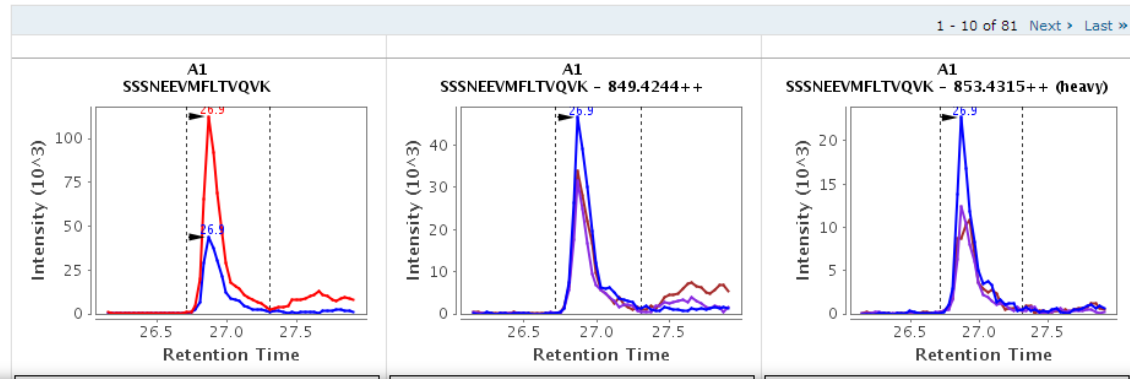


Viewing Peptides

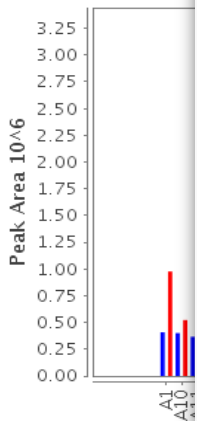
Peptide Summary

Name	Human_plasma.zip
Group	A2MG
Sequence	SSSNEEVMFLTVQVK
NeutralMass	1696.8342
Avg. RT	26.6774
Predicted RT	26.3900
Precursors	SSSNEEVMFLTVQVK - 8 SSSNEEVMFLTVQVK - 8

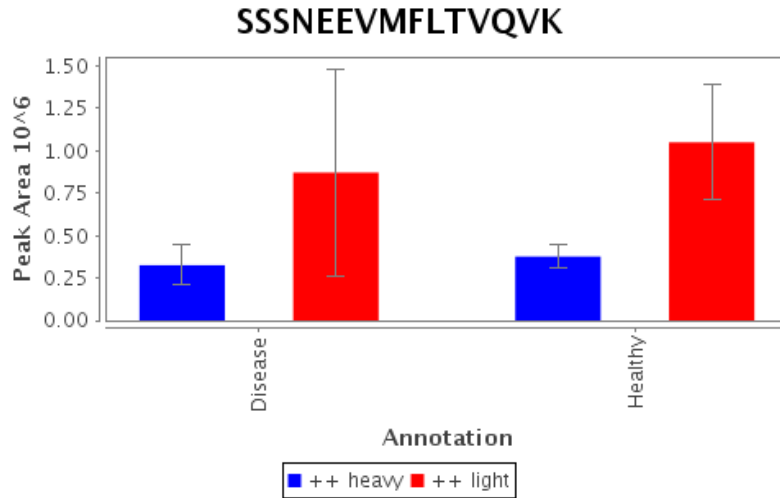
Chromatograms



Peak Areas



Peak Areas



Group By:

Condition

CV Values:

Width:

500

Height:

300

UPDATE

Protein and Peptide Search

Mass Spec Search

Protein Search Peptide Search Modification Search

Protein Name * ?

Search in subfolders?

SEARCH

Mass Spec Search

Protein Search Peptide Search Modification Search

Peptide sequence * ? MQILVSSFEEQLNR

Exact matches only?

Search in subfolders?

