



PROJECT DELIVERABLE REPORT

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** Type: R – Report; P – Prototype; D – Demonstrator; - O - Other



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JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Dash-in – Do you have any comments on the Web based Dash-in federated analyses? 83



INTRODUCTION

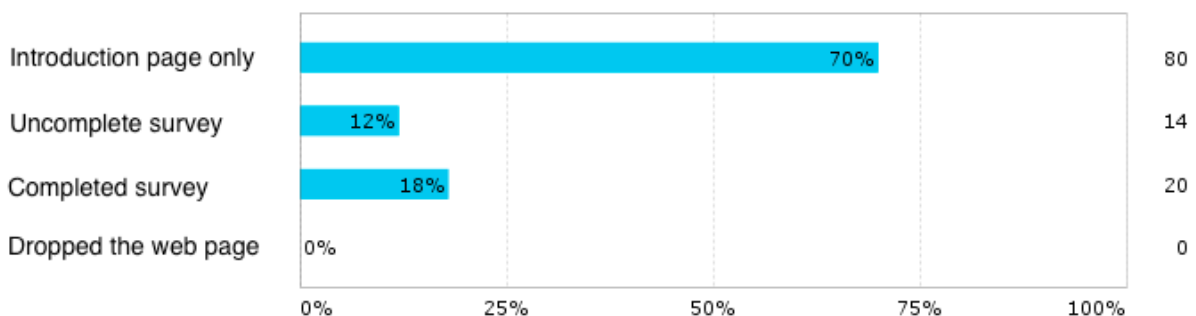
This task has executed a survey of the designed interface of the common access point to the infrastructure. The focus of the design and the user survey actions was on usability and easiness of use for the end-user.

Initial plan didn't match with project-wide availability of resources. This survey is based on the interfaces of the Phenotype database (dbNP), the BioShare OPAL DataShield tools, in the scope of this consortium and the initial part of this federated analysis. Moreover, the newly developed extensions were tested. Brand new design and implementation of the value-added DASH-IN visual analytical platform was conducted (T3.2, T3.5) and surveyed.

A first survey call was launched from the 30th of November to the 31st December 2016 with a reminder sent on the 19th of December 2016. Survey data was collected from the system on 16th January 2017 (which was left accessible all the time). A total of 21 surveys were taken (counted as those that had entered at least the email addresses). Two persons, identified by email address, took the survey twice, therefore a total of 19 unique people took the survey.

Due to some requests the survey, on January 31st 2017 the survey was extended until the 15th of February 2017. The system was left accessible until the 27th of February 2017 when survey data was collected. Seven more answers were provided by the partners (counted as a partner who at least entered the email address), two of which already took the survey during the first session. Therefore, finally a total of 24 unique people answered the survey for a total of 28 times.

Globally the first introduction survey page was visited 114 times. In 70% of the cases the user took no further action, while in 12% of the cases some pages were went through but without finishing and 18% (exactly 20 survey answers, possibly provided by fewer than 20 people) of the persons completed the entire survey. No one was dropped out of the web pages.



Given the heterogeneous background of the partners, the survey aimed at assessing whether:

- The systems have been used at all.
- How easy is perceived the instructed visual process (which of course depends on whether the person has actually used it or is just looking at the pictures).



- The user has suggestions for improvements.

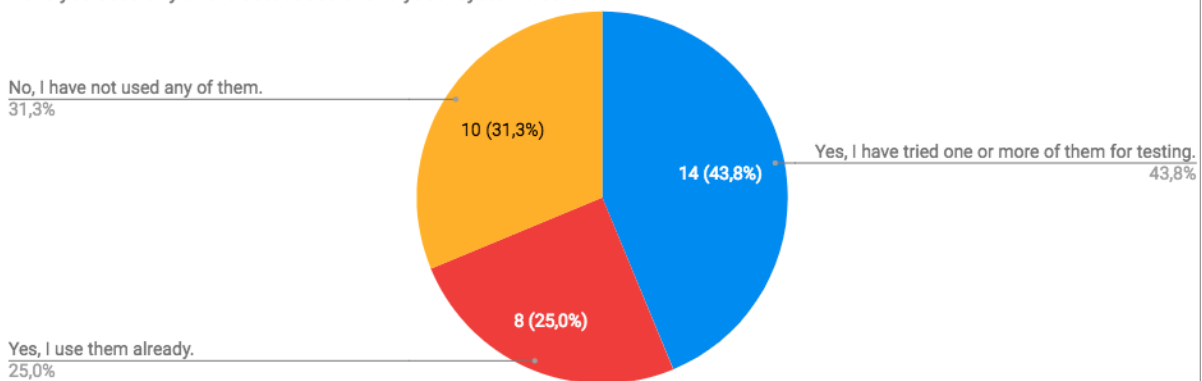
Each surveyed section is reported in this document with all the screenshots and texts along with the statistics of responders. Overall it is noticeable a quite high rate of responders who answered the survey without trying at all the system or started trying it without actually completing the real session.

All survey data described in the remaining is relative to the overall answers globally provided during the two survey sessions.

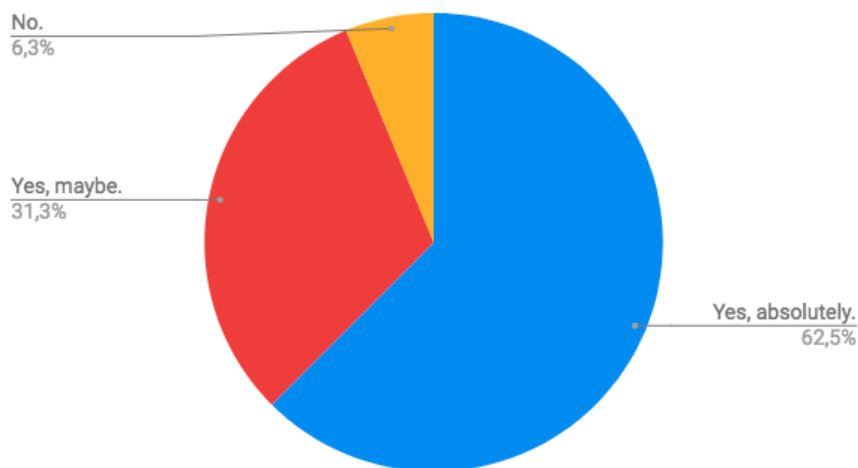
RESULTS OF THE SURVEY

See below a representation of the survey and the analysis of the responses.

Have you used any of the databases or analytical systems before?



Would you or your institution be interested in using the ENPADASI/DASH-IN?





OPAL – ADD A NEW PROJECT

Projects

A project is a repository of data and dictionaries. In a project, data can be imported, exported, analysed and transformed. Projects also offer access controls to the dictionaries, the summaries and the individual data.

[+ Add Project](#) Total 1

Name	Title	Description	Last updated
LifeLines	LifeLines		23 days ago

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Add Project

Name *
Finn prj 01
A project must have a unique name.

Title
Finn's Project 1
A project title for the human beings.

Database
opal_data (default storage)
None
opal_data (default storage)

Save Cancel



These images show part of the process to add a new project to OPAL

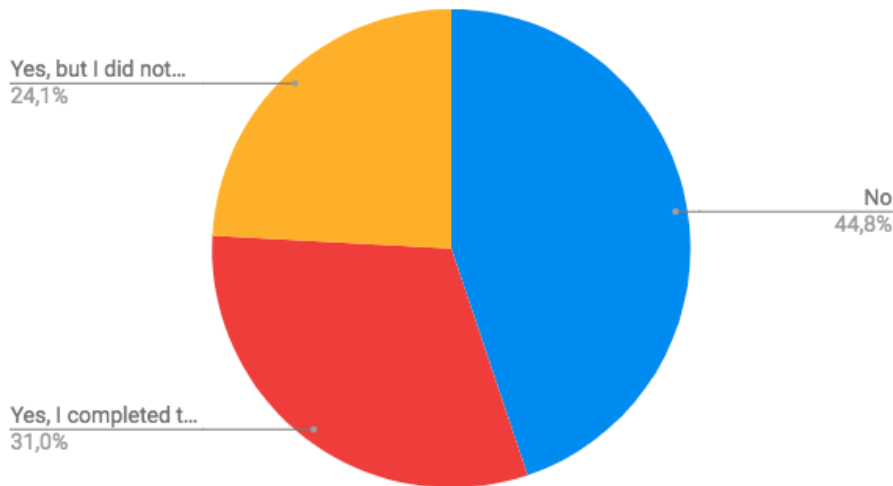
Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

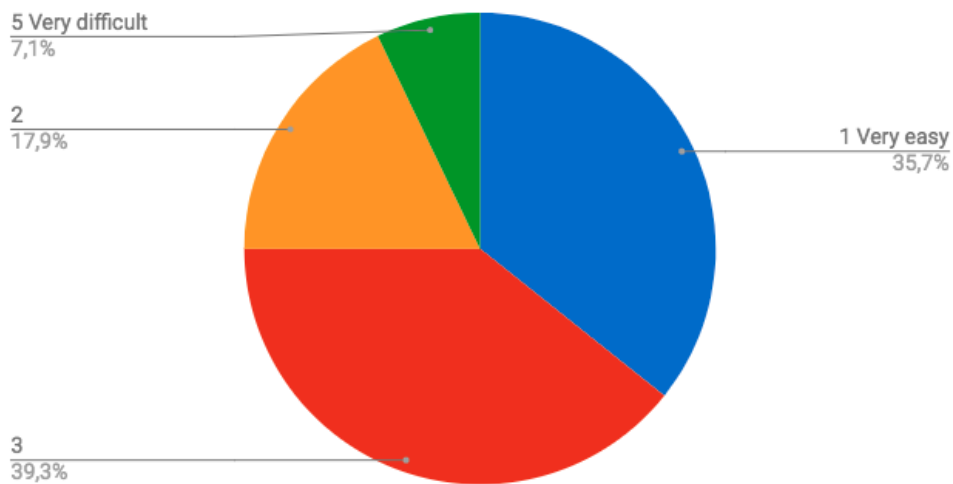
How easy or difficult is it to add a new project to OPAL

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

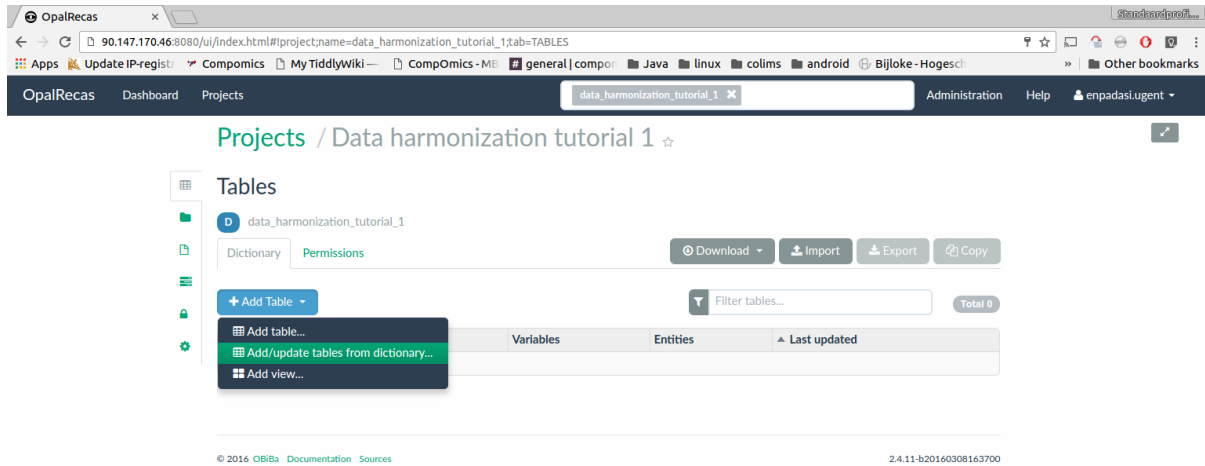
Add a new project to OPAL



Add a new project to OPAL - How easy or difficult is it to add a new project to OPAL

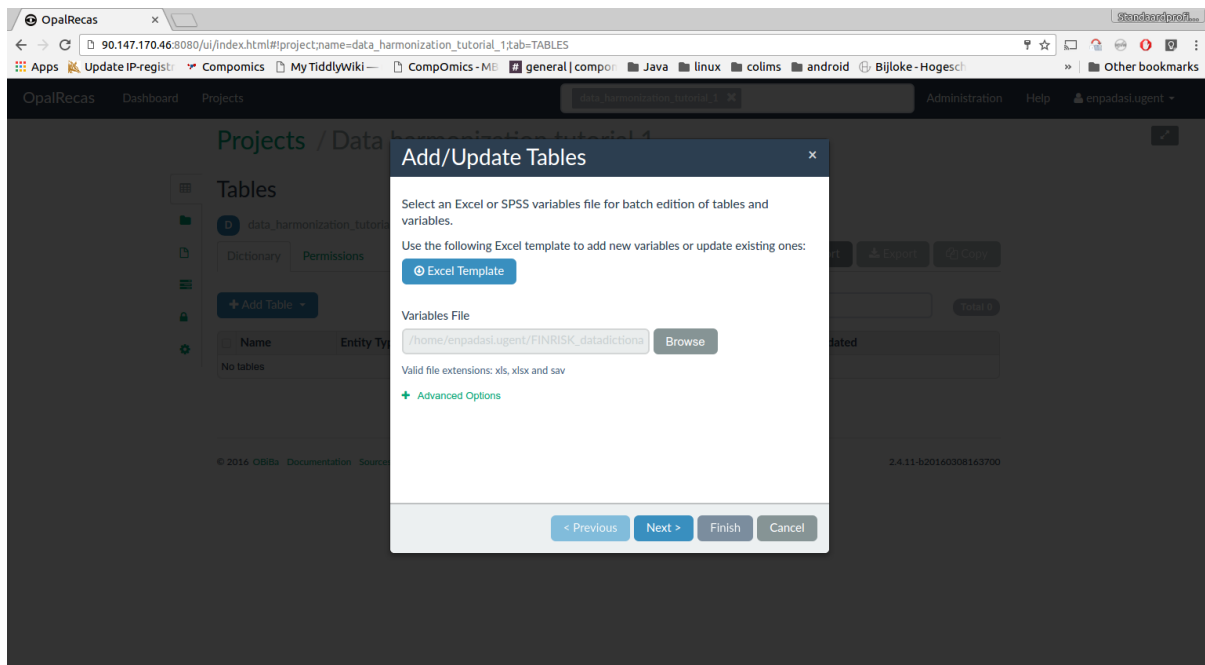


OPAL - ADD TABLE DICTIONARY

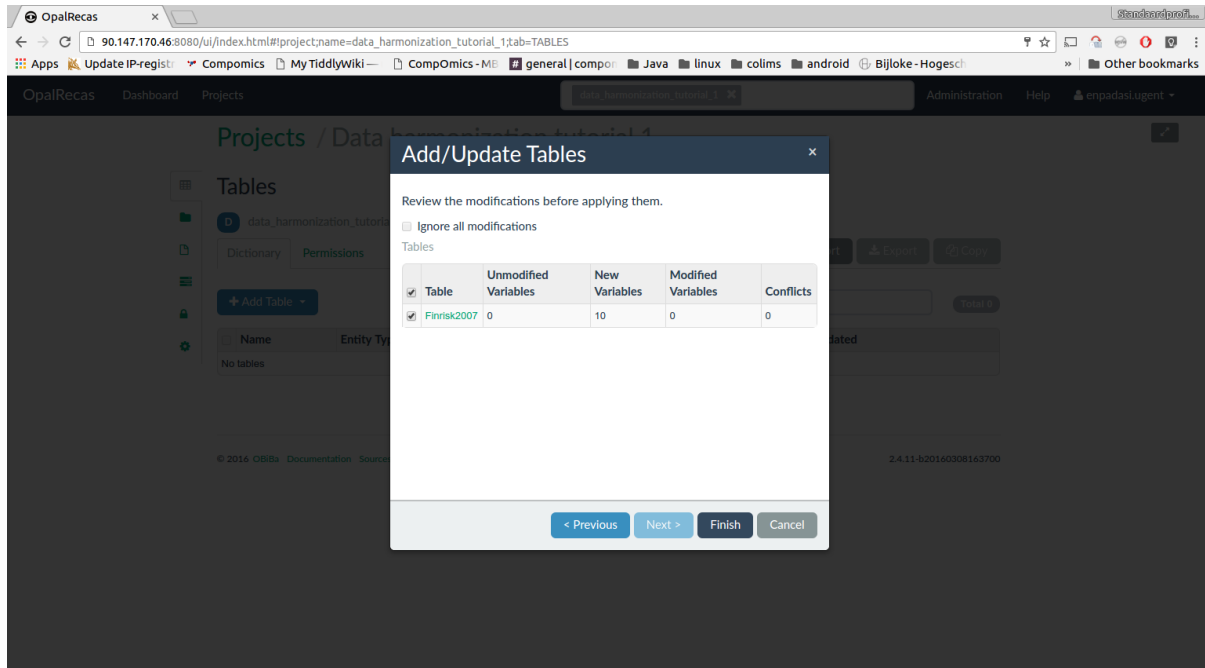


javascript,;

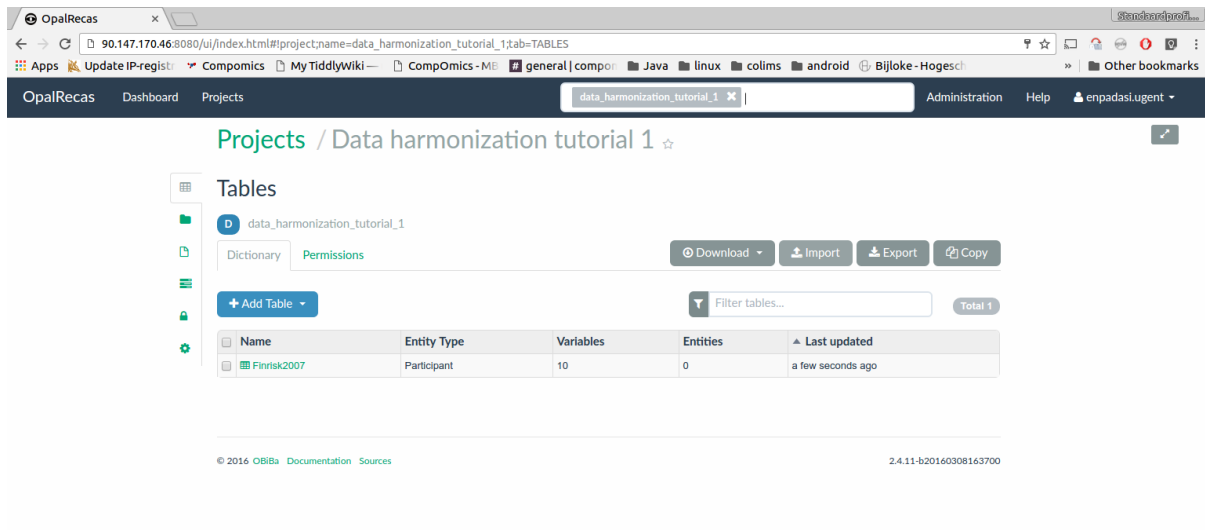
Before uploading table data in .csv format, a data dictionary needs to be added. In case of data in .sav format, both the data dictionary and the data are included in the same file.



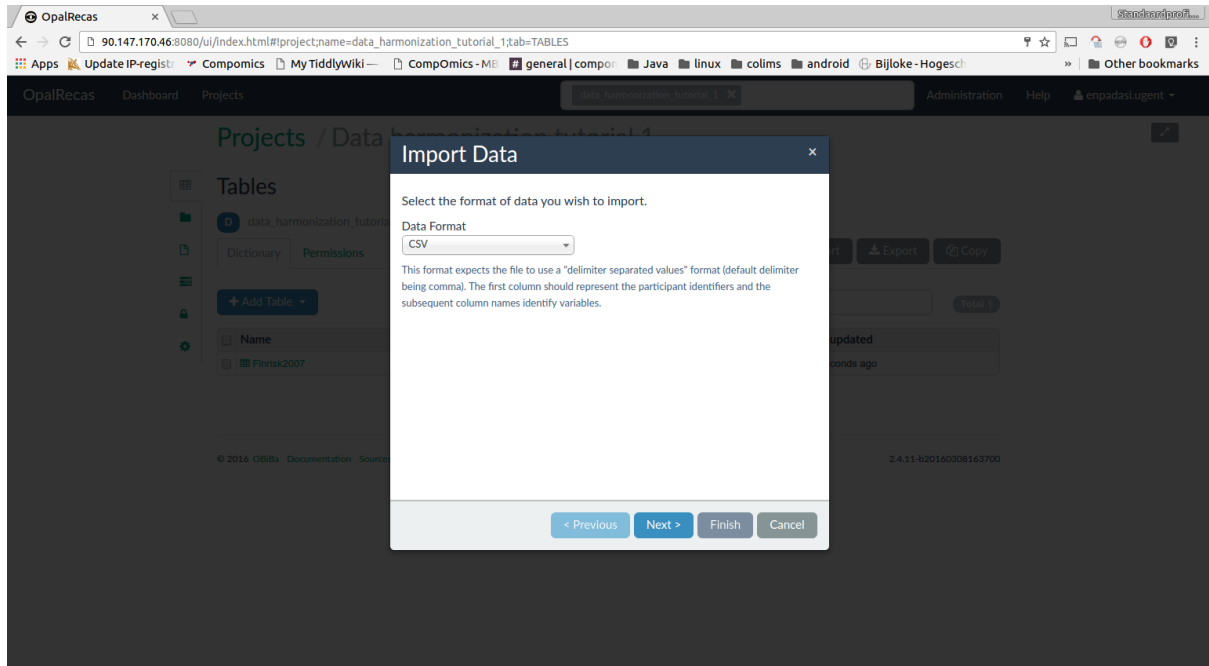
After uploading the data dictionary to the Opal server, the user can select it (.xls, .xlsx and .sav are supported).



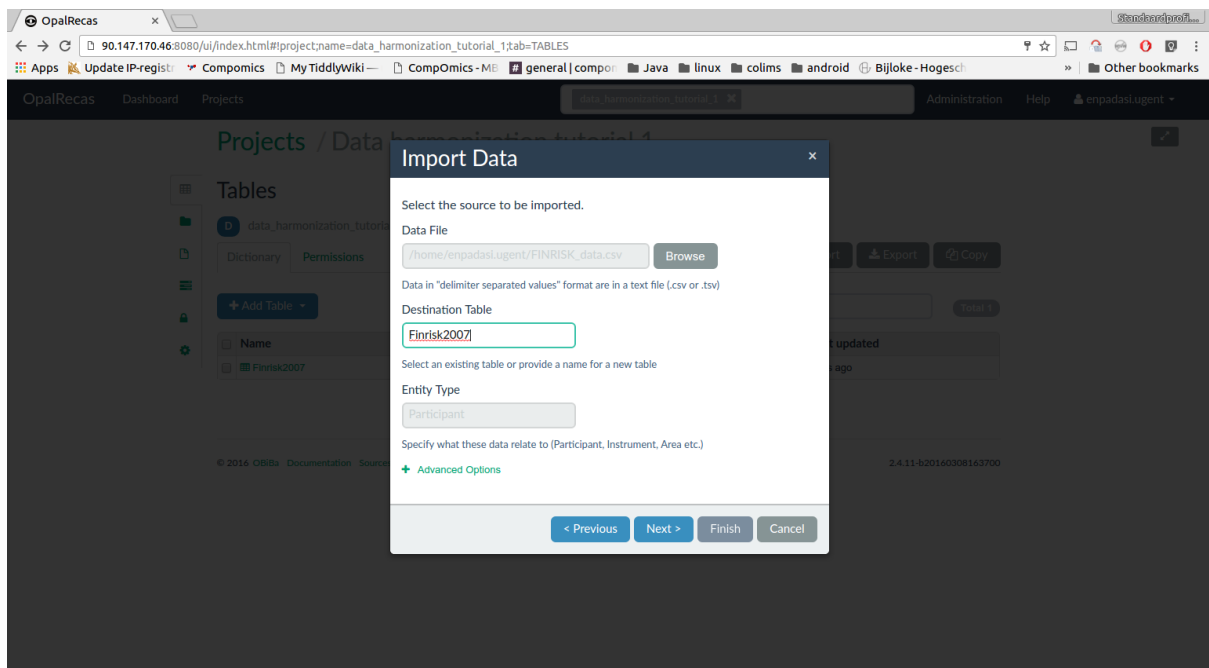
Select the table of interest and click on the "Finish" button.



Once the data dictionary has been added, click on the "Import" button to import the table data.



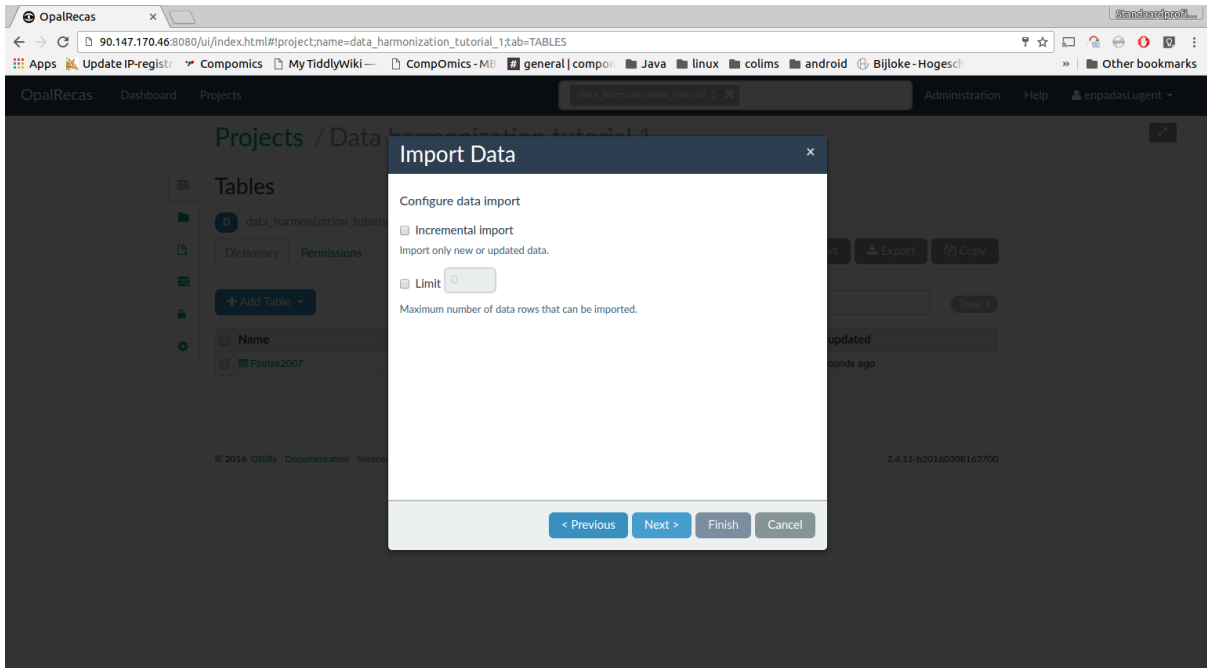
Leave the default "CSV" setting and click on the "Next" button.



