

Tutorial for logging a sample requisition into YPED

Go To:

[YPED Results Login](#) or <http://yped.med.yale.edu/>

Your account is:

Login ID:

Password:

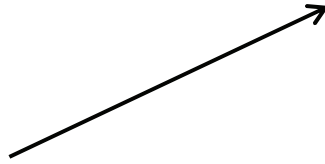
**** Note if you do not have a YPED account, select “Apply for account”, and fill out the appropriate information, and your account will be activated within 24 hrs.**



User Functions User:

- [Compare LCMS Peptides](#)
- [Compare MUDPIT/ITRAQ](#)
- [Compare Results \(ProteinPilot/LCMS\)](#)
- [Compare DIGE ITRAQ\(ProteinPilot\)](#)
- [Add Sample Requisition](#)
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- [Project Management help](#)
- [Project Management](#)
- [Repository Management](#)
- [Protein ID Peptide Report](#)
- [YPED Usage Statistics](#)
- [Log Out](#)

Pick





Select Sample Requisition USER:

Select Project

You can also create a project

Select Analysis Type Select Funding Source Template Sample(optional)

- Gel-Protein ID
- Solution-Protein ID
- ITRAQ4plex
- ITRAQ8plex
- Label Free Quant
- SILAC
- SmallMoleculeQuant
- MRM Protein Quant
- TMTduplex
- TMTsixplex

Pick

Projects are used to group samples that are from the same experiment. A Project will also allow you to do some comparisons of protein identifications between samples once the analysis is complete.



Select Sample Requisition USER:

Select Project

Select Analysis Type Select Funding Source Select Number of Samples Template Sample(optional)

Gel-Protein ID 1

For questions or comments c
Updated 06 Dec 2013 9:58

- PTAEO
- NHLBI
- NIDA
- NBC
- Credit Card
- PO
- Other

[hifman](#)

Pick payment option
And # of samples submitted

If you already have submitted
a sample previously, you will
be given the option to choose
a past sample as a template to
use for your new sample set



Add Sample Requisition USER:

Analysis Type:

Funding Source:

Type Billing Information*

Special Billing Instructions

Number of Samples to add

Sample #	1
Sample Name *	<input type="text"/>
Stain *	<input type="radio"/> silver <input type="radio"/> Coomassie <input type="radio"/> none
Organism *	<input type="text"/>
Tissue	<input type="text"/>
Cell Line	<input type="text"/>
Estimated Total Amount (ug) *	<input type="text"/>
Estimated Total Amount (picomoles)	<input type="text"/>
Accession ID suspected protein	<input type="text"/>
Digest	<input checked="" type="radio"/> Trypsin <input type="radio"/> LysC/Trypsin <input type="radio"/> LysC <input type="radio"/> GluC <input type="radio"/> other (Please specify) <input type="text"/>
Sample Description/Comment *	<input type="text"/>
Growth Conditions	<input type="text"/>
Extraction Procedure	<input type="text"/>
Post Translation Modification	<input checked="" type="radio"/> none <input type="radio"/> Phospho <input type="radio"/> other (Please specify) <input type="text"/>
Key Words *	<input type="text"/>

Fill out the information
And then hit

* Are required fields



Add Sample Requisition USER:

Sample Requisition created. req_num: r_37810

Analysis Type: Gel-Protein ID

Funding Source: Other

Type Billing Information* PO or PTAE0 #

Special Billing Instructions

Number of Samples to add 1

Sample #	1
Sample Name *	test
Stain *	<input type="radio"/> silver <input checked="" type="radio"/> Coomassie <input type="radio"/> none
Organism *	Homo sapiens
Tissue	
Cell Line	
Estimated Total Amount (ug) *	0.1
Estimated Total Amount (picomoles)	
Accession ID suspected protein	
Digest	<input checked="" type="radio"/> Trypsin <input type="radio"/> LysC/Trypsin <input type="radio"/> LysC <input type="radio"/> GluC <input type="radio"/> other (Please specify) <input type="text"/>
Sample Description/Comment *	protein ID
Growth Conditions	
Extraction Procedure	
Post Translation Modification	<input checked="" type="radio"/> none <input type="radio"/> Phospho <input type="radio"/> other (Please specify) <input type="text"/>
Key Words *	protein ID

Once the form is completed,
Print the submission form and place
With your sample.

****Your online sample
submission is now complete,
and you can press "Done".
Make sure to exit out of the
browser if you are using a
public computer, so that your
login information is secure.**