

Search

Search

Type: Location

Folder

Filters

Group by

Save

X Clear

< > 1-50 of 1159 items

Barcode	Name	Location	Modified	Schema
4C002	4C EE&SB fridge transient storage	DTU Buildi...	10/12/2020	4°C Fridge
4C002	4C Fridge 00271	DTU Buildi...	09/08/2018	4°C Fridge
4C009	4C Fridge 01223			4°C Fridge
4C008	4C Fridge 01233			4°C Fridge
4C014	4C Fridge 01871			4°C Fridge
4C015	4C Fridge Aaron	BioInnovati...	15/04/2021	4°C Fridge
4C016	4C Fridge Adam			4°C Fridge
4C005	4C Fridge ANALYTICS			4°C Fridge
4C011	4C Fridge CFB00266			4°C Fridge
4C01478	4C Fridge CFB01478	DTU Buildi...	19/11/2018	4°C Fridge
4C01653	4C Fridge CFB01653	DTU Buildi...	19/11/2018	4°C Fridge
4C003	4C Fridge DSP1	DTU Buildi...	09/08/2018	4°C Fridge

Part 1

The Basics of Benchling

An introduction to our **Laboratory Information Management System (LIMS)**

Reach out when struggling with Benchling:

Biosustain Benchling support

lims_support@biosustain.dtu.dk

Access Benchling:

biosustain.benchling.com

(login with DTU credentials)

Agenda

Introduction to Benchling
and best practices

~ 30 min

Hands-on

~ 15 min

Agenda

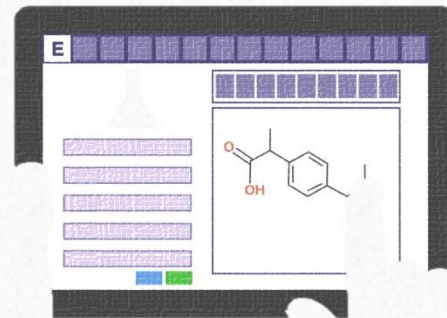
Introduction to Benchling
and best practices

~ 30 min

Hands-on

~ 15 min

What is a LIMS?



What is a LIMS?

L.I.M.S = Laboratory Information Management System

- It **keeps track of laboratory data** associated with samples and experiments
- At Biosustain, we use **Benchling**, a Cloud-based platform



biosustain.benchling.com

Main functionalities

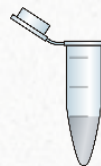
ELN

- Experiment notes
- Registration of samples



Samples storage

- Samples transfer in boxes/plates



Molecular Biology tool

- Import of sequences
- Plasmid design and annotation



Requests

- Samples submission for analysis
- Lab material order request



Working with Benchling

Benefits

- Facilitates **passing over** of projects
- Foster **collaboration**
- Promotes data capture in a **defined place** preventing its loss
- Make knowledge **findable**
- Promotes the adoption of **common practices** across research groups

Working with Benchling

Obstacles

- Adapting to a **new way of working**
- **Software limitations** and “pain points”
- **Learning curve**
 - Many functionalities
 - Complex / confusing data model

Good practices

- ✓ Record all necessary information to make your experiment **clear to others** and reproducible
- ✓ Register important data (strains, media, plasmids...)
- ✓ Keep your project folders organized and use **clear names** and **descriptions**

Good practices

- ✓ Make sure **not** to save relevant data in the **Biosustain Training** project folder



Projects / Biosustain Training /
Inventory Saved Searches

Search

< > 1-100 of 2227 items

We noticed that the list include real data



Good practices

- ✓ Reach out when struggling using the platform
- ✓ There might be a **quicker** and **easier** way to do what you are doing !

Submit your questions to
lims_support@biosustain.dtu.dk

Get hands-on Benchling support

Mondays 13:00 -14:00 (Room 222)

DTU
Research Data
Management Team

DROP-IN HOURS

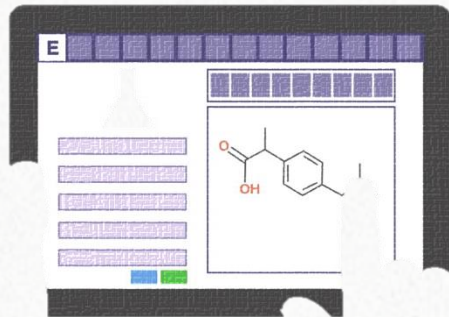
Get hands-on support for **Benchling** and
other **data management** tasks.

MONDAYS
13:00 – 14:00

3rd FLOOR

Image by Joamp on Freepik

Getting started



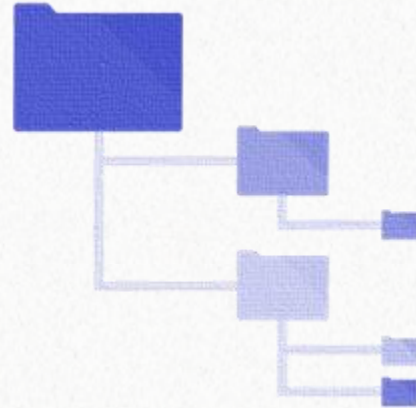
Step 1: Create your project folder

Step 2: Create your experiment ELN & register samples

Step 3: Navigate through your data

Step 1:

Create your project folder



Project folder

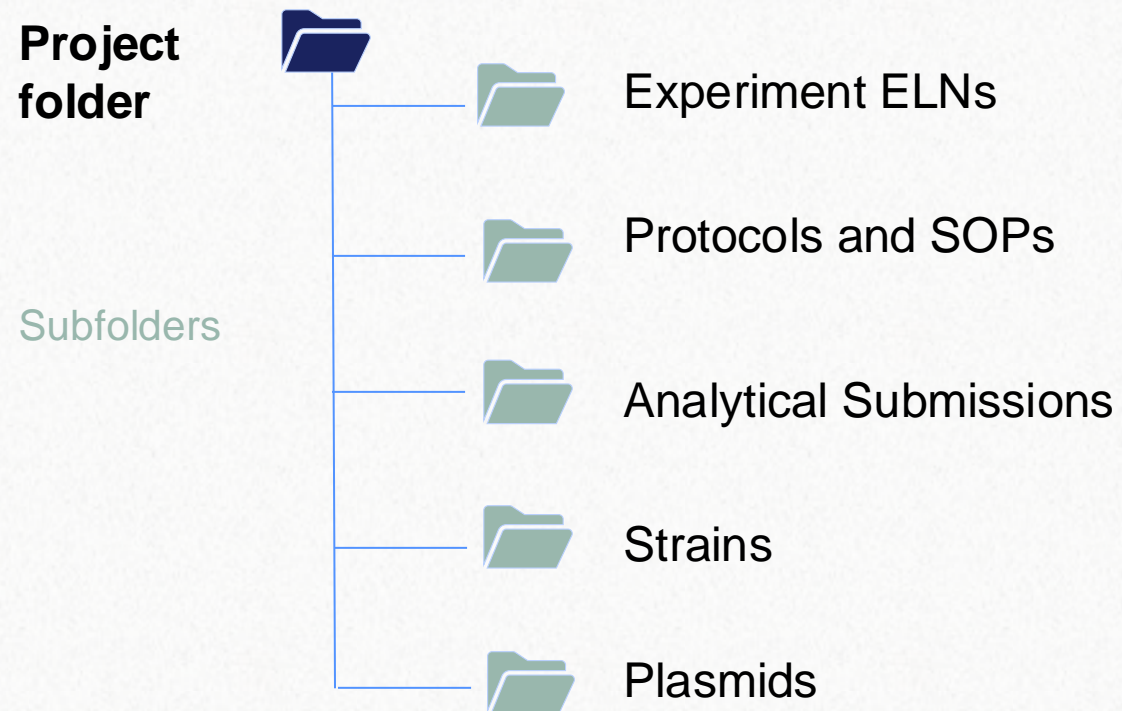
Your Project folder will contain:

✓ your Electronic Notebook pages (called “**Entries**”) 

✓ your registry items (called “**Entities**”) strains, plasmids,...



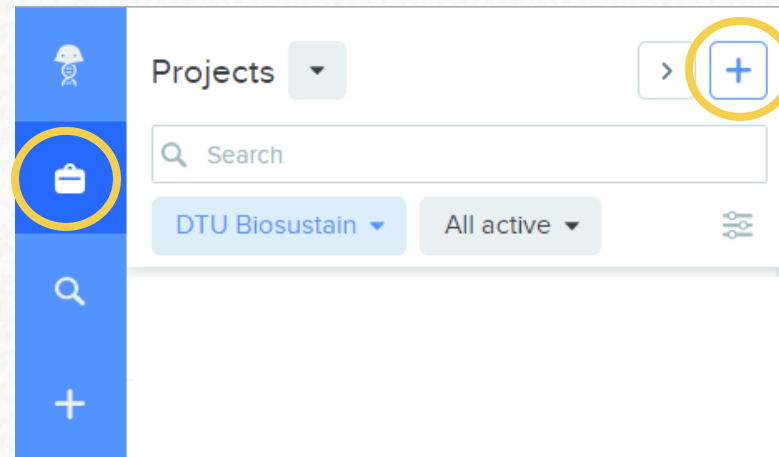
Example of folder structure



Project folder

To create a New Project:

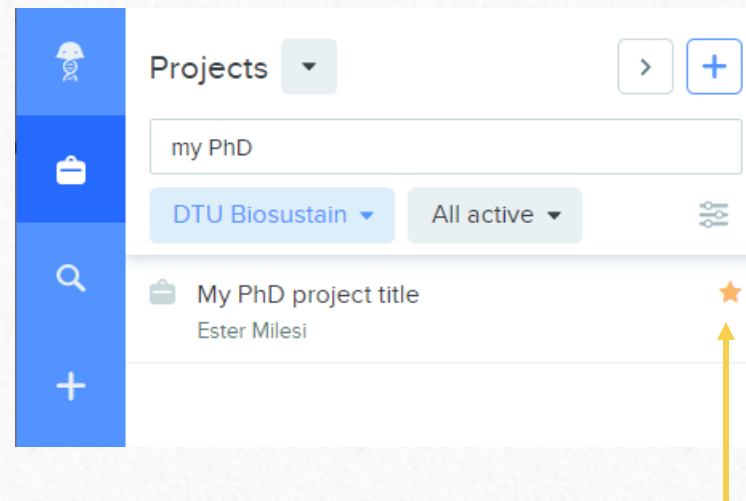
- Click on the “Project” icon
- Click on the “+” icon
- Give the folder a clear name (e.g., your PhD project title)



Project folder

Star ★ your Project:

- This way, it will appear on top of every other Project that you have access to



Project folder

Give access to your team

- Almost all Research groups have a **Benchling Team**
- When possible, add **your Team** among the collaborators of the Project

The screenshot shows the Benchling interface. At the top, there is a navigation bar with a blue sidebar containing icons for home, search, and other functions. The main content area shows a project titled "My PhD project title" with a settings gear icon. A yellow arrow points from this gear icon to the "Project Settings" dialog box.

The "Project Settings" dialog box has a "Name" field containing "My PhD project title". Below this is a "Manage collaborators" section with a search bar and an "Add collaborator" button. The "Collaborators" section is a table with the following data:

User / App / Organization / Team	Access policies	Auditor
Members of DTU Biosustain	NONE	<input type="checkbox"/>
Ester Milest	ADMIN	<input type="checkbox"/>
DNA Foundry (NGS)	READ	<input type="checkbox"/>
Admins of DNA Foundry (NGS)	ADMIN	<input type="checkbox"/>

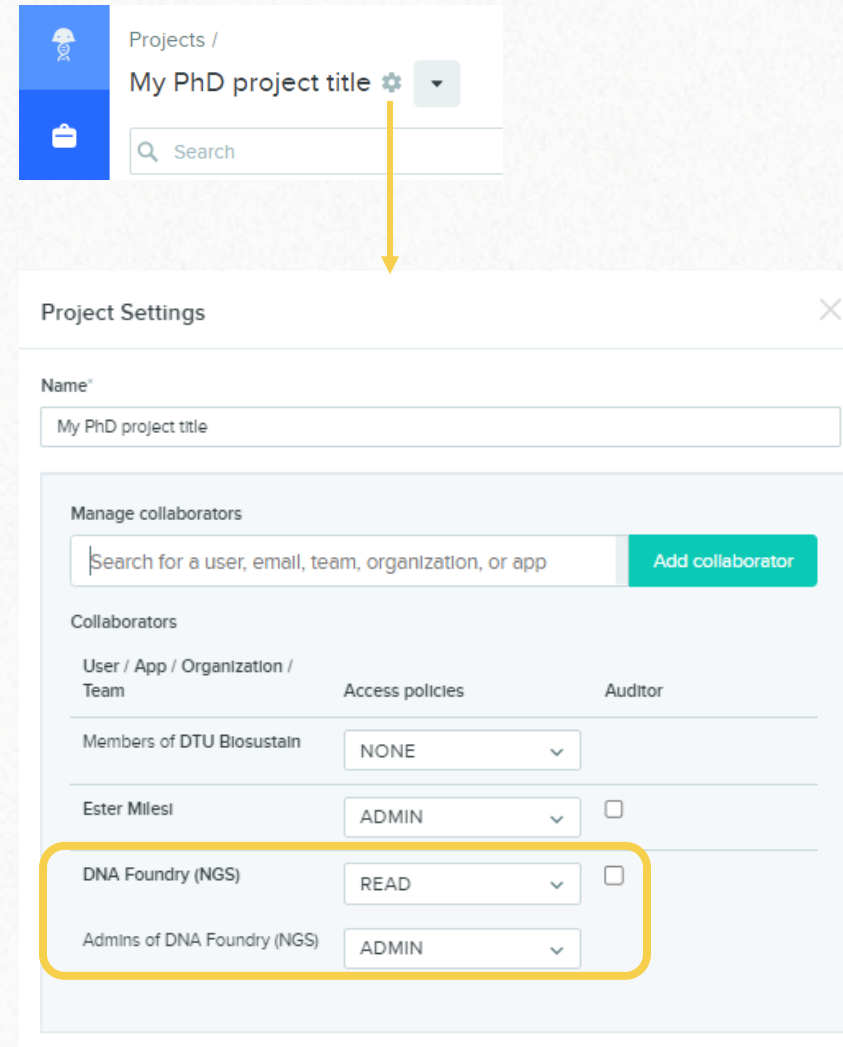
The row for "DNA Foundry (NGS)" is highlighted with a yellow border.