SEARCH

Targeted MS Peptides

VIEWS CHARTS EXPORT PRINT PAGE SIZE

	Protein / Label	Peptide	Path	File
DETAILS	TAF11	MQILVSSFEEQLNR	/MacCoss/brendanx/Test1/Libraries/Human	Stergachis - Supplementary Data 2_2013-05-30_10-54-36.zip
DETAILS	TAF11	MQILVSSFEEQLNR	/MacCoss/brendanx/Test1/Libraries/Human	Stergachis - Supplementary Data 2_2013-06-02_17-14-10.zip
Distinct Count:		1		2

Modification Search

Targeted MS Modification Search

Search By: Delta Mass Modification Name

Type: Names used in imported experiments
 Common Unimod modifications
 All Unimod modifications

Include: Structural Isotope Label

Custom Name: Label: 13C(6)15N(2)

Search in Subfolders: Carbamidomethyl (C)
 Carbamidomethyl C

SEARCH

Heavy K
 Heavy R
 Label: 13C(6)15N(2)
 Label: 13C(6)15N(4)
 Phospho (S,T)
 Phospho (Y)

Targeted MS Peptides

VIEWS CHARTS EXPORT PRINT PAGE SIZE

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	Protein / Label	Peptide	Precursor	File
DETAILS	Rv3133c devR	TLLGLLSEGLTNK	TLLGLLSEGLTNK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Rv2626c hrp1	HLPEHAIVQFVK	HLPEHAIVQFVK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Decoys	HLIDDALK	HLIDDALK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Decoys	TLLGLLSEGLTNK	TLLGLLSEGLTNK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Rv2623 TB31.7	HLIDDALK	HLIDDALK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Rv2031c hspX	TVSLPVGAEDEDDIK	TVSLPVGAEDEDDIK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Rv2031c hspX	GILTVSVAVSEGKPTK	GILTVSVAVSEGKPTK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Decoys	TVSLPVGAEDEDDIK	TVSLPVGAEDEDDIK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Decoys	GILTVSVAVSEGKPTK	GILTVSVAVSEGKPTK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	Decoys	HLPEHAIVQFVK	HLPEHAIVQFVK	SRMCourse_DosR-hDP__20130501_forBrendan.zip
DETAILS	B2MG	VNHVTLSPQK	VNHVTLSPQK	Human_plasma.zip
DETAILS	CO4A	GSFEFPVGDVSK	GSFEFPVGDVSK	Human_plasma.zip
DETAILS	CO4A	LGQYASPTAK	LGQYASPTAK	Human_plasma.zip
DETAILS	CO5	FQNSAILTIQPK	FQNSAILTIQPK	Human_plasma.zip
DETAILS	CFAB	YGLVTYATYPK	YGLVTYATYPK	Human_plasma.zip
DETAILS	B2MG	VEHSDLFSK	VEHSDLFSK	Human_plasma.zip
DETAILS	APOA1	DLATVYVDVLK	DLATVYVDVLK	Human_plasma.zip
DETAILS	CFAB	QLNEINYEDHK	QLNEINYEDHK	Human_plasma.zip
DETAILS	GELS	AGALNSNDAFVLK	AGALNSNDAFVLK	Human_plasma.zip
DETAILS	C1R	YTTTGMGVNTYK	YTTTGMGVNTYK	Human_plasma.zip

VIEWS CHARTS EXPORT PRINT PAGE SIZE

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Customizing Views, Sorts, Filters

Targeted MS Peptides

VIEWS ▾ CHARTS ▾ EXPORT ▾ PRINT PAGE SIZE ▾ 1 - 20 of 48 Next ▸ Last ▹

Columns Available Fields Selected Fields

Filter

Sort (1)

- Peptide
- Group
- Peptide
- Start Index
- End Index
- Peptide Neutral Mass
- Missed Cleavages
- Rank
- RT Score
- Predicted RT
- Avg. Measured RT
- Decov