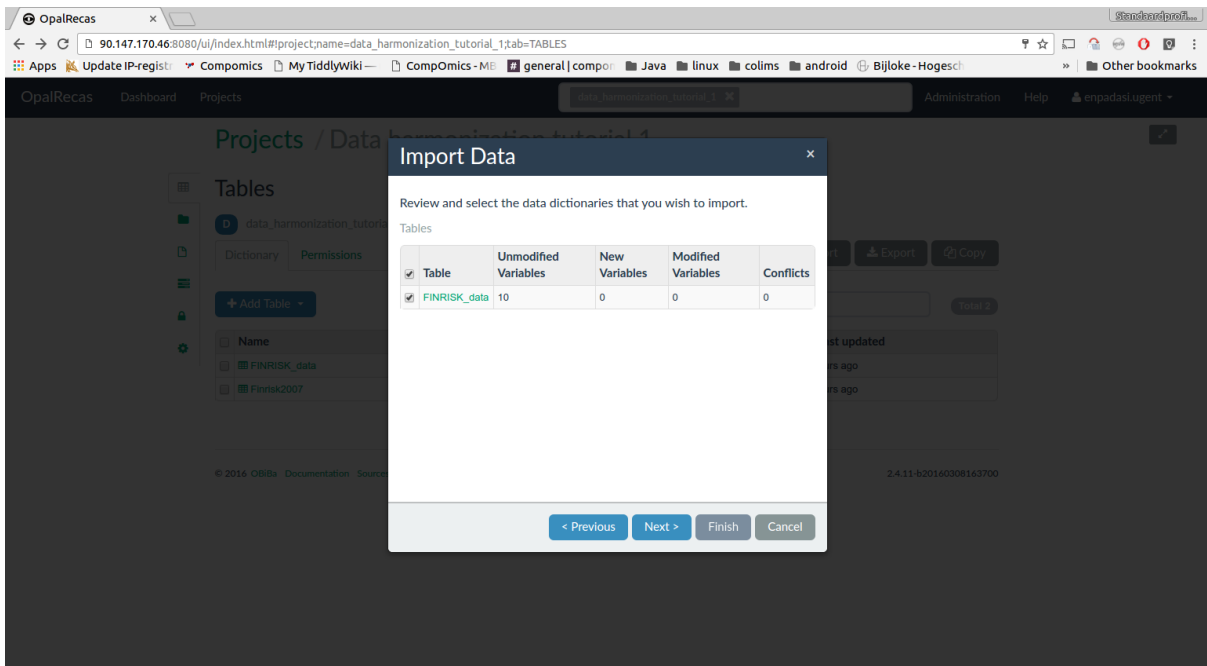
Select the data file and the destination table and click on the "Next" button.



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Keep the default settings and click on the "Next" button.



Select the appropriate table and click on the "Next" button.



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Review the data that will be imported.

Table: FINRISK_data

1-20 of 5,024

ID	SUKUP	PAINO	PITUUS	FR
1000948357610215153	2	89.38351149082827	168.2201360071805	1
1002400312040470929	2	94.28620400965093	172.98381515846853	1
1004168722520747805	1	58.05548910267279	168.59543649669538	1
1007800521831721882	1	81.44858356305257	174.25781292787622	1
1010975981543513267	2	79.9910303261611	181.97687069030263	1
1013334134400726098	1	47.11228817596795	145.27003920443656	1
1013735151969205356	2	79.80130160335341	169.52511608291368	1
1015429109149314392	1	52.32337587441517	166.3847009852287	1

< Previous Next > Finish Cancel

Click on the "Next" button.

Specify whether the data file is to be archived.

Once a file has been successfully imported:

- leave it where it is
- move it to another folder: Browse

< Previous Next > Finish Cancel

Click on the "Finish" button.



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Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

How easy or difficult is it to add a table dictionary in OPAL?

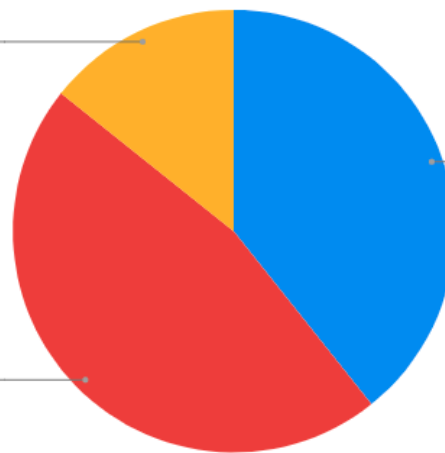
- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Opal - Add table dictionary - Have you used this functionality

Yes, but I did not complete the process
14,3%

No
46,4%

Yes, I completed the process
39,3%



Opal - Add table dictionary - How easy or difficult did you find it?

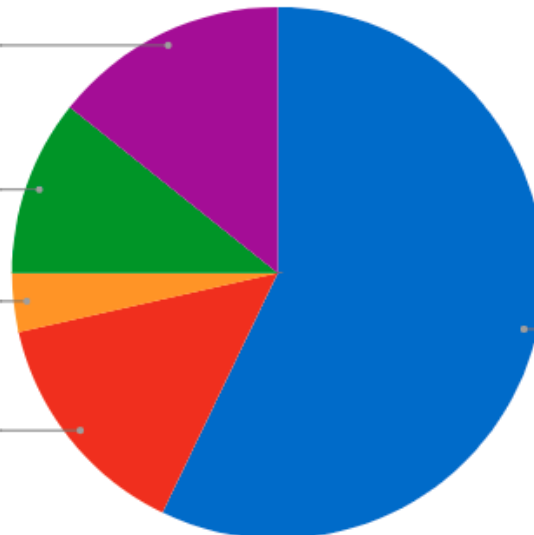
2
14,3%

5 Very difficult
10,7%

4
3,6%

1 Very easy
14,3%

3
57,1%



OPAL - ADD HARMONIZED SCHEMA

The screenshot shows the OpalRecas web interface. The browser address bar shows the URL: `90.147.170.46:8080/ui/index.html#/project,name=data_harmonization_tutorial_1,tab=TABLES`. The page title is "Projects / Data harmonization tutorial 1". The main content area is titled "Tables" and shows a table with the following data:

Name	Variables	Entities	Last updated
Finrisk2007	10	0	7 hours ago

A context menu is open over the table, with the following options: "Add table...", "Add/update tables from dictionary...", and "Add view...". The "Add view..." option is highlighted.

Click on the "Add table" button and select "Add view..."

The screenshot shows the "Add View" dialog box in the OpalRecas web interface. The dialog has the following fields and buttons:

- Name ***:
- Table References ***:
- Variables**:
- Buttons**:

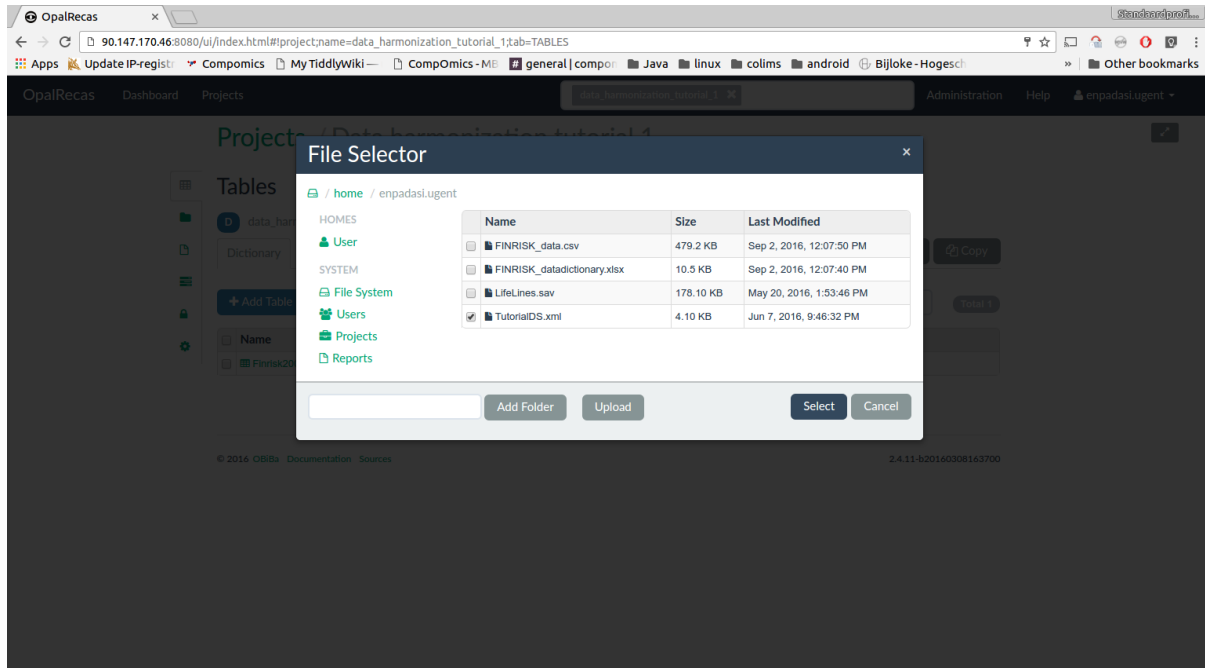
The dialog also contains the following text:

- "A view must have a unique name."
- "The table(s) from which the entities are extracted."
- "Provide derived variables from a view XML or a table dictionary in Excel, optional."

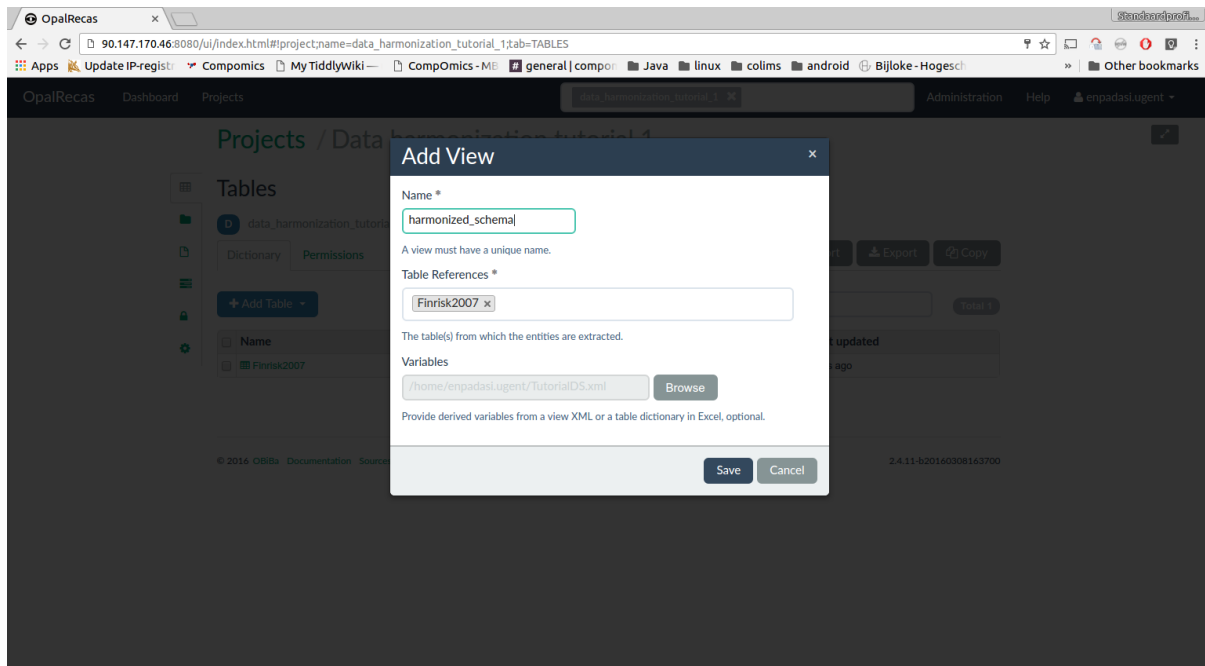
Select the previously added data table in the "Table References" field. Click on the "Browse" button next to the "Variables" field for selecting the harmonized variables.



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Upload the harmonized variables in view XML format or table dictionary Excel format and the select it.



Click on the "Save" button to add the harmonized view.



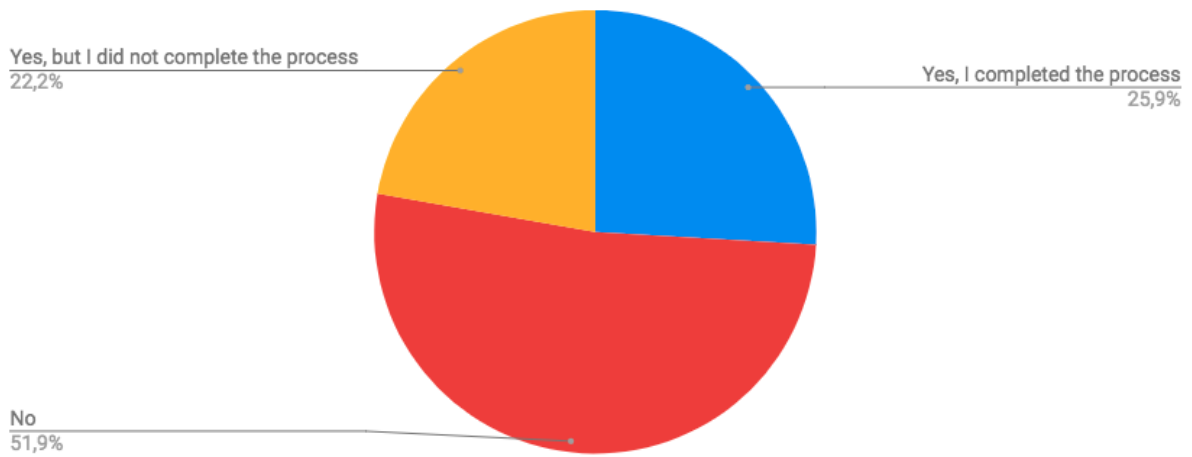
Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

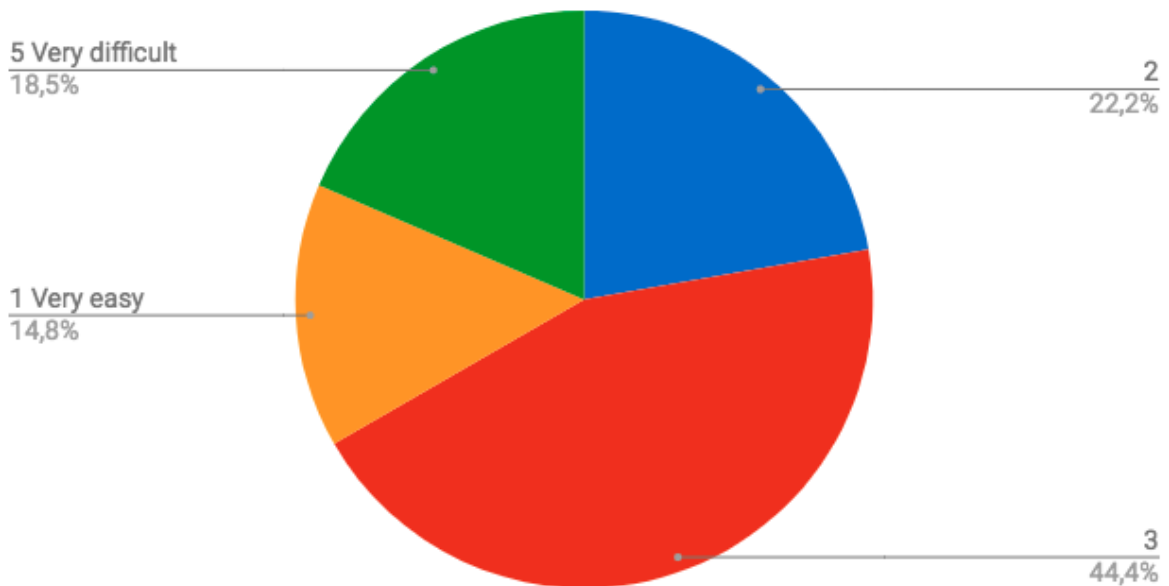
How easy or difficult is it to add a harmonized schema in OPAL?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Opal - Add harmonized schema - Have you used this functionality



Opal - Add harmonized schema - How easy or difficult did you find it?





OPAL - VARIABLE HARMONIZATION - ADVANCED

The screenshot shows the 'Tables' view in the OpalRecas application. The breadcrumb is 'Projects / Data harmonization tutorial 1'. The 'Tables' section shows a list of tables:

Name	Entity Type	Variables	Entities	Last updated
harmonized_schema	Participant	5	0	a few seconds ago
Finrisk2007	Participant	10	0	7 hours ago

At the bottom of the page, the copyright information reads: © 2016 OBIBA Documentation Sources 2.4.11-b20160308163700

Click on the "Save" button to add the harmonized view.

The screenshot shows the 'Properties' view for the 'harmonized_schema' table. The breadcrumb is 'Projects / data_harmonization_tutorial_1 / harmonized_schema'. The 'Properties' section shows:

Name	harmonized_schema
Entity Type	Participant
Table References	data_harmonization_tutorial_1.Finrisk2007

The 'Entity Filter' section shows: // no filter

The 'Variables' section shows a list of variables:

Name	Label	Value Type	Categories
Gender		integer	0, 1, 9
PM_BMI_CAT		integer	1, 2, 3, 9
DIS_DIAB		integer	0, 1, 9
PM_DIASTOLIC		integer	999
SMK_CIG_CURRENT		integer	0, 1, 9

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Click on the "Gender" variable. It will be used as an example of an "easy" variable to harmonize.



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The screenshot shows the OpalRecas web interface. The main content area displays the 'Properties' section for a variable named 'Gender'. A 'Derive' button is visible, and a dropdown menu is open, showing the following options:

- Add variable to view...
- Categorize this variable to another...
- Categorize another variable to this...** (highlighted)
- Custom derivation...

Below the 'Properties' section, there is a 'Categories' table:

Name	Label	Missing
0	Male en Male	
1	Female en Female	
9	Missing	✓

Click on the "Derive" button and select "Categorize another variable to this..."

The screenshot shows the 'Derive Variable' dialog box open over the web interface. The dialog box contains the following text and fields:

Select from which variable derivation will be performed. The resulting derived variable will be the current variable.

Table:

Variable:

At the bottom of the dialog box, there are four buttons: '< Previous', 'Next >', 'Finish', and 'Cancel'.

Select the "Finrisk2007" table and the "SUKUP" variable.



Original Value	Frequency	Original Label	New Value	Missing
1	0	Male	0	<input type="checkbox"/>
2	0	Female	1	<input type="checkbox"/>
N/A	0	Empty value	9	<input checked="" type="checkbox"/>
*	0	Other value	9	<input checked="" type="checkbox"/>

Give each original value a new value in the "New Value" column using the drop-down boxes. Click on the "Next" button to review the variable derivation.

Value	Frequency	Percentage	
		Subtotal	Total
Non-Missing			
0	0	0%	0%
1	0	0%	0%

Click on the "Finish" button to finalize the harmonization of the "Gender" variable.

OpalRecas Dashboard Projects data_harmonization_tutorial_1 harmonized_schema Administration Help enpadasI.ugent

Projects / Data harmonization tutorial 1

Tables

data_harmonization_tutorial_1 / harmonized_schema / Gender

Dictionary Script Summary Values Permissions

Derive Edit History

```
$(*SOKUP*).map({
  *1*: '0',
  *2*: '1'
},
*9*);
```

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Click the "Script" tab of the "Gender" variable to see the generated harmonization script.

OpalRecas Dashboard Projects data_harmonization_tutorial_1 harmonized_schema Administration Help enpadasI.ugent

data_harmonization_tutorial_1 / harmonized_schema

Dictionary Summary Values Permissions Download Export Copy

Properties

Name	harmonized_schema
Entity Type	Participant
Table References	data_harmonization_tutorial_1.Finrisk2007

Entity Filter

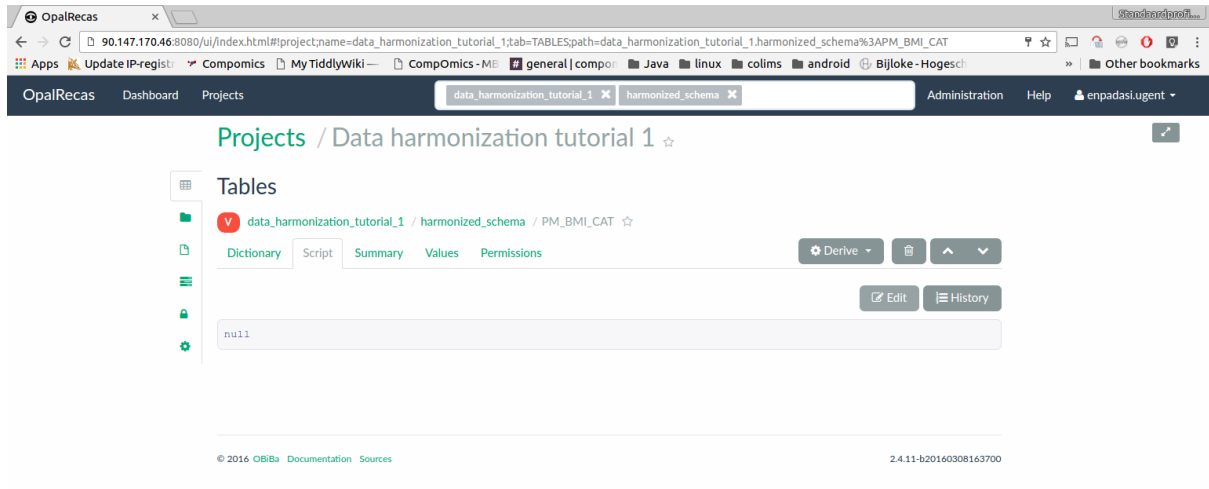
// no filter

Variables

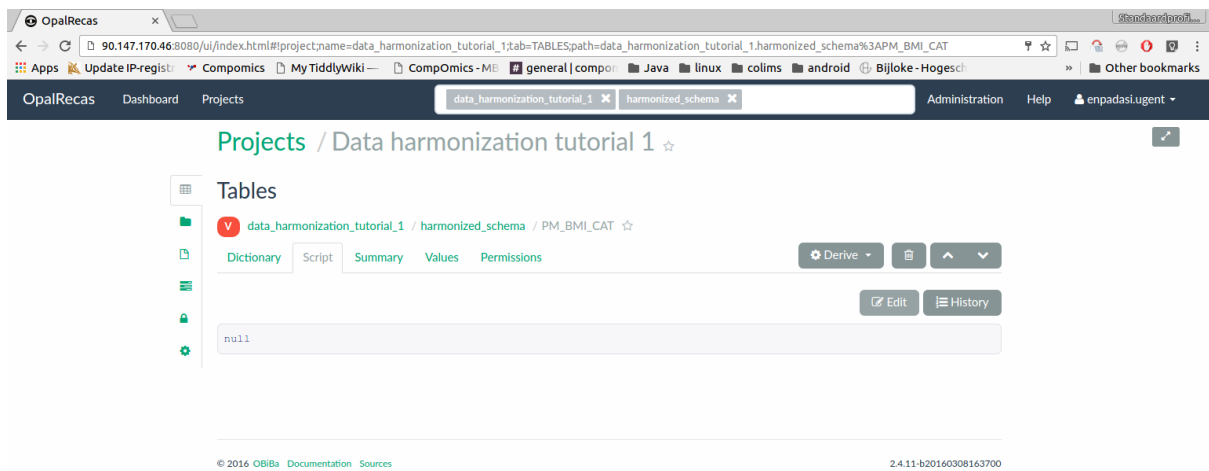
+ Add Variables Filter variables... Total 5

Name	Label	Value Type	Categories
Gender		integer	0, 1, 9
PM_BMI_CAT		integer	1, 2, 3, 9
DIS_DIAB		integer	0, 1, 9
PM_DIASTOLIC		integer	999
SMK_CIG_CURRENT		integer	0, 1, 9

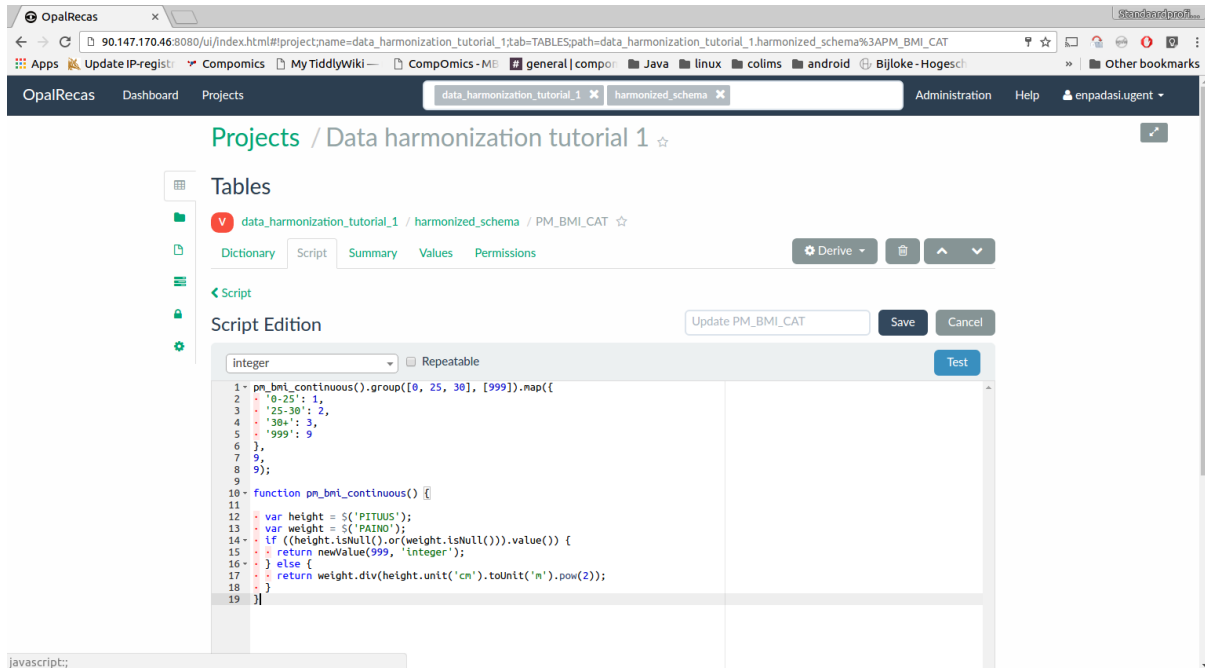
Click on the "PM_BMI_CAT" variable (Body Mass Index). It will be used as an example of a more complex transformation.



