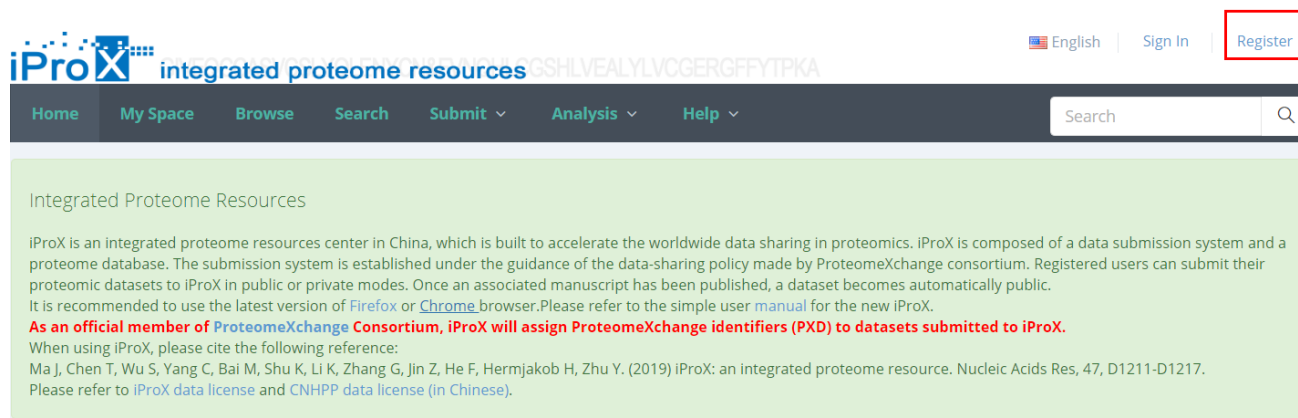


第一步：进入 iprox 主页

<https://www.iprox.org/>

第二步：注册账号

选择主页右上角的“Register”



新出现的页面中填写个人信息

🏠 REGISTRATION WIZARD - STEP 1 OF 4

1 Account Setup 2 User Profile Setup 3 Institute/Organization Setup 4 Confirm

Provide your account details.

Username * ① 不得超过12个字符
Provide your username.

Password * ②
Provide your password.

Confirm Password * ③
Confirm your password.

Email * ④
Provide your email address.

⑤ ☒ I agree to the [Terms of Service](#)

⑥

一次填完 2、3 步，星号*标记的是必填内容
第 4 步确定填写的信息

Back

Submit

核实填写内容正确时，点击“Submit”

然后去填写的邮箱，激活账号

Registration is completed. Please check your email to activate your account.
If you haven't received this email in two minutes, please click the following button to get it again.
In addition, you can send your 'Username' to iprox@iprox.org, we'll active it as quickly as possible.

89 seconds later can resend

进入自己的邮箱，在收件箱找到邮件

iprox

Registration for iProX - Thanks for registering iProX! Please click the following link to activate your account in 48 hours.

点击邮件正文的链接，完成邮件激活

Thanks for registering iProX!

Please click the following link to activate your account in 48 hours.

<https://www.iprox.org/page/URV001.html?userName=december&password=dai88336539>

Following is the same content in Chinese.

以下是与上面英文信息相同的中文信息。

感谢您注册iProX!

请点击下面链接激活账号，链接有效期为48小时，超过期限请重新注册。

<https://www.iprox.org/page/URV001.html?userName=december&password=dai88336539>

出现这个提示

Email has been activated, please log in

注册就完成了，就可以在网站进行登陆

第三步：登陆账号

点击右上角“Sign In”