Show Hidden

DELETE REVERT

	Protein / Label	Peptide	Precursor
DETAILS ▸	Rv3133c devR ▾	TLLGLLSEGLTNK	TLLGLLSI
DETAILS ▸	Rv2626c hrp1 ▾	HLPEHAIVQFVK	HLPEHAIV
DETAILS ▸	Decoys ▾	HLIDDALK	HLIDDALI
DETAILS ▸	Decoys ▾	TLLGLLSEGLTNK	TLLGLLSI
DETAILS ▸	Rv2623 TB31.7 ▾	HLIDDALK	HLIDDALI
DETAILS ▸	Rv2031c hspX ▾	TVSLPVGAEDDIK	TVSLPVG
DETAILS ▸	Rv2031c hspX ▾	GILTVSVAVSEGKPTK	GILTVSV
DETAILS ▸	Decoys ▾	TVSLPVGAEDDIK	TVSLPVG
DETAILS ▸	Decoys ▾	GILTVSVAVSEGKPTK	GILTVSV
DETAILS ▸	Decoys ▾	HLPEHAIVQFVK	HLPEHAIV
DETAILS ▸	B2MG ▾	VNHVTLSQLPK	VNHVTL
DETAILS ▸	CO4A ▾	GSFEFPVGDVSK	GSFEFPV
DETAILS ▸	CO4A ▾	LGQYASPTAK	LGQYASF
DETAILS ▸	CO5 ▾	FQNSAILTIQPK	FQNSAIL
DETAILS ▸	CFAB ▾	YGLVTYATYPK	YGLVTYA
DETAILS ▸	B2MG ▾	VEHSDLSFSK	VEHSDLS
DETAILS ▸	APOA1 ▾	DLATVYVDVLK	DLATVYV
DETAILS ▸	CFAB ▾	QLNEINYEDHK	QLNEINYI
DETAILS ▸	GELS ▾	AGALNSNDAFVLK	AGALNSN
DETAILS ▸	C1R ▾	YTTMTGMVNTYK	YTTMTGM

VIEWS ▾ CHARTS ▾ EXPORT ▾ PRINT PAGE SIZE ▾

Targeted MS Peptides

VIEWS ▾ CHARTS ▾ EXPORT ▾ PRINT PAGE SIZE ▾ 1 - 20 of 37 Next ▹

View: With RT

Filter: (PredictedRetentionTime > 15)

	Protein / Label	Peptide	Precursor	File	Predicted RT ▾ ▲
DETAILS ▸	CO6 ▾	IGESIELTCPK	IGESIELTCPK	Human_plasma.zip	17.6500
DETAILS ▸	PON1 ▾	IQNILTEEPK	IQNILTEEPK	Human_plasma.zip	17.7100
DETAILS ▸	Rv2626c hrp1 ▾	HLPEHAIVQFVK	HLPEHAIVQFVK	SRMCourse_DosR-hDP__20130501_forBrendan.zip	17.9000
DETAILS ▸	Decoys ▾	HLPEHAIVQFVK	HLPEHAIVQFVK	SRMCourse_DosR-hDP__20130501_forBrendan.zip	17.9000
DETAILS ▸	THBG ▾	EGQMSVEEAMSSK	EGQMSVEEAMSSK	Human_plasma.zip	18.0700
DETAILS ▸	FN1 ▾	SYTITGLQPGTDYK	SYTITGLQPGTDYK	Human_plasma.zip	19.3800
DETAILS ▸	Rv2031c hspX ▾	GILTVSVAVSEGKPTK	GILTVSVAVSEGKPTK	SRMCourse_DosR-hDP__20130501_forBrendan.zip	20.2700
DETAILS ▸	Decoys ▾	GILTVSVAVSEGKPTK	GILTVSVAVSEGKPTK	SRMCourse_DosR-hDP__20130501_forBrendan.zip	20.2700
DETAILS ▸	HPT ▾	VTSIQDWVQK	VTSIQDWVQK	Human_plasma.zip	20.4300
DETAILS ▸	GELS ▾	AGALNSNDAFVLK	AGALNSNDAFVLK	Human_plasma.zip	20.5000
DETAILS ▸	C1QB ▾	LEQGENVFLQATDK	LEQGENVFLQATDK	Human_plasma.zip	20.6600
DETAILS ▸	CFAB ▾	YGLVTYATYPK	YGLVTYATYPK	Human_plasma.zip	20.6800
DETAILS ▸	CO5 ▾	FQNSAILTIQPK	FQNSAILTIQPK	Human_plasma.zip	20.6900
DETAILS ▸	Rv2031c hspX ▾	TVSLPVGAEDDIK	TVSLPVGAEDDIK	SRMCourse_DosR-hDP__20130501_forBrendan.zip	20.7000
DETAILS ▸	Decoys ▾	TVSLPVGAEDDIK	TVSLPVGAEDDIK	SRMCourse_DosR-hDP__20130501_forBrendan.zip	20.7000
DETAILS ▸	AACT ▾	EIGELYLPK	EIGELYLPK	Human_plasma.zip	21.3600
DETAILS ▸	THY ▾	AADDTWEPFASGK	AADDTWEPFASGK	Human_plasma.zip	21.4600
DETAILS ▸	RET4 ▾	FSGTWYAMAK	FSGTWYAMAK	Human_plasma.zip	21.8700
DETAILS ▸	CO4A ▾	GSFEFPVGDVSK	GSFEFPVGDVSK	Human_plasma.zip	22.9400
DETAILS ▸	IC1 ▾	VTTSQDMLSIMEK	VTTSQDMLSIMEK	Human_plasma.zip	23.8500

VIEWS ▾ CHARTS ▾ EXPORT ▾ PRINT PAGE SIZE ▾ 1 - 20 of 37 Next ▹

Export and Client APIs

The image displays the PanoramaWeb interface for managing targeted MS peptides. It includes a sidebar for navigation, a main content area with a search bar, and a table of peptide data. The interface is overlaid with a Microsoft Excel window showing the exported data and a R script window with the code used for data selection.