Click on the "PM_BMI_CAT" variable (Body Mass Index). It will be used as an example of a more complex transformation.



After selecting the "Script" tab, click on the "Edit" button.



The screenshot shows the 'Script Edition' interface for the 'PM_BMI_CAT' table. The script is as follows:

```

1 pm_bmi_continuous().group([0, 25, 30], [999]).map({
2   '18-25': 1,
3   '25-30': 2,
4   '30+': 3,
5   '999': 9
6 },
7 9,
8 9);
9
10 function pm_bmi_continuous() {
11
12   var height = S('FITBUS');
13   var weight = S('PAINO');
14   if ((height.isNull().or(weight.isNull()).value()) {
15     return newValue(999, 'integer');
16   } else {
17     return weight.div(height.unit('cm').toUnit('m').pow(2));
18   }
19 }

```

Copy the transformation script for the "PM_BMI_CAT" variable from the "Tutorial_Algorithms.doc" file, paste it into the script editor and click on the "Save" button. Do the same thing for the remaining variables.

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

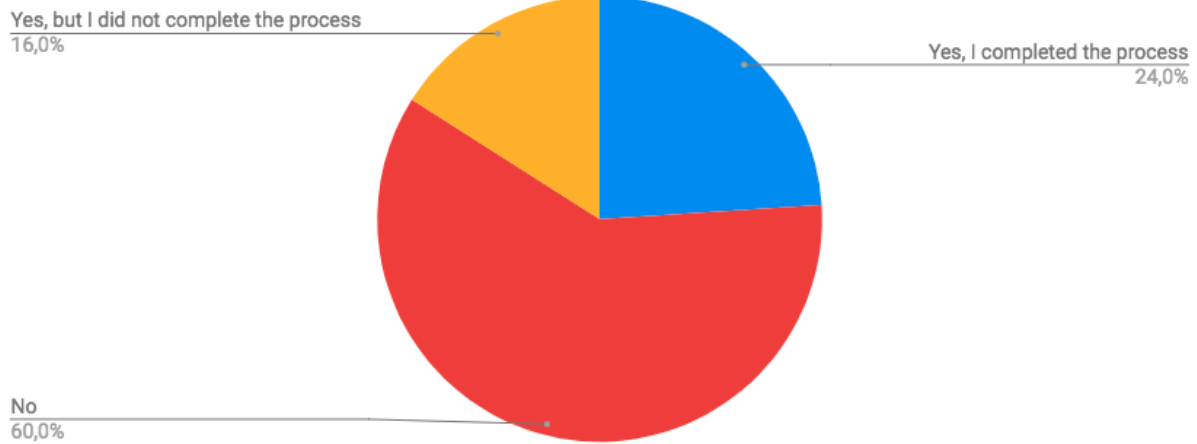
How easy or difficult is it to harmonize variables in OPAL?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

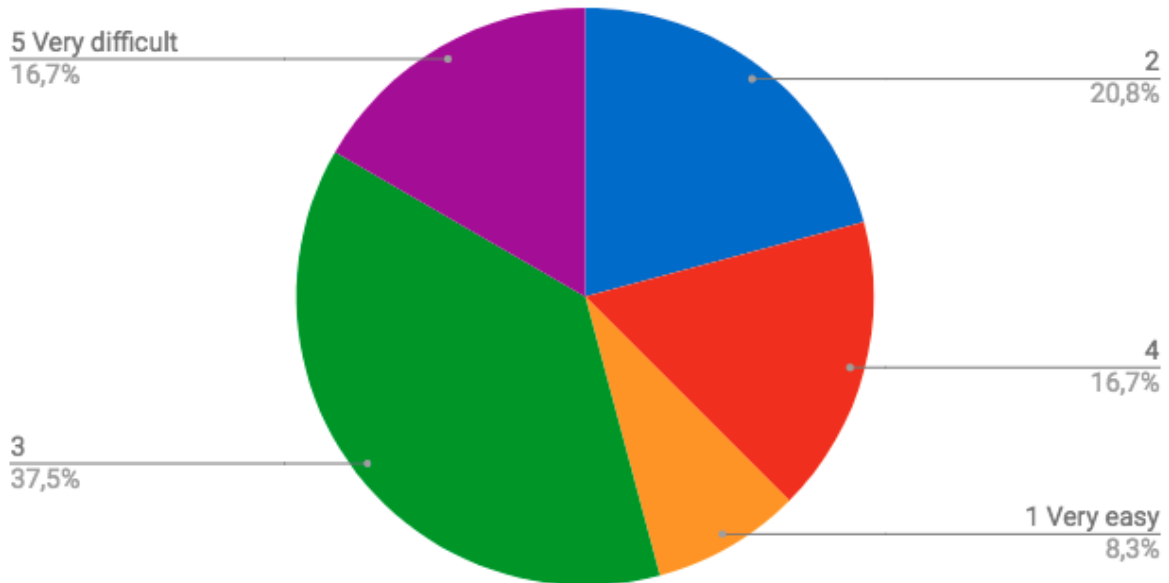


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Opal - Variable harmonization - Have you used this functionality



Opal - Variable harmonization - How easy or difficult did you find it?



OPAL - DO YOU HAVE ANY COMMENTS ON THE OPAL SERVER?

From the tutorial pages, is not obvious how to add tables from my own PC. To better explain, the fact that there is already a file browser displayed on the "add table" page, which browse the file of the OPAL instance and not the files from my local hard drive, can cause some minor confusion

I has to work with a virtual machine than don't work properly.

Simply leaving the requested information "by default" is not very complicated, but understanding the meaning of each choice is complex. For example Default Storage for database - Difference between "Add



Table/Add table for dictionary/Add View" - Different options of derive variables - Magma JavaScript
data format from missing values leads to difficulties - problems with certificate based authentication
I'm sorry to say, but I find it weird to be forced to assess the difficulties related to a "functionality" that I have not used at all!
I did not use it.
Maybe it is not so immediate for people from 'bio' area.
Who will define Harmonized data ? Will each of the partners use the same Harmonized data on each project on each OPAL server ?
More Information regarding the structure of the dictionary is needed(we had trouble with categories) - Project Name should be modifiable
I did not use it.
when harmonizing the variables (the simple way without "Script Edition"), if the type of variables (e.g. Integer, etc.) can be selected, it would be better.
A complete tutorial should always be available.

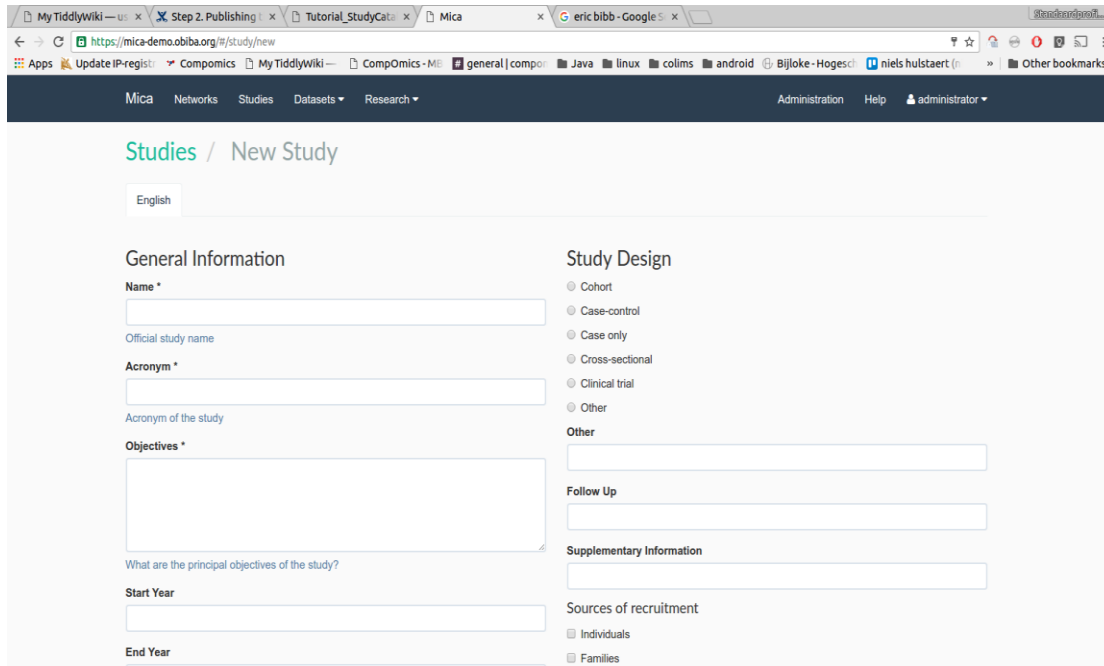
MICA - ADD STUDY

ID	Acronym	Name	Published	Status	Actions
atlantic-path	en Atlantic PATH	en Atlantic Partnership for Tomorrow's Health (The)	★	Draft	✓
bcgp	en BCGP	en BC Generations Project	★	Draft	✓
cag	en CaG	en CARTaGENE	★	Draft	✓
cls	en CLS	en Canberra Longitudinal Study	★	Draft	✓
clsa	en CLSA	en Canadian Longitudinal Study on Aging	★	Draft	✓
finrisk-2007	en FINRISK 2007	en National FINRISK Study 2007	★	Draft	✓
frele	en FR&LE	en FR&LE: Fragilité, une étude longitudinale de ses expressions [Frailty, a longitudinal study of its expressions]	★	Draft	✓
lassa	en LASA	en Longitudinal Aging Study Amsterdam	★	Draft	✓
lbis	en LBLS	en Long Beach Longitudinal Study	★	Draft	✓
lifelines	en LifeLines	en LifeLines Cohort Study & Biobank	★	Draft	✓
nuage	en NuAge	en Quebec Longitudinal Study on Nutrition and Successful Aging	★	Draft	✓
ohs	en OHS	en Ontario Health Study	★	Draft	✓
shin	en SHIP	en Study of Health in Pomerania	★	Draft	✓

After navigating to the studies tab in the main menu, click on the "Add Study" button.



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My TiddlyWiki — us x Step 2. Publishing x Tutorial_StudyCat: x Mica x eric bibb - Google S x

https://mica-demo.obiba.org/#/study/new

Apps Update IP-Regist: Compomics My TiddlyWiki — CompOmics - MBI general | compo Java linux colims android Bijloke - Hogescl niels hulstaert (Other bookmarks

Mica Networks Studies Datasets Research Administration Help administrator

Studies / New Study

English

General Information

Name *
Official study name

Acronym *
Acronym of the study

Objectives *
What are the principal objectives of the study?

Start Year

End Year

Study Design

Cohort
 Case-control
 Case only
 Cross-sectional
 Clinical trial
 Other

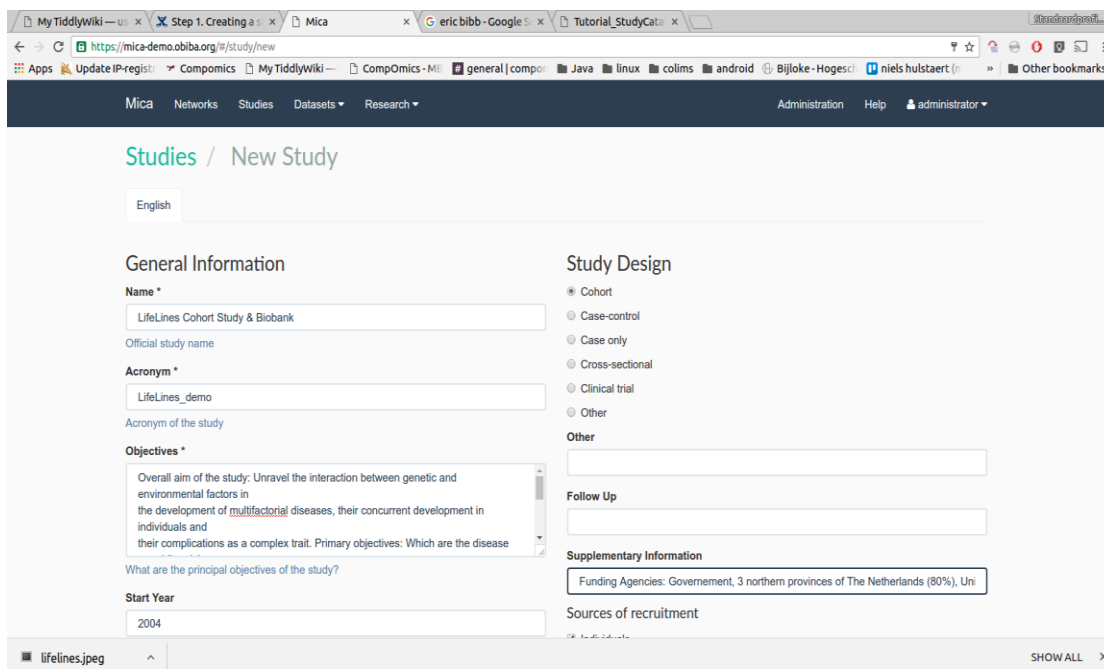
Other

Follow Up

Supplementary Information

Sources of recruitment

Individuals
 Families



My TiddlyWiki — us x Step 1. Creating a s x Mica x eric bibb - Google S x Tutorial_StudyCat: x

https://mica-demo.obiba.org/#/study/new

Apps Update IP-Regist: Compomics My TiddlyWiki — CompOmics - MBI general | compo Java linux colims android Bijloke - Hogescl niels hulstaert (Other bookmarks

Mica Networks Studies Datasets Research Administration Help administrator

Studies / New Study

English

General Information

Name *
LifeLines Cohort Study & Biobank
Official study name

Acronym *
LifeLines_demo
Acronym of the study

Objectives *
Overall aim of the study: Unravel the interaction between genetic and environmental factors in the development of multifactorial diseases, their concurrent development in individuals and their complications as a complex trait. Primary objectives: Which are the disease
What are the principal objectives of the study?

Start Year
2004

Study Design

Cohort
 Case-control
 Case only
 Cross-sectional
 Clinical trial
 Other

Other

Follow Up

Supplementary Information
Funding Agencies: Government, 3 northern provinces of The Netherlands (80%), Un

Sources of recruitment

lifelines.jpeg SHOW ALL X



JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Fill in all the required fields and click on the "Save" button on the bottom of the page

On the study main page, click on the "Add contact" or "Add investigator" button. Fill in the required fields and click on the "Save" button.

These images show part of the process to add a study to MICA.



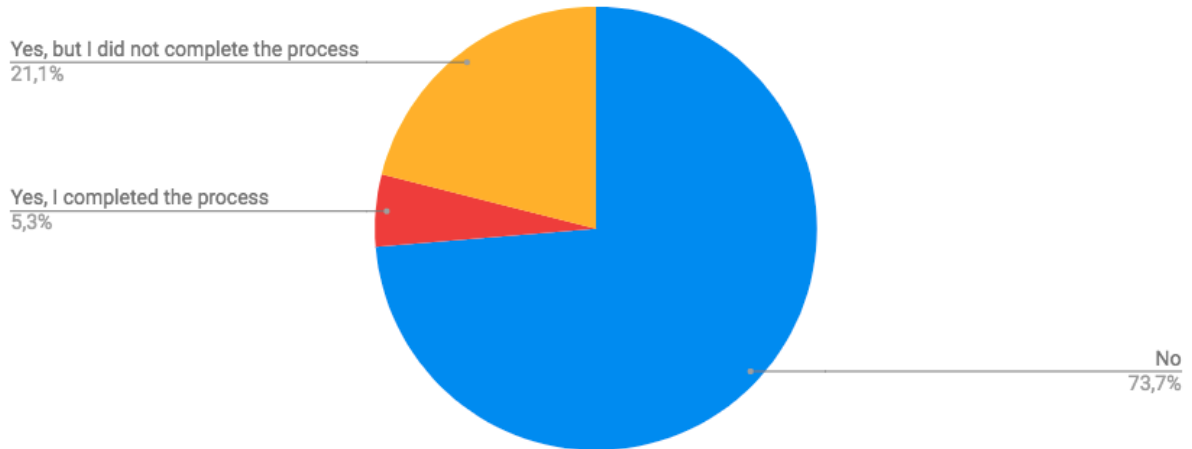
Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

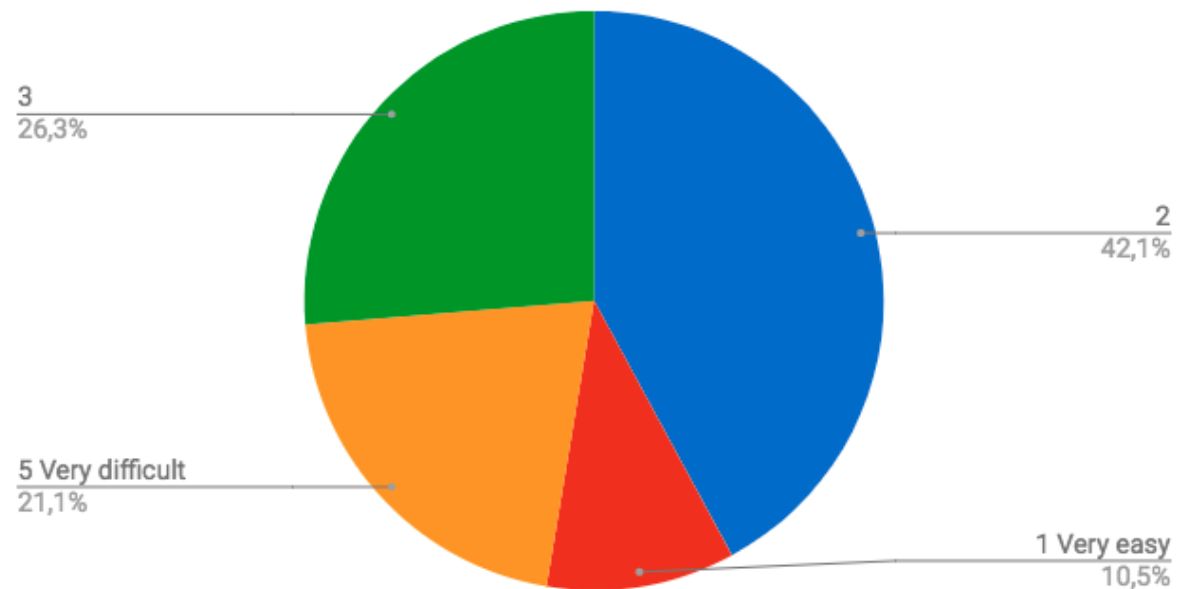
How easy or difficult is it to add a study to MICA?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

MICA - add study - Have you used this functionality



MICA - add study - How easy or difficult did you find it?





MICA - ADD POPULATION

The screenshot shows the MICA web application interface. The browser address bar indicates the URL is https://mica-demo.obiba.org/#/study/lifelines_demo. The main navigation bar includes 'Mica', 'Networks', 'Studies', 'Datasets', and 'Research'. The 'Populations' section is highlighted, showing a blue button labeled '+ Add Population'. The 'Marker Paper' section contains a citation: 'Stolk RP, Rosmalen JG, Postma DS, de Boer RA, Navis G, Slaets JP, Ormel J, Woffenbuttel BHR. Universal risk factors for multifactorial diseases: LifeLines: a three-generation population-based study. Eur J Epidemiol 2008; 23(1): 67-74' with a Pubmed ID of [PUBMED 18075776](#).

On the study main page, click on the "Add Population" button.

The screenshot shows the 'Add Population' form in the MICA application. The breadcrumb trail is 'Studies / lifelines_demo'. The form is titled 'Add Population' and includes a sidebar with a menu icon. The 'General Information' section contains:

- ID ***:
- Name ***:
- Description**:

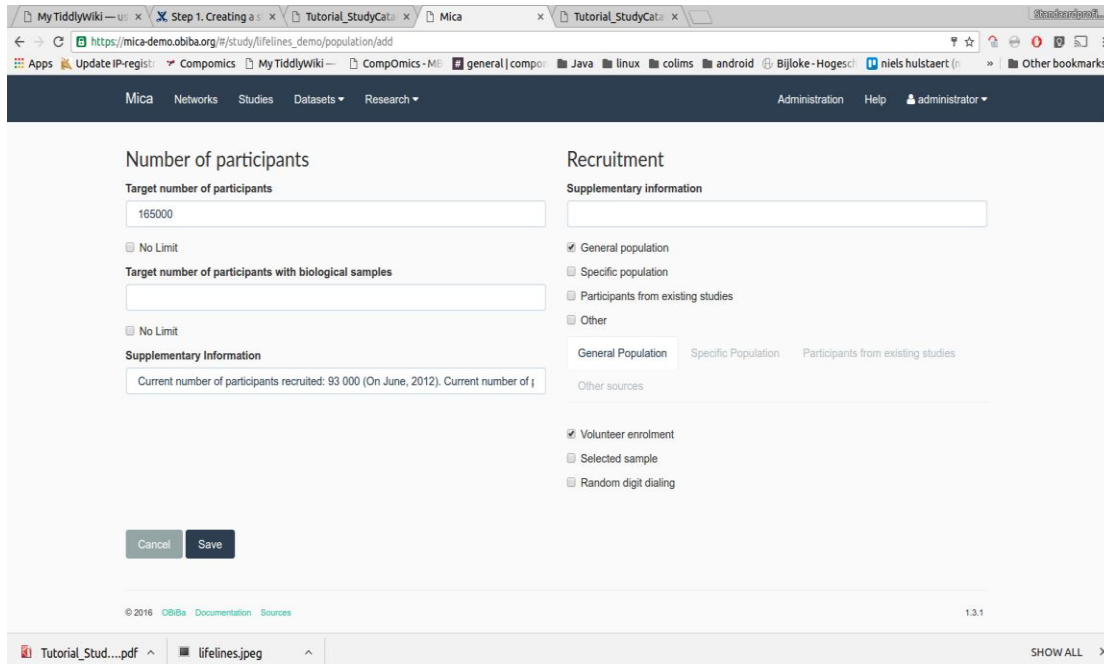
 The 'Selection criteria' section includes:

- Gender**:
- Minimum age**:
- Maximum age**:

 On the right side, there are two buttons: '+Add health status' and '+Add ethnic origin'. The bottom of the page shows a file manager with 'Tutorial_Stud....pdf' and 'lifelines.jpeg'.



JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE



Number of participants

Target number of participants: 165000

No Limit

Target number of participants with biological samples

No Limit

Supplementary Information

Current number of participants recruited: 93 000 (On June, 2012). Current number of [

Recruitment

Supplementary information

General population

Specific population

Participants from existing studies

Other

General Population | Specific Population | Participants from existing studies

Other sources

Volunteer enrolment

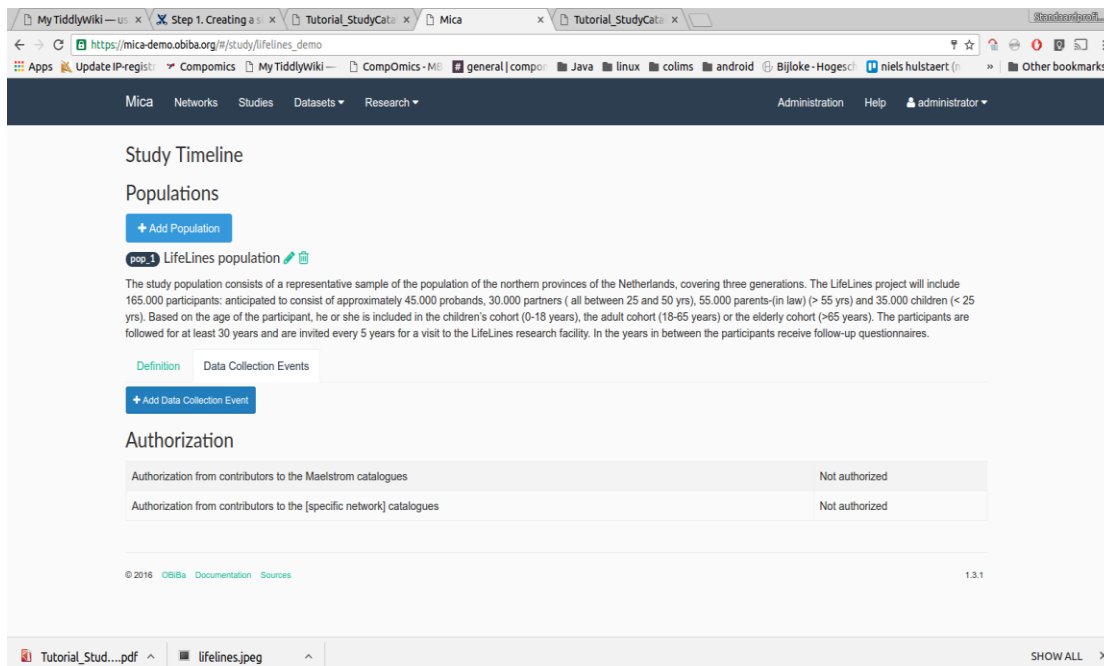
Selected sample

Random digit dialing

Cancel Save

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Fill in the required fields and click on the "Save" button.



Study Timeline

Populations

+ Add Population

pop_1 LifeLines population

The study population consists of a representative sample of the population of the northern provinces of the Netherlands, covering three generations. The LifeLines project will include 165,000 participants: anticipated to consist of approximately 45,000 probands, 30,000 partners (all between 25 and 50 yrs), 55,000 parents-(in law) (> 55 yrs) and 35,000 children (< 25 yrs). Based on the age of the participant, he or she is included in the children's cohort (0-18 years), the adult cohort (18-65 years) or the elderly cohort (>65 years). The participants are followed for at least 30 years and are invited every 5 years for a visit to the LifeLines research facility. In the years in between the participants receive follow-up questionnaires.

Definition | Data Collection Events

+ Add Data Collection Event

Authorization

Authorization from contributors to the Maelstrom catalogues	Not authorized
Authorization from contributors to the [specific network] catalogues	Not authorized

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On the study main page in the "Population" section, click on the "Data Collection Events" tab. Then click on the "Add Data Collection Event" button.