Project folder

Benefits

- ✓ You don't have to add each member one-by-one
- ✓ When new researchers join the Team, **they get automatically access to all shared Project folders**



The screenshot shows a 'Project Settings' dialog box. At the top, the project name is 'My PhD project title'. Below this is a 'Manage collaborators' section with a search bar and an 'Add collaborator' button. The 'Collaborators' table lists several users and their access levels:

User / App / Organization / Team	Access policies	Auditor
Members of DTU Biosustain	NONE	<input type="checkbox"/>
Ester Milest	ADMIN	<input type="checkbox"/>
DNA Foundry (NGS)	READ	<input type="checkbox"/>
Admins of DNA Foundry (NGS)	ADMIN	<input type="checkbox"/>

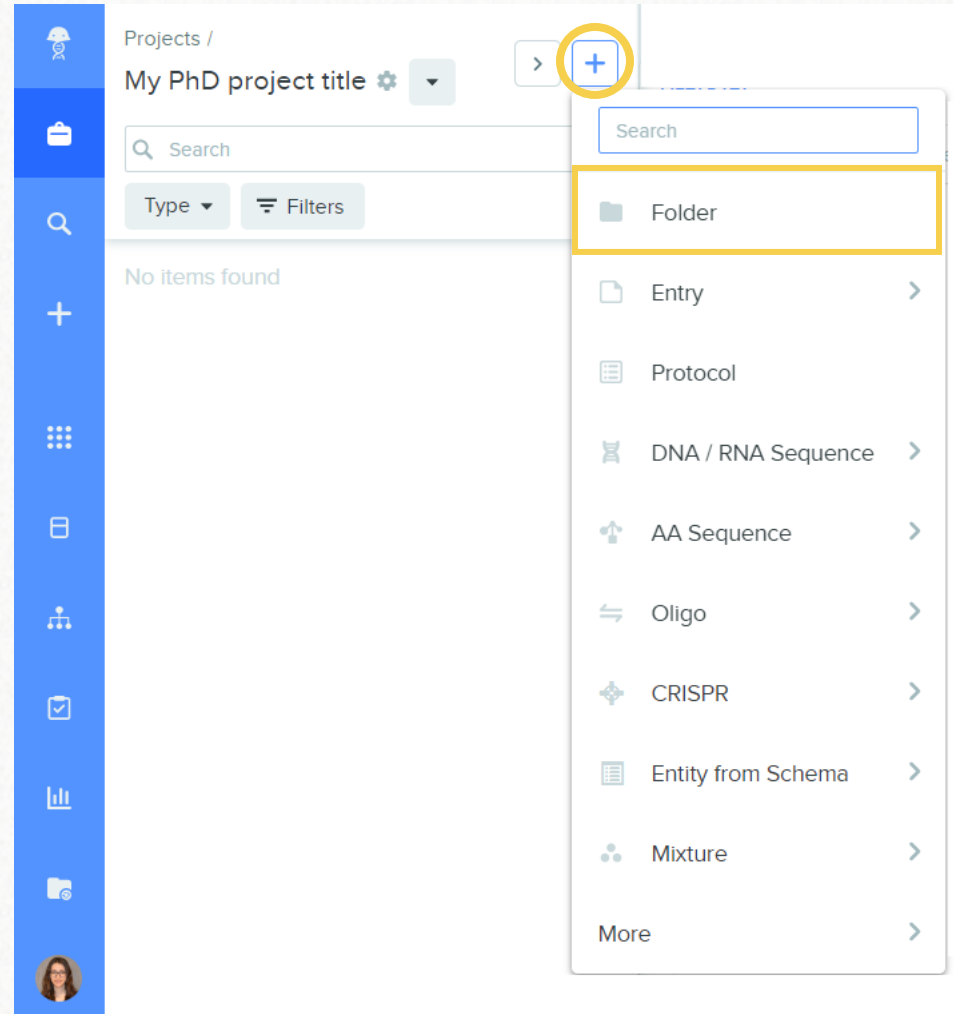
The 'DNA Foundry (NGS)' row is highlighted with a yellow box.



Creating subfolders

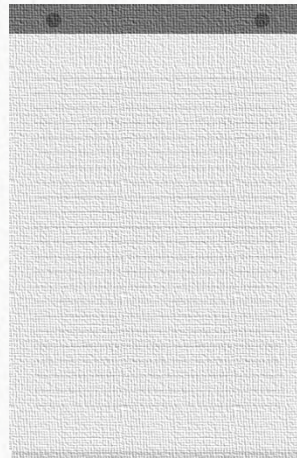
To create a subfolder:

- Enter your newly created Project
- Click on the “+” icon
- Select “Folder”



Step 2:

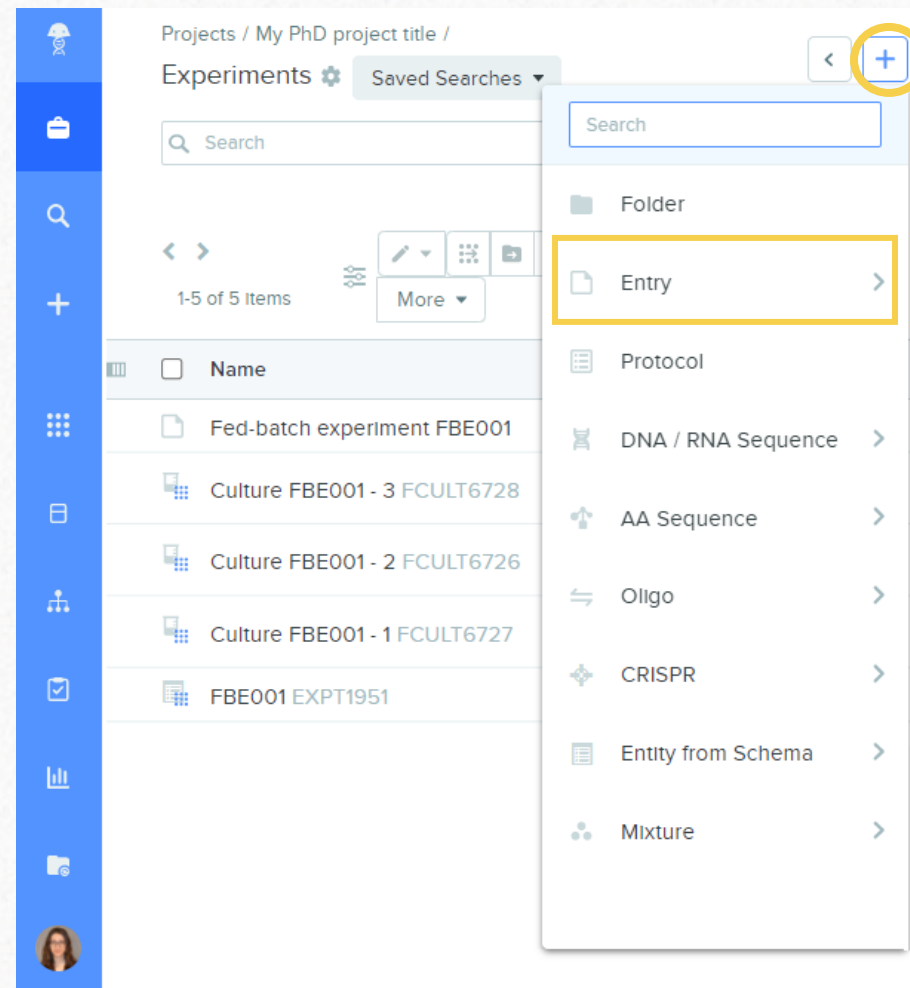
Create your experiment ELN & create samples



Electronic notebook

To create a new Entry:

- Go to the relevant folder (e.g., “Experiments”)
- Click on the “+” icon
- Select “Entry”



Electronic notebook

Option 1:

Blank entry

Option 2:

New Entry from Template

--- you can create your own!

The screenshot displays the interface of an electronic notebook. At the top, the breadcrumb path is 'Projects / My PhD project title / Experiments'. A search bar is visible. Below the search bar, there are navigation arrows and a 'More' button. A list of experiments is shown with columns for 'Name' and 'ID'. The list includes:

- Fed-batch experiment FBE001
- Culture FBE001 - 3 FCULT6728
- Culture FBE001 - 2 FCULT6726
- Culture FBE001 - 1 FCULT6727
- FBE001 EXPT1951

A 'More' menu is open, showing various options for creating new entries. The options are:

- Folder
- Entry
- Protocol
- DNA / RNA Sequence
- AA Sequence
- Oligo
- CRISPR
- Entity from Schema
- Mixture

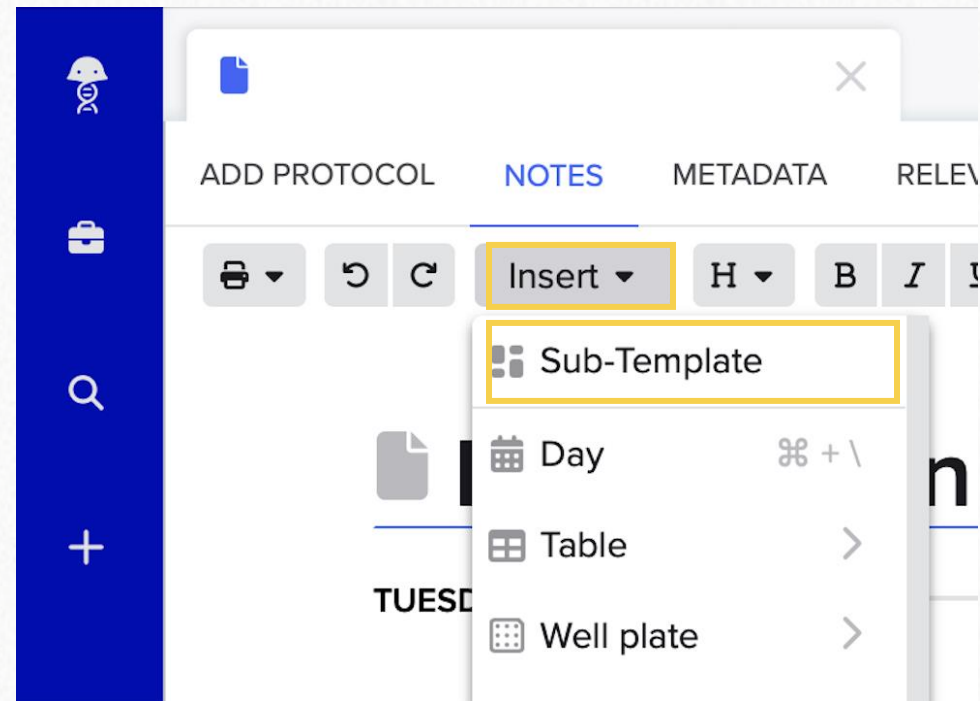
Two options are highlighted with yellow boxes: 'Blank entry' and 'Entry from template'. A yellow circle highlights the '+' icon in the top right corner of the interface.

Electronic notebook

Option 3:

Blank entry + Sub-Template

--- you can create your own!