Targeted MS Peptides Table:

Protein / Label	Peptide	Precursor
CO6	IGESIELTCPK	IGESIELTCPK
PON1	IQNILTEEPK	IQNILTEEPK

Excel Export Options:

- Excel 2007 File (.xlsx) Maximum 1,048,576 rows and 16,384 columns
- Excel 97 File (.xls) Maximum 65,536 rows and 256 columns
- Refreshable Web Query (.iqy)

R Script:

```
mydata <- labkey.selectRows(
  colFilter=makeFilter(c("PeptideId/PredictedRetentionTime", "GREATER_THAN", "15")),
  containerFilter=NULL)
```

Permissions

https://panoramaweb.org x
https://panoramaweb.org:9443/labkey/security/MacCoss/brendanx/project.view?

Admin ▾ Help ▾ jeckels ▾

Permissions for /MacCoss/brendanx

SAVE AND FINISH SAVE CANCEL

Folders

- MacCoss
 - absterga
 - brendanx**
 - genn
 - maccoss
 - maccosstester
 - mbereman
 - nicksh
 - spionin
 - vsharma

Permissions Project Groups Site Groups Impersonate Group Diagram

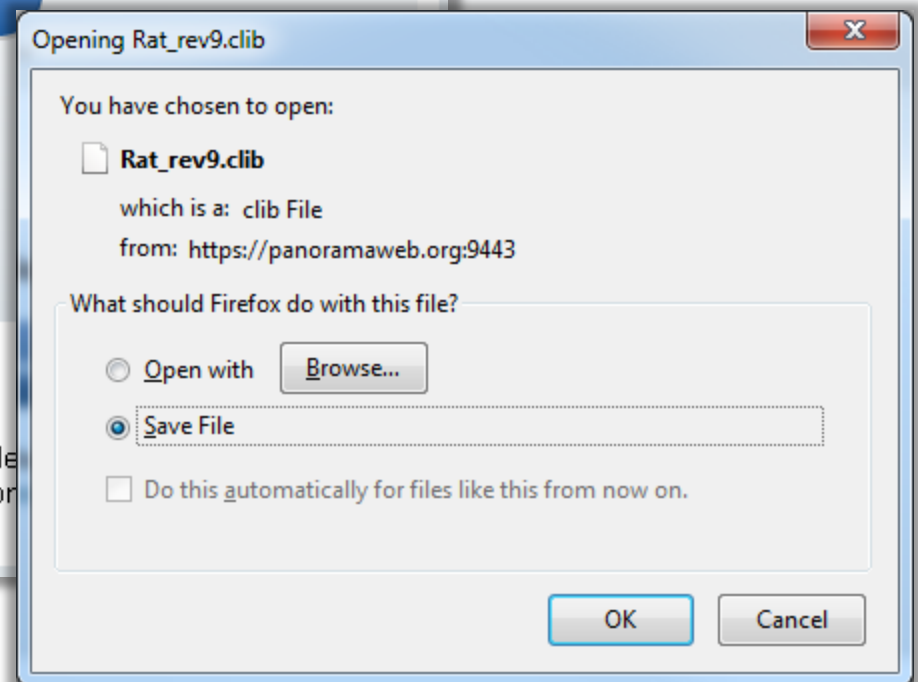
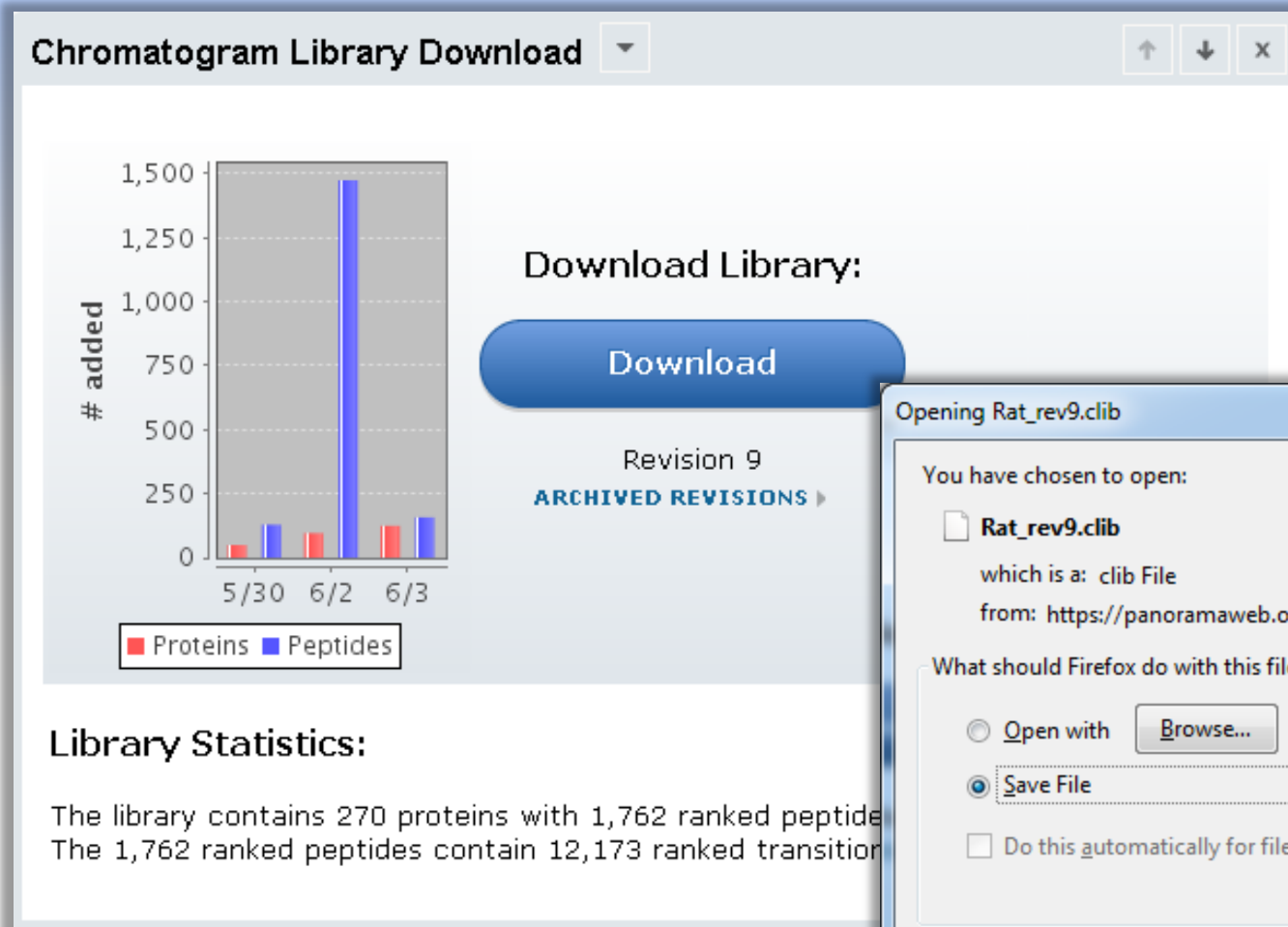
VIEW PERMISSIONS REPORT ▾