The screenshot shows the 'General Information' form in the MICA application. The form includes the following fields:

- ID ***: data_coll_1
- Start Year ***: 2007
- Name ***: Baseline recruitment / initial data collection
- Start Month**: (empty)
- Description**: (empty text area)
- End Year**: 2013
- End Month**: (empty)

Under the 'Data sources' section, the following options are checked:

- Questionnaires
- Physical measures
- Biosamples
- Administrative databases
- Others

Buttons for 'Cancel' and 'Save' are located at the bottom left of the form.

The screenshot shows the 'Study Timeline' and 'Populations' sections of the MICA application. The 'Study Timeline' shows a red bar representing the study duration from 2007 to 2018. The 'Populations' section includes a table of data collection events:

#	Name	Description	Start	End	Actions
data_coll_1	Baseline recruitment / initial data collection		2007 (January)	2013 (December)	
data_coll_2	First Follow-up of participants		2011 (January)	2017 (December)	

Fill in the required fields and click on the "Save" button on the bottom of the page. The added date collection events are shown in a time-line on the study main page.

These images show part of the process to add population to MICA



Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

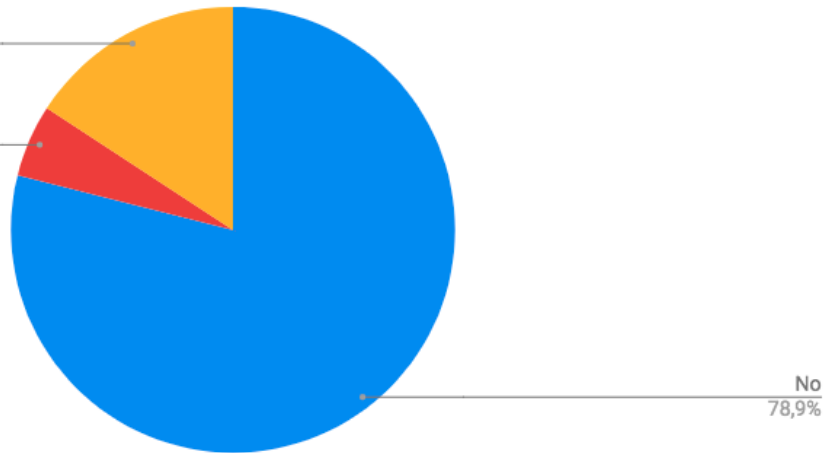
How easy or difficult is it to add new population to MICA?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

MICA - add population - Have you used this functionality

Yes, but I did not complete the process
15,8%

Yes, I completed the process
5,3%



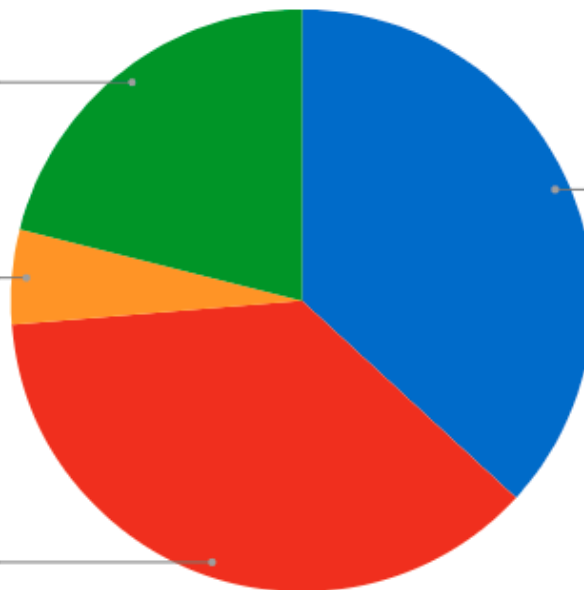
MICA - add population - How easy or difficult did you find it?

5 Very difficult
21,1%

1 Very easy
5,3%

3
36,8%

2
36,8%

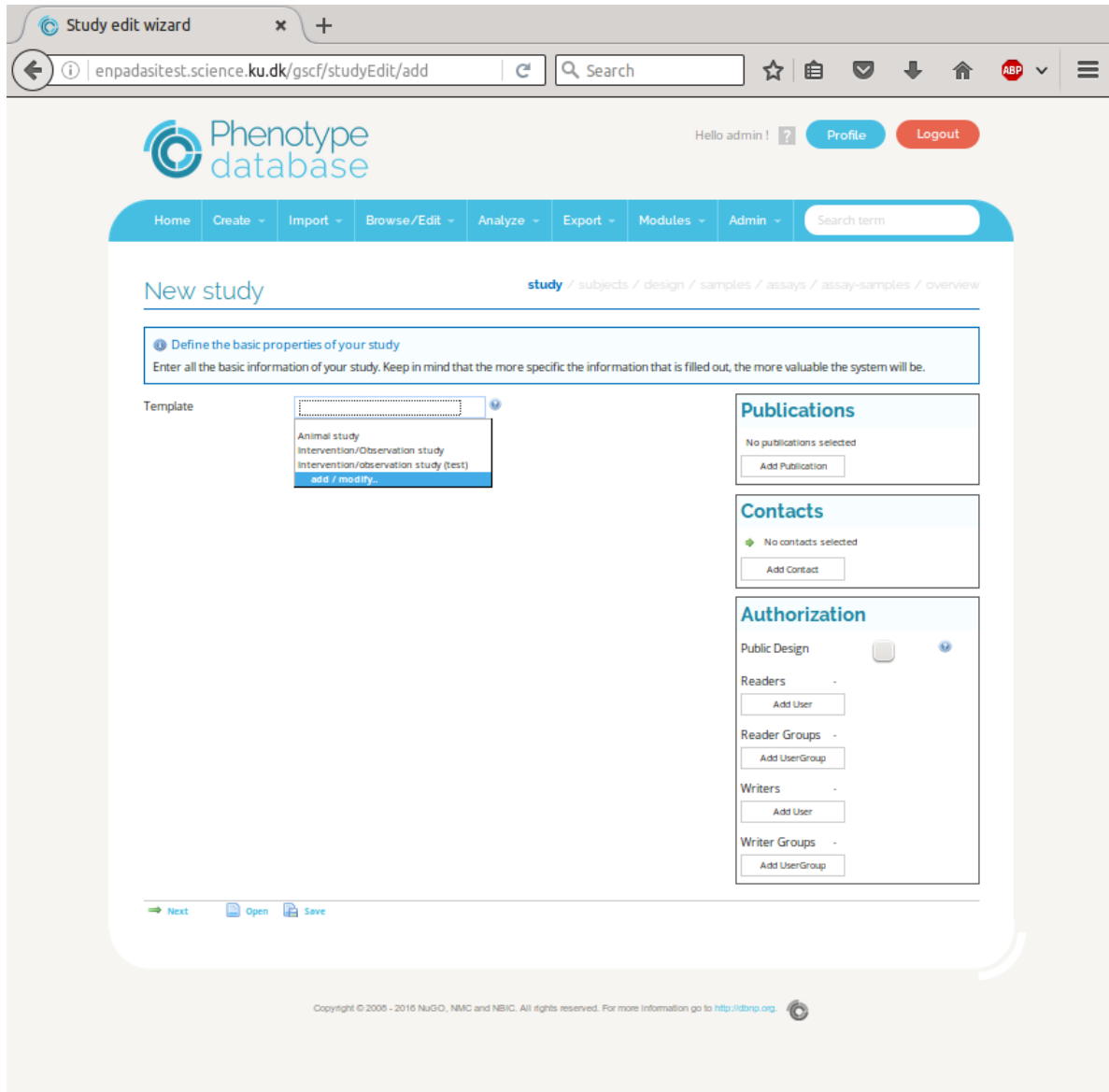




MICA - DO YOU HAVE ANY COMMENTS ON THE MICA SERVER?

Was not able to access the MICA instance at INFN of Bari for testing
Unable to do anything cause the account seems to have no admin privileges.
No access to the study creation on the mica server http://mica.cloud.ba.infn.it So I could not test anything.
I'm sorry to say, but I find it weird to be forced to assess the difficulties related to a "functionality" that I have not used at all!
no
I have not found the possibility to test MICA, even if it seems that it is not difficult to use.
Give a hint on how to add multiple population and multiple samples within the same population. This could be a tutorial part.
no
Please, provide a practical test.

PHENOTYPE DATABASE (DBNP) - CREATE A NEW STUDY

Study edit wizard

enpadasitest.science.ku.dk/gscf/studyEdit/add

Phenotype database

Hello admin! Profile Logout

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

New study

study / subjects / design / samples / assays / assay-samples / overview

Define the basic properties of your study
Enter all the basic information of your study. Keep in mind that the more specific the information that is filled out, the more valuable the system will be.

Template

- Animal study
- Intervention/Observation study
- Intervention/observation study (test)
- add / modify...

Publications

No publications selected

Add Publication

Contacts

No contacts selected

Add Contact

Authorization

Public Design

Readers -

Add User

Reader Groups -

Add UserGroup

Writers -

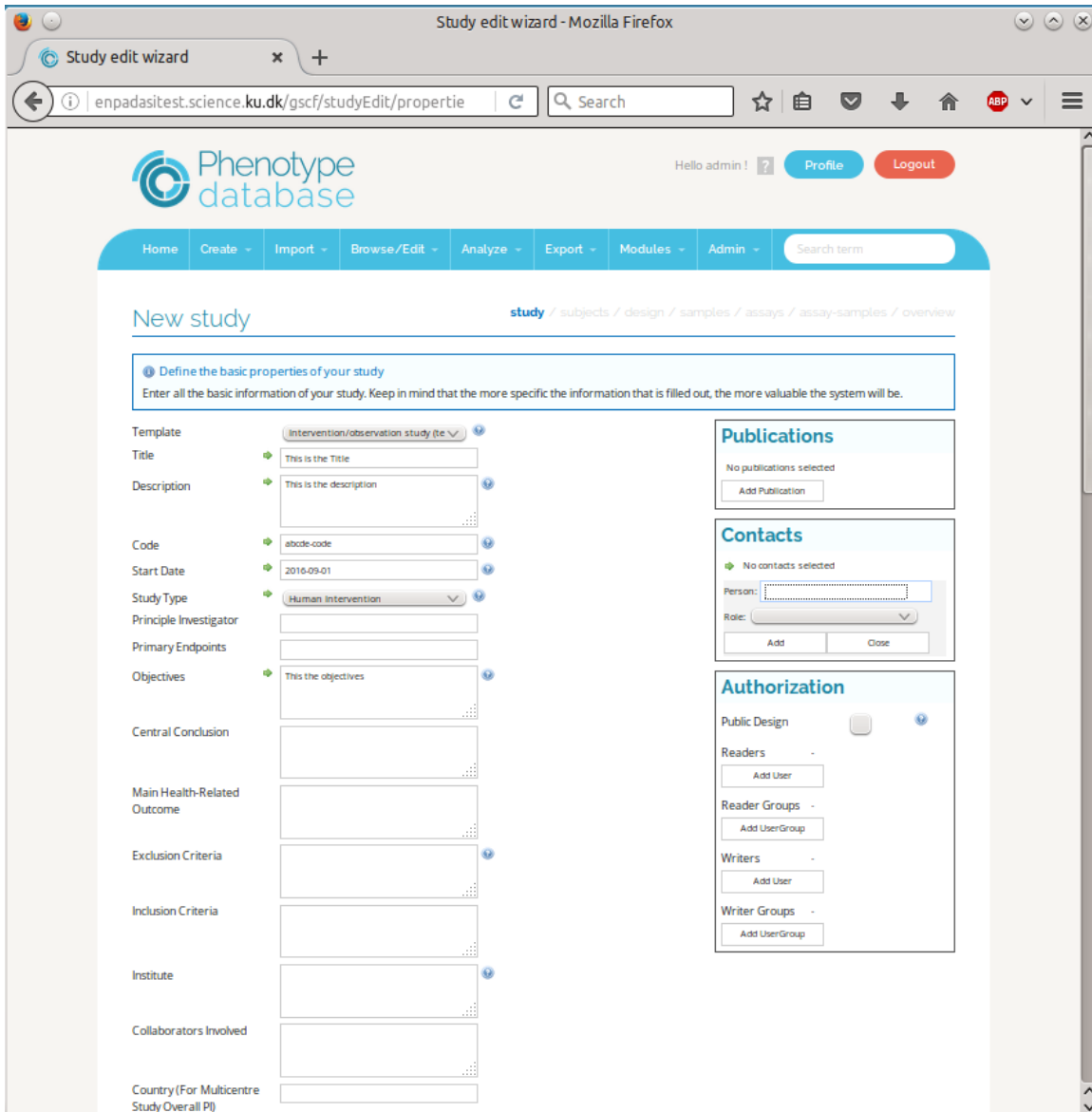
Add User

Writer Groups -

Add UserGroup

Next Open Save

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The screenshot shows the 'Study edit wizard' interface in Mozilla Firefox. The browser address bar shows the URL: `enpadasitest.science.ku.dk/gscf/studyEdit/propertie`. The page header includes the 'Phenotype database' logo, a user greeting 'Hello admin!', and buttons for 'Profile' and 'Logout'. A navigation menu contains: Home, Create, Import, Browse/Edit, Analyze, Export, Modules, Admin, and a search bar.

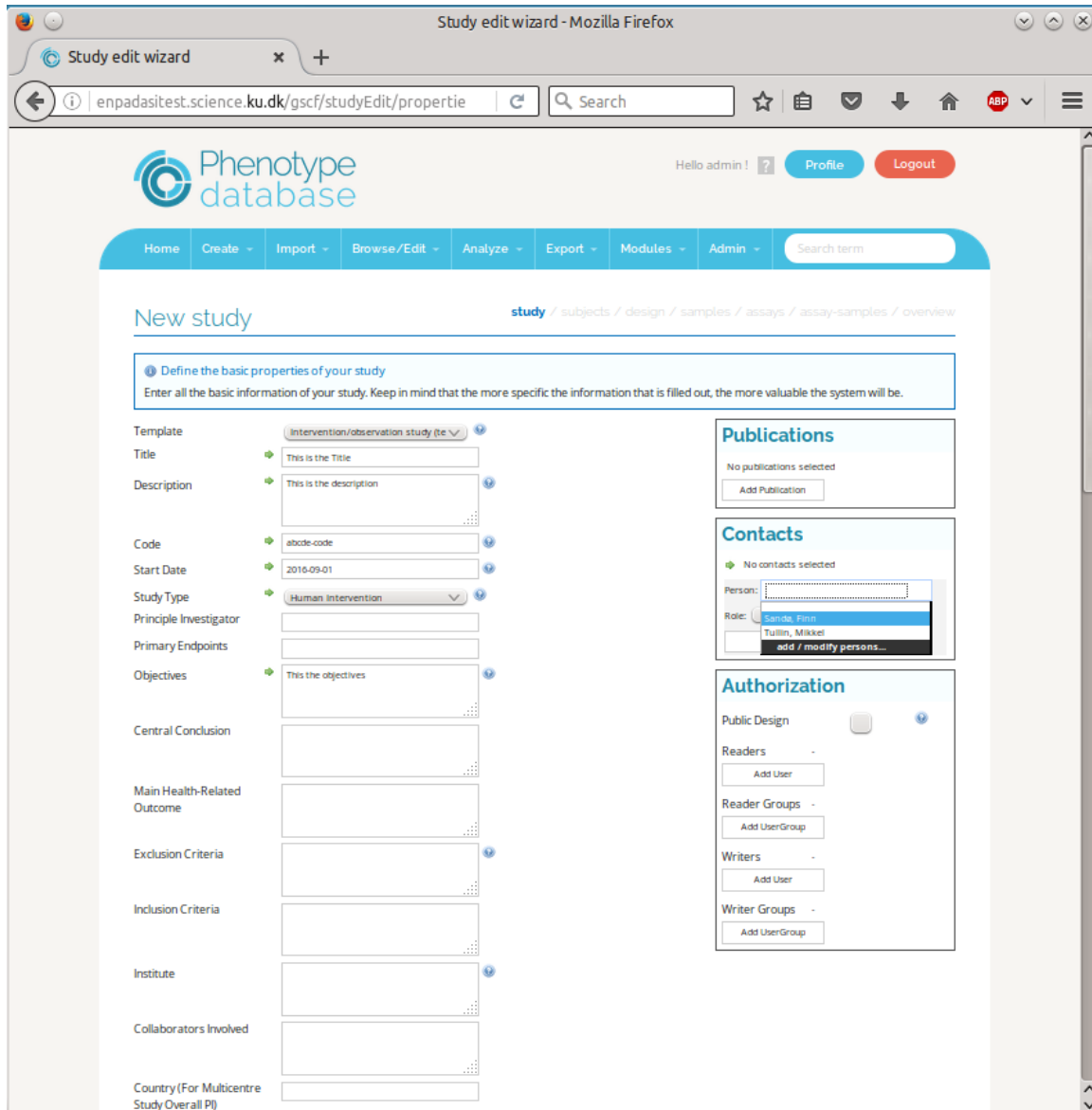
The main content area is titled 'New study' and includes a breadcrumb trail: `study / subjects / design / samples / assays / assay-samples / overview`. A blue box contains the instruction: 'Define the basic properties of your study. Enter all the basic information of your study. Keep in mind that the more specific the information that is filled out, the more valuable the system will be.'

The form fields are as follows:

- Template: `Intervention/observation study (b` (dropdown)
- Title: `This is the Title` (text input)
- Description: `This is the description` (text input)
- Code: `abcde-code` (text input)
- Start Date: `2016-09-01` (date input)
- Study Type: `Human Intervention` (dropdown)
- Principle Investigator: (empty text input)
- Primary Endpoints: (empty text input)
- Objectives: `This the objectives` (text input)
- Central Conclusion: (empty text input)
- Main Health-Related Outcome: (empty text input)
- Exclusion Criteria: (empty text input)
- Inclusion Criteria: (empty text input)
- Institute: (empty text input)
- Collaborators Involved: (empty text input)
- Country (For Multicentre Study Overall PI): (empty text input)

On the right side, there are three panels:

- Publications:** 'No publications selected', with an 'Add Publication' button.
- Contacts:** 'No contacts selected', with 'Person:' and 'Role:' dropdowns, and 'Add' and 'Close' buttons.
- Authorization:** 'Public Design' toggle, and sections for 'Readers', 'Reader Groups', 'Writers', and 'Writer Groups', each with 'Add User' and 'Add UserGroup' buttons.

Study edit wizard - Mozilla Firefox

enpadasitest.science.ku.dk/gscf/studyEdit/propertie

Phenotype database

Hello admin! Profile Logout

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

New study

study / subjects / design / samples / assays / assay-samples / overview

Define the basic properties of your study
Enter all the basic information of your study. Keep in mind that the more specific the information that is filled out, the more valuable the system will be.

Template: intervention/observation study (be)

Title: This is the Title

Description: This is the description

Code: abcde-code

Start Date: 2016-09-01

Study Type: Human Intervention

Principle Investigator: [empty]

Primary Endpoints: [empty]

Objectives: This the objectives

Central Conclusion: [empty]

Main Health-Related Outcome: [empty]

Exclusion Criteria: [empty]

Inclusion Criteria: [empty]

Institute: [empty]

Collaborators Involved: [empty]

Country (For Multicentre Study Overall PI): [empty]

Publications

No publications selected

Add Publication

Contacts

No contacts selected

Person: [empty]

Role: Sando, Finn
Tullin, Mikkel

add / modify persons...

Authorization

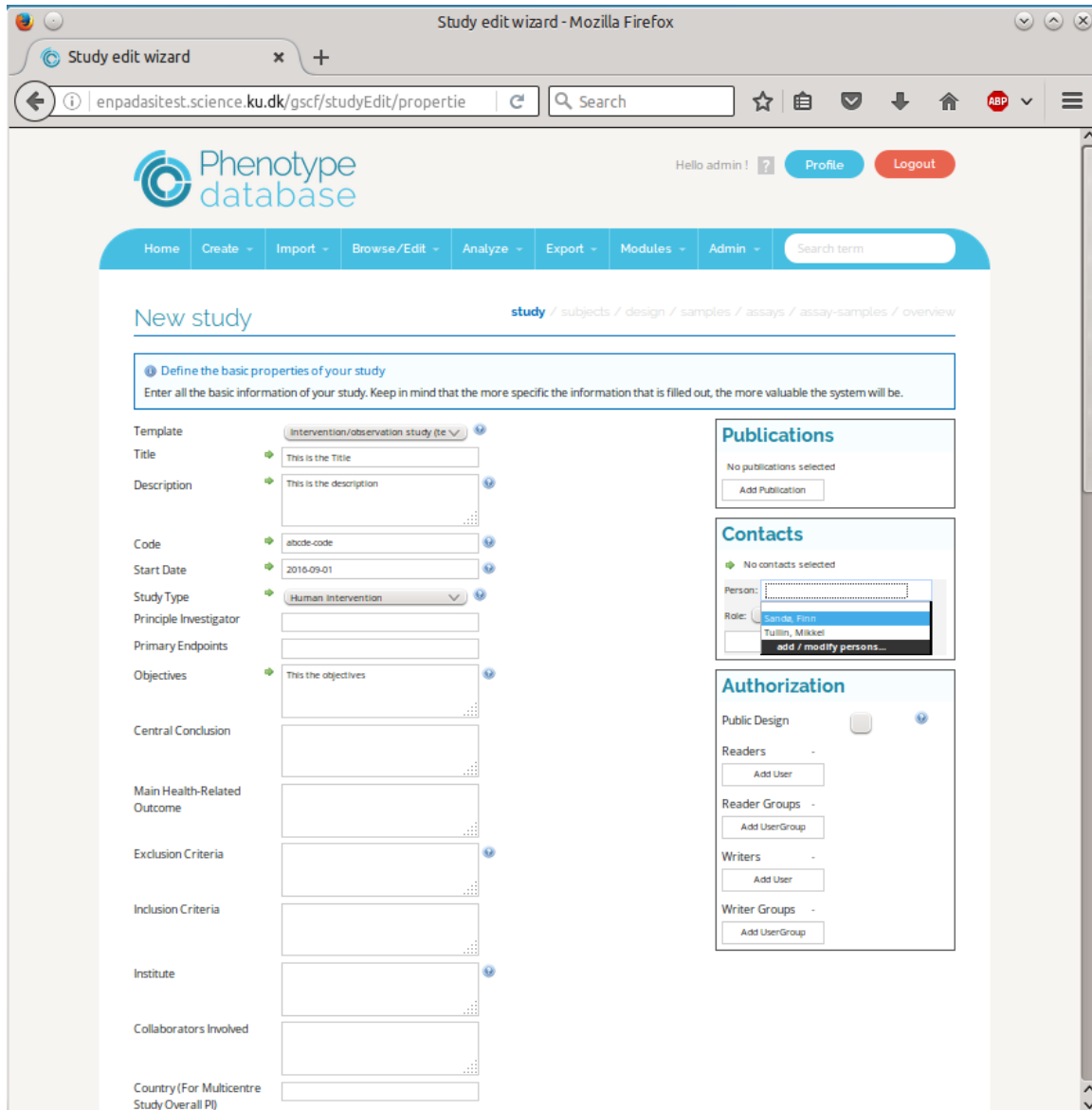
Public Design: [checkbox]

Readers: Add User

Reader Groups: Add UserGroup

Writers: Add User

Writer Groups: Add UserGroup



These images show part of the process to create a new study in the Phenotype Database (dbNP).