integrated proteome resources

English

Sign In

Register

Home

My Space

Browse

Search

Submit

Analysis

Help

Search

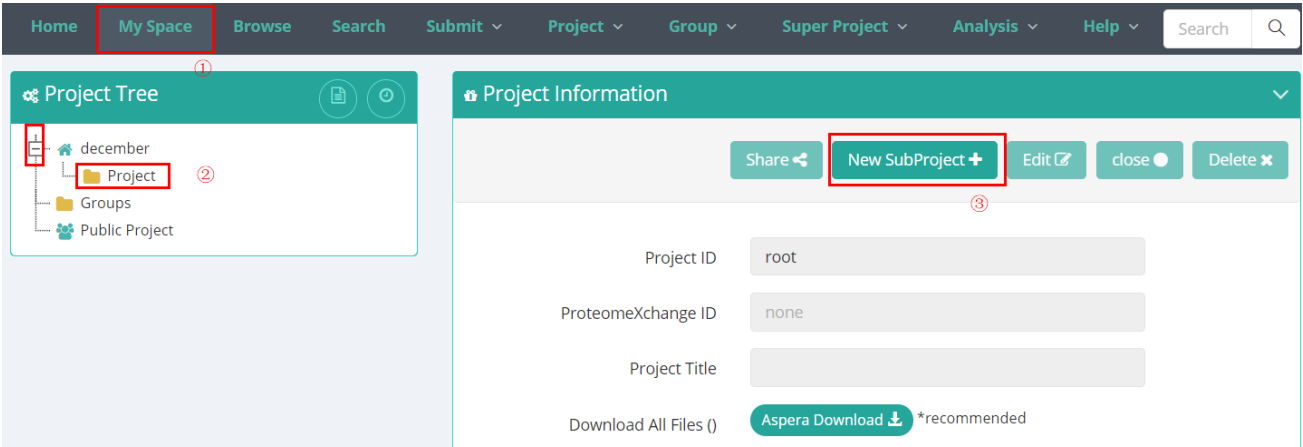
Q

Activation code has been expired

填入用户名、密码、验证码，登陆个人账号

第四步：提交数据

点击导航栏——My Space，
左部 Project Tree 里。点击 “Project”
点击右侧 “New SubProject”



填写项目信息，
共需要四步：

1 Add New Project 添加新项目

Add New Project

Project ID	<input type="text" value="IPX0001717000"/>	Add link 不用填，自动生成
	<small>Project ID will be generated automatically.</small>	<small>Provide external links for your dataset.</small>
ProteomeXchange ID	<input type="text"/>	
Project Title *	<input type="text"/>	项目名称
	<small>Provide title of the project (required).</small>	
Project Tag (key words) *	<input type="text"/>	项目标签
	<small>Provide key words of the project (recommended).</small>	
Description *	<input type="text"/>	项目描述 (不少于50个字符)
	<small>Provide description of the project (recommended).</small>	
Institution *	<input type="text"/>	机构名
	<small>Provide your institution (required).</small>	

Submitter *

Add New +

一般自动生成，如没有，可点击“Add New+”添加，填入后“save”

Name	Email	Institution	Edit	Delete
december			Edit	Delete

Lab Head *

Add New +

点击“Add New+”添加，填入后“save”

Name	Email	Institution	Edit	Delete
			Edit	Delete

Publication

Add New +

PubMed ID	References	Edit	Delete
No data available in table			

Access Level *

Public

三个选项，“Public”公开、“Plan to public”计划公开（在后面提示的日期栏选择将要公开的日期）、“Private”保密。

Data release note: the data will be released to public when the paper associated with this iProX ID is published, no matter 'Plan to public' or 'Private' is chosen here.

没有标注选择为可选内容，根据具体情况选择填与不填

2 Add New Subproject 添加新子项目

Add New SubProject

Subproject ID

Subproject ID will be generated automatically.

ProteomeXchange ID

Subproject Title *

项目名称

Provide title of the subproject (required).

Subproject Tag

Provide key words of the subproject (recommended).

Species *

Homo sapiens

← Choose

+ New

通过“choose”选择物种

Provide species information for your dataset (required).

MS instrument *

Q Exactive HF

≡ more

Add New MS instrument +

质谱仪型号

The instrument name is required, and your can click the 'More' option to provide further instrument information.