Inherit permissions from parent

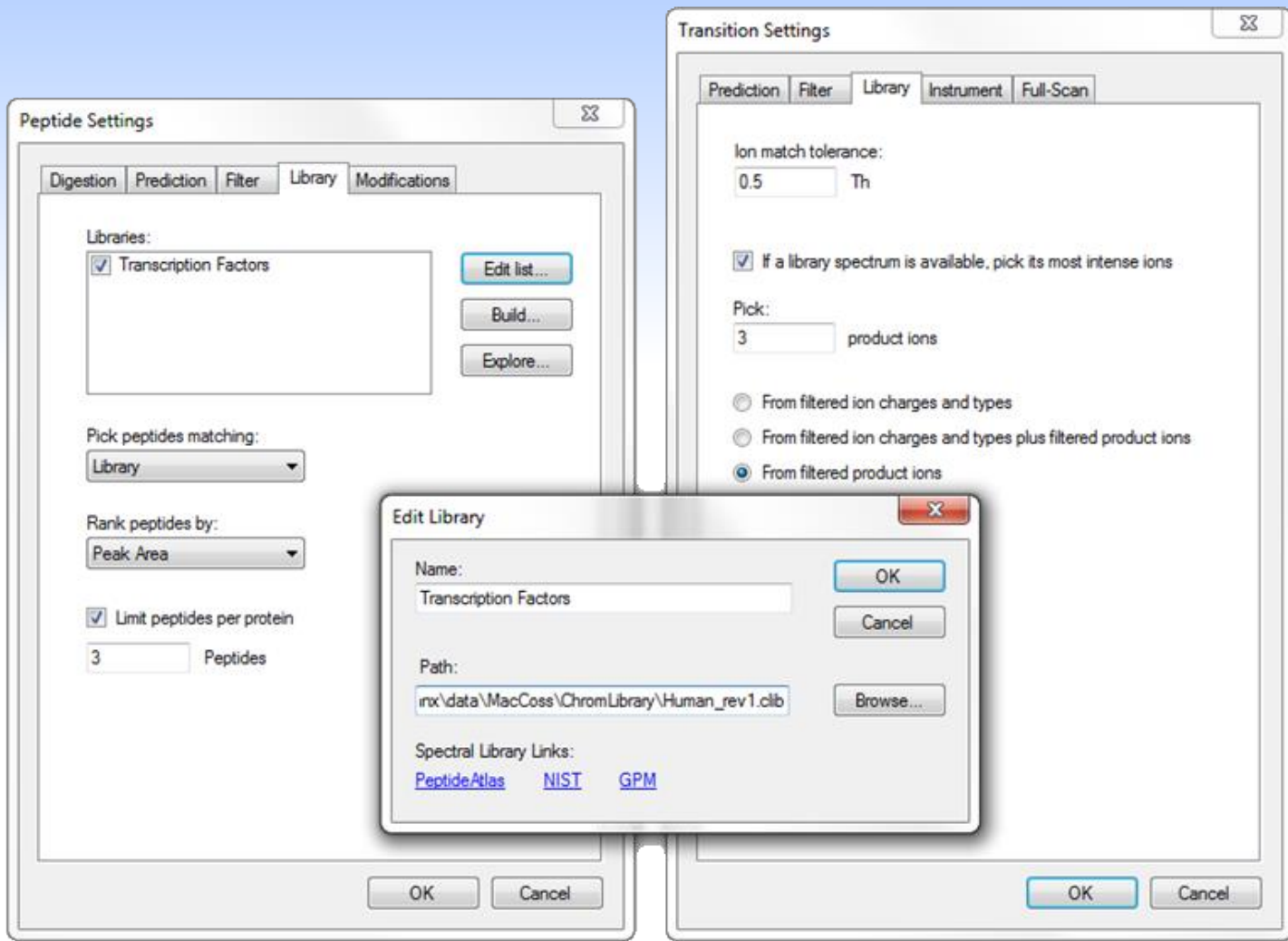
Roles	Groups
Folder Administrator Folder Administrators have full control over their particular folder, but not others.	BRENDANX@PROTEINMS.NET <input type="button" value="x"/> Add user or group...
Editor Editors may read, add, update and delete information.	Add user or group...
Author Authors may read and add information in some cases, but may update and delete only information they added.	MACCOSS MEMBERS <input type="button" value="x"/> ABSTERGA@UW.EDU <input type="button" value="x"/> Add user or group...
Reader Readers may read information but may not change anything.	MACCOSS MEMBERS <input type="button" value="x"/> GUESTS <input type="button" value="x"/> Add user or group...
Submitter Submitters may submit new information, but may not read or change anything.	Add user or group...
Assay Designer Assay designers may perform several actions related to designing assays.	Add user or group...