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

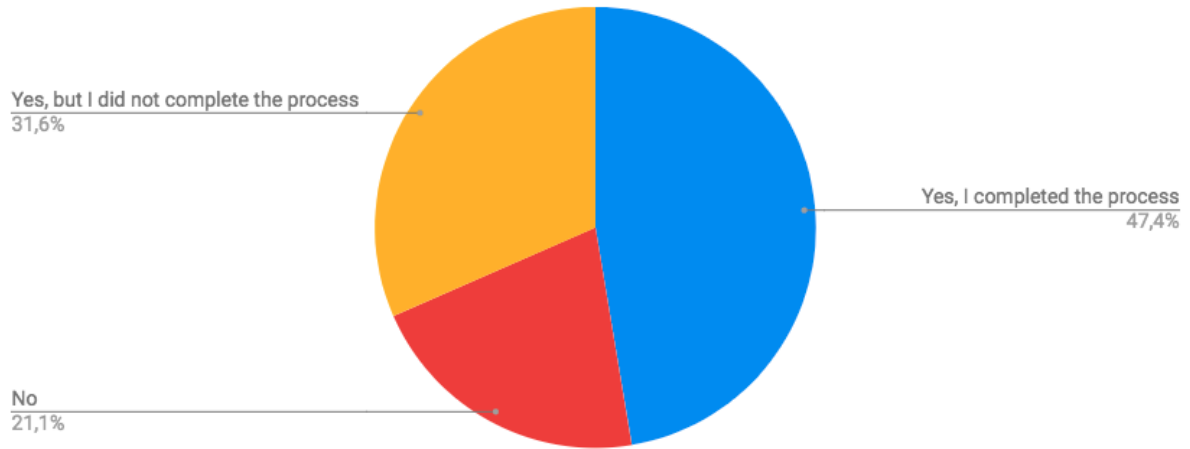
How easy or difficult is it to create a new study to the Phenotype Database (dbNP)?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

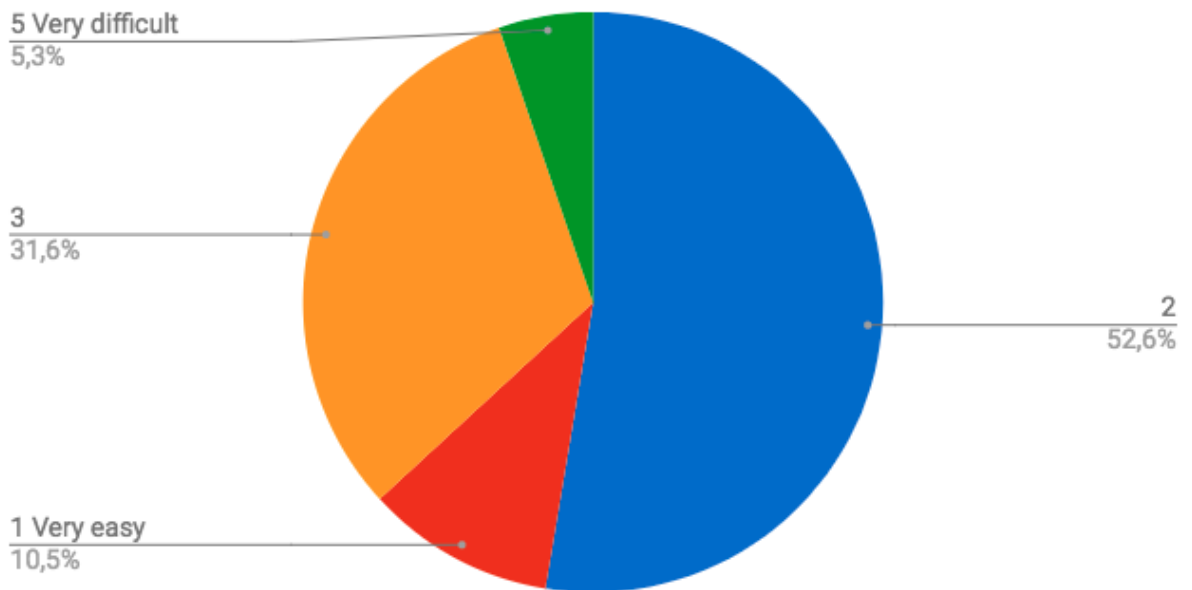


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Phenotype Database (dbNP) - create a new study - Have you used this functionality

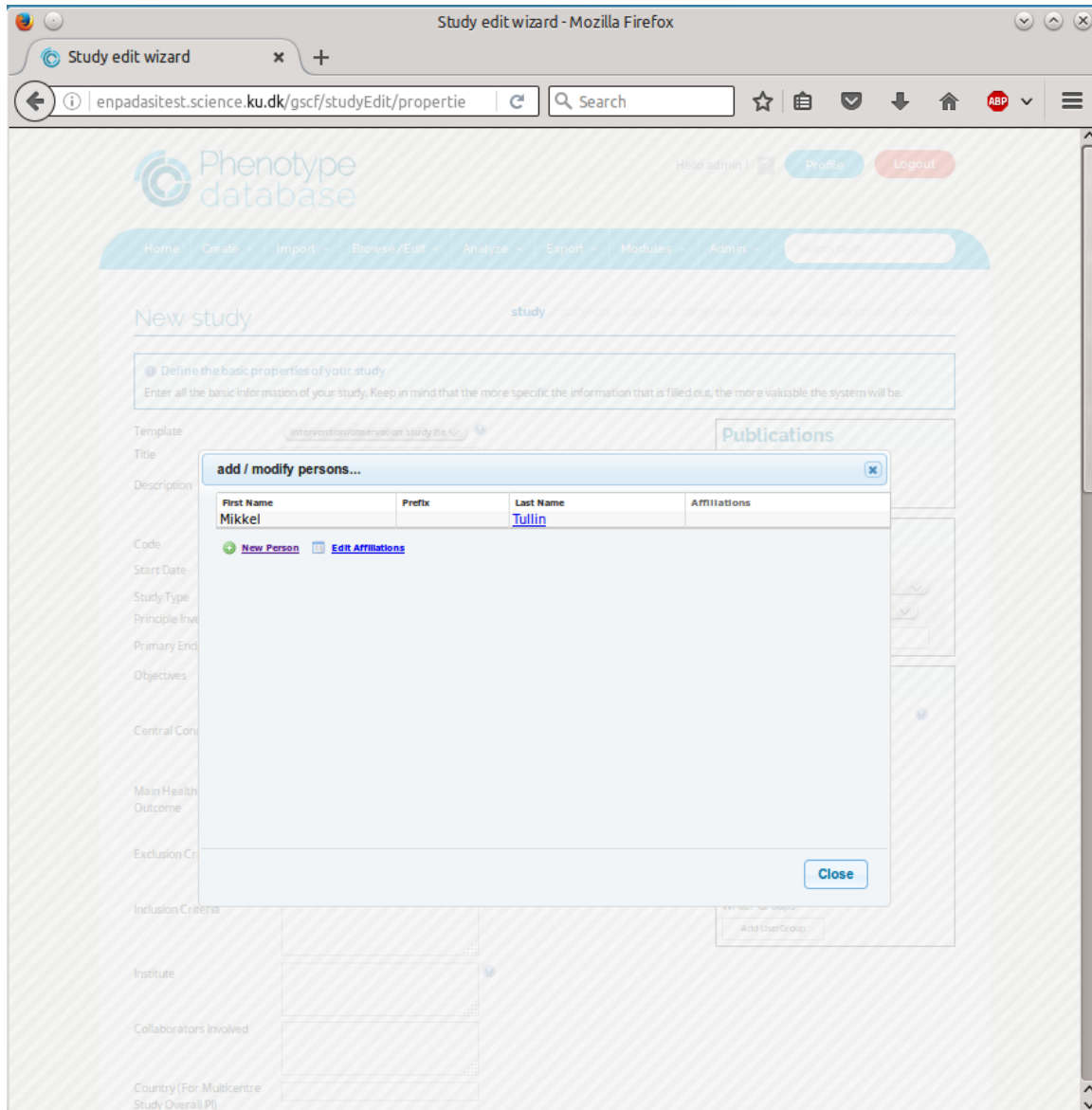


Phenotype Database (dbNP) - create a new study - How easy or difficult did you find it?





PHENOTYPE DATABASE (DBNP) - ADD A PERSON TO THE PERSON LIST

Study edit wizard - Mozilla Firefox

enpadasitest.science.ku.dk/gscf/studyEdit/propertie

Phenotype database

Home Create Import Browse/Edit Analyze Export Modules Admin

New study

Define the basic properties of your study. Enter all the basic information of your study. Keep in mind that the more specific the information that is filled out, the more valuable the system will be.

Template: [intervention/observation study \(6\)](#)

add / modify persons...

First Name	Prefix	Last Name	Affiliations
Mikkel		Tullin	

[New Person](#) [Edit Affiliations](#)

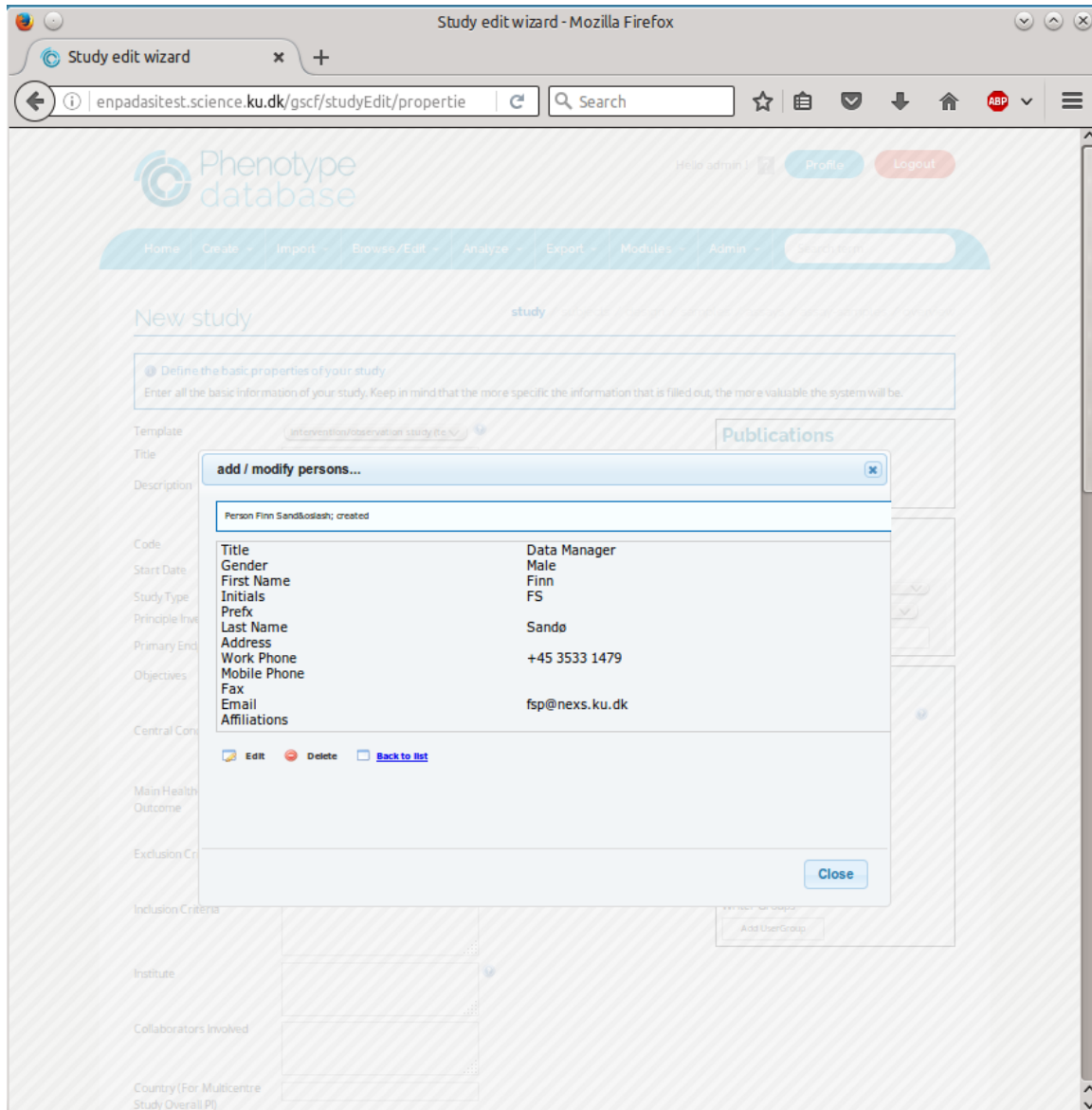
Close

Add UserGroup

Institute

Collaborators Involved

Country (For Multicentre Study Overall PI)



These images show the process to add a person to the person list in the Phenotype Database (dbNP)

Have you used this functionality?

- No
 Yes, but I did not complete the process
 Yes, I completed the process

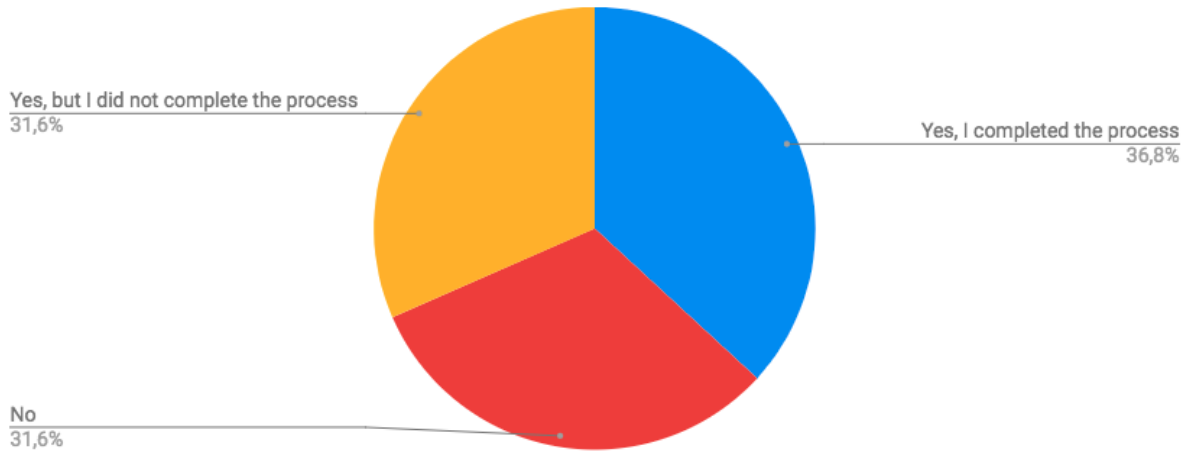
How easy or difficult is it to add a person to person list in the Phenotype Database (dbNP)?

- 1 Very easy
 2
 3
 4
 5 Very difficult

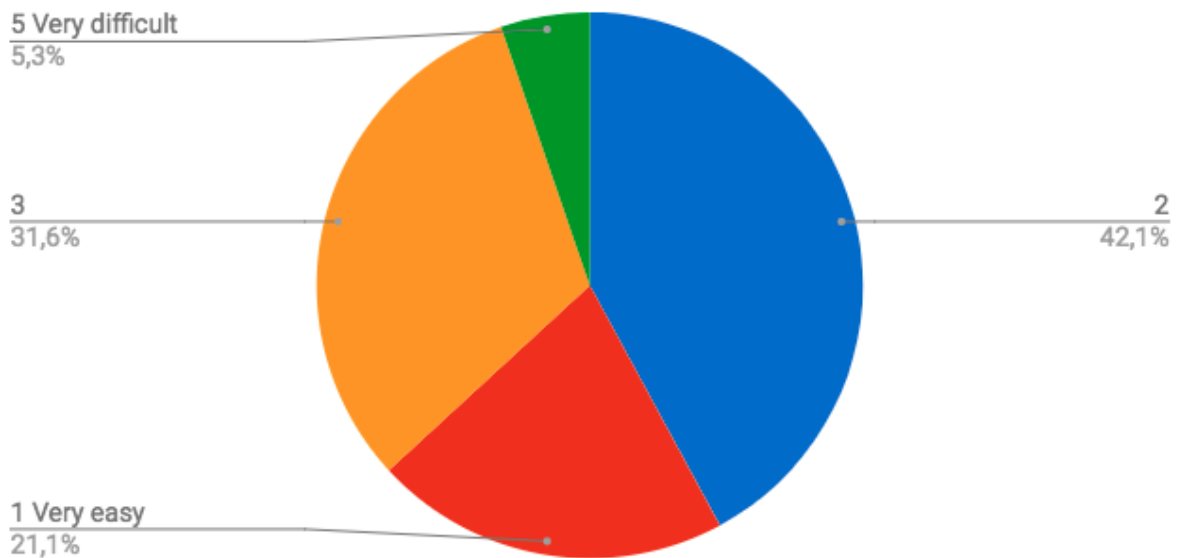


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

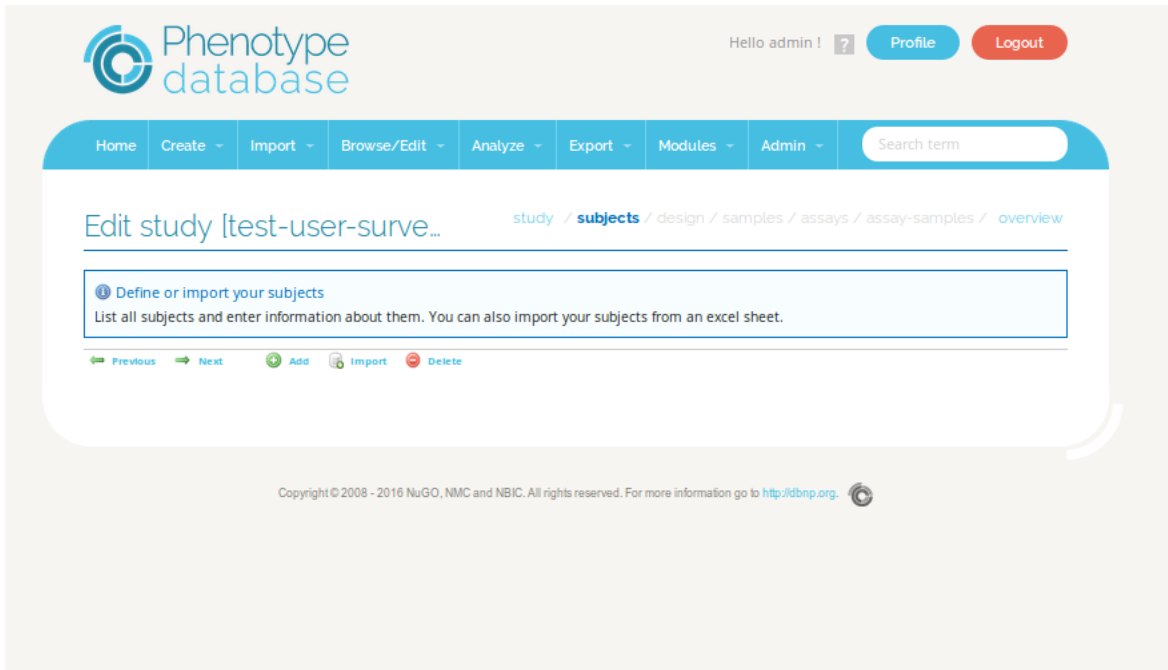
Phenotype Database (dbNP) - add a person to the person list - Have you used this functionality



Phenotype Database (dbNP) - add a person to the person list - How easy or difficult did you find it?



PHENOTYPE DATABASE (DBNP) – UPLOAD SUBJECTS

Phenotype database

Hello admin! [Profile](#) [Logout](#)

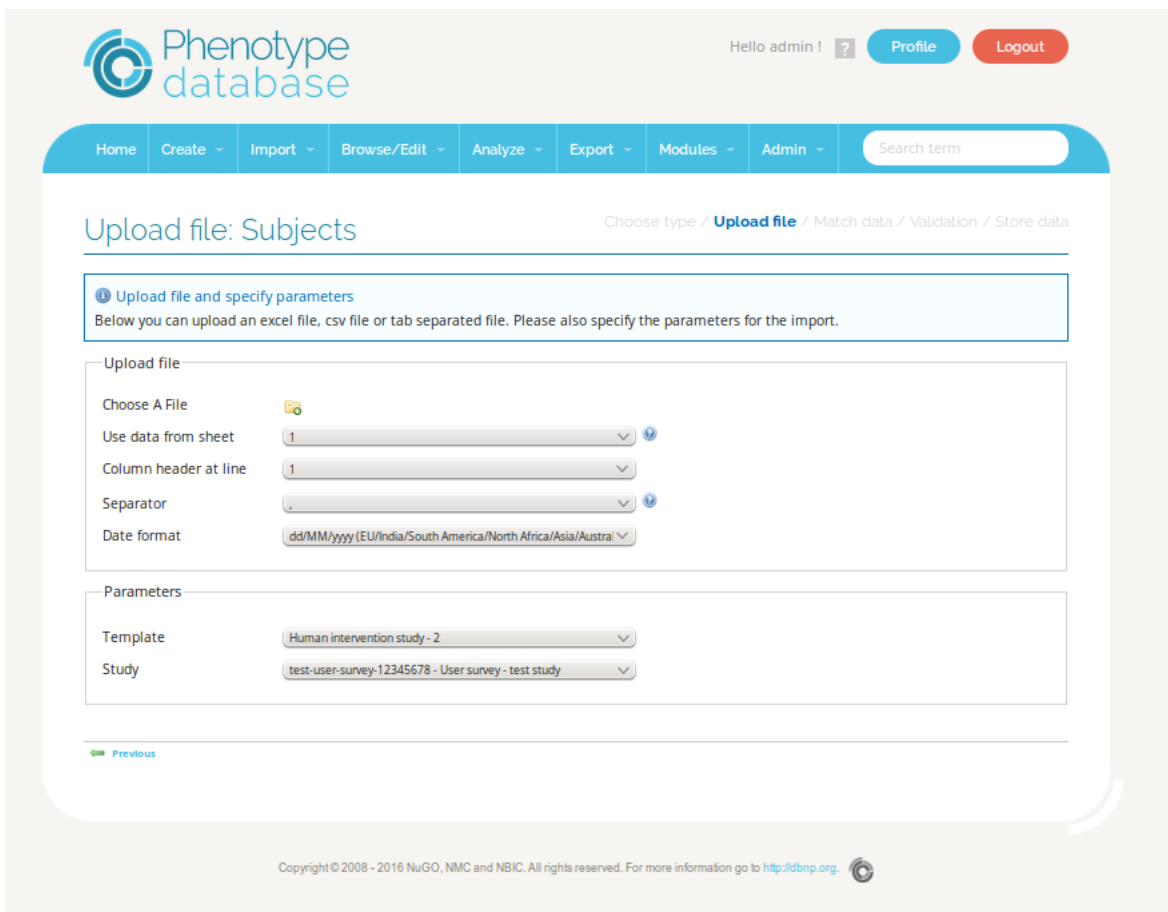
Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Edit study | test-user-surve... study / **subjects** / design / samples / assays / assay-samples / overview

Define or import your subjects
List all subjects and enter information about them. You can also import your subjects from an excel sheet.

← Previous → Next + Add 📄 Import - Delete

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Phenotype database


Hello admin! [Profile](#) [Logout](#)

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Upload file: Subjects Choose type / **Upload file** / Match data / Validation / Store data

Upload file and specify parameters
Below you can upload an excel file, csv file or tab separated file. Please also specify the parameters for the import.

Upload file

Choose A File 

Use data from sheet 1

Column header at line 1

Separator ,

Date format dd/MM/yyyy (EU/India/South America/North Africa/Asia/Austral)

Parameters

Template Human intervention study - 2

Study test-user-survey-12345678 - User survey - test study

← Previous

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The screenshot shows a web browser window titled 'Import data - Mozilla Firefox' with the URL 'enpadasitest.science.ku.dk/gscf/importer/upload?ini'. The page displays the 'Phenotype database' interface with a user profile for 'admin'. A 'File Upload' dialog box is open, showing a file explorer view of the path 'media > sf_H_DRIVE > Documents > Opgaver > ENPADASI > dash-in > survey-prepare'. The file list includes:

Name	Size	Modified
dbnp		Tuesday
dbnp-fs-2016-09-26		14:09
enpadas_i_survey		Tuesday
enpadas_i_survey-1		Tuesday
opal-1		08/31/2016
revised docs		Tuesday
enpadas_i-logo	424 bytes	07/06/2016
ENPADASi-logo.jpg	32.5 kB	07/06/2016
enpadas_i_survey.zip	4.2 MB	09/18/2016
enpadas_i_survey-1.zip	2.7 MB	09/07/2016
revised docs.zip	1.8 MB	09/06/2016
SUBJECTS_DATA-enpadasitutorial_study.xlsx	23.8 kB	14:57

The 'SUBJECTS_DATA-enpadasitutorial_study.xlsx' file is highlighted. Below the file list are 'All Files', 'Open', and 'Cancel' buttons. The background interface shows a 'Template' dropdown set to 'Human intervention study' and a 'Study' dropdown set to 'abode-code - This is the Title'.





Hello admin! [?](#) [Profile](#) [Logout](#)

Home
Create
Import
Browse/Edit
Analyze
Export
Modules
Admin

Search term

Upload file: Subjects

Choose type / **Upload file** / Match data / Validation / Store data

Upload file and specify parameters

Below you can upload an excel file, csv file or tab separated file. Please also specify the parameters for the import.

Upload file

Choose A File: Uploaded SUBJECTS_DATA-Sea bu.

Use data from sheet:

Column header at line:

Separator:

Date format:

Data preview

Name	Start group	Species	Country (humans)	Ethnicity (or strain)	Gender	Age	Initial bodyweight	Initial Body Height	BMI	Initial waist circumference	Commer
5493	1	Homo sapiens	Denmark	Caucasian	Male	26	118.2	1.855	34	110.5	Test d seque 1Sea
5502	1	Homo sapiens	Denmark	Caucasian	Male	44	89.9	1.76	29	111	Test d seque 1Sea
5506	1	Homo sapiens	Denmark	Caucasian	Male	23	111.3	1.83	33	102.5	Test d seque 1Sea
5491	2	Homo	Denmark	Caucasian	Male	20	106.1	1.79	33	104.5	Test d seque 1Sea

Showing rows 1 to 5 out of a total of 9 example rows

Parameters

Template:

Study:

[← Previous](#) [Next →](#)



Phenotype database

Hello admin! ? Profile Logout

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Match data

Choose type / Upload file / **Match data** / Validation / Store data

Match the rows and columns from your file
Below you can specify where and how to store the data you provided.

Match Clear mapping

Name	Start group	Species	Country (humans)
(Don't import)	(Don't import)	(Don't import)	(Don't import)
5493	1	Homo sapiens	Denmark
5502	1	Homo sapiens	Denmark
5506	1	Homo sapiens	Denmark
5491	2	Homo sapiens	Denmark
5492	2	Homo sapiens	Denmark

Showing rows 1 to 5 out of a total of 9 example rows

Previous Validate Import

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Phenotype database

Hello admin! ? Profile Logout

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Match data

Choose type / Upload file / **Match data** / Validation / Store data

Match the rows and columns from your file
Below you can specify where and how to store the data you provided.

Match Clear mapping

Name	Start group	Species	Country (humans)
name	Start group	species	Country
5493	1	Homo sapiens	Denmark
5502	1	Homo sapiens	Denmark
5506	1	Homo sapiens	Denmark
5491	2	Homo sapiens	Denmark
5492	2	Homo sapiens	Denmark

Showing rows 1 to 5 out of a total of 9 example rows

Previous Validate Import

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Phenotype database

Hello admin! [?](#) [Profile](#) [Logout](#)

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Validation results [Choose type](#) / [Upload file](#) / [Match data](#) / **Validation** / [Store data](#)

✔ **Your data looks OK**
Your data was validated and no errors were found. You can safely continue importing this data.

[Back to upload](#) [Back to mapping](#) [Import](#)

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Phenotype database

Hello admin! [?](#) [Profile](#) [Logout](#)

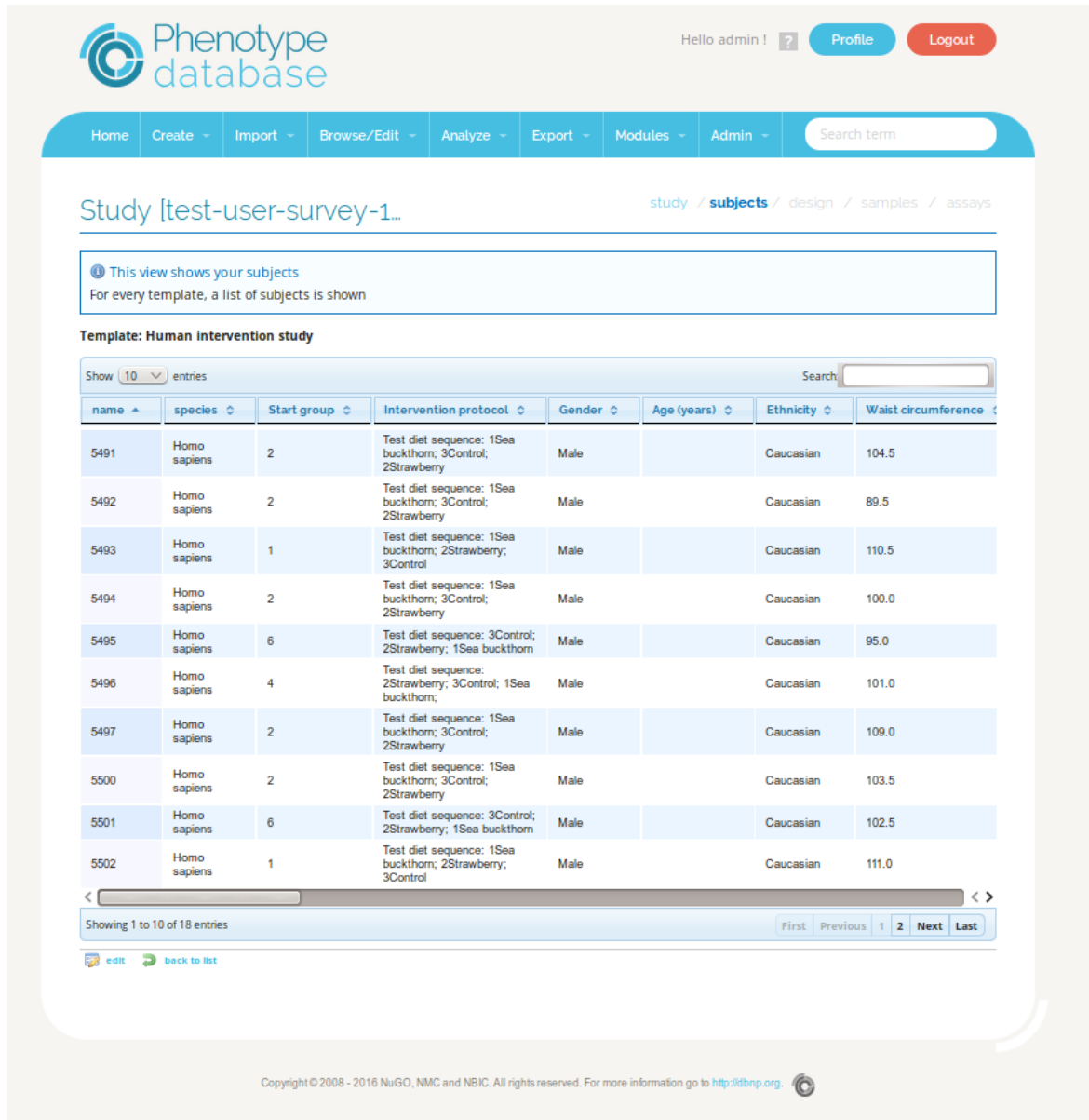
Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Store data [Choose type](#) / [Upload file](#) / [Match data](#) / [Validation](#) / **Store data**

✔ **Data imported successfully**
18 line(s) from your file have been successfully imported and no errors were found.