没有标注选择为可选内容，根据具体情况选择填与不填

3 Upload Data File: 上传文件

上传文件的方式有两种：

“Upload by Browser”：当上传的文件不超过 4GB 的时候，可以选择

否则可选择 “Upload by Aspera”，可上传大于 4GB 文件

iProX 支持两种不同的提交方式（“完整提交”和“部分提交”）。
在“完全提交”中，需要具有支持格式的原始文件和结果文件，
而在“部分提交”中，原始文件和搜库文件都是必需的，但可以是非标准格式。
表 1 列出了两种提交方式支持的详细文件类型。

Files types required in complete submission	
Raw file	Raw file refers to MS instrument raw output, such as: 1) *.raw 2) *.wiff 3) *.wiff.scan 4) *.baf 5) *.mzMXL 6) *.mzML
Result file	Result file could be one of the following types: 1) mzTab file 2) mzIdentML file with pealists/spectra files 3) prideXML file converted by PRIDE Converter .
Files types required in partial submission	
Raw file	MS instrument raw output
Search file	Search engine results (protein/peptide identifications)

目前大多采用 **Partial submission** 的形式来上传数据。

Upload your data file

Two different submission ways ('Complete submission' and 'Partial submission') are supported by iProX. In 'Complete submission', both raw files and result files with supported formats (mzTab, mzIdentML files with Peak files, or prideXML files) are required, while in 'Partial submission', both raw files and search files are required, but can be in non-standard formats.

Upload by Browser

Upload by Aspera

①

Upload files by web browser without any plugin or third-party software.
The single file size is limited to 4G. Please choose Aspera to upload large files.

☐ Complete submission

☒ Partial submission

②

+ Add File

Start Upload

Stop Upload

Cancel Upload

File Type	File Name	Source/Result File	Remarks	Size	Operation
RAW	HFX2_FFRA19H000562-1A.raw	Select related file		64K	✖
		进度: <div></div>			
OTHERS	propep.zip	Select related file		29.08K	✖
		进度: <div></div>			

Raw 文件来自释放文件夹 Raw_file 中*.raw 文件

Search file 文件来自 Match_result ，可将*PeptideGroups.txt/*Proteins.txt 文件进行准备好文件后，

点击“Add File”按钮，选中要上传的文件，然后点击“打开”按钮，如果有必要需要在 file type 一栏的下拉菜单中修改一下“file type”。

Buttons: + Add File, Start Upload, Stop Upload, Cancel Upload

File Type	File Name	Source/Result File	Remarks	Size	Operation
RAW	HFX2_FFRA19H000562-1A.raw 进度: [Progress Bar]	Select related file		64K	✕
SEARCH	propep.zip 进度: [Progress Bar]	Select related file		29.08K	✕

Complete submission 需要同时上传 RAW 文件和 Result 文件；Partial submission 需要同时上传 RAW 文件和 Search 文件。添加完之后点击“Start upload”即可开始上传。

Upload by Browser | Upload by Aspera

Upload files by web browser without any plugin or third-party software. The single file size is limited to 4G. Please choose Aspera to upload large files.

☐ Complete submission ☒ Partial submission

Buttons: + Add file, Start upload, Stop upload, Cancel upload

File Type	File Name	Source/Result File	Remarks	Size	Operation
RAW	CHPP_LM3_RP1_1.raw 进度: 已上传:580.64M/580.64M(100%)	✕ F030023.dat		580.64M	
RAW	CHPP_LM3_RP2_1.raw 进度: [Progress Bar]	✕ F030024.dat		601.17M	✕
SEARCH	F030023.dat 进度: [Progress Bar]	✕ CHPP_LM3_RP1_1.raw		66.14M	✕
SEARCH	F030024.dat 进度: [Progress Bar]	✕ CHPP_LM3_RP2_1.raw		67.96M	✕

4 Submit

核实上传内容，点击 Sumit 就完成了整个提交数据的流程。管理者会对上传数据进行审核。

STEP 4 OF 4

- 1 ✓ Add New Project
- 2 ✓ Add New Subproject
- 3 ✓ Upload Data File
- 4 Submit

Submission Summary

Please double-check the information before the final submission. Choose the project and its subproject(s), and click the "submit" button to finally submit your dataset.

Show 5 entries

Search:

	<input type="checkbox"/>	Project ID	Project Title	disease	Data Size	1	Status				
<input type="checkbox"/>	<input type="checkbox"/>	IPX0001717000					save	<input data-bbox="1098 674 1121 707" type="button" value="+"/>	<input data-bbox="1153 674 1177 707" type="button" value="📄"/>	<input data-bbox="1209 674 1233 707" type="button" value="✖"/>	

< 1 >

- ⏪ Back
- Submit ⏪
- Add New Subproject ⏪