** indicates permissions are inherited*

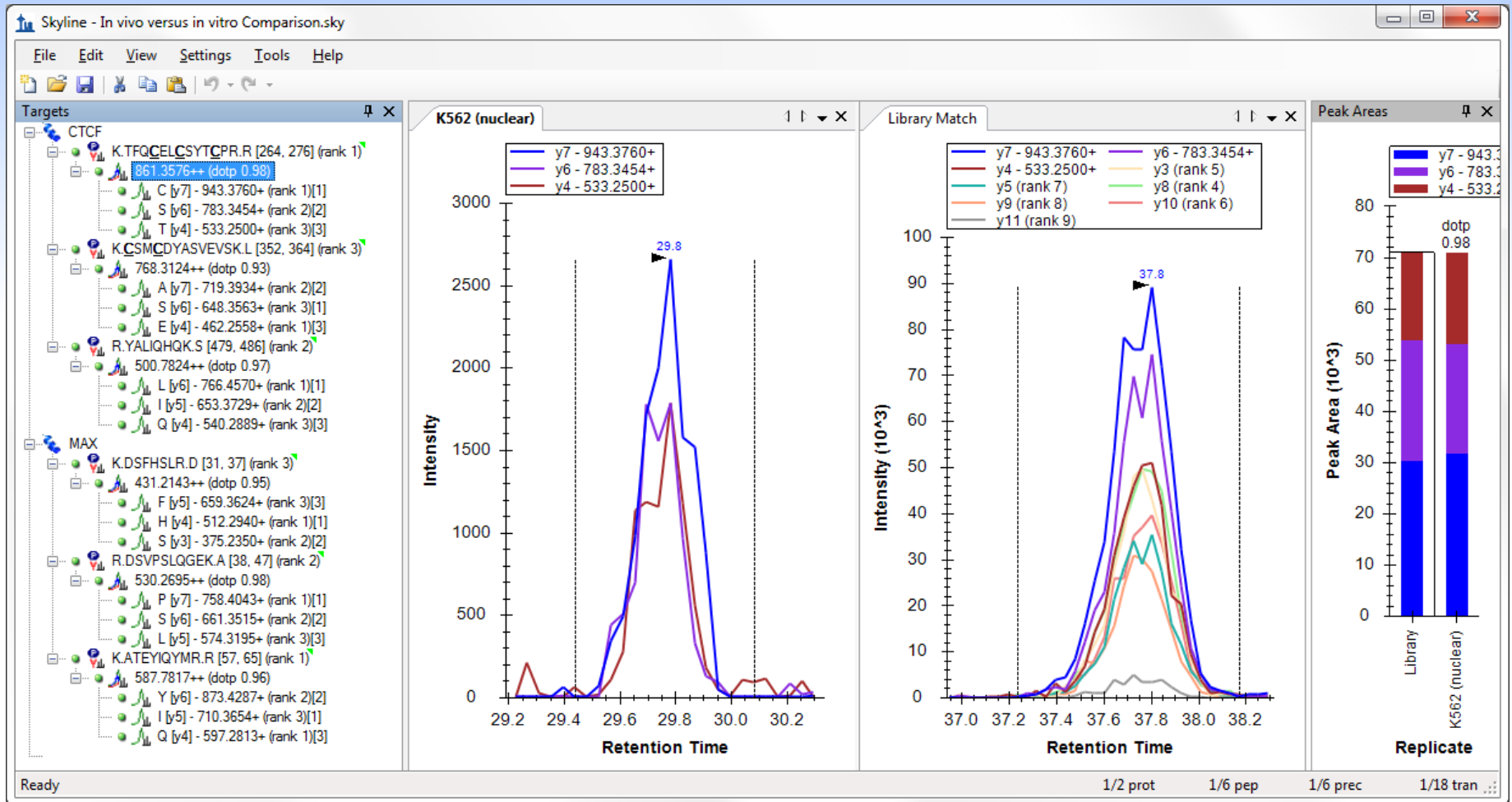
Chromatogram Library Export



Chromatogram Library Import



Chromatogram Library Usage



How to Get Panorama

- PanoramaWeb (panoramaweb.org)
 - Hosted at University of Washington
 - Request access for a lab or organization
 - Data can be private or public
- Create your own installation
 - Download and documentation at labkey.com
 - Install on Windows, Linux, OSX, etc
- Use LabKey Software's private hosting

Future Plans

- Release chromatogram library support
 - July 2013
 - Will coincide with new Skyline release
- Store and calculate normalized retention times (iRT)
- Store and view QC runs
- Show more Skyline graphs
- Support response curves with QuaSAR

More Information

- Poster tomorrow
 - **Sharing Targeted Proteomics Assays using Skyline and Panorama**
 - MP 382
- panoramaweb.org
 - View recorded webinar
 - Request a project
- labkey.org/labkey.com
 - General information
 - Downloads for installers and source code



Questions

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Skyline Users Group - June 9, 2013