You can choose to: [view subjects for study User survey - test study](#), [edit subjects for study User survey - test study](#) or [restart the importer](#).

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The screenshot shows the Phenotype Database interface. At the top, there is a navigation bar with 'Home', 'Create', 'Import', 'Browse/Edit', 'Analyze', 'Export', 'Modules', and 'Admin'. A search bar is also present. The main content area displays 'Study [test-user-survey-1...' with a breadcrumb trail: 'study / subjects / design / samples / assays'. A message box states: 'This view shows your subjects. For every template, a list of subjects is shown.' Below this, the 'Template: Human intervention study' is selected. A table lists 18 subjects with columns for name, species, start group, intervention protocol, gender, age, ethnicity, and waist circumference. The table shows subjects 5491 through 5502. At the bottom, there are 'edit' and 'back to list' buttons, and a copyright notice: 'Copyright © 2008 - 2016 NuGO, NMC and NBIC. All rights reserved. For more information go to http://dbnp.org.'

name	species	Start group	Intervention protocol	Gender	Age (years)	Ethnicity	Waist circumference
5491	Homo sapiens	2	Test diet sequence: 1Sea buckthorn; 3Control; 2Strawberry	Male		Caucasian	104.5
5492	Homo sapiens	2	Test diet sequence: 1Sea buckthorn; 3Control; 2Strawberry	Male		Caucasian	89.5
5493	Homo sapiens	1	Test diet sequence: 1Sea buckthorn; 2Strawberry; 3Control	Male		Caucasian	110.5
5494	Homo sapiens	2	Test diet sequence: 1Sea buckthorn; 3Control; 2Strawberry	Male		Caucasian	100.0
5495	Homo sapiens	6	Test diet sequence: 3Control; 2Strawberry; 1Sea buckthorn	Male		Caucasian	95.0
5496	Homo sapiens	4	Test diet sequence: 2Strawberry; 3Control; 1Sea buckthorn;	Male		Caucasian	101.0
5497	Homo sapiens	2	Test diet sequence: 1Sea buckthorn; 3Control; 2Strawberry	Male		Caucasian	109.0
5500	Homo sapiens	2	Test diet sequence: 1Sea buckthorn; 3Control; 2Strawberry	Male		Caucasian	103.5
5501	Homo sapiens	6	Test diet sequence: 3Control; 2Strawberry; 1Sea buckthorn	Male		Caucasian	102.5
5502	Homo sapiens	1	Test diet sequence: 1Sea buckthorn; 2Strawberry; 3Control	Male		Caucasian	111.0

These images show the process to upload subjects to the Phenotype Database (dbNP)

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

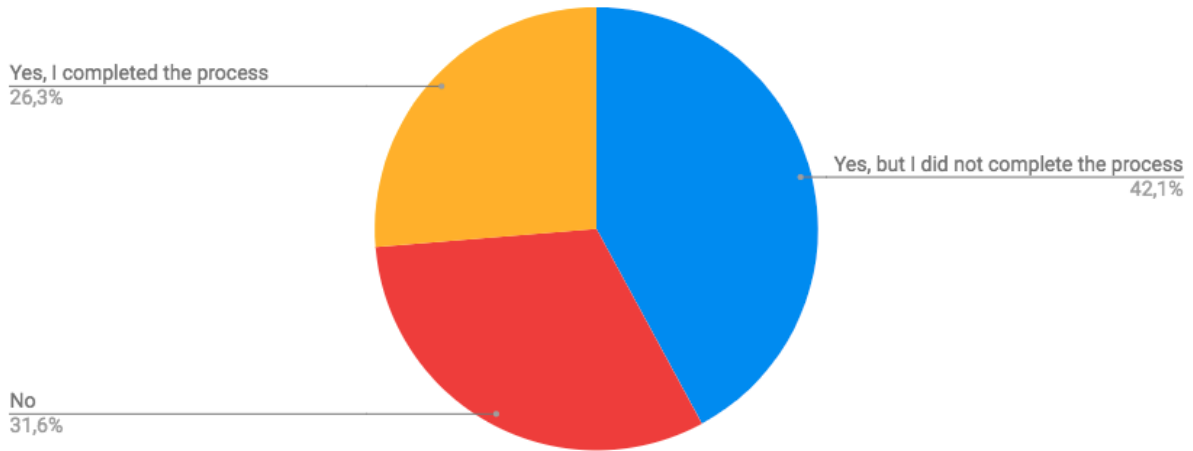
How easy or difficult is it to upload subject to the Phenotype Database (dbNP)?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

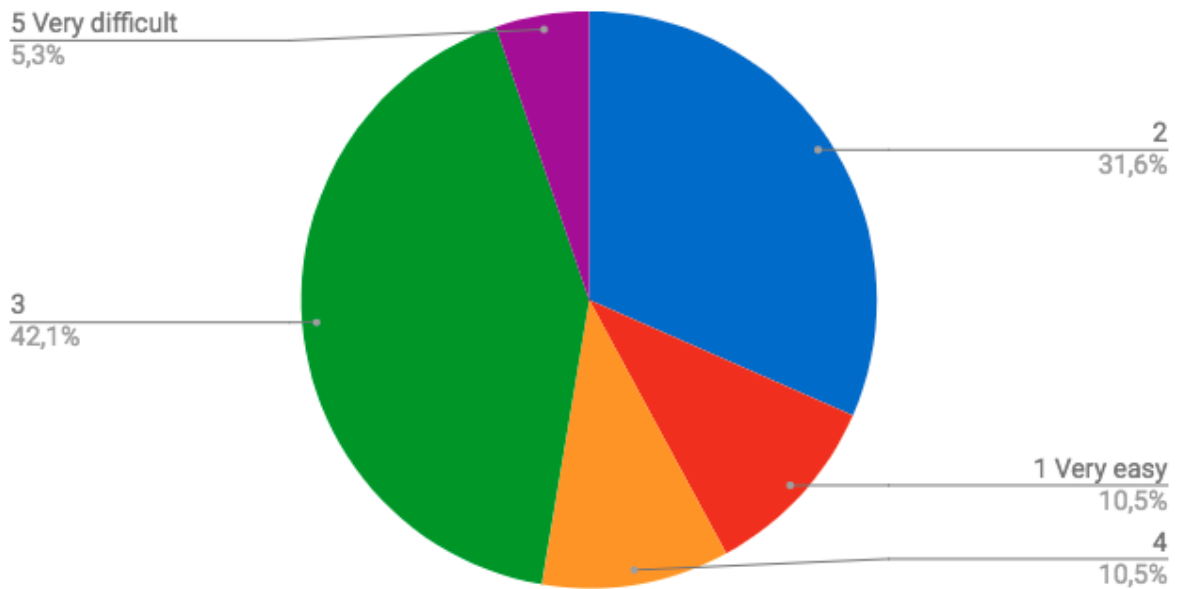


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Phenotype Database (dbNP) - upload subjects - Have you used this functionality

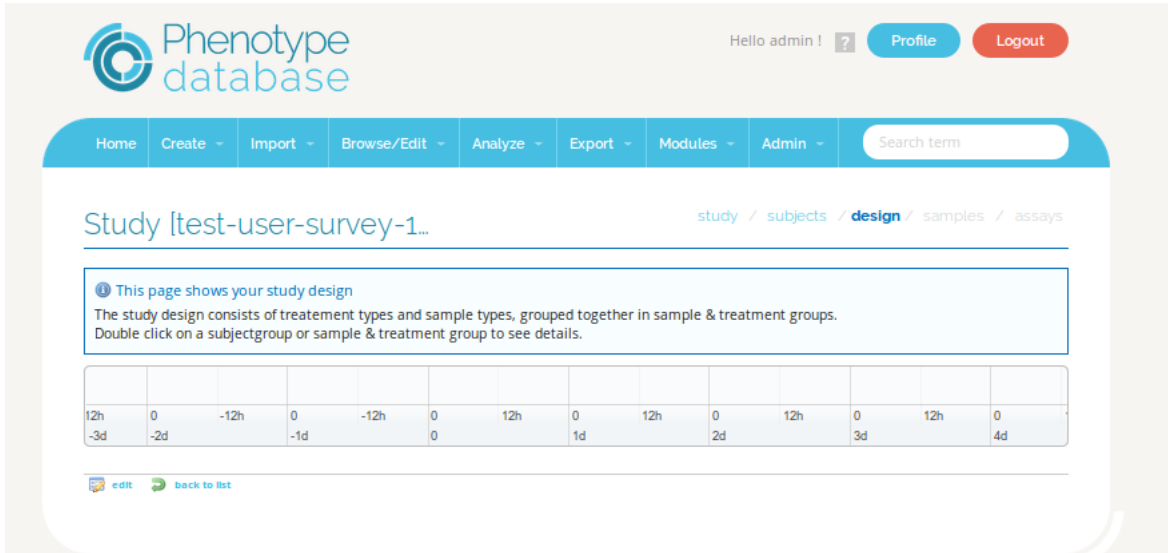


Phenotype Database (dbNP) - upload subjects - How easy or difficult did you find it?





PHENOTYPE DATABASE (DBNP) – STUDY DESIGN, ADD SUBJECT GROUPS



Phenotype database Hello admin! [?](#) [Profile](#) [Logout](#)

Home Create Import Browse/Edit Analyze Export Modules Admin

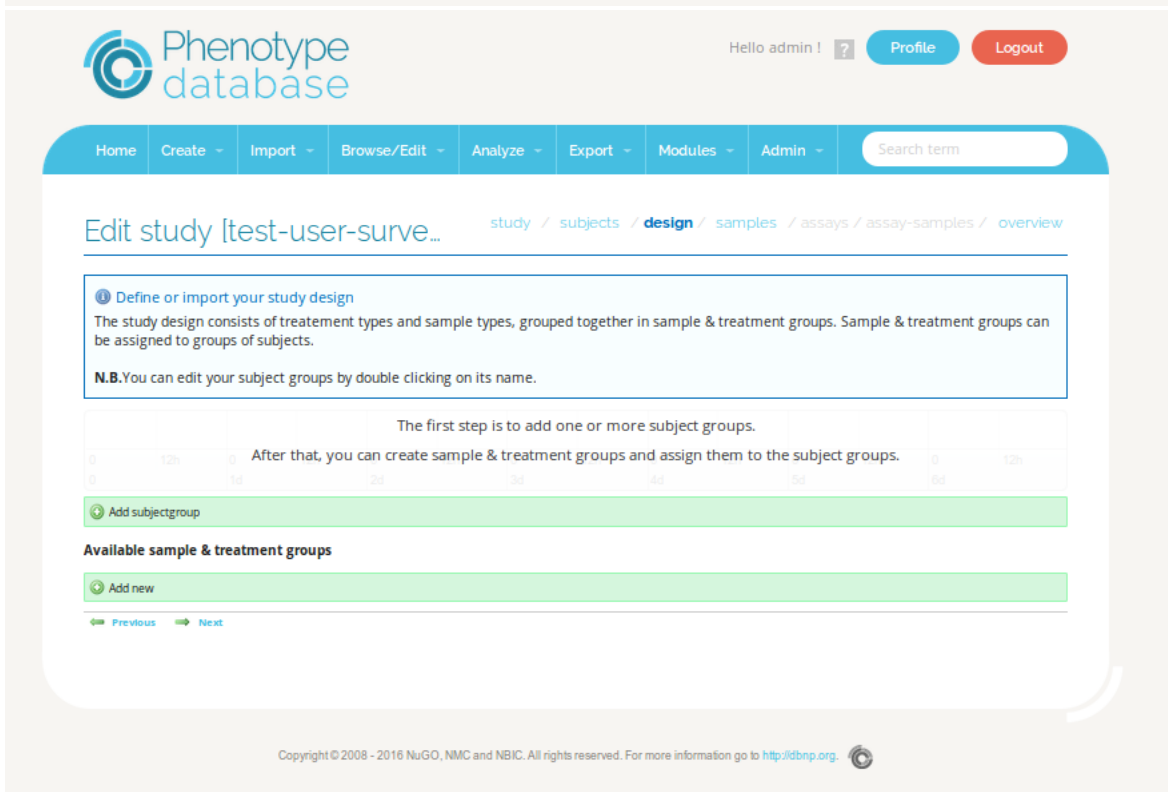
Study [test-user-survey-1...] study / subjects / **design** / samples / assays

1 This page shows your study design
The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Double click on a subjectgroup or sample & treatment group to see details.

12h	0	-12h	0	-12h	0	12h	0	12h	0	12h	0	12h	0
-3d	-2d	-1d	0	1d	2d	3d	4d						

[edit](#) [back to list](#)

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Phenotype database Hello admin! [?](#) [Profile](#) [Logout](#)

Home Create Import Browse/Edit Analyze Export Modules Admin

Edit study [test-user-surve...] study / subjects / **design** / samples / assays / assay-samples / overview

1 Define or import your study design
The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be assigned to groups of subjects.
N.B. You can edit your subject groups by double clicking on its name.

The first step is to add one or more subject groups.
After that, you can create sample & treatment groups and assign them to the subject groups.

0	12h	0	0	1d	2d	3d	4d	5d	6d	0	12h
---	-----	---	---	----	----	----	----	----	----	---	-----

[Add subjectgroup](#)

Available sample & treatment groups

[Add new](#)

[Previous](#) [Next](#)

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Add subjectgroup ✕

[Edit the details of the subjectgroup](#)
Check the subjects that are in this group

Name:

Show 100 entries Search:

	Name	Template	Species
<input type="checkbox"/>	5491	Human intervention study	Homo sapiens
<input type="checkbox"/>	5492	Human intervention study	Homo sapiens
<input checked="" type="checkbox"/>	5493	Human intervention study	Homo sapiens
<input type="checkbox"/>	5494	Human intervention study	Homo sapiens
<input type="checkbox"/>	5495	Human intervention study	Homo sapiens
<input type="checkbox"/>	5496	Human intervention study	Homo sapiens
<input type="checkbox"/>	5497	Human intervention study	Homo sapiens
<input type="checkbox"/>	5500	Human intervention study	Homo sapiens
<input type="checkbox"/>	5501	Human intervention study	Homo sapiens
<input checked="" type="checkbox"/>	5502	Human intervention study	Homo sapiens
<input type="checkbox"/>	5503	Human intervention study	Homo sapiens
<input type="checkbox"/>	5504	Human intervention study	Homo sapiens
<input type="checkbox"/>	5505	Human intervention study	Homo sapiens
<input type="checkbox"/>	5506	Human intervention study	Homo sapiens
<input checked="" type="checkbox"/>	5508	Human intervention study	Homo sapiens
<input type="checkbox"/>	5509	Human intervention study	Homo sapiens
<input type="checkbox"/>	5510	Human intervention study	Homo sapiens
<input type="checkbox"/>	5511	Human intervention study	Homo sapiens

Showing 1 to 18 of 18 entries (3 selected) [clear selection](#) First Previous 1 Next Last



Home Create Import Browse/Edit Analyze Export Modules Admin

Edit study [test-user-surve... [study](#) / [subjects](#) / **[design](#)** / [samples](#) / [assays](#) / [assay-samples](#) / [overview](#)

Define or import your study design

The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be assigned to groups of subjects.

N.B. You can edit your subject groups by double clicking on its name.

Group6															
Group5															
Group4															
Group2															
Group1															
	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h	
	0		1d		2d		3d		4d		5d		6d		

➕ Add subjectgroup

Available sample & treatment groups

➕ Add new

⬅ Previous Next ➡

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Add eventgroup ✕

Edit the details of the sample & treatment group

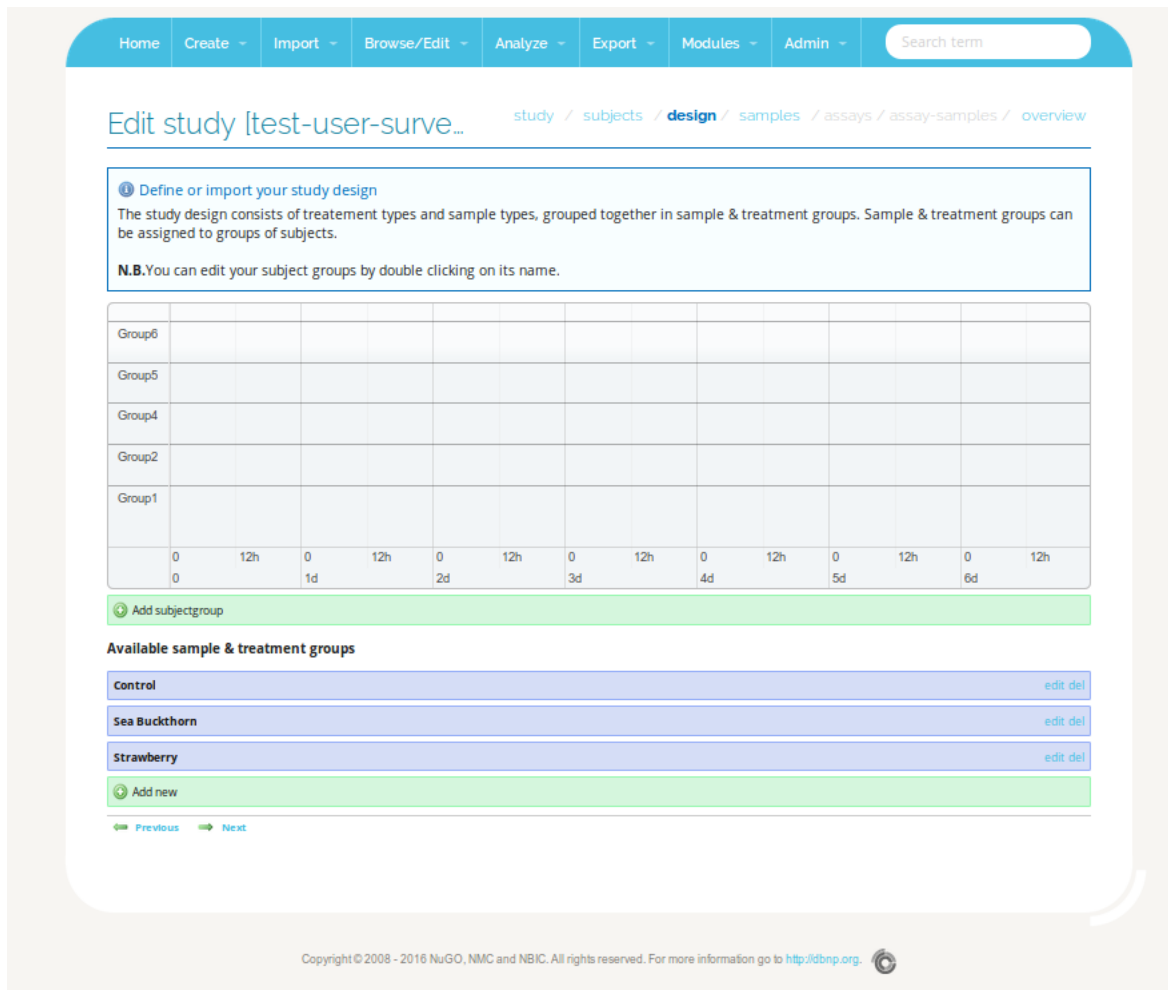
Drag treatment types and sample types into the group. Changes will be saved immediately. However, changes in the name require a click on the 'save name' button.

Please note: changes to this sample & treatment group will affect all instances of the group.

Please note: the timing of treatment and sample types also depends on the timing of the group within the study.

Name:

Available sample & treatment groups

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Edit study ttest-user-surve... study / subjects / design / samples / assays / assay-samples / overview

Define or import your study design
 The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be assigned to groups of subjects.
 N.B. You can edit your subject groups by double clicking on its name.

Group6														
Group5														
Group4														
Group2														
Group1														
	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h
	0		1d		2d		3d		4d		5d		6d	

Add subjectgroup

Available sample & treatment groups

Control	edit del
Sea Buckthorn	edit del
Strawberry	edit del
Add new	

Previous Next

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These images show the process to study design, add subject groups to a study design in the Phenotype Database (dbNP)

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

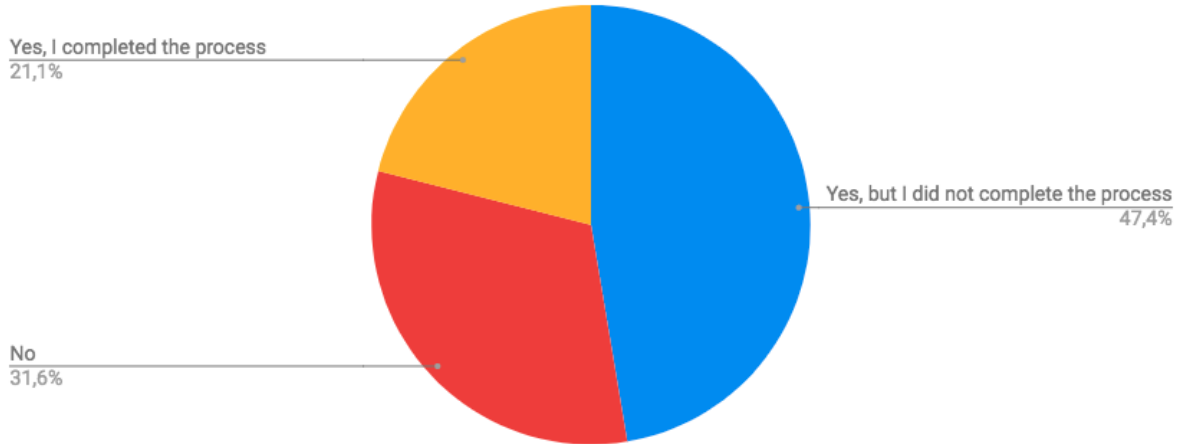
How easy or difficult is it to add subject groups to a study design in the Phenotype Database (dbNP)?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

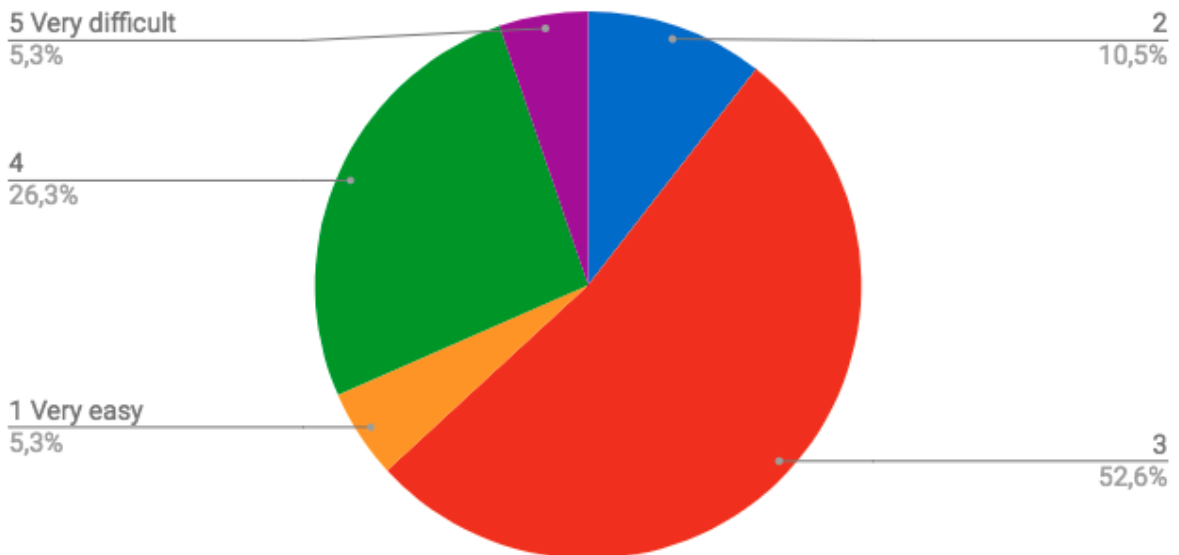


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Phenotype Database (dbNP) - study design, add subject groups - Have you used this functionality



Phenotype Database (dbNP) - study design, add subject groups - How easy or difficult did you find it?





PHENOTYPE DATABASE (DBNP) – ADD TREATMENT TYPES AND SAMPLE TYPES

Edit sample & treatment group

① Edit the details of the sample & treatment group
 Drag treatment types and sample types into the group. Changes will be saved immediately. However, changes in the name require a click on the 'save name' button.
Please note: changes to this sample & treatment group will affect all instances of the group.
Please note: the timing of treatment and sample types also depends on the timing of the group within the study.

