



Quick Guide Module PMS

Core Facility Proteomics Mass Spectrometry

https://www.dbmr-mare.ch/mare/

Support manfred.heller@unibe.ch, Head PMS sophie.lagache@unibe.ch, PMS Lab ilker.romann@unibe.ch, IT DBMR The MARE system has been introduced in order to provide core facilities with a digitalized system for the management of projects. Please note the following:

- Every user of the core facility, as well as their PI, must have registered once with the MARE system before submitting samples (section 1 and 2).
- PIs are specifically requested to report their PI status (section 2) in order to be listed as such and get permission to oversee all the user projects associated with them.
- Every user must register specifically with PMS (section 3).
- Each registered user can then create a PMS project (section 4): this involves stating a PI and giving a few details about your project; you will notice that we suggest meeting with you and your PI, as we believe this improves the likelihood of success of your experiment.
- Once the project is accepted by PMS, all the necessary information about the experiment can be entered, such as the type of samples submitted, their names, the samples taxonomy etc., and for UniBe or Inselspital users the corresponding REF number or cost center, respectively (section 5).
- Once your samples are processed, both user and PI will be informed, and the result files will be available in the user project space for the next 30 days.

Workflow in a Nutshell





In the first step, each user must create a user account: Click «Sign Up». If you have already performed this step, click Sign In and jump to section 4.

LU	ogin
Username	
Password	
🗸 Sign In	Password forgotten
Don't have an acc	count yet? Sign Up!
Sign Firstna	Up me *
Charles	
Lastna	me *
Darwin	
E-Mail Address (University	, Hospital or Business) *
charles.darwin@unibe.ch	
Passwo	ord *
Repea	at *
I agree with terms of use	
✓ I'm not a robot	reCAPTCHA Privacy - Terms
Sign Up	
Already have an a	ccount? Sign In!



4

Please fill in all fields of your personal data.

MARE		Darwin Charles
DBMR Management Software	命	은 Change user data
		🚨 Notifications
合 Home	A Welcome Darwin Charles	A Change Password
O Profile		[→ Logout
	PMS	
	Proteomics & Mass Spectrometry	
	Request Access	

If the personal data is incomplete, access to the module in step 3 will not be granted.

Personal In	formation	Contact Detail	s
Username *	charles.darwin@unibe.ch	E-Mail *	charles.darwin@unibe.ch
Salutation *	Mr. ~	Phone	+41 000 00 00
Title	Prof. Dr. med., Dr. phil. nat.		
Lastname *	Darwin		
Firstname *	Charles		
Address		Affiliation/Con	npany
Street	Murtenstrasse	University/Company	University of Bern
Number	28	Department	Department for BioMedical Research
Zip Code	3008	Institute/Group	Proteomics Mass Spectrometry Core Facility
City	Bern		
State	BE		
Country	Switzerland V		
Save	< Back		

immediately after registering, stating that you are a PI requesting PI permission. This is very important, as otherwise your lab members will not be able to capture projects. 3 Request Access

In order to use PMS services, you need to specifically request it:

DBMR Management	命	[→	
☆ Home ◇ Profile	A Welcome Darwin Charles		
	PMS Proteomics & Mass Spectrometry		
	Request Access		

New request for access

	t projects	ear Mani vould like to su est, Charles	De I w Be	Message *	
			< Back	Send	7
					6

PMS is informed and activates access.

PMS	PMS
Proteomics & Mass Spectrometry	Proteomics & Mass Spectrometry
Request Pending	Projects

Steps 1 to 3 are performed only once.



Select Project Request in the menu:

合 Home	ి Welcome Charles Darwin
A Profile	
ee PMS ^	
Project Requests	PMS
4	Proteomics & Mass Spectrometry
	Projects

Create a new project application.

🗅 Project Requests		
0 Project Requests		New project request +
Request Date 1	Status ↑↓	

Please select the responsible PI from the list. The PI must already be registered in the system and requires special authorization (steps 1 and 2).