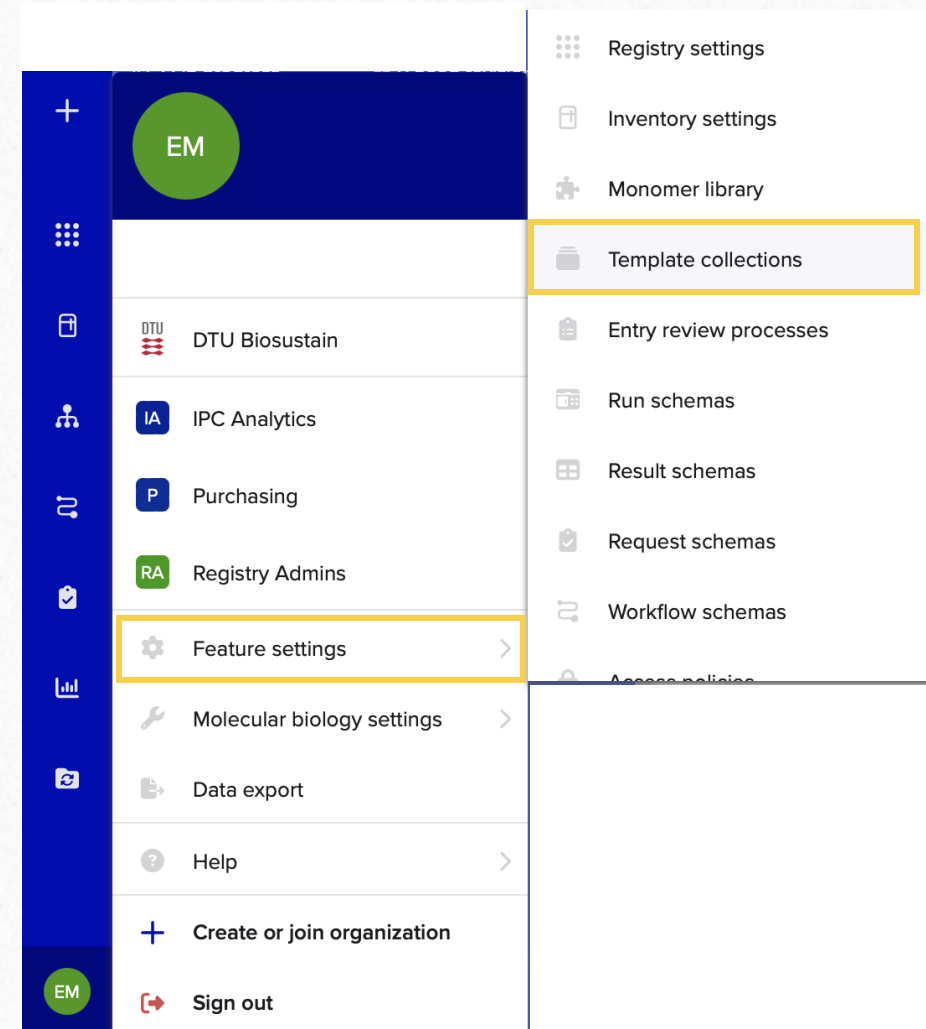


Electronic notebook

To create a Template or a Sub-template:

- Go to your profile
- Go to Feature settings
- Go to Template collections

Create your own!



Electronic notebook

To create a Template or a Sub-template:

- Go to your profile
- Go to Feature settings
- Go to Template collections

Create your own!

Template Collections > My templates

Search All items Filtering by: Not archived Create

<< < > >> 1-4 of 4 items

NAME	TYPE	AUTHOR	CREATED

Template
Sub-template

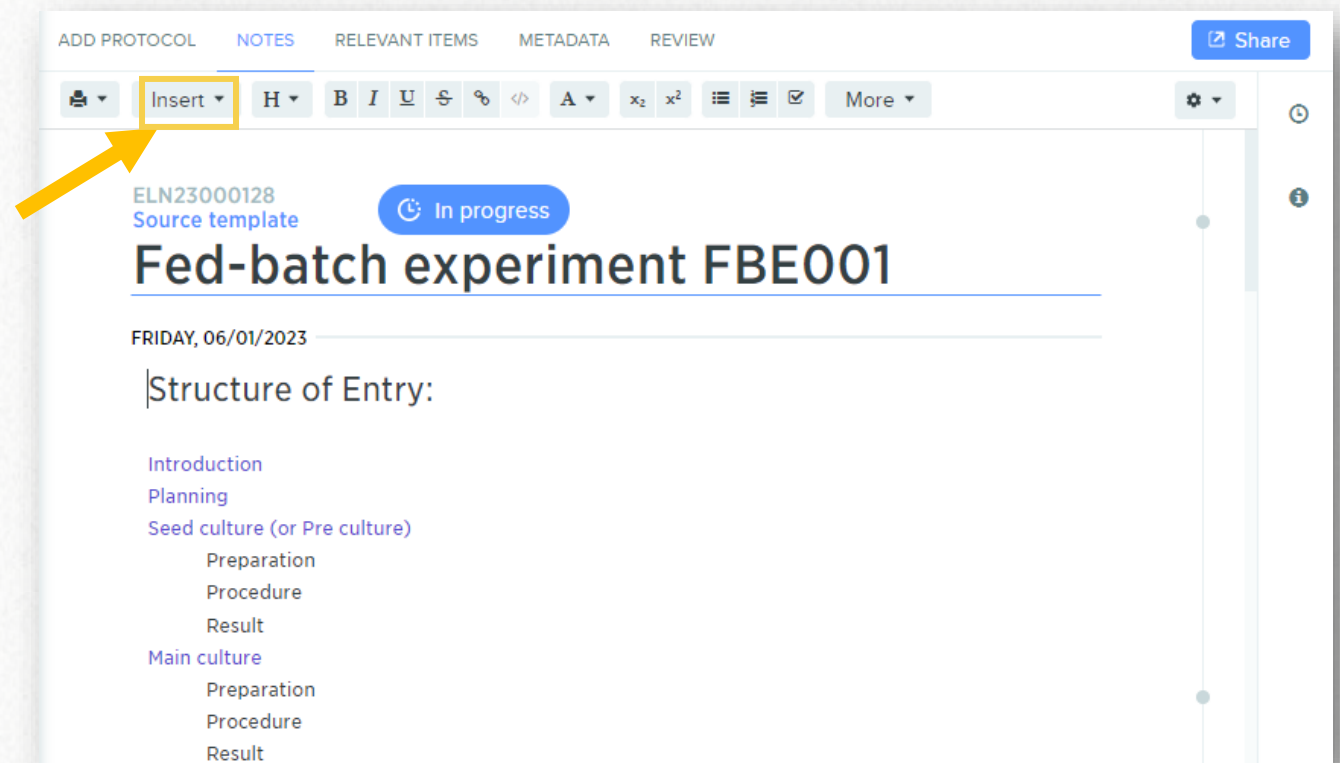


Electronic notebook

In your Entry you can:

- Take **notes**
- Add **attachments/files**
- Create **tables**

Click on **Insert** to see the whole list of options



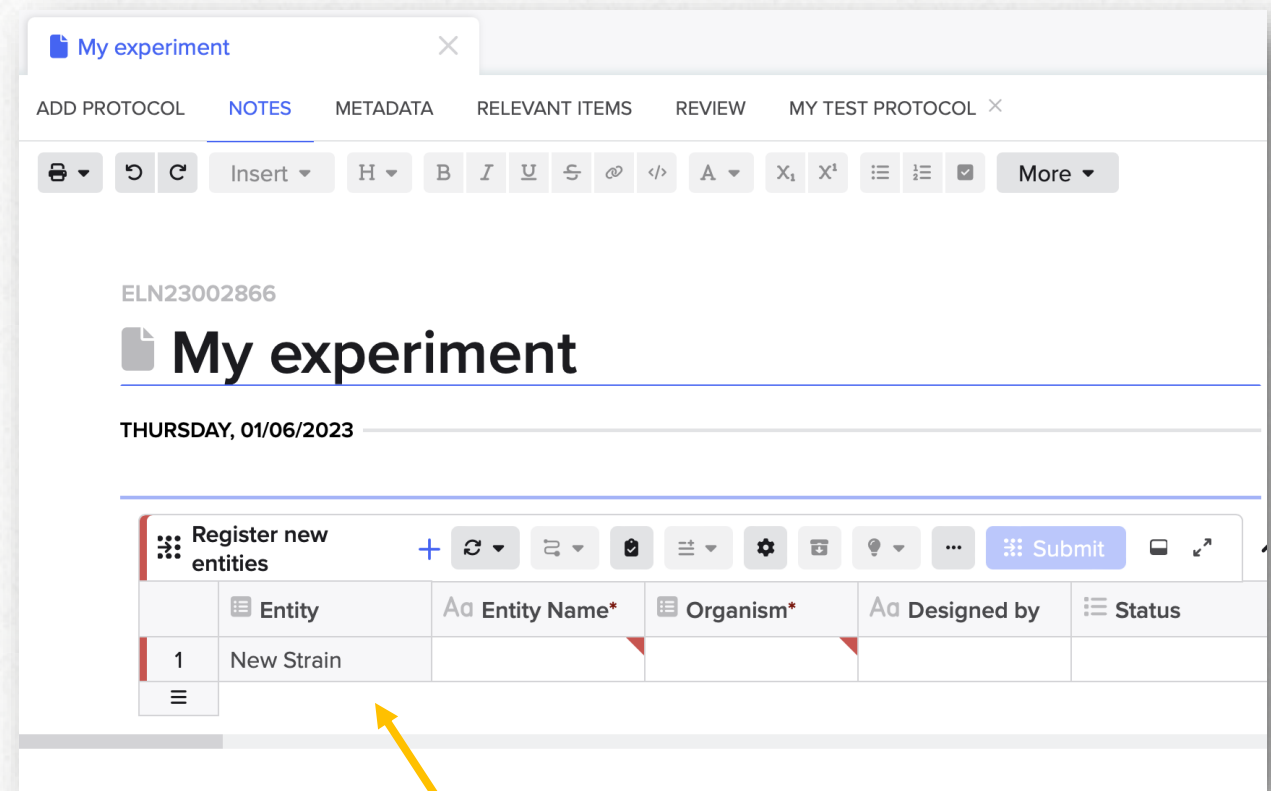
The screenshot displays the interface of an electronic notebook. At the top, there are navigation tabs: 'ADD PROTOCOL', 'NOTES', 'RELEVANT ITEMS', 'METADATA', and 'REVIEW'. A 'Share' button is located in the top right corner. Below the navigation is a rich text editor toolbar with various icons for text formatting (bold, italic, underline, strikethrough, link, unlink, code), alignment, and other functions. The 'Insert' menu is highlighted with a yellow box, and a yellow arrow points to it. The main content area shows an entry titled 'ELN23000128 Source template' with a status 'In progress'. The entry title is 'Fed-batch experiment FBE001'. Below the title, the date 'FRIDAY, 06/01/2023' is displayed. The main text area contains the text 'Structure of Entry:' followed by a list of sections: 'Introduction', 'Planning', 'Seed culture (or Pre culture)', 'Preparation', 'Procedure', 'Result', 'Main culture', 'Preparation', 'Procedure', and 'Result'.

Electronic notebook

In your Entry you can:

- Register strains, media, etc. using **Registration tables**
- Assign **storage location** to registered entities

= more of this in the *Hands-on*



The screenshot shows a web interface for an electronic notebook. At the top, there's a tab labeled "My experiment" and a navigation bar with options: "ADD PROTOCOL", "NOTES", "METADATA", "RELEVANT ITEMS", "REVIEW", and "MY TEST PROTOCOL". Below this is a rich text editor toolbar with various icons for text formatting and insertion. The main content area displays the entry ID "ELN23002866" and the title "My experiment" with a date "THURSDAY, 01/06/2023". A "Register new entities" table is visible, which is a registration table for strains. The table has columns for "Entity", "Entity Name*", "Organism*", "Designed by", and "Status". A single row is shown with "New Strain" in the "Entity" column. A yellow arrow points to this row.

	Entity	Aa Entity Name*	Organism*	Aa Designed by	Status
1	New Strain				

A Registration table for strains

Electronic notebook

In your Entry you can:

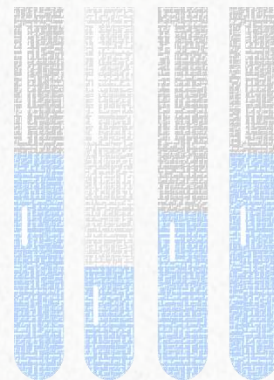
- Tag your plasmids, sequences, etc. (everything that you create)
- Tag another ELN or SOP

To tag an object, type @...

The screenshot shows a web-based interface for an electronic notebook. At the top, there is a tab labeled 'My experiment' with a close button. Below the tab are navigation options: 'ADD PROTOCOL', 'NOTES' (which is selected), 'METADATA', 'RELEVANT ITEMS', and 'REVIEW'. A rich text editor toolbar is visible, containing icons for undo, redo, insert, heading, bold, italic, underline, strikethrough, link, code, and text color. The main content area displays the entry ID 'ELN23002866' and the title 'My experiment' underlined. Below the title is the date 'THURSDAY, 01/06/2023'. Two tags are present: 'pUC18' with a plasmid icon and 'SOP' with a document icon. Two yellow arrows point from text labels to these tags: 'Link to a plasmid' points to 'pUC18' and 'Link to a SOP' points to 'SOP'.