Name:

0	12h	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h
0		1d		2d		3d		4d		5d		6d	

Available treatments types **Available sample types**

Add treatment type

New treatment type

Template

- Blood sampling
- Body fluid sampling
- Bone scan prep
- Compound challenge
- Compound intervention
- Diet challenge**
- Diet intervention
- Diet+Physical challenge
- Humanization
- Observation
- Physical activity challenge
- SHAM-UG-procedure
- Screening visit
- Surgery
- Urine Sample
- add / modify..



Add treatment type

New treatment type

Template: Diet challenge

Template for fasting or feeding intervention or challenge.

Name: Ad Libitum

Route: Oral

Event Name (STRING):

Intervention/Challenge: Challenge

Event-Type: Diet

Description:

Diet Description:

Diet Carbohydrate-Level:

Diet Fat-Level:

Diet Protein-Level:

Migration:

Save name Cancel

Edit sample & treatment group

Edit the details of the sample & treatment group

Drag treatment types and sample types into the group. Changes will be saved immediately. However, changes in the name require a click on the 'save name' button.

Please note: changes to this sample & treatment group will affect all instances of the group.

Please note: the timing of treatment and sample types also depends on the timing of the group within the study.

Name: Control

0	12h	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h	0	12h
0		1d		2d		3d		4d		5d		6d			

Available treatments types: Ad Libitum (edit del), Add new

Available sample types: Add new

Save name Cancel

Add sample type

New sample type

Template:

- Body composition
- Body fluid
- Dietary intake
- Excretion
- Genetic variation (SNPs)
- Imaging
- Organ/tissue
- Physiology
- Questionnaire
- Tissue
- add / modify.

Save name Cancel



Add sample type ✕

New sample type

Template ▼ ⓘ
 Name ▶ Anthropometry ⓘ
 Sample Template ▶ Anthropometrics ⓘ

Save name
Cancel

▶ Edit the details of the sample & treatment group

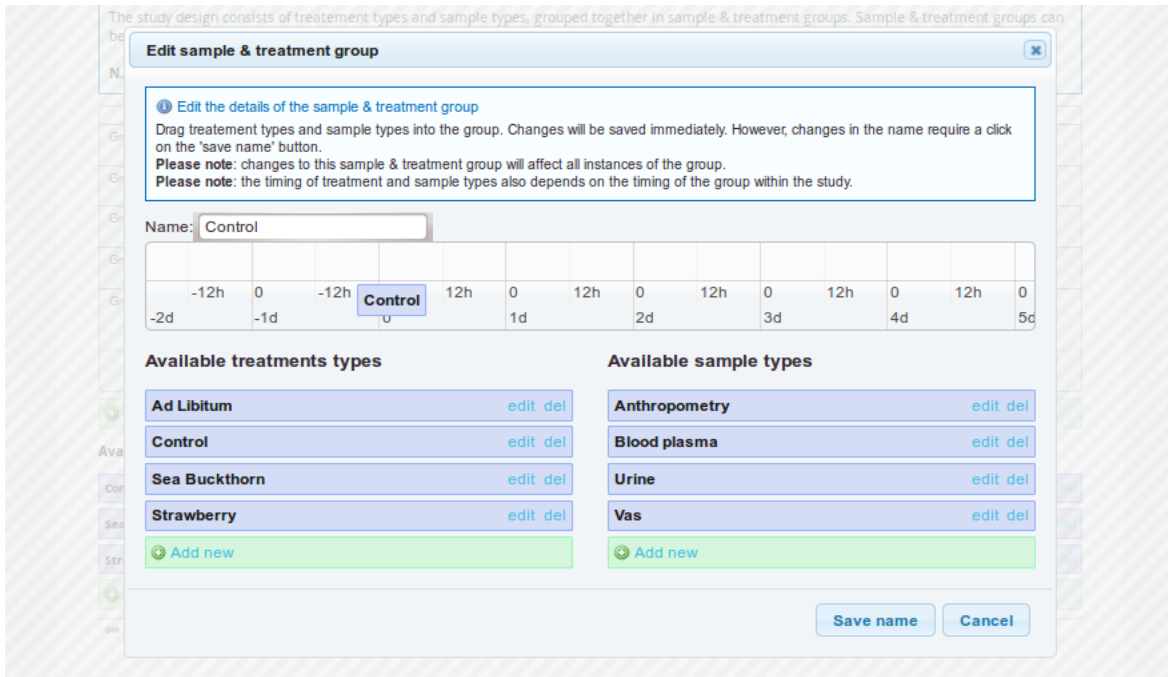
Edit sample type ✕

Edit sample type [Blood plasma]

Template ▼ ⓘ
▶ **Body fluid**
▶ **Template for Body fluids (e.g: blood, lymph).**

Name ▶ Blood plasma ⓘ
 Sample Template ▶ Blood sample ⓘ
 Sampling Name Short ▶ ⓘ
 Sampling-Type ▼ ⓘ
 Body Fluid ▼ ⓘ
 Related Event/Chall. ▼ ⓘ
 Relative Time In Related Event 0s ⓘ
 Sampling Method ▼ ⓘ
 Sample Fraction ▼ ⓘ
 Sampling Description ▶ ⓘ
 Fasting Period 0s
 External Treatment ▶
 Migration ▶

Save name
Cancel



These images show the process to add treatment types and sample types to a study design in the Phenotype Database (dbNP)

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

How easy or difficult is it to add treatment types and sample types to a study design in the Phenotype Database (dbNP)?

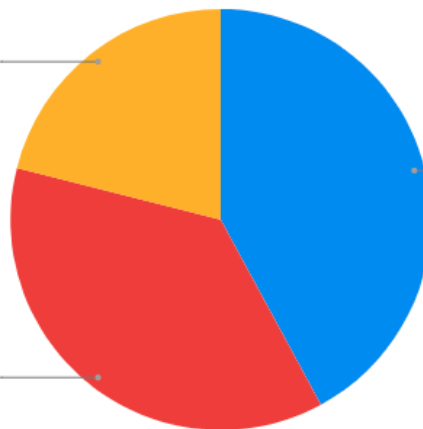
- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Phenotype Database (dbNP) - study design: add treatment types and sample types - Have you used this functionality

Yes, I completed the process
21,1%

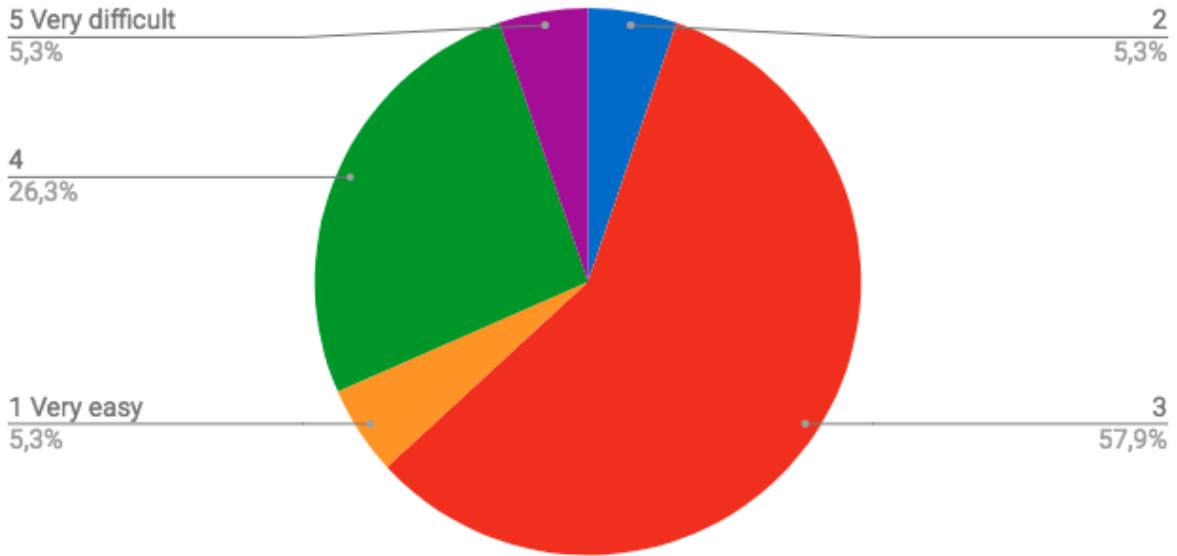
Yes, but I did not complete the process
42,1%

No
36,8%





Phenotype Database (dbNP) - study design: add treatment types and sample types - How easy or difficult did you find it?



PHENOTYPE DATABASE (DBNP) – CREATE STUDY DESIGN FOR A SINGLE SUBJECT GROUP

The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be...

Edit sample & treatment group

ⓘ Edit the details of the sample & treatment group
 Drag treatment types and sample types into the group. Changes will be saved immediately. However, changes in the name require a click on the 'save name' button.
Please note: changes to this sample & treatment group will affect all instances of the group.
Please note: the timing of treatment and sample types also depends on the timing of the group within the study.

Name:

	-12h	0	-12h	Control	12h	0	12h	0	12h	0	12h	0	12h	0
	-2d	-1d		u		1d		2d		3d		4d		5d

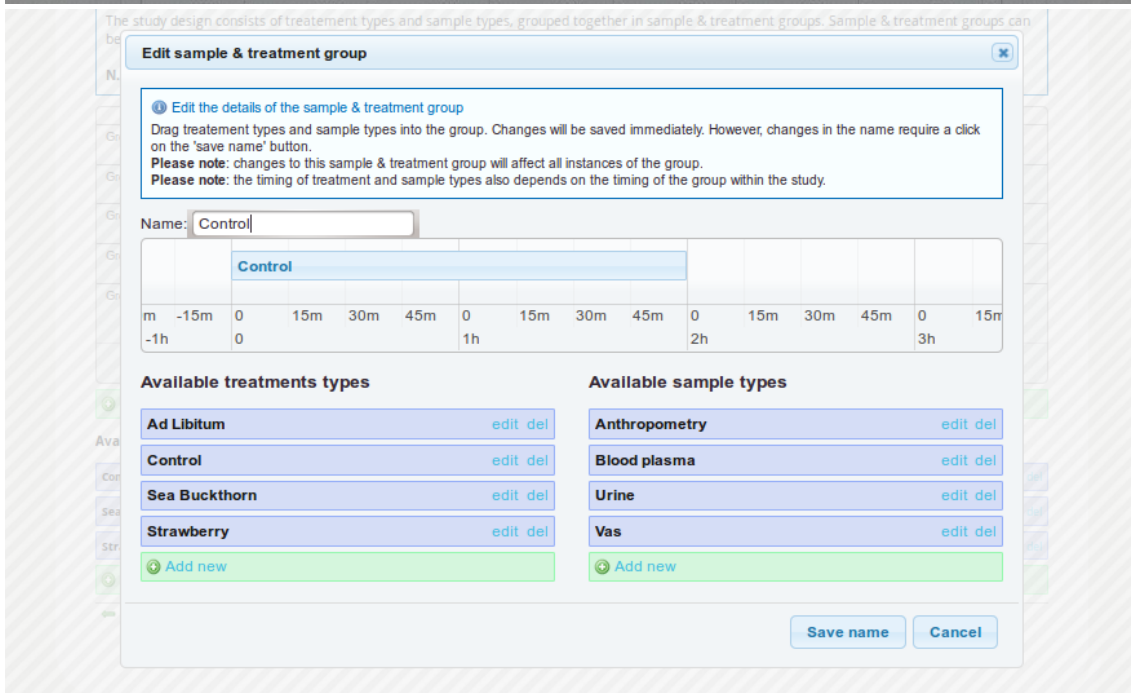
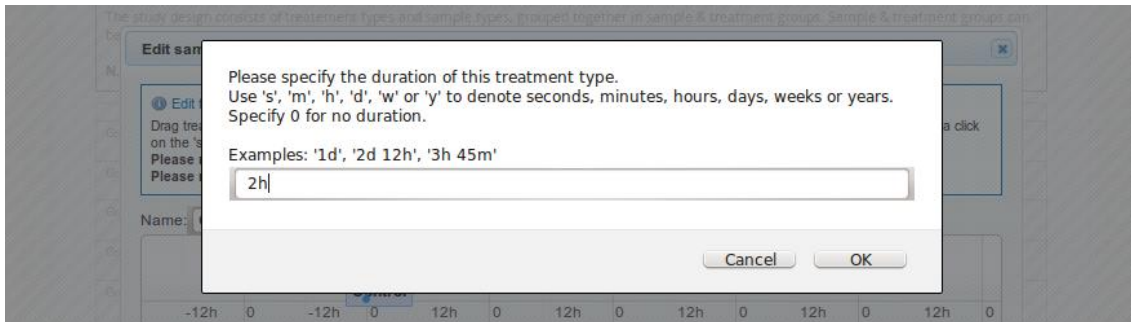
Available treatments types

- Ad Libitum [edit del](#)
- Control [edit del](#)
- Sea Buckthorn [edit del](#)
- Strawberry [edit del](#)
- [Add new](#)

Available sample types

- Anthropometry [edit del](#)
- Blood plasma [edit del](#)
- Urine [edit del](#)
- Vas [edit del](#)
- [Add new](#)

[Save name](#) [Cancel](#)





The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be edited.

Edit sample & treatment group

1 Edit the details of the sample & treatment group
 Drag treatment types and sample types into the group. Changes will be saved immediately. However, changes in the name require a click on the 'save name' button.
Please note: changes to this sample & treatment group will affect all instances of the group.
Please note: the timing of treatment and sample types also depends on the timing of the group within the study.

Name:

Time	Event
0m	Vas, Urine, Blood plasma, Anthropometry
15m	Vas
30m	Blood plasma, Vas
45m	Blood plasma

Available treatments types

- Ad Libitum edit del
- Control edit del
- Sea Buckthorn edit del
- Strawberry edit del
- + Add new

Available sample types

- Anthropometry edit del
- Blood plasma edit del
- Urine edit del
- Vas edit del
- + Add new

Save name Cancel

The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be...

Edit sample & treatment group

1 Edit the details of the sample & treatment group
 Drag treatment types and sample types into the group. Changes will be saved immediately. However, changes in the name require a click on the 'save name' button.
Please note: changes to this sample & treatment group will affect all instances of the group.
Please note: the timing of treatment and sample types also depends on the timing of the group within the study.

Name:

Available treatments types

Ad Libitum	edit del
Control	edit del
Sea Buckthorn	edit del
Strawberry	edit del
Add new	

Available sample types

Anthropometry	edit del
Blood plasma	edit del
Urine	edit del
Vas	edit del
Add new	

These images show the process to create a study design for a single subject group in the Phenotype Database (dbNP)

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

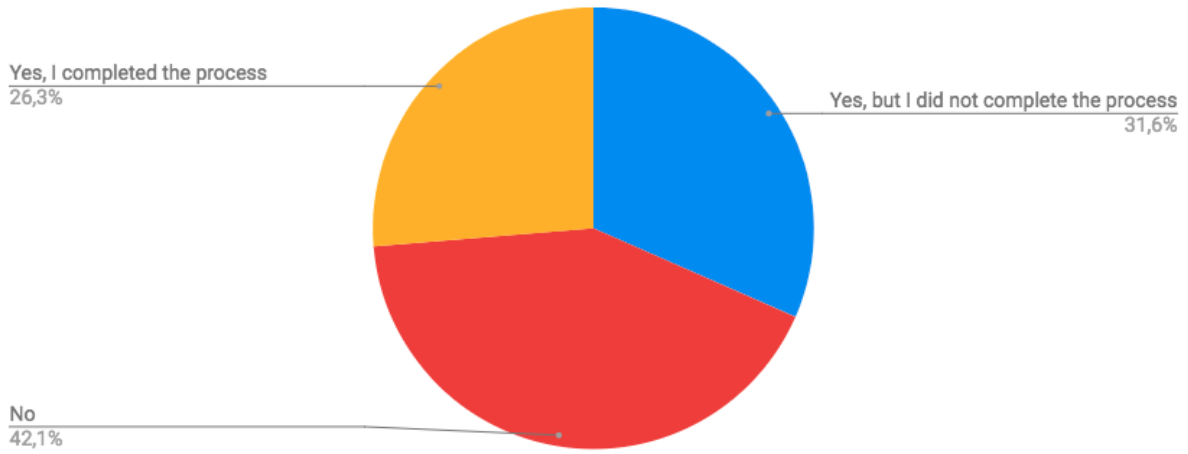
How easy or difficult is it to create a study design for a single subject group in the Phenotype Database (dbNP)?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

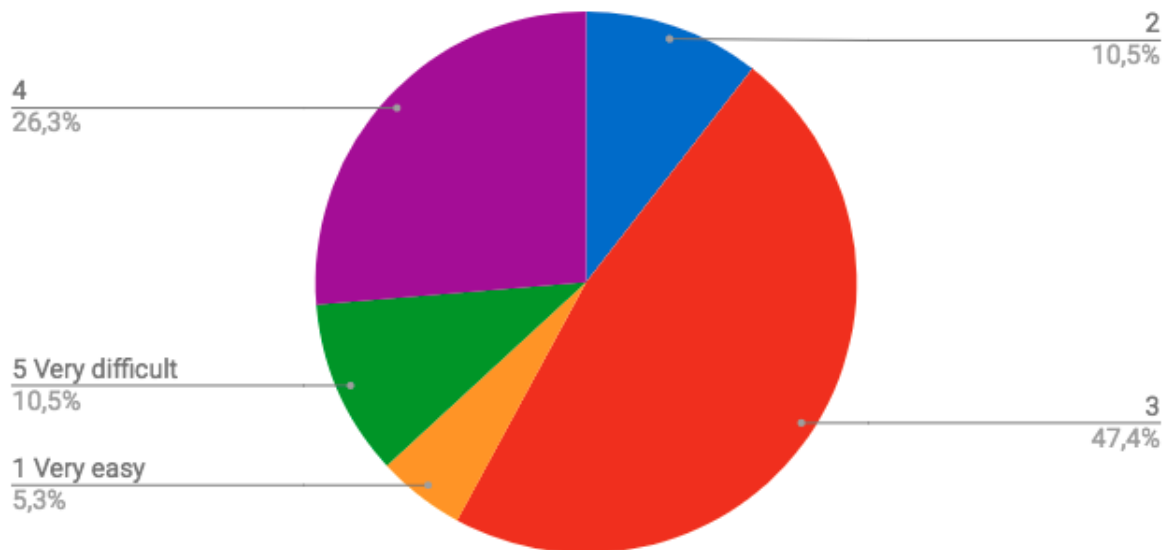


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Phenotype Database (dbNP) - create study design for a single subject group - Have you used this functionality



Phenotype Database (dbNP) - study design: add treatment types and sample types - How easy or difficult did you find it?



PHENOTYPE DATABASE (DBNP) – CREATE OVER-ALL STUDY DESIGN



Home Create Import Browse/Edit Analyze Export Modules Admin

Edit study [test-user-surve...

study / subjects / **design** / samples / assays / assay-samples / overview

1 Define or import your study design

The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be assigned to groups of subjects.

N.B. You can edit your subject groups by double clicking on its name.

Group6														
Group5														
Group4														
Group2														
Group1														
	-2d	-12h	0	-12h	0	12h	0	12h	0	12h	0	12h	0	12h
			-1d		0		1d		2d		3d		4d	

+ Add subjectgroup

Available sample & treatment groups

Control 8x Vas, 6x Blood plasma, 4x Urine...
edit del

Sea Buckthorn 8x Vas, 6x Blood plasma, 4x Urine...
edit del

Strawberry 7x Vas, 6x Blood plasma, 4x Urine...
edit del

+ Add new

[← Previous](#) [Next →](#)

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Home
Create
Import
Browse/Edit
Analyze
Export
Modules
Admin

Search term

Edit study [test-user-surve...]

[study](#) / [subjects](#) / **[design](#)** / [samples](#) / [assays](#) / [assay-samples](#) / [overview](#)

1 Define or import your study design

The study design consists of treatment types and sample types, grouped together in sample & treatment groups. Sample & treatment groups can be assigned to groups of subjects.

N.B. You can edit your subject groups by double clicking on its name.

Group6							
	Control						
Group5							
Group4							
Group2							
Group1							
	0	4h	8h	12h	16h	20h	0
-1d	0						1d

[Add subjectgroup](#)

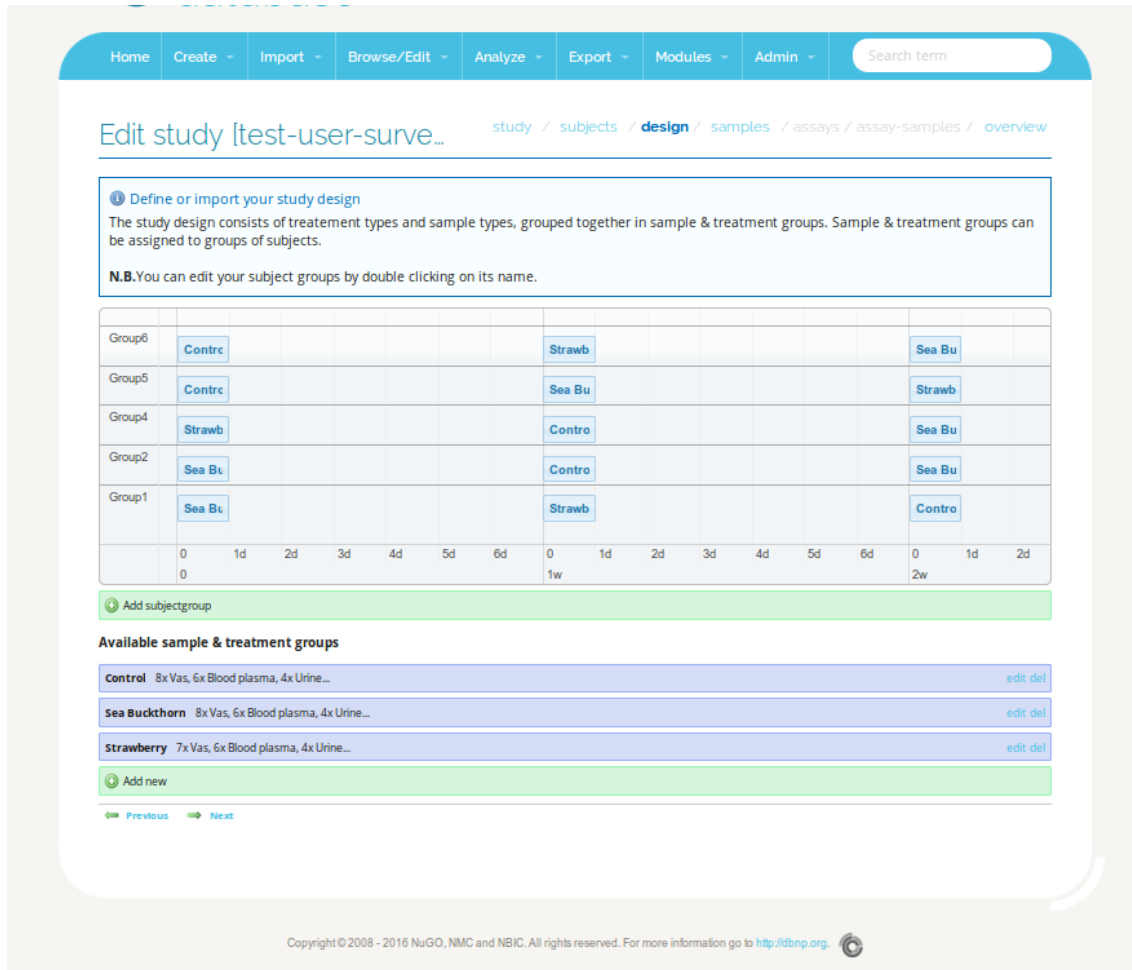
Available sample & treatment groups

Control	8x Vas, 6x Blood plasma, 4x Urine...	edit del
Sea Buckthorn	8x Vas, 6x Blood plasma, 4x Urine...	edit del
Strawberry	7x Vas, 6x Blood plasma, 4x Urine...	edit del

[Add new](#)

[Previous](#) [Next](#)

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These images show the process to create the over-all study design in the Phenotype Database (dbNP)

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

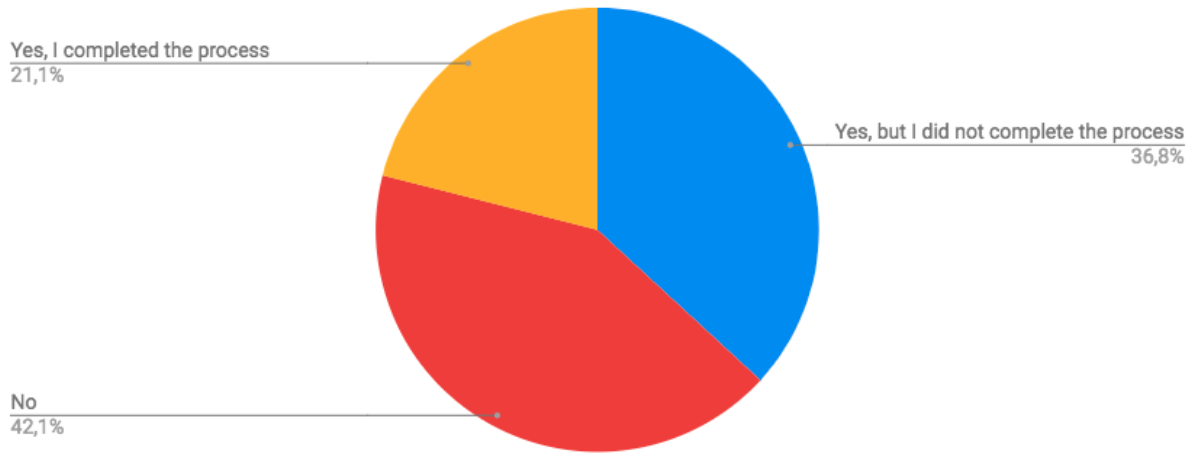
How easy or difficult is it to create the over-all study design in the Phenotype Database (dbNP)?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

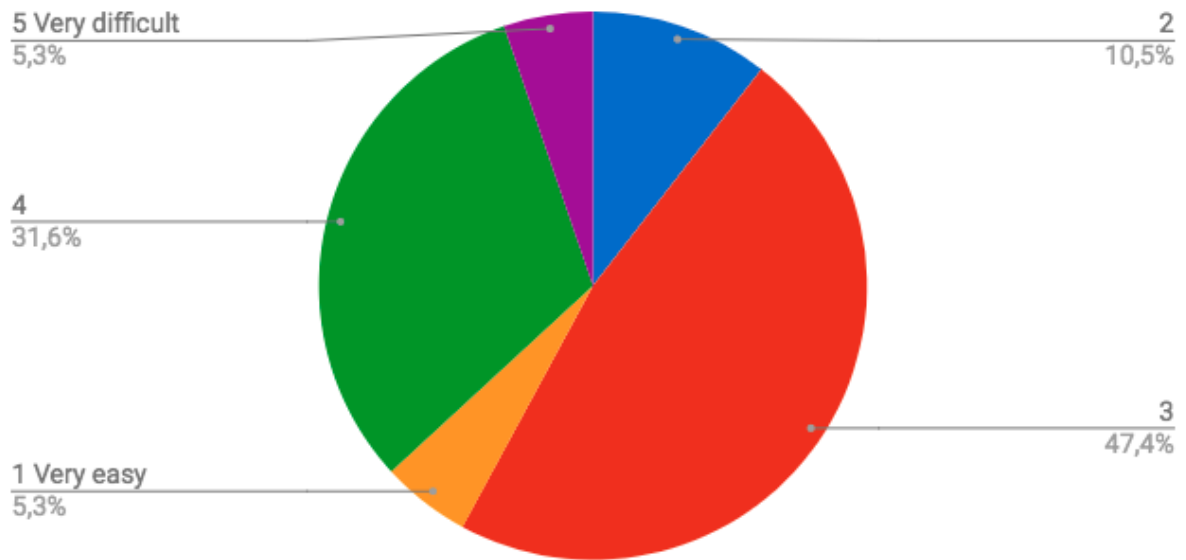


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Phenotype Database (dbNP) - create over-all study design - Have you used this functionality



Phenotype Database (dbNP) - create over-all study design - How easy or difficult did you find it?



