

Import Data

Choose file

Step 1: Choose Data File Step 2: Map Fields Step 3: Commit Data

Files Templates **View saved mapping templates**

Choose a file to upload: Secondary CRC Assay.zip Project: Internal data

Uploaded Files Show files being imported and: Files you uploaded All files

Filename	Date	Status
▶ Secondary CRC Assay.zip	January 23, 2018	Partially mapped Continue mapping
▶ Secondary CRC Assay.zip	December 14, 2017	Imported Explore data Report Import again

Revisit previous imports

Select file and project to import data

Revisit previous imports

View imported data in table

View import data report

Map fields

Step 1: Choose Data File **Step 2: Map Fields** Step 3: Commit Data

File: Secondary CRC Assay.zip

 Use a saved mapping template, if available

Add readouts (protocol data) only Update existing molecules Register new molecules / batches with structures

Add data to existing molecules

A	B	C	D	E	F
Molecule Name	Plate	Well	Conc (uM)	Raw (RLU)	Microscopy
Sample-10-001	Sample Pl._20151108	A10	0.01	2.67E+03	MicroscopyImage
Sample-10-001	Sample Pl._20151108	B10	0.03	2.84E+03	
Sample-10-001	Sample Pl._20151108	C10	0.1	2.04E+03	

Map fields Note: System will map columns with similar names to fields

Map fields Note: System will map columns with similar names to fields

Issues to be fixed before importing can occur

Save mapping as template for future use

Type of information to be registered
Note: To register records without structures, select "without structure" option

Use a saved mapping template, if available

Add data to existing molecules

Map fields
Note: System will map columns with similar names to fields

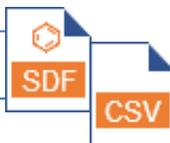
Issues to be fixed before importing can occur

Save mapping as template for future use

File format

Molecules

File types



CSV format

Unique batch field

Note: this can be used for uploading data

	A	B	C	D	E	F	G	H
1	Canonical_Smiles	CAS #	Purity	Vendor	Synonym	Catalog #	Amt	External ID
2	<chem>CC(NC(C)(C)C(=O)c1cccc(Cl)c1</chem>	34911-55-2	95	NIH	SAM001246723	CPD000058423	5	Sample-10-2
3	<chem>CNCC[C@H](Oc1cccc2ccccc12)c3cccs3</chem>		95	NIH	SAM001247059	CPD000449282	5	Sample-10-3
)cc3	128196-01-0	95	NIH	SAM001246668	CPD000469191	5	Sample-10-4

Use SMILES or mol file for structures

Note: Salts and solvents will be stripped by default

File type



Single point data

First row – column headers

Each type of data entered into a separate column

	A	B	C
1	SampleID	Inhibition	SEM
2	Sample-10-001	-10.336685	4.596999925
3	Sample-10-001	4.142833609	6.56848643
4	Sample-10-001	0.047860794	0.137099726
5	Sample-10-002	-4.29821	4.29821
6	Sample-10-002	8.835689174	0.158027477
7	Sample-10-002	3.668918207	4.560620974

A batch of a molecule may be uniquely identified by:

- Molecule name and Batch name
- Synonym and Batch name
- Batch field that is unique
- Plate and Well location – plate preregistered
- A combination of one of the first 3 options and Plate and Well location

e.g. Molecule name, Batch name, Plate and Well location

Dose response

Controls – if applicable

Note: do not need to be associated with a batch

	A	B	C	D	E	F
1	Molecule ID	Batch	Plate	Well	Conc	RLU
2			Plate 20180418	A01		2.82E+03
3			Plate 20180418	A02		30
4	DV-0000054	1	Plate 20180418	A04	0.01	3.37E+03
5	DV-0000048	1	Plate 20180418	A10	0.01	2.67E+03
6	DV-0000052	1	Plate 20180418	A12	0.01	3.34E+03
7	DV-0000053	1	Plate 2018041			

Samples

- 1 row per measurement
- Location of samples specified in file

Importing attachments

File format

File type ZIP

Name	Type
MicroscopyImage1.jpg	JPG File
MicroscopyImage2.jpg	JPG File
MicroscopyImage3.jpg	JPG File
MicroscopyImage4.jpg	JPG File
Secondary CRC Assay.csv	Microsoft Excel C

Create zip file containing attachments and CSV or SDF file.