Step 3:

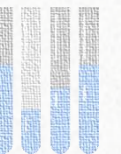
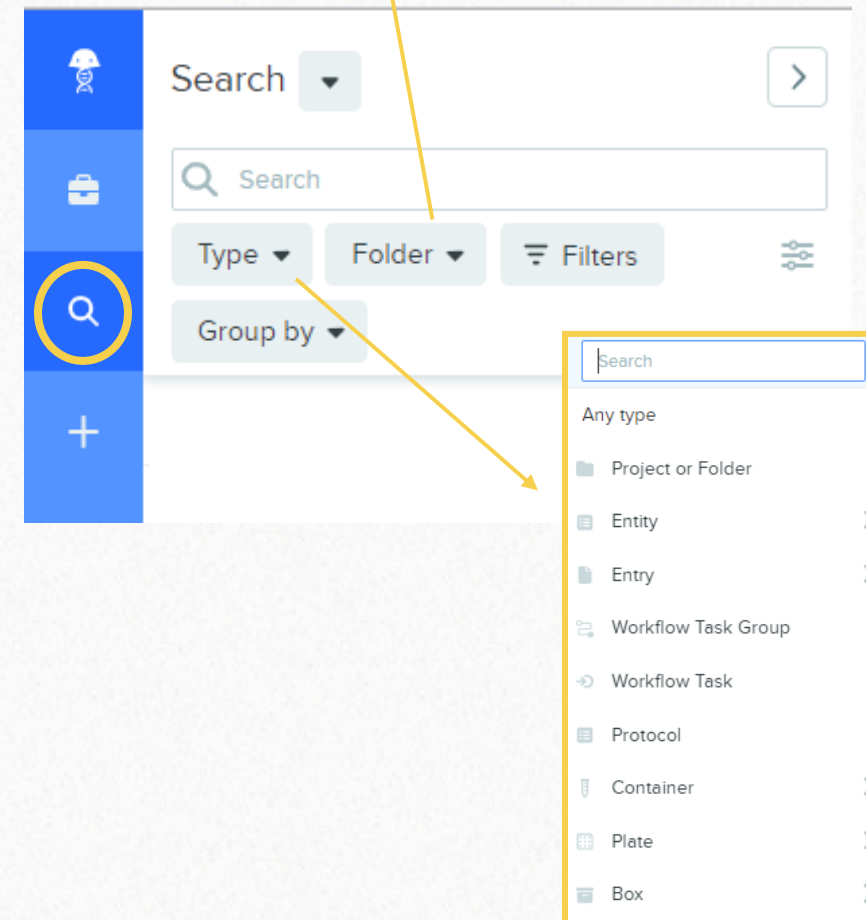
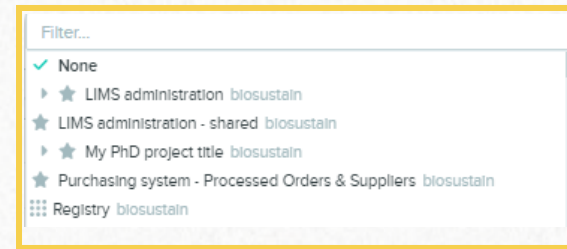
Navigate through your data



Global search

It allows to search through all your data and filter by:

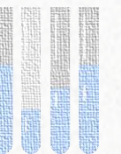
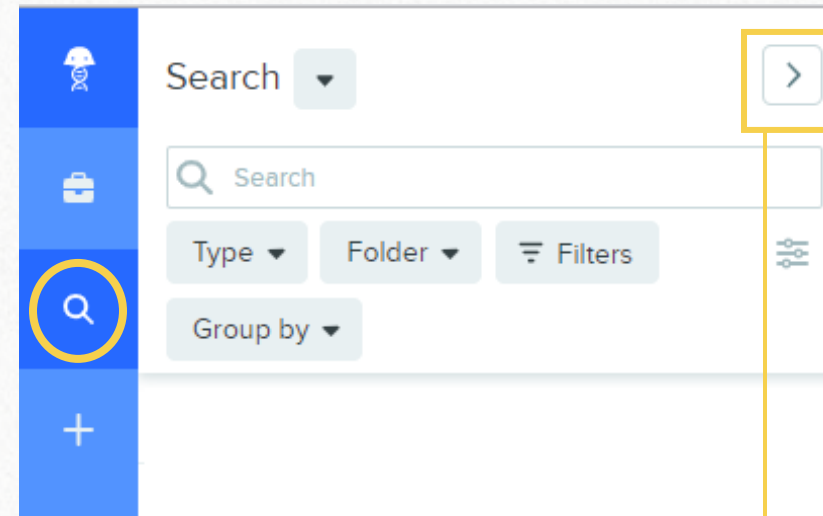
- Data type (samples, boxes...)
- Folder
- Metadata field



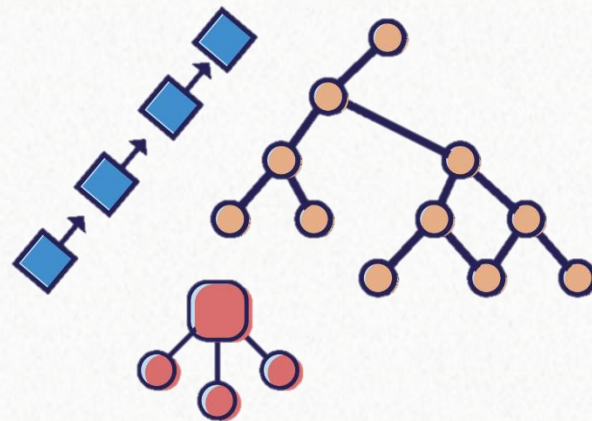
Global search

Expand view to do bulk actions on your data:

- Bulk **edit** of metadata fields
- Bulk **registration** of imported sequences
- Bulk **transfer** to a different folder
- Bulk **archive**



Benchling entities: what you need to know



Registering entities: what to consider

1. Entities can store different information

Entity types that can store:

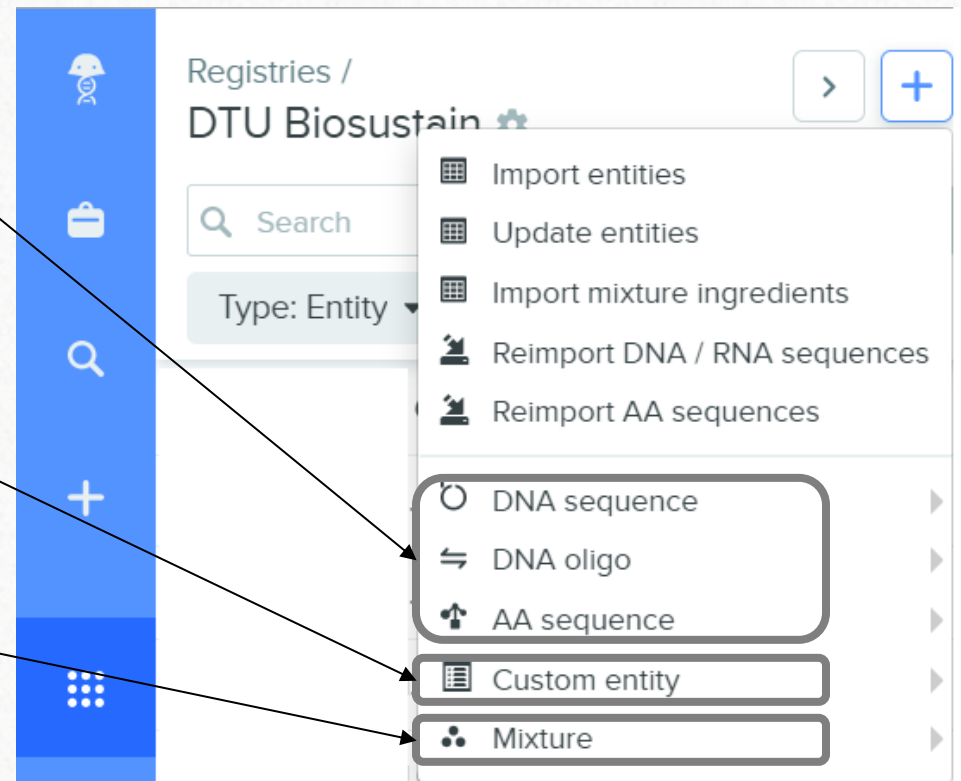
- metadata
- a sequence

Entity type that can store:

- metadata




Entity type that can store:

- metadata
- media ingredients and recipe

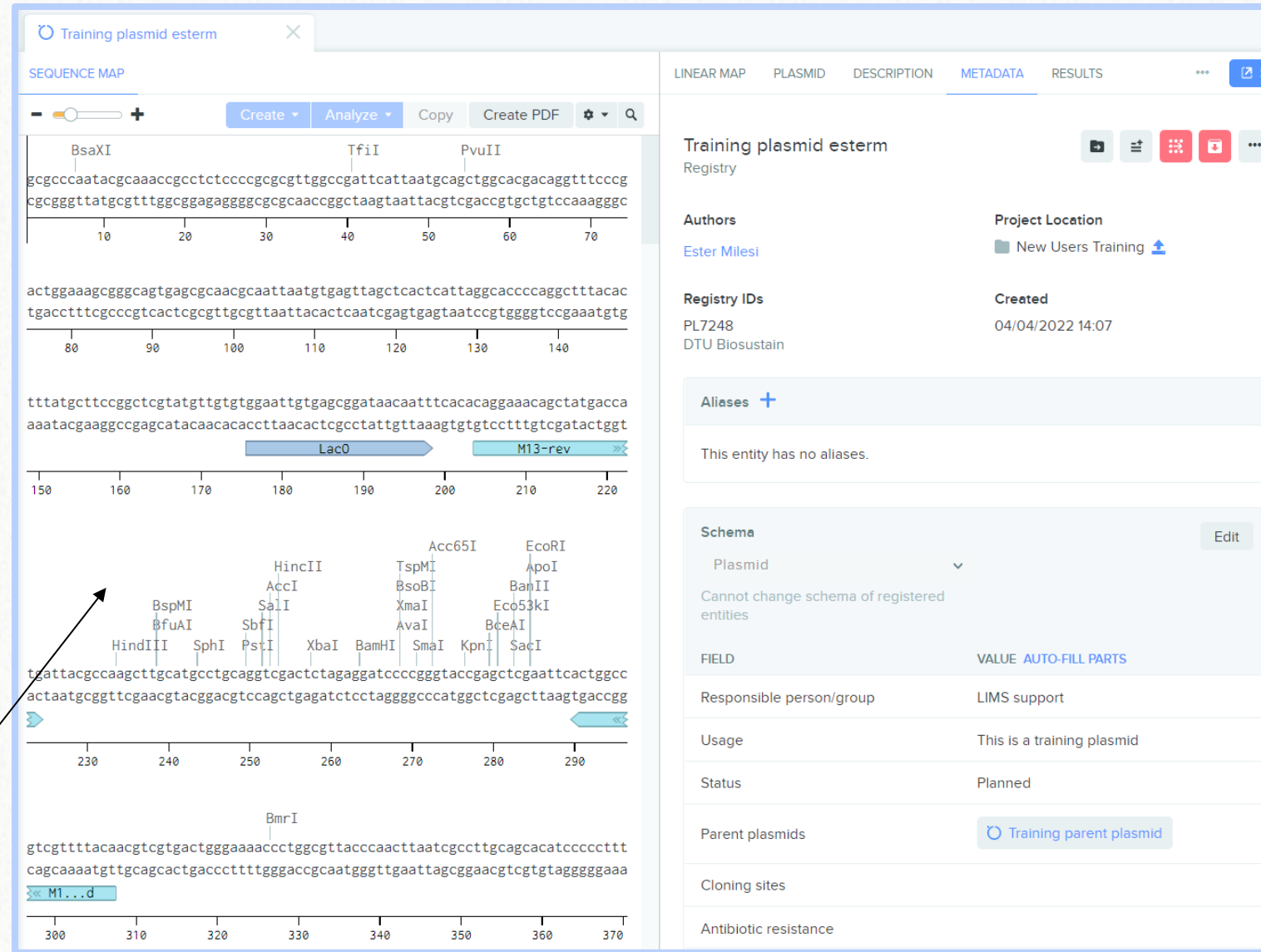


Entity types that can store:

- metadata
- a sequence

-  DNA sequence
-  DNA oligo
-  AA sequence

Sequence



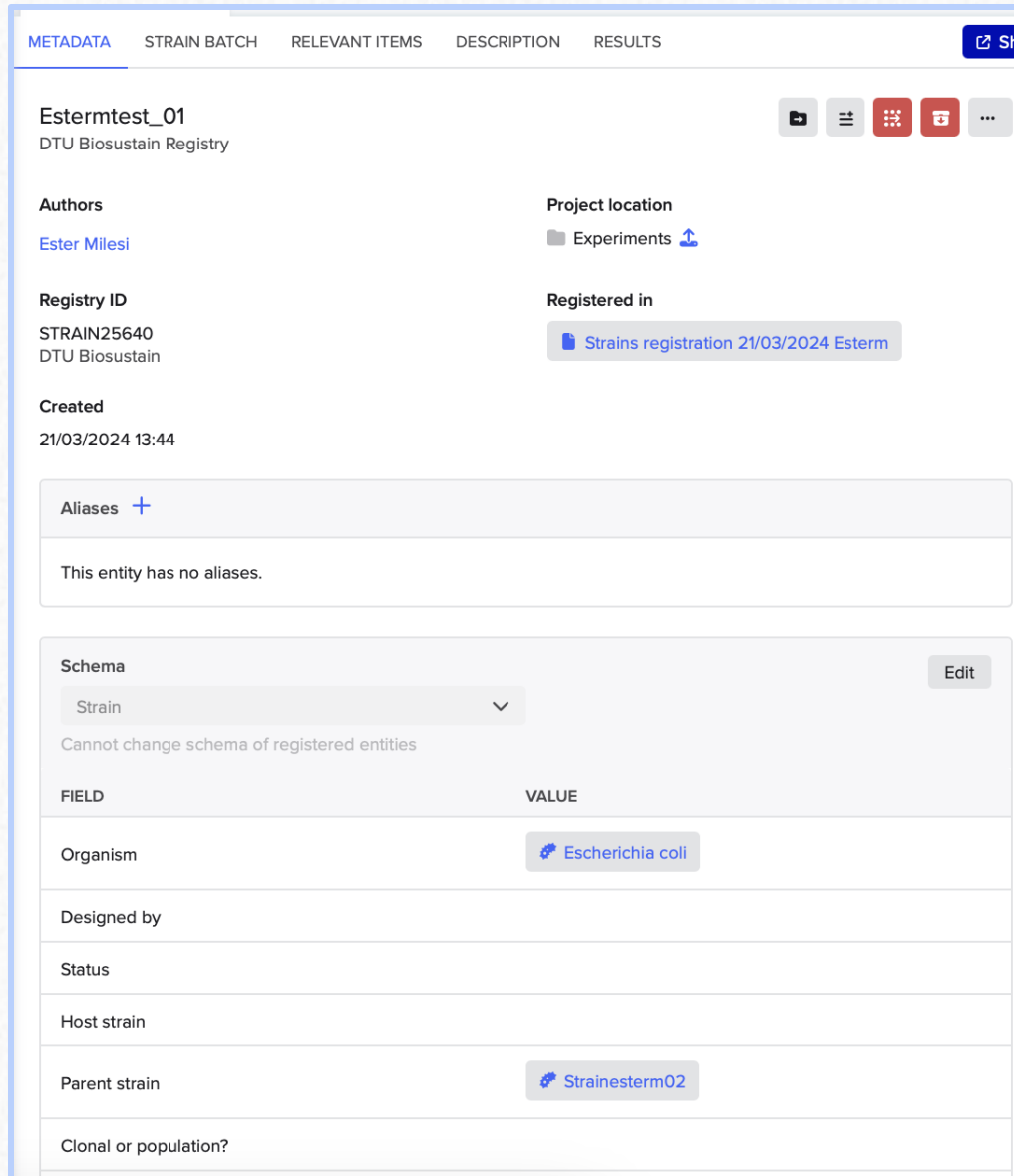
The screenshot shows a web interface for a plasmid registry. The left pane displays a 'SEQUENCE MAP' for a plasmid named 'Training plasmid esterm'. The map shows a DNA sequence with various restriction enzyme sites (BsaXI, TfiI, PvuII, LacO, M13-rev, etc.) and their positions along the sequence. The right pane shows the 'METADATA' for this plasmid, including authors (Ester Milesi), project location (New Users Training), registry IDs (PL7248, DTU Biosustain), and creation date (04/04/2022 14:07). A table below the metadata lists fields like 'Responsible person/group' (LIMS support), 'Usage' (This is a training plasmid), 'Status' (Planned), and 'Parent plasmids' (Training parent plasmid).



Entity types that can store:

- metadata

Custom entity



The screenshot displays the 'METADATA' tab for the entity 'Estermtest_01' in the DTU Biosustain Registry. The interface includes a top navigation bar with tabs for METADATA, STRAIN BATCH, RELEVANT ITEMS, DESCRIPTION, and RESULTS. The entity name 'Estermtest_01' is shown with its registry ID 'DTU Biosustain Registry'. Below this, there are sections for Authors (Ester Milesi), Registry ID (STRAIN25640), Created date (21/03/2024 13:44), Project location (Experiments), and Registered in (Strains registration 21/03/2024 Esterm). An Aliases section indicates that the entity has no aliases. A Schema section shows the entity is of type 'Strain' and cannot be changed. A table below lists fields and their values: Organism (Escherichia coli), Parent strain (Strainesterm02), and Clonal or population? (empty).

FIELD	VALUE
Organism	Escherichia coli
Designed by	
Status	
Host strain	
Parent strain	Strainesterm02
Clonal or population?	



Entity types that can store:

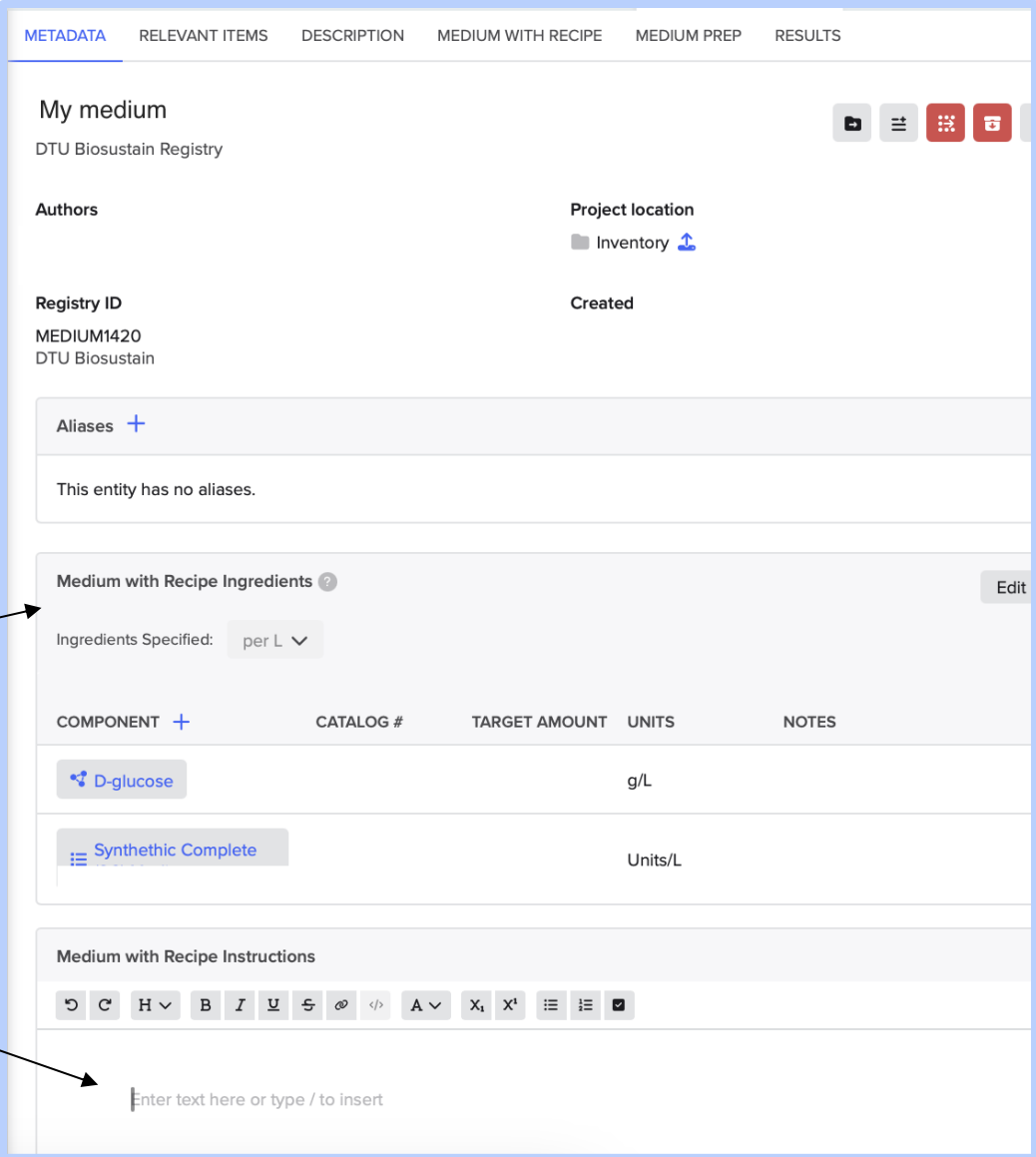
- metadata
- media ingredients and recipe

 Mixture

Component list
(with amount)

Recipe

Metadata



METADATA RELEVANT ITEMS DESCRIPTION MEDIUM WITH RECIPE MEDIUM PREP RESULTS

My medium
DTU Biosustain Registry

Authors Project location
Inventory

Registry ID Created
MEDIUM1420
DTU Biosustain

Aliases +
This entity has no aliases.

Medium with Recipe Ingredients Edit

Ingredients Specified: per L

COMPONENT +	CATALOG #	TARGET AMOUNT	UNITS	NOTES
D-glucose			g/L	
Synthetic Complete			Units/L	

Medium with Recipe Instructions

Enter text here or type / to insert

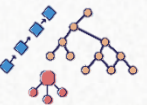
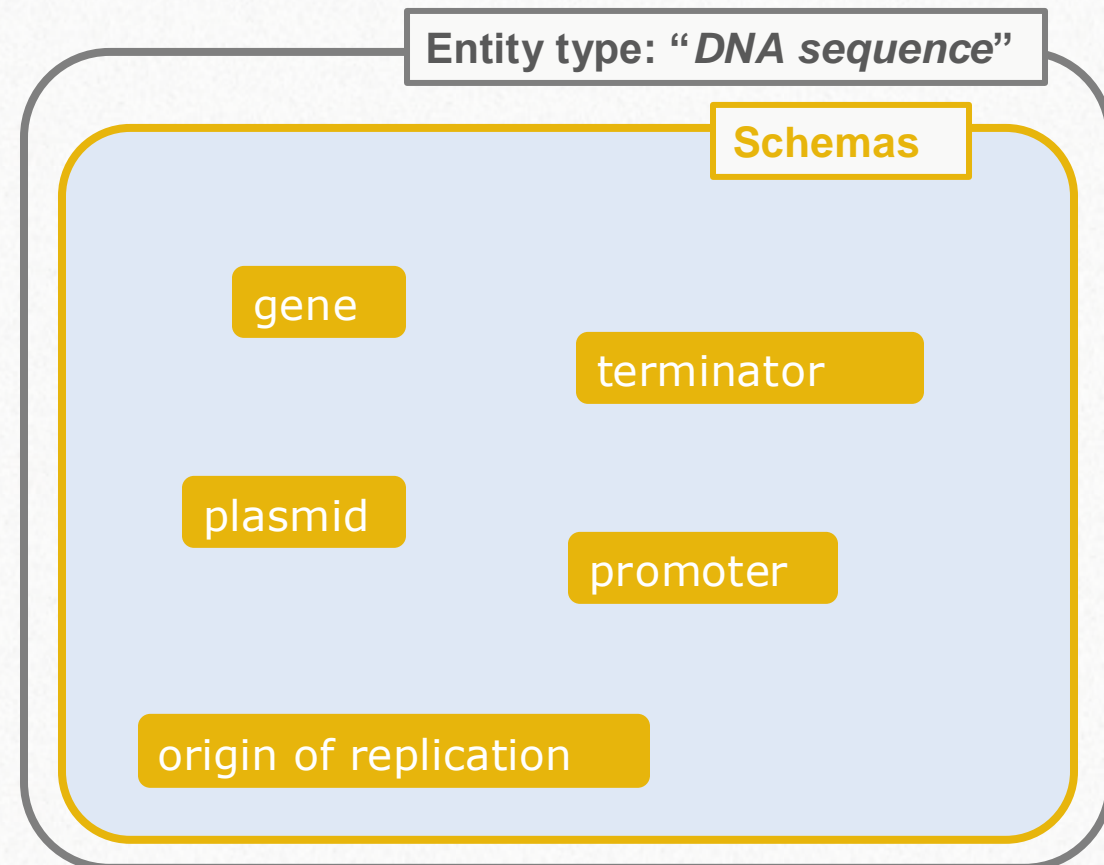


Registering entities: what to consider

2. Entities are assigned a “schema”

→ The “**type**” only tells you which information the entity stores

The various sample types are described by “**schemas**”

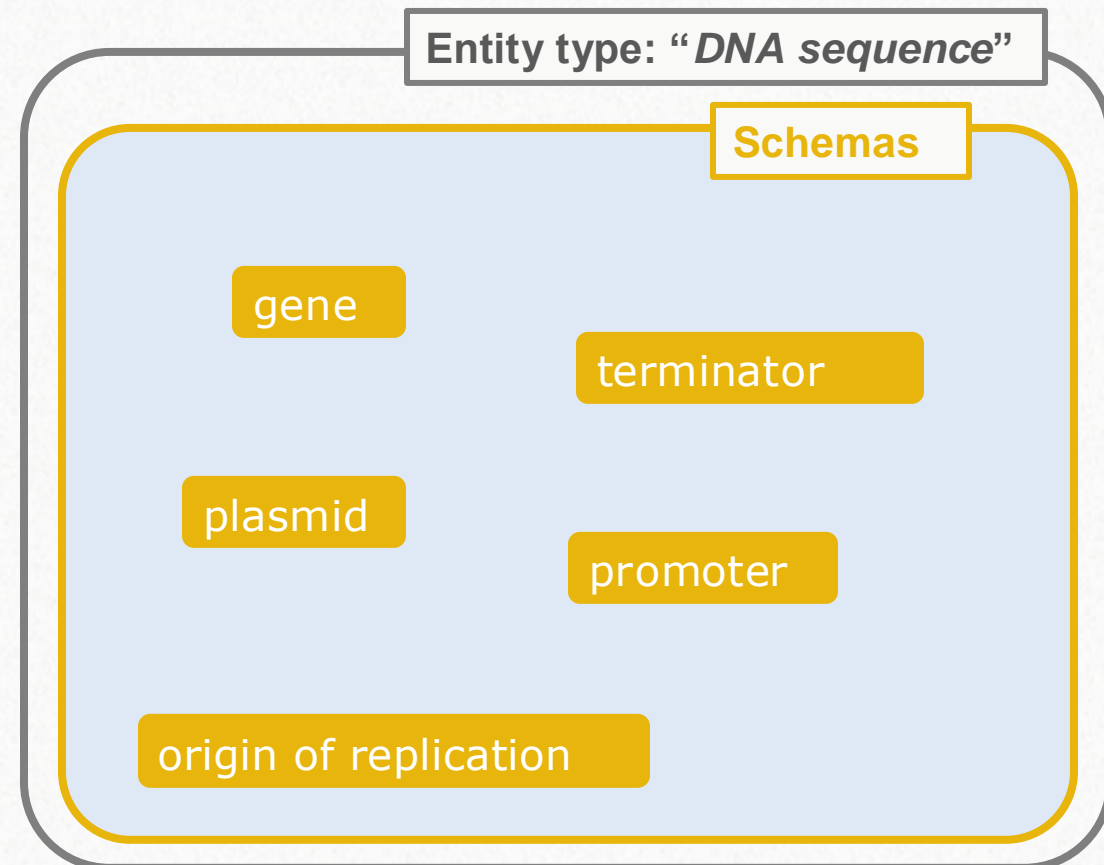


Registering entities: what to consider

2. Entities are assigned a “schema”

A schema specifies:

- sample type
- required information to fill-in
- **links** to other schemas

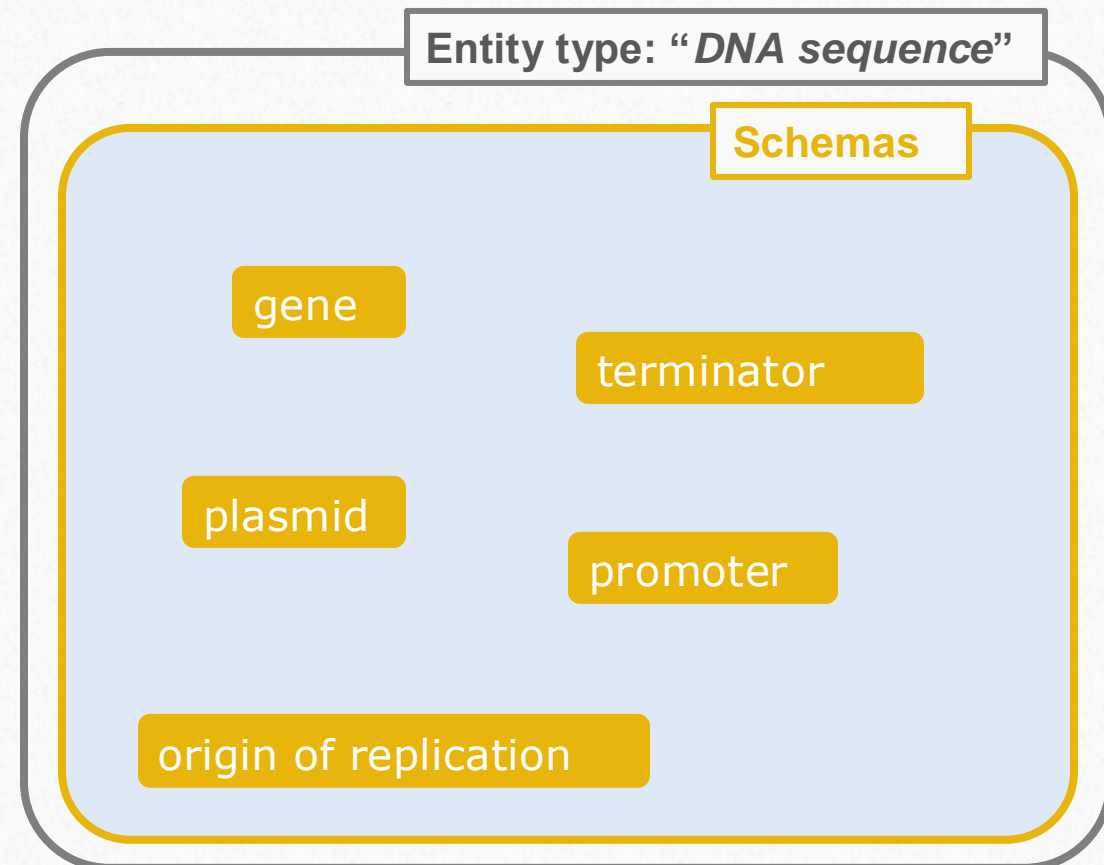


Registering entities: what to consider

2. Entities are assigned a “schema”

A schema specifies:

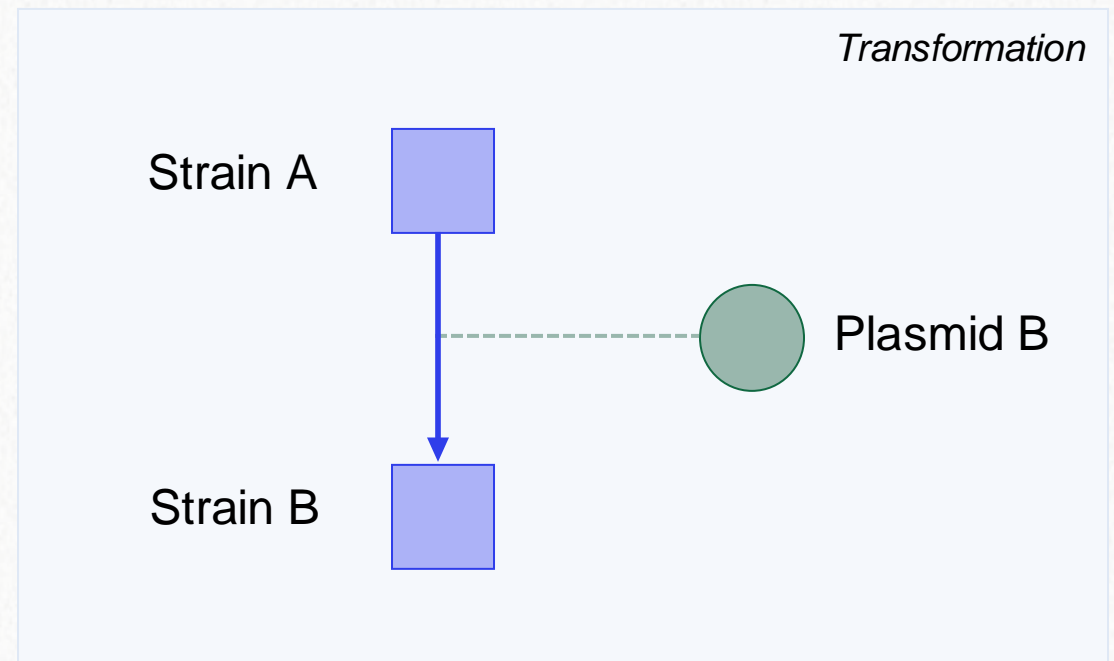
- sample type
- required information to fill-in
- links to other schemas



Registering entities: what to consider

2. Entities are assigned a “schema”

For example, the schema “**Strain**” can link to another strain (parent) and to a plasmid



Links  allow to track the sample “history”



Registering entities: what to consider

3. You can create entities 1-by-1 or in bulk

To create entities 1-by-1:

- Go to Registry > Click on the “+” icon
- Select the **entity type** e.g., “DNA sequence”
- Select the **schema** e.g., “Plasmid”

The screenshot shows the 'Registries / DTU Biosustain' interface. A search bar is visible at the top. A vertical blue sidebar on the left contains several icons, with the grid icon (representing 'Add') circled in yellow. A dropdown menu is open, listing various entity types: DNA Fragment, Gene, gRNA, Marker, Origin of Replication, Plasmid, Promoter, Tag, and Terminator. A second dropdown menu is open over 'DNA sequence', listing schemas: DNA sequence, DNA oligo, AA sequence, Custom entity, and Mixture.



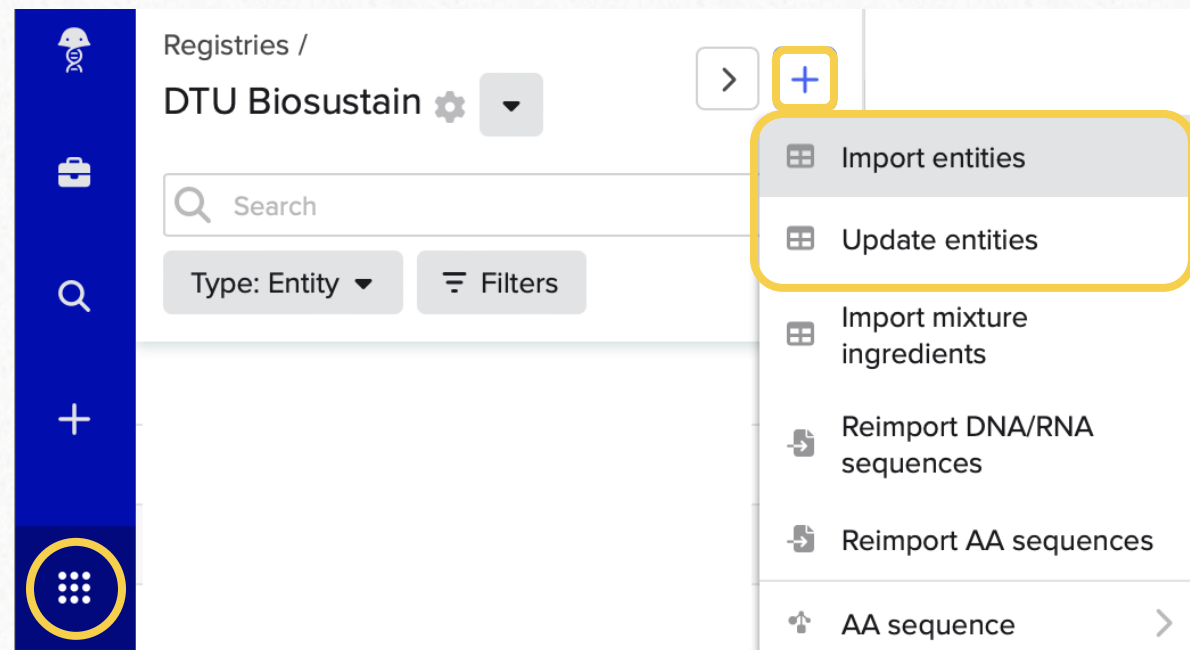
Registering entities: what to consider

3. You can create entities 1-by-1 or in bulk

To create entities in bulk:

- Upload a **spreadsheet**
Select *import* or *update entities*

(make sure that the values in the cells are the one that Benchling expects)



The screenshot shows the Benchling web interface. On the left is a dark blue sidebar with icons for home, search, add, and a grid icon (highlighted with a yellow circle). The main content area is titled 'Registries / DTU Biosustain' with a settings gear and a dropdown arrow. Below the title is a search bar, a 'Type: Entity' dropdown, and a 'Filters' button. A yellow box highlights a '+' icon in the top right corner, which has opened a dropdown menu. The menu contains the following options: 'Import entities' (highlighted with a grey background), 'Update entities', 'Import mixture ingredients', 'Reimport DNA/RNA sequences', 'Reimport AA sequences', and 'AA sequence' with a right-pointing arrow.



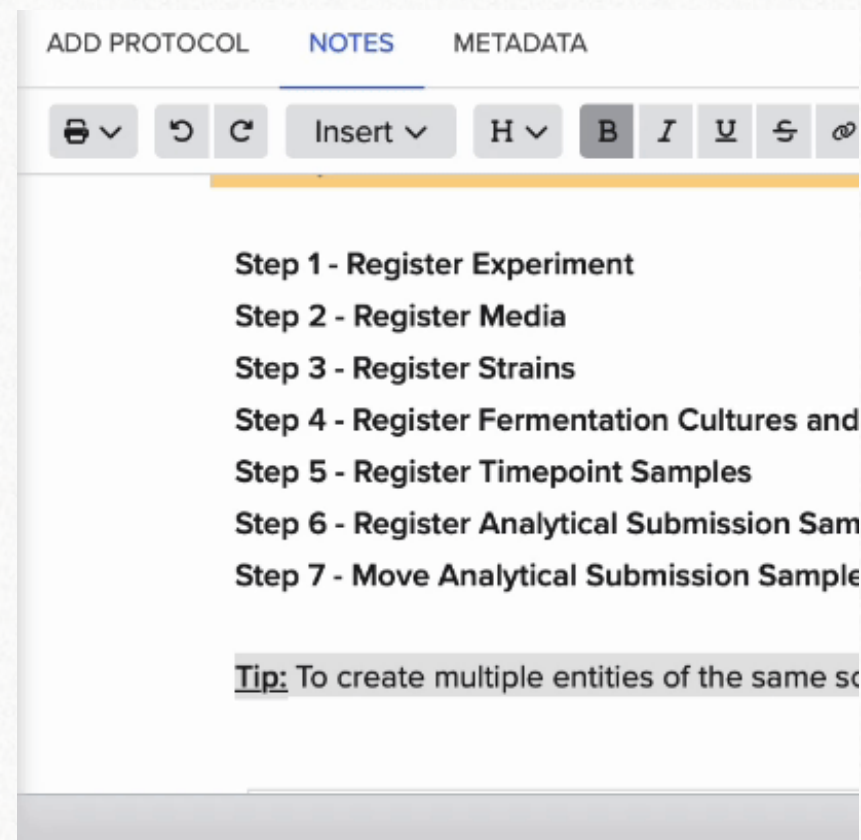
Registering entities: what to consider

3. You can create entities 1-by-1 or in bulk

To create entities in bulk:

- Use a registration table directly in the **Electronic Notebook**

Click “**Insert**” > “Registration table” > Select Schema



The screenshot shows the 'NOTES' tab of the Electronic Notebook interface. The top navigation bar includes 'ADD PROTOCOL', 'NOTES', and 'METADATA'. Below this is a toolbar with icons for undo, redo, insert, heading, bold, italic, underline, strikethrough, and link. The main content area displays a list of seven steps for registering an experiment:

- Step 1 - Register Experiment
- Step 2 - Register Media
- Step 3 - Register Strains
- Step 4 - Register Fermentation Cultures and
- Step 5 - Register Timepoint Samples
- Step 6 - Register Analytical Submission Sam
- Step 7 - Move Analytical Submission Sample

At the bottom, a tip is displayed: Tip: To create multiple entities of the same sc



Registering entities: what to consider

4. Some entities have “batches” schemas

Batches = physical samples

- When storing your sample long-term, create **batches in Benchling**
- This helps your team to track **where samples are stored**

Plasmid

Strain

Plasmid batch

Strain batch

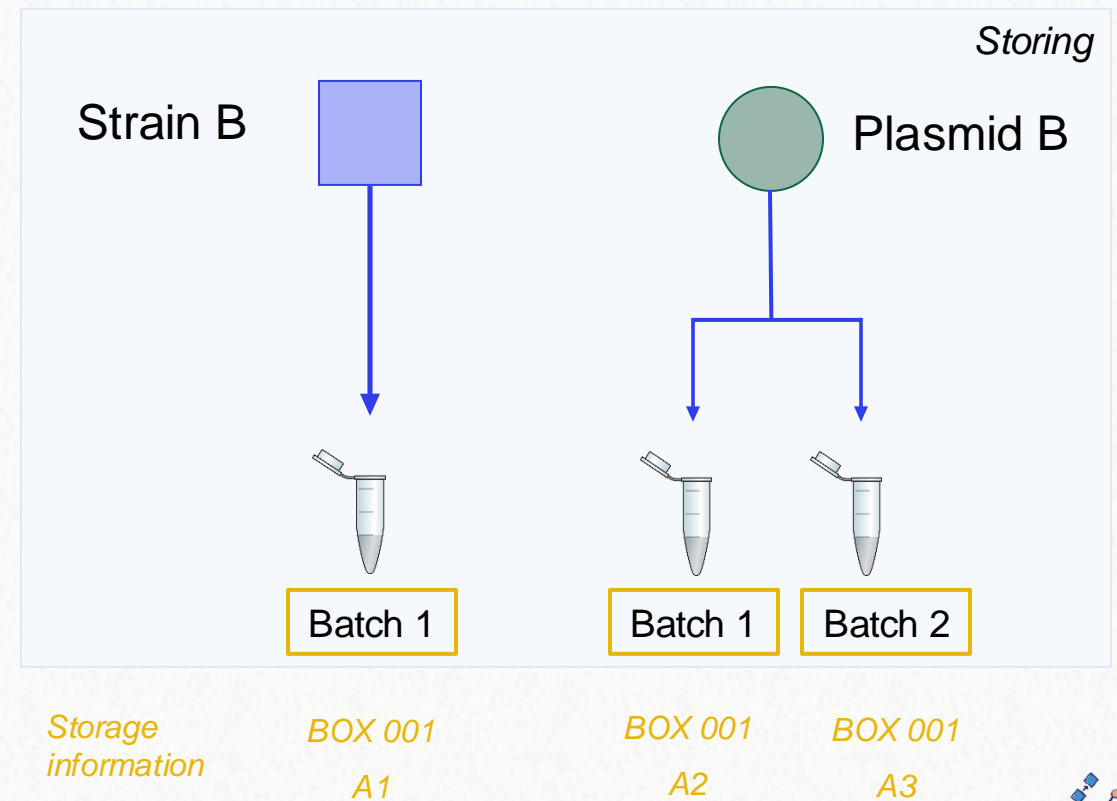


Registering entities: what to consider

4. Some entities have “batches” schemas

Batches = physical samples

- When storing your sample long-term, create **batches in Benchling**
- This helps your team to track **where samples are stored**

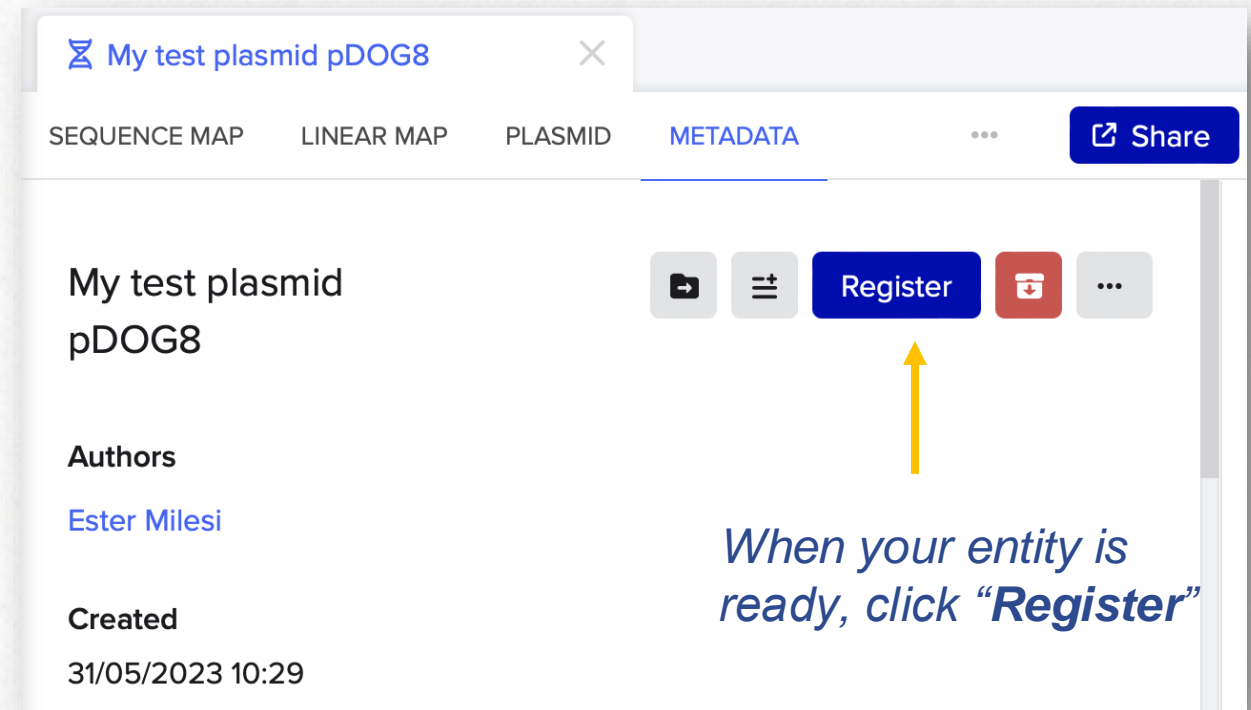


Registering entities: what to consider

5. Entities can exist outside of the Registry

Sometime, entities are not automatically registered

- In order to register an entity, you need to **select the Schema** (entity type)
- Registering the entity will add a **unique identifier** to your sample



My test plasmid pDOG8

SEQUENCE MAP LINEAR MAP PLASMID METADATA Share

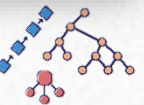
My test plasmid pDOG8

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Created
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Register

When your entity is ready, click “Register”



Good practices

(If you haven't done this consistently during your project)

At the end of your project:

- ✓ **Re-organize** your registered samples and entries and **give access** to your team
- ✓ Register **important strains/other samples** and their **location**, and print the label for the Box before moving it in the freezer



Storage:

track your samples



The Inventory

Benchling allows you to track the location of your samples

Room > Fridge > Box > Vial

If your fridge/location is not registered, let RDM support know

Create new box/plate/vial

Storage / DTU Biosustain

Search

Type: Location Barcode

Filters

1159 results X Clear

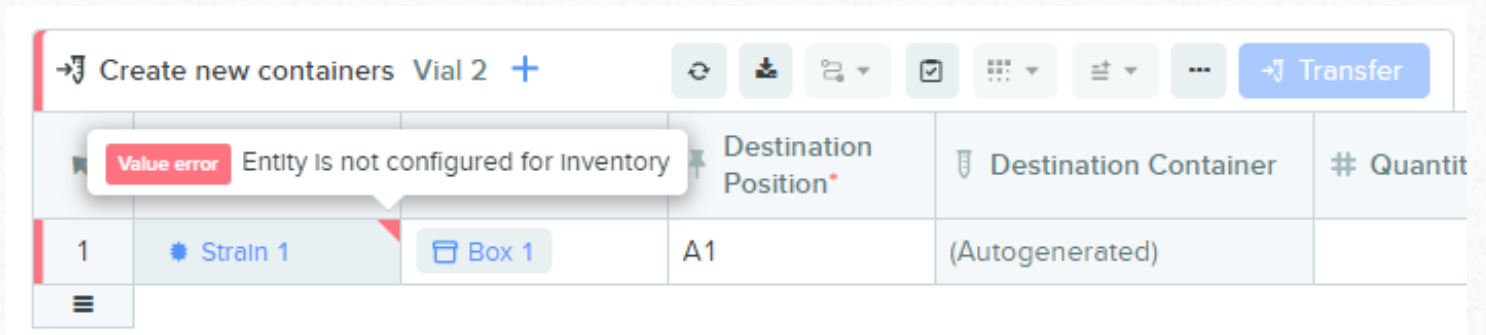
4C Fridge 00271 4C002
4C Fridge DSP1 4C003
4C Fridge DSP2 4C004
4C Fridge ANALYTICS 4C005



The Inventory

Storable samples include:

- Batches
(e.g., “Strain batch”)
- Fermentation cultures
- Submission samples
(e.g., for analysis)



The screenshot shows a software interface for managing inventory. At the top, there is a header bar with the text "Create new containers Vial 2 +" and several action icons, including a refresh icon, a download icon, a dropdown menu, a checkmark icon, a grid icon, a list icon, and a "Transfer" button. Below the header is a table with the following columns: "Destination Position*", "Destination Container", and "# Quantit". The table has one row with the following data: "1", "Strain 1", "Box 1", "A1", and "(Autogenerated)". A red tooltip with the text "Value error Entity is not configured for inventory" is pointing to the "Strain 1" cell.

			Destination Position*	Destination Container	# Quantit
1	Strain 1	Box 1	A1	(Autogenerated)	

Error showed in an Inventory table **in the Notebook** when trying to move a “Strain” in a Vial instead of a “Strain batch”



The Inventory

Benchling allows you to track the volume or concentration in of each vial




- Example of a **Box** in Benchling



Update quantity ✕

Current quantity	New quantity*	New units*
Not specified	<input type="text" value="20"/>	<input style="border: none; border-bottom: 1px solid #ccc; background-color: #f0f0f0; width: 100%;" type="text" value="uL"/>








Training box esterm ✕


METADATA RESULTS






Training box esterm
Move   

Barcode 81BOX984
Location  DTU Building 220 / Training Location  ✕


	1	2	3	4	5	6	7	8	9
A	1	2	3	4	5	6	7	8	9
B	10	11	12	13	14	15	16	17	18
C	19	20	21	22	23	24	25	26	27
D	28	29	30	31	32	33	34	35	36
E	37	38	39	40	41	42	43	44	45
F	46	47	48	49	50	51	52	53	54
G	55	56	57	58	59	60	61	62	63
H	64	65	66	67	68	69	70	71	72
I	73	74	75	76	77	78	79	80	81

Fill containers       

Actions 

	Position 	Container	Quantity 
	1 A1	 Training vial e	20 uL
<input checked="" type="checkbox"/>	2 A2	 Training vial fc	40 uL

[← Back](#)

Position 2 (A2) 

Barcode VIAL25349 Quantity 40 uL

No contents in Position 2 (A2).



Lastly: Archiving

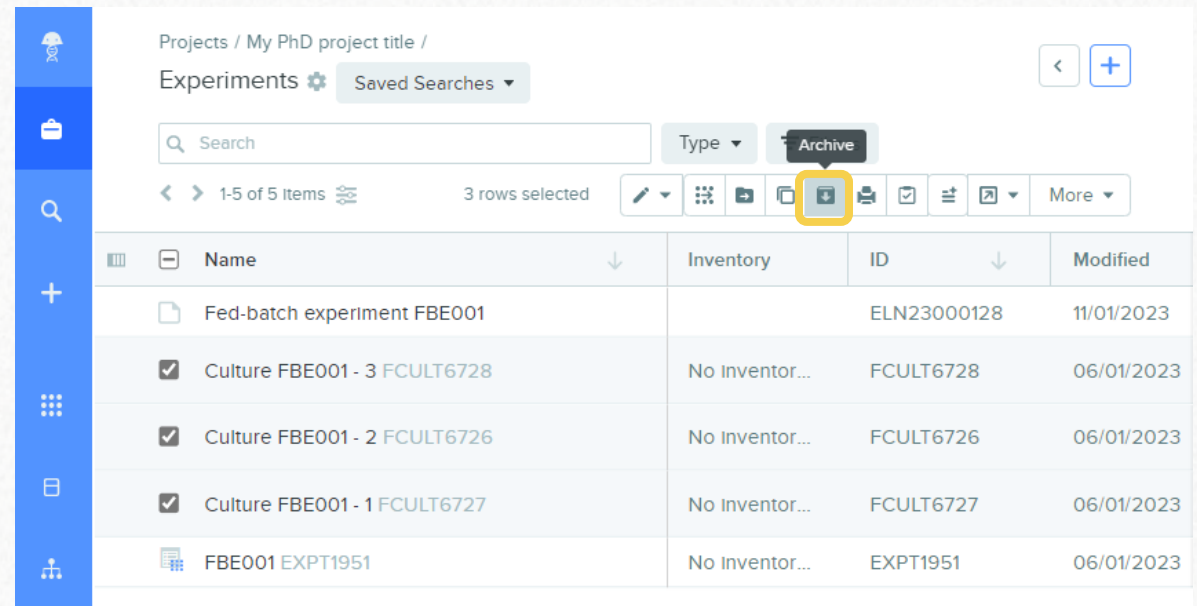


Archiving entities

Nothing can be deleted but only archived

- You can archive
 - ✓ Project folders
 - ✓ ELN entries
 - ✓ Entities

If you created them **by mistake** or if they are not relevant




The screenshot shows a web application interface for managing experiments. The breadcrumb path is "Projects / My PhD project title / Experiments". There is a search bar and a "Type" dropdown menu. A toolbar contains various icons, with the "Archive" icon (a trash can) highlighted by a yellow box. Below the toolbar is a table with 5 rows. The first row is "Fed-batch experiment FBE001". The next three rows are "Culture FBE001 - 3 FCULT6728", "Culture FBE001 - 2 FCULT6726", and "Culture FBE001 - 1 FCULT6727", each with a checked checkbox in the first column. The last row is "FBE001 EXPT1951". The table columns are "Name", "Inventory", "ID", and "Modified".

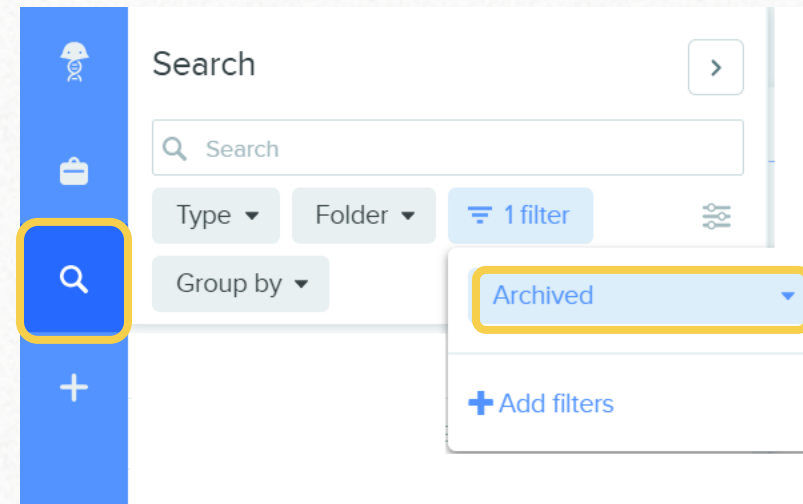
<input type="checkbox"/>	Name	Inventory	ID	Modified
<input type="checkbox"/>	Fed-batch experiment FBE001		ELN23000128	11/01/2023
<input checked="" type="checkbox"/>	Culture FBE001 - 3 FCULT6728	No Inventor...	FCULT6728	06/01/2023
<input checked="" type="checkbox"/>	Culture FBE001 - 2 FCULT6726	No Inventor...	FCULT6726	06/01/2023
<input checked="" type="checkbox"/>	Culture FBE001 - 1 FCULT6727	No Inventor...	FCULT6727	06/01/2023
<input type="checkbox"/>	FBE001 EXPT1951	No Inventor...	EXPT1951	06/01/2023



Archiving entities

It is still possible to go through archived items and unarchive them 

- In the search tab, filter by “Archive” status



Questions?



Agenda

Introduction to Benchling
and best practices

~ 30 min

Hands-on

~ 15 min

Agenda

Introduction to Benchling
and best practices

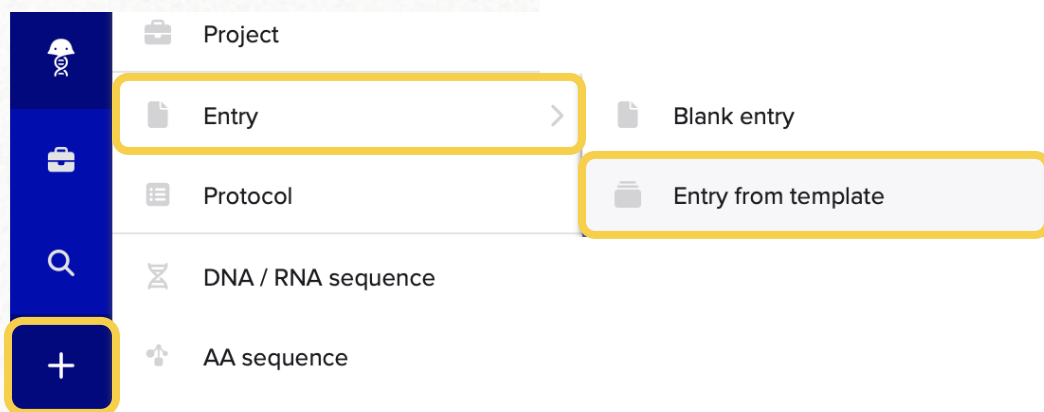
~ 30 min

Hands-on

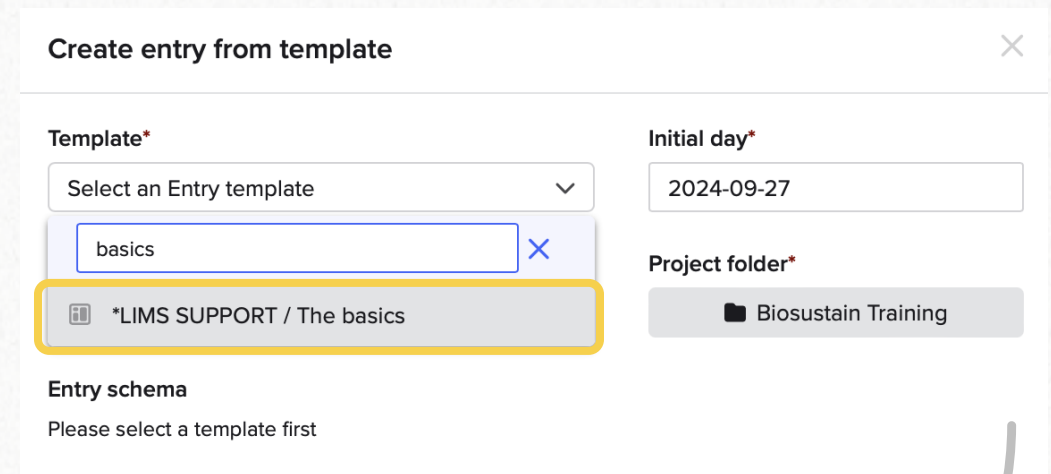
~ 15 min

Explore the Notebook functionalities

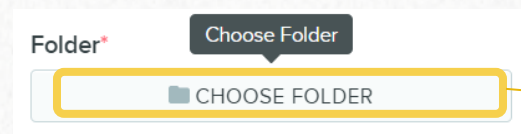
1 CREATE ENTRY FROM TEMPLATE



2 CHOSE "THE BASICS" TEMPLATE



3 SAVE IT IN THE BIOSUSTAIN TRAINING FOLDER



Projects

Filter...

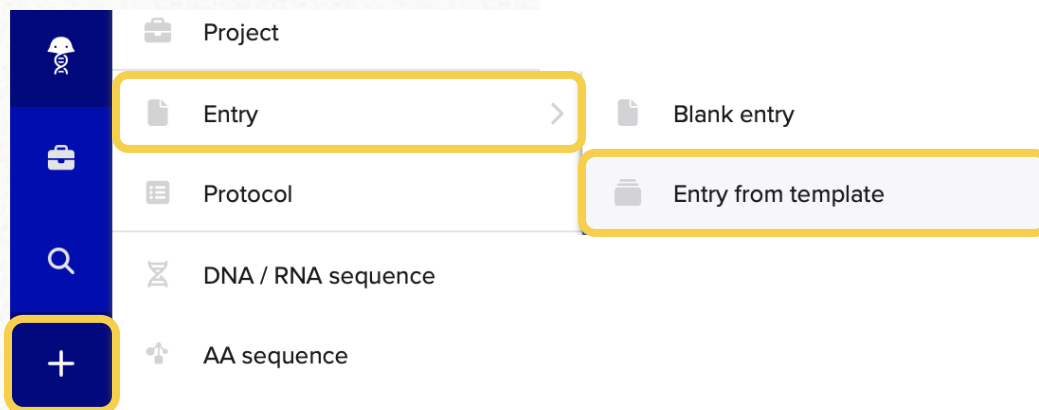
- ▼ ★ Biosustain Training biosustain
- ▶ ✓ Notebook



Additional useful template to use:

Submit samples for analysis to the Analytics team

1 CREATE ENTRY FROM TEMPLATE

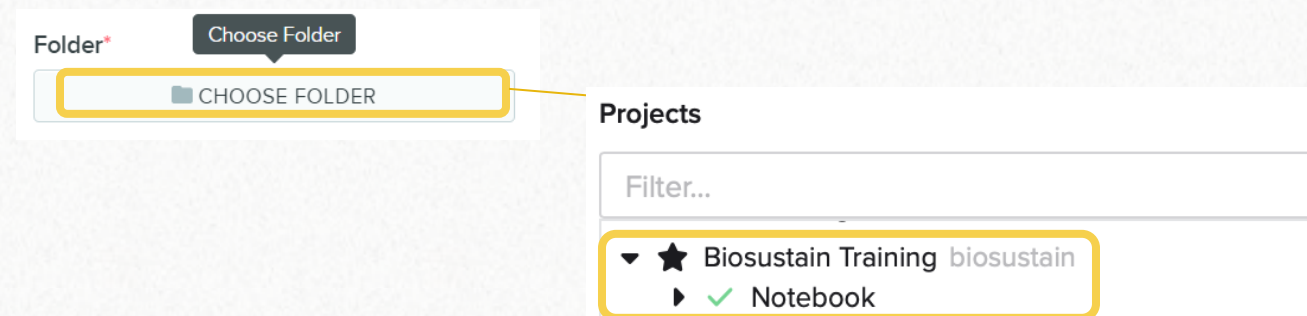


2 CHOSE "AC Analytical Sample Submission" TEMPLATE

The dialog box 'Create entry from template' contains the following fields:

- Template***: *Submission templates / AC Analytical Sample Submission [YOUR USERNAME]
- Initial day***: 2024-10-02
- Name***: AC Analytical Sample Submission [YOUR USERNAME]
- Project folder***: Biosustain Training

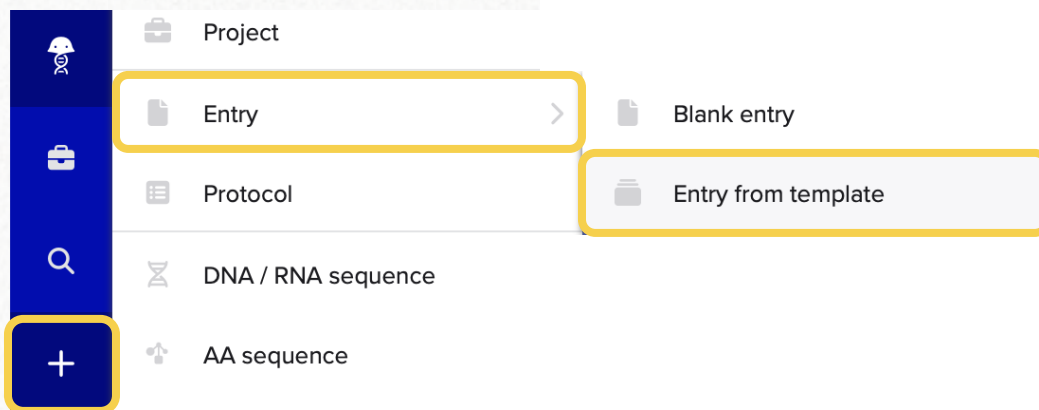
3 SAVE IT IN THE BIOSUSTAIN TRAINING FOLDER



Additional useful template to use:

Registration and storage of Strain

1 CREATE ENTRY FROM TEMPLATE



2 CHOSE "Strains registration" TEMPLATE

A screenshot of the 'Create entry from template' dialog box. The dialog has a title bar with a close button. It contains four fields: 'Template*' with a dropdown menu showing '*LIMS SUPPORT / Template - Strains registration' (highlighted in yellow), 'Initial day*' with a text input field containing '2024-10-02', 'Name*' with a text input field containing 'Template - Strains registration', and 'Project folder*' with a button labeled 'Biosustain Training' (highlighted in yellow). A blue circle with the number '2' is positioned to the left of the dialog.

3 SAVE IT IN THE BIOSUSTAIN TRAINING FOLDER