New project request

Basedata Co	sulting Overview	
Principle Investigate	r / Group	
Please give us a sho	t description Please enter only a short description	of the project.

Please specify 3 possible dates for a consultation.

Tip: Copying and pasting from the first appointment is helpful.

New project request

Basedata	Consulting	Overview	
Please choose 3	dates/times for a	neeting:	
First meeting o	date/time *	15.07.2024 09:00	Ċ
Second meeti	ng date/time *	16.07.2024 10:00	Ċ
Third meeting	;date/time *	17.07.2024 11:00	*
Please choose	an option *	Online (Teams, Zoom, Slack etc.)	~
← Back			Next →

New project request

Basedata	Consulting	Overview
Basedata)	
Principle Investigator /		Albert Einstein

Group Leader	
Customer Name	Darwin Charles
Customer Email Address	charles.darwin@unibe.ch
Customer Phone	+41 000 00 00
Project description	Please enter only a short description of the project.

Consulting

First meeting date/time	15.07.2024 09.00
Second meeting date/time	16.07.2024 10.00
Third meeting date/time	17.07.2024 11.00
Meeting type	Online (Teams, Zoom, Slack etc.)

Send

Project Requests

1 Project Requests		× +	
Request Date 1ें≓	Status ↑↓		
10.07.2024 16:37:18	Request Sent		

You will be contacted by PMS for a consultation.

Project Requests

1 Project Requests		B +	
Request Date 1	Status ↑↓		
10.07.2024 16:37:18	Consulting Pending	$\boxed{\bigcirc}$	

After the consultation, a project is generated.

Project Requests

1 Project Requests		B +
Request Date 17	Status ↑↓	
10.07.2024 16:37:18	Request Accepted	

Once all the above is completed, you will see it as "accepted" under PMS-Project Requests.



Now click on Home, then Projects under PMS. You should see your created project:

Project over	view			
0 / 1 Project				
Project Id 1े₹ 3	UNI REF 17	PI/PL ↑₹ 2	Status ↑₹ 🕚	
20240710_12		Einstein Albert	Created	

Note the Project id that has been automatically generated: it will be useful for reference in future communications.

You are first in the "Basedata" widget. Fill in everything, then click next:

\equiv Project Details

Affiliate *	University of Bern				
Project Category *	Customer Project				
Uni REF Number	Important! Enter correct REF number!				
Type of request analysis Protein identification, short gradient, per sample					
Protein identification with PTM analysis, per sample					
	Comparative proteomics, shot gun, per sample				
	Comparative proteomics, in gel, per gel bande				
MW determination (Infusion or LCMS) / intact mass determination					
Deschaproteomics with phosphopentide enrichment					

Note:

If affiliate is University of Bern

- Select University of Bern
- Important! Enter correct REF number!

If affiliate is Inselspital

- Select University Hospital Inselspital
- Important! Enter correct Cost Center number!

• Enter Insel Advisor

\equiv Project Details

Basedata Samples Sar	nple Groups Result files Overview		
Affiliate *	University Hospital Berne - Inselspital		
Project Category *	Customer Project		
Cost center	Important! Enter correct Cost Center number!		
Insel advisor	firstname.name@insel.ch		
Type of request analysis	Protein identification, short gradient, per sample		
	Protein identification with PTM analysis, per sample		
	Comparative proteomics, shot gun, per sample		
	Comparative proteomics, in gel, per gel bande		
MW determination (Infusion or LCMS) / intact mass determination			
	Phosphoproteomics with phosphopentide enrichment		
	Next →		

Fill in next details about the samples you will submit:

Basedata	Samples	Sample Groups	Result files	Overview	
🧹 Gel band	S			Pull-down on beads	
Membrar	ie pieces			Cell Lysate pellet	
In solutio	n			Other	
Taxonomy *		Taxonomy			
Sample descri	ption	Sample descri	ption		
Processing Re	quest	Optional infor	mations		
	4	optionation			
					_
← Back					Ne
ease not	e that it i	s essential fo	r us to kno	w the taxonomy or the organ	ism