Create SDF or CSV file as usual. Add a column for the attachment filename

	A	B	C	D	E	F	
1	SampleID	Plate	Well	Conc (uM)	Raw (RLU)	Microscopy	
2	Sample-10-001	Sample Plate 20151108	A10	0.01	2.67E+03	MicroscopyImage3.jpg	
3	Sample-10-001	Sample Plate 20151108	B10	0.03	2.84E+03		
4	Sample-10-001	Sample Plate 20151108	C10	0.1	2.04E+03		
5	Sample-10-001	Sample Plate 20151108	D10	0.3	1.25E+03		
6	Sample-10-001	Sample Plate 20151108	E10	1	775		
7	Sample-10-001	Sample Plate 20151108	F10	3	255		
8	Sample-10-001	Sample Plate 20151108	G10	10	1		
9	Sample-10-001	Sample Plate 20151108	H10	30	63		
10	Sample-10-001	Sample Plate 20151108	I10	0.01	2.44E+		
11	Sample-10-001	Sample Plate 20151108	J10	0.03	3.22E+03		
			08	K10	0.1	2.53E+03	
			08	L10	0.3	1.68E+03	
			08	M10	1	628	
			08	N10	3	267	
			08	O10	10	109	
			08	P10	30	84	
18	Sample-10-002	Sample Plate 20151108	A20	0.01	2.73E+03	MicroscopyImage2.jpg	
19	Sample-10-002	Sample Plate 20151108	B20	0.03	3.00E+03		

Enter filename of attachment
Note: Attachment only needed once per batch

A batch of a molecule may be uniquely identified by:

- Molecule name and Batch name
- Synonym and Batch name
- Batch field that is unique
- Plate and Well location – plate preregistered
- A combination of one of the first 3 options and Plate and Well location
e.g. Molecule name, Batch name, Plate and Well location

Import Data

Commit Data

Step 1: Choose Data File | Step 2: Map Fields | **Step 3: Commit Data**

File: Benzodiazepines.sdf (Review mapping) | Project: Bayesian | Owner: Heather Arnaiz

This data import is ready for review

- 7 records will be imported - Only records that are not associated with any rejected event will be imported.
- 2 records will be rejected - You will be able to download these records and their specific issues once you commit.

Noteworthy Events - Usually fine. Associated records will be imported except if you choose otherwise.

- 7 New Batches (2 indirectly rejected)

Suspicious Events - Usually unexpected. Associated records will not be imported except if you choose otherwise.

- 2 Existing Molecules Associated with New Project

Line #	Record #	Molfile	Event Description
1	1	BENZOEXAM...M.END	REJECTED: Molecule HA-0000962 associated with project Bayesian
58	2	BENZOEXAM...M.END	REJECTED: Molecule HA-0000156 associated with project Bayesian

Export

Commit Data Import | Reject Data Import

View mapping

- Create template from mapping

Noteworthy events

- Things to be aware of
- Decide to Accept or Reject

Suspicious events

- Problems in registration
- Decide to Accept or Reject

Click down arrow to see details of noteworthy or suspicious events

Export problematic data to fix and reimport, as desired

Commit or Reject Import

Report

Explore Data | ELN | Import Data | Reports | Settings | [Email]

Your Current Import: Benzodiazepines.sdf Processed (2 Suspicious Events) Email me updates

+ Create a new ... | Search | Saved Searches | Col

Step 1: Choose Data File | Step 2: Map Fields | **Step 3: Commit Data**

File: Benzodiazepines.sdf (Review mapping) | Project: Bayesian | Owner: Heather Arnaiz

Import details for Benzodiazepines.sdf: Queued (#1) Email me updates

Cancel Import

Explore Data | ELN | **Import Data** | Reports | Settings | [Email] | Heather Arnaiz

File: Benzodiazepines.sdf (Review mapping) | Project: Bayesian | Owner: Heather Arnaiz

This data import was committed

- 7 records imported | View or export registered molecules
- 2 records rejected | Export rejected records to fix and re-import the problematic data

Explore imported data

View imported records

View or export records

Noteworthy Events - Usually fine. Associated records will be imported except if you choose otherwise.

- 7 New Batches (2 indirectly rejected)

Suspicious Events - Usually unexpected. Associated records will not be imported except if you choose otherwise.

- 2 Existing Molecules Associated with New Project

Import status

I changed my mind – Cancel the import

View mapping

- Create template from mapping

View imported records

View or export records