PHENOTYPE DATABASE (DBNP) – GENERATE SAMPLES



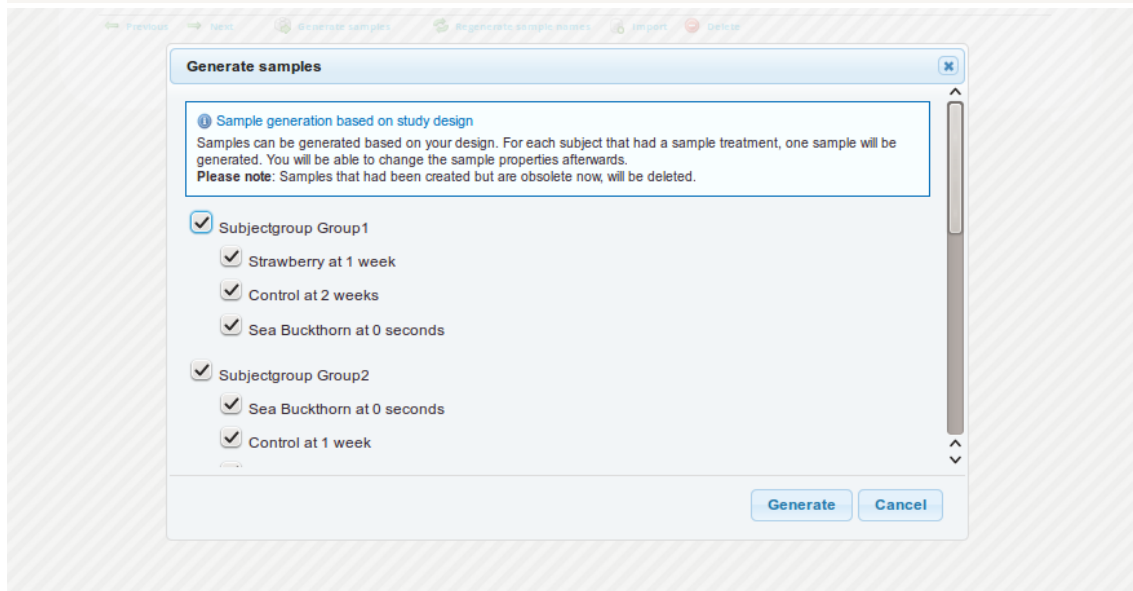

Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Edit study [test-user-surve... study / subjects / design / **samples** / assays / assay-samples / overview

Generate samples
Click 'Generate samples' to initiate the sample generation based on your study design.

Previous Next Generate samples Regenerate sample names Import Delete

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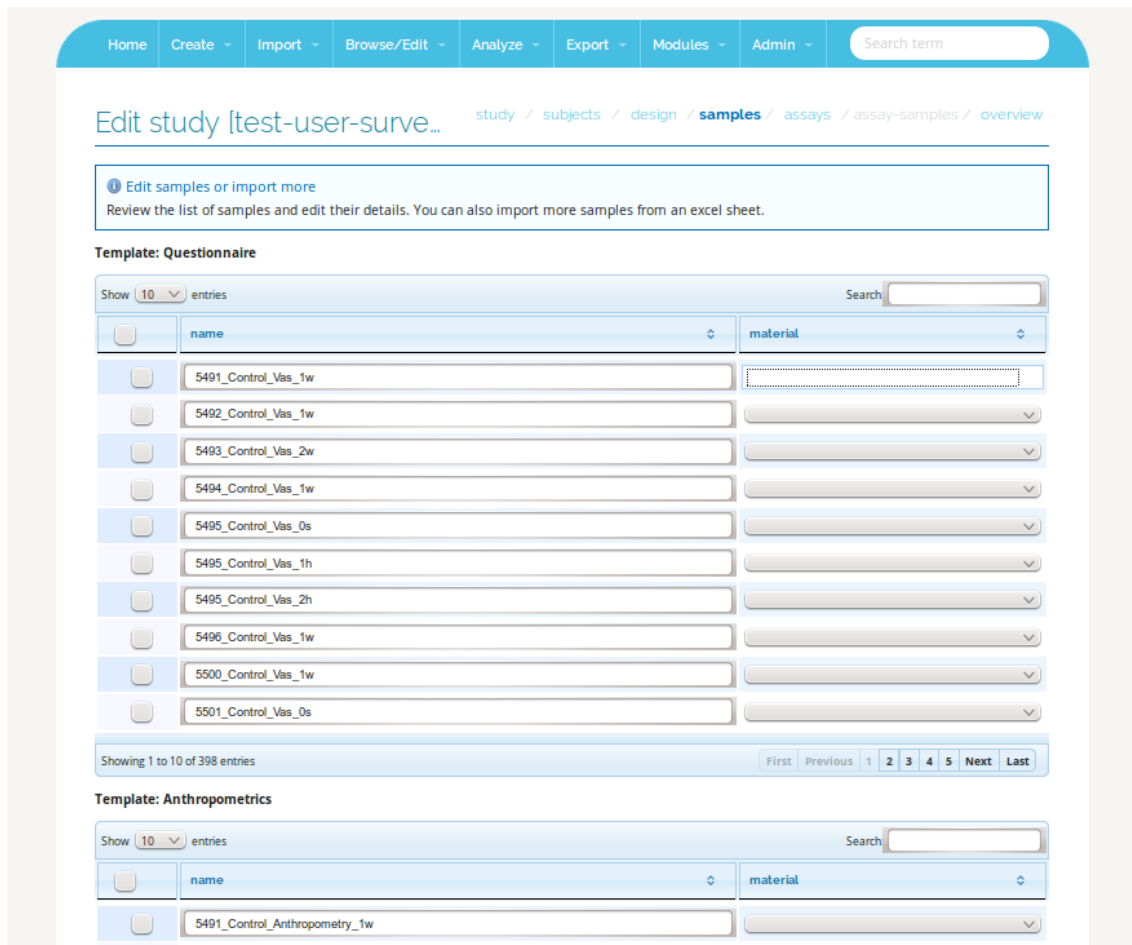
Previous Next Generate samples Regenerate sample names Import Delete

Generate samples

Sample generation based on study design
Samples can be generated based on your design. For each subject that had a sample treatment, one sample will be generated. You will be able to change the sample properties afterwards.
Please note: Samples that had been created but are obsolete now, will be deleted.

- Subjectgroup Group1
 - Strawberry at 1 week
 - Control at 2 weeks
 - Sea Buckthorn at 0 seconds
- Subjectgroup Group2
 - Sea Buckthorn at 0 seconds
 - Control at 1 week

Generate Cancel



Home Create Import Browse/Edit Analyze Export Modules Admin Search term

Edit study [test-user-surve... study / subjects / design / **samples** / assays / assay-samples / overview

1 Edit samples or import more
Review the list of samples and edit their details. You can also import more samples from an excel sheet.

Template: Questionnaire

Show 10 entries Search

<input type="checkbox"/>	name	material
<input type="checkbox"/>	5491_Control_Vas_1w	
<input type="checkbox"/>	5492_Control_Vas_1w	
<input type="checkbox"/>	5493_Control_Vas_2w	
<input type="checkbox"/>	5494_Control_Vas_1w	
<input type="checkbox"/>	5495_Control_Vas_0s	
<input type="checkbox"/>	5495_Control_Vas_1h	
<input type="checkbox"/>	5495_Control_Vas_2h	
<input type="checkbox"/>	5496_Control_Vas_1w	
<input type="checkbox"/>	5500_Control_Vas_1w	
<input type="checkbox"/>	5501_Control_Vas_0s	

Showing 1 to 10 of 398 entries First Previous 1 2 3 4 5 Next Last

Template: Anthropometrics

Show 10 entries Search

<input type="checkbox"/>	name	material
<input type="checkbox"/>	5491_Control_Anthropometry_1w	

These images show the process to generate samples in the Phenotype Database (dbNP)

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

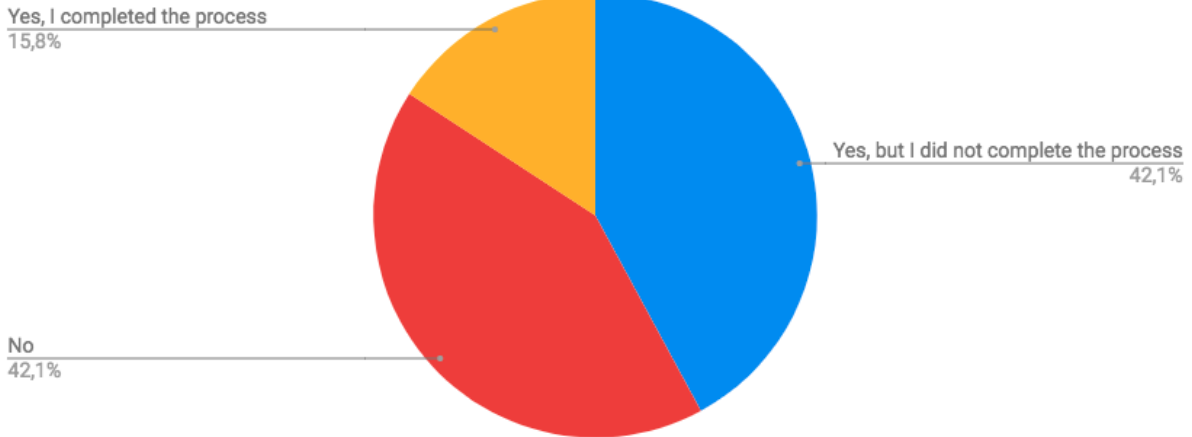
How easy or difficult is it to generate samples in the Phenotype Database (dbNP)?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

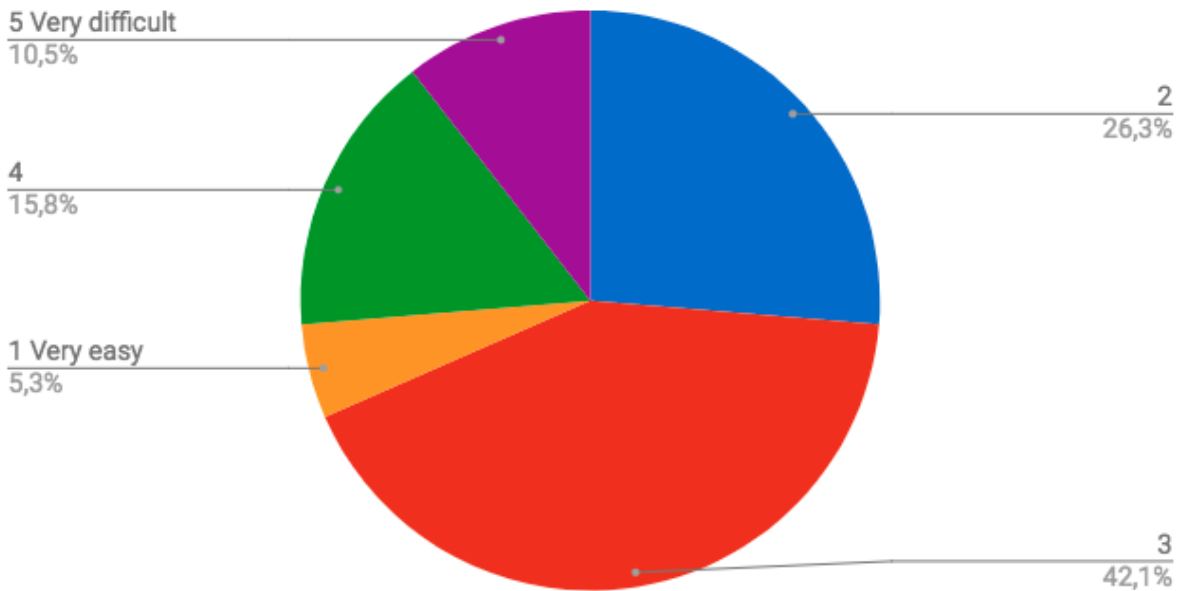


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Phenotype Database (dbNP) - generate samples - Have you used this functionality



Phenotype Database (dbNP) - generate samples - How easy or difficult did you find it?



PHENOTYPE DATABASE (DBNP) – DO YOU HAVE ANY COMMENTS ON THE PHENOTYPE DATABASE (DBNP) SERVER?

It is difficult to understand the meaning of the concepts "sample treatment", "treatments types" and "sample types". But, above all, to apprehend all the differences between many templates.

Quite lengthy process in comparison to DataShield and Mica. However more visual than a harmonization script.

we not used the dbnp and have no experience with it

It's about 1 and 1/2 years ago that I used the database. It was during the meeting in Barcelona. So I don't



remember very well.
no
Provide easier and clearer templates. Sometimes treatment and sample templates can overlap. To use dbNP, all its instances should have the same synchronized templates and a clear user-guide on how to create the study design.
No

DASH-IN TEXT BASED FEDERATED ANALYSES – INSTALLING AND RUNNING THE OPAL/DATASHIELD R PACKAGES

```
R version 3.2.4 (2016-03-10) -- "Very Secure Dishes"
Copyright (C) 2016 The R Foundation for Statistical Computing
Platform: x86_64-apple-darwin13.4.0 (64-bit)

R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.

Natural language support but running in an English locale

R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> |
```

In case you want to launch a federated analysis you need to have already running the R environment. Within an active R session the installation of the DataShield client (the piece of software that enables to access the data and run the analysis) is performed in a console environment issuing the commands listed above.

Have you installed the functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

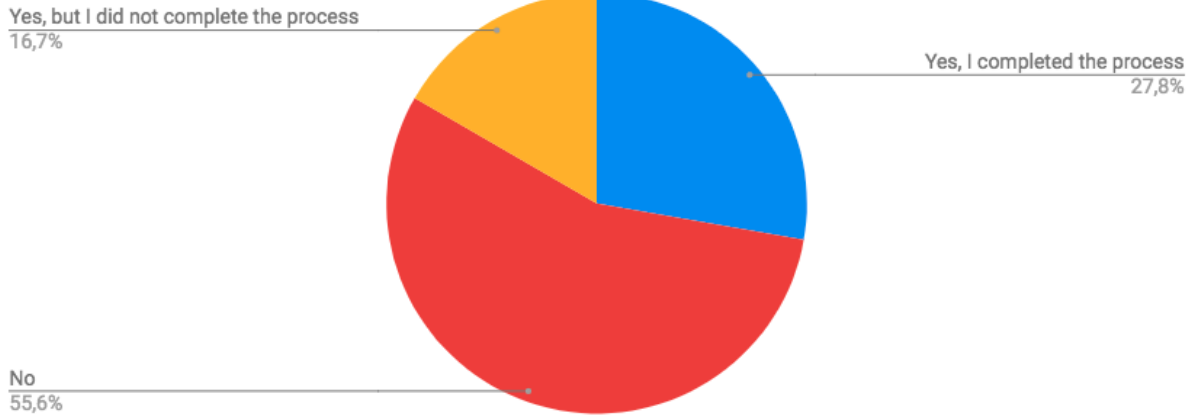
How easy or difficult is the installation of R and the Datashield pre-requisites?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

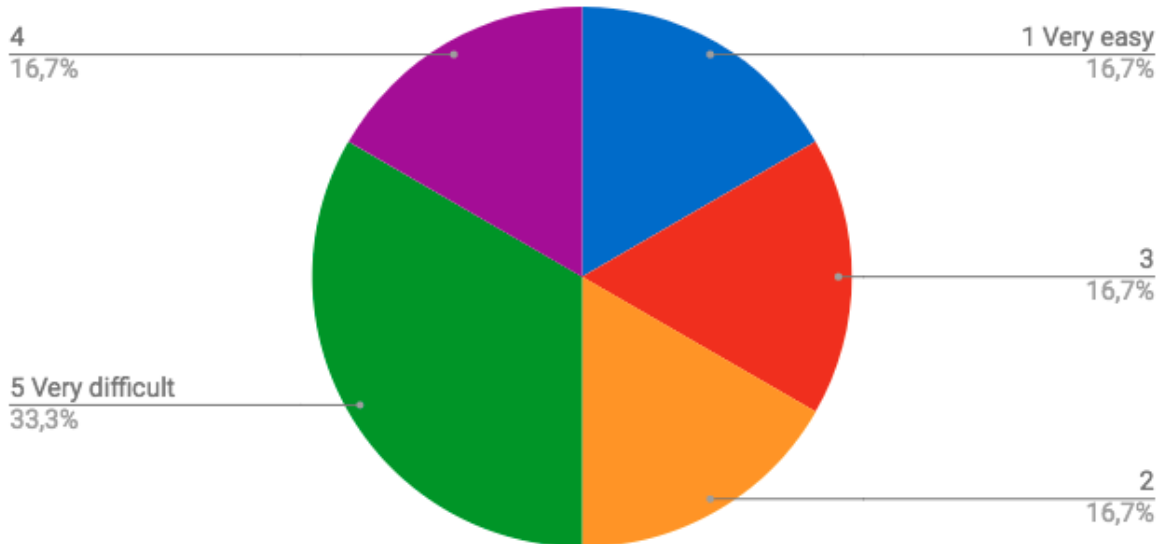


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Dash-in text based federated analyses - Installing and running the Opal/DataShield R packages - Have you used this functionality



Dash-in text based federated analyses - Installing and running the Opal/DataShield R packages - How easy or difficult did you find it?



DASH-IN TEXT BASED FEDERATED ANALYSES – RUN A FEDERATED ANALYSIS

```
> # load libraries
library(opal)
library(dsBaseClient)
library(dsStatsClient)
library(dsGraphicsClient)
library(dsModellingClient)

# load the login file
my_login<-read.table('../logins.txt', sep=" ", header=TRUE)

# log in to the remote servers
opals <- datashield.login(logins=my_login, assign=TRUE, symbol = 'D')

# analyse the outcome variable DIS_DIAB (diabetes status)
# and the covariates PM_BMI_CONTINUOUS (continuous BMI), LAB_HDL (HDL cholesterol)
# and GENDER (gender), with an interaction between the latter two.
ds.glm(formula=D$DIS_DIAB ~ D$PM_BMI_CONTINUOUS + D$LAB_HDL * D$GENDER, family='binomial')
```

```
Iteration 1...
CURRENT DEVIANCE:      5772.52971970323
Iteration 2...
CURRENT DEVIANCE:      1316.4695853419
....
SUMMARY OF MODEL STATE after iteration 9
Current deviance 534.369101004808 on 4159 degrees of freedom
Convergence criterion TRUE (1.38346480863211e-12)
beta: -6.90641416691308 0.142256253558181 -0.96744073975236 -1.40945273343361 0.646007073975006
Information matrix overall:
              (Intercept) PM_BMI_CONTINUOUS  LAB_HDL  GENDER1  LAB_HDL:GENDER1
(Intercept)    52.47803      1624.8945    71.78440   16.77192     25.91140
PM_BMI_CONTINUOUS 1624.89450    51515.6450  2204.90085  503.20484    774.18028
LAB_HDL          71.78440     2204.9008   109.48855  25.91140    42.87626
GENDER1         16.77192     503.2048   25.91140  16.77192    25.91140
LAB_HDL:GENDER1  25.91140     774.1803   42.87626  25.91140    42.87626
              [,1]
(Intercept) -3.610710e-10
PM_BMI_CONTINUOUS -9.867819e-09
LAB_HDL        -6.308052e-10
GENDER1        -2.910556e-10
LAB_HDL:GENDER1 -5.339693e-10
Current deviance: 534.369101004808
$formula
[1] "D$DIS_DIAB ~ D$PM_BMI_CONTINUOUS + D$LAB_HDL * D$GENDER"
$family
Family: binomial
Link function: logit

$coefficients
              Estimate Std. Error  z-value  p-value low0.95CI.LP
(Intercept) -6.9064142  1.08980103  -6.3373166  2.338013e-10  -9.04238494
PM_BMI_CONTINUOUS  0.1422563  0.02932171   4.8515676  1.224804e-06   0.08478676
```

To run a federated analysis it is required to enter a set of commands. The bare minimum set of commands are presented in the first screen while the second screen shows the results. The type of analysis involves a Generalized Linear Model regression using the binomial link function.



JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

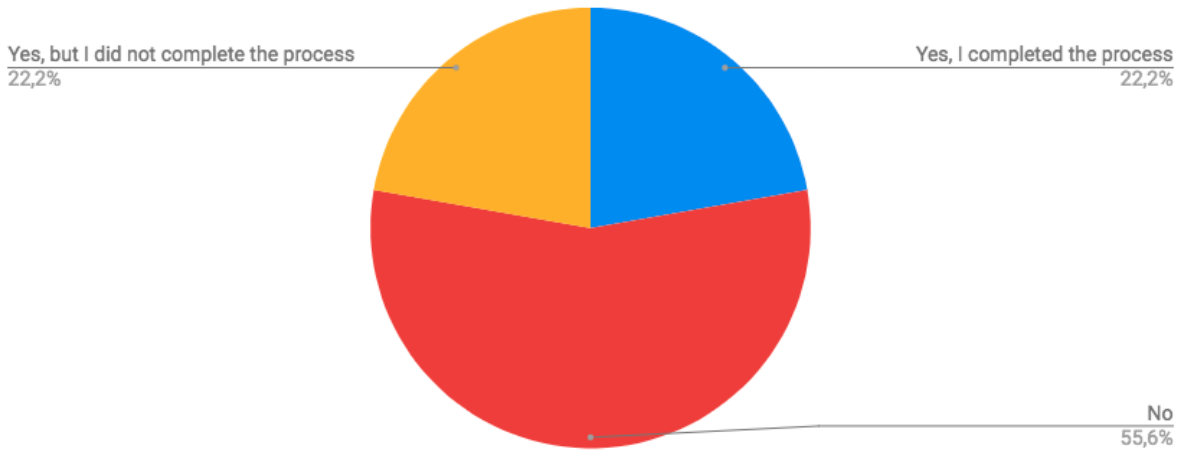
Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

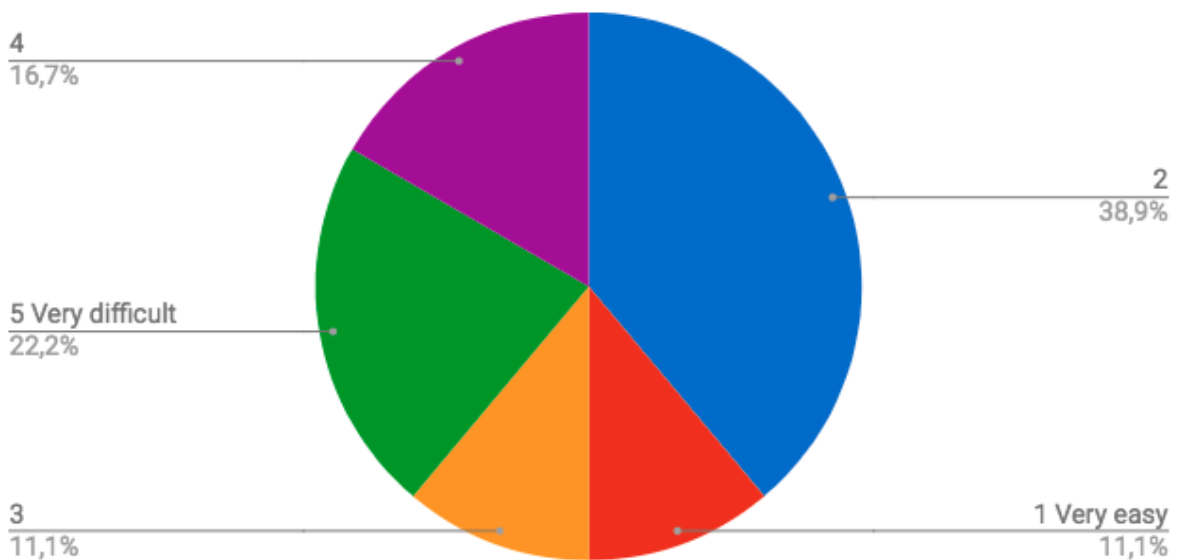
How easy or difficult are the analyses of this kind?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Dash-in text based federated analyses - Run a federated analysis - Have you used this functionality



Dash-in text based federated analyses - Run a federated analysis - How easy or difficult did you find it?





No, the process it perfect like it is now
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a standalone application (like excel) could be considered
Yes, a simple web page allowing to select data and explore results could be considered
Yes, a simple web page allowing to select data and explore results could be considered
unable to install the opal packages on mac osx.
Most of the commands in the previous pages do not work. Impossible for a non-expert to succeed in using these tools (R command lines to connect and to analyze OPAL and DbNP datas).
A combination of both a web application and the normal R procedures is appropriate. Those more familiar with commands and scripts, can do it the traditional way. Others who are used to for example SPSS might like a visual interface.
I'm sorry to say, but I find it weird to be forced to assess the difficulties related to a "functionality" that I have not used at all!
no
Tutorials to connect to PhenotypeDatabase R-Client, OPAL Datashield.
no

DASH-IN – SETTING UP DATA SOURCES



Selection of the Projects into OPAL servers.

Selection of tables from OPAL projects



The screenshot shows the 'Data Servers' configuration page. It includes instructions: (1) Select the desired DBNP / OPAL studies. (2) Press the 'Setup remote data tables' button below. The interface features four data server sections: DEMO, OpalRecas, UniLiege, and TNO. Each section has a dropdown for 'Opal Projects' and a dropdown for 'Data tables'. At