≡ Project Details

Please note that it is essential for us to know the taxonomy or the organism corresponding to your samples; examples are Human, Mouse, Rat, Toxoplasma gondii (strain ATCC 50611 / Me49), etc. We will use this information to find the appropriate protein database which we will use to search your samples. You must tell us in the next widget ("Sample Groups") the names of the samples as they appear on your vial labels. In order to facilitate the listing of samples, we ask you first how many conditions ("Number of Experiment Groups") you have. Ex: if you have 3 replicates of 2 different conditions, put 2 here. If you have 4 different samples, put 4 here. The replicates will be given automatic names which you can change. See below an example if you have 3 control samples called CTRL_1, CTRL_2, CTRL_3, and 4 treated samples called Treat_1, Treat_2, Treat_3, and Treat_D.

The number of conditions is 2, so enter 2 and click update:

\equiv Project Details

Basedata Samples	Sample Groups	Result files	Overview
Number of Experiment Groups	2 Update		
← Back			Next →

The number of replicates is 3 and 4; click update and modify each sample name individually as needed:

≔ Project Detai	ls				
Basedata Sample	s Sample Groups	Result files Ove	rview		
Number of Experiment Gro	pups 2 Upd	ate			
Group Name	CTRL	# of replicas	3 Update		
Replicate/Sample Name	1	Protein Amount		Buffer Composition	Number of Fractions/Sample *
Replicate/Sample Name	2	Protein Amount		Buffer Composition	Number of Fractions/Sample *
Replicate/Sample Name	3	Protein Amount		Buffer Composition	Number of Fractions/Sample *
Group Name	Treat	# of replicas	4 Update		
Replicate/Sample Name	1	Protein Amount		Buffer Composition	Number of Fractions/Sample *
Replicate/Sample Name	2	Protein Amount		Buffer Composition	Number of Fractions/Sample *
Replicate/Sample Name	3	Protein Amount		Buffer Composition	Number of Fractions/Sample *
Replicate/Sample Name	D	Protein Amount		Buffer Composition	Number of Fractions/Sample *
← Back					Next $ ightarrow$

If the buffer composition or protein amount is known and relevant, fill in the fields.

Important: press Next to register your input.

The next widget is ""Result files".

\equiv Project Details

 \equiv Project Details

Basedata Samples Sa	ample Groups Result file	os Overview		
Document Name ↑↓	Document Size ↑↓	Upload Date ↑↓	Expired Date 1	
No records found.				
← Back				Next ->

After completion of the project, you can download the results here, within 30 days.

Basedata	Samples	Sample Groups	Result files	Overview
Basedata				
PI/GL		Albert Einstein		
Customer nam	e	Charles Darwin		
Customer Ema	il	charles.darwin@	@unibe.ch	
Customer Pho	ne	+41 000 00 00		
Id		20240710_12		
Affiliate		University Hosp	ital Berne - Inselsp	spital
Uni REF Numb	er			
Company Orde	ering Number			
Funding				
Project Catego	ry	Customer Proje	ct	
Type of reques	t analysis			
Samples				
Gel bands	5			Pull-down on beads
Membran	e pieces			Cell Lysate pellet
In solutio	n			Other
Taxonomy		Taxonomy		
Sample descri	ption	Sample descript	tion	
Processing Rec	quest	Optional inform	ations	
Sample Grou	ps			
20240710_12_	Gr 1_R 1_1			
20240710_12_	Gr 2_R 1_1			
I agree to	the Tern	ns and Conditions		
← Back				🔊 Send

Important: Your input won't be saved until you press the "Send" button. So don't forget that you need to click to the last widget and press "Send" in order to save your input or changes.



Done.

For easy navigation, use the arrow keys on the keyboard.

	F16	F17	F18	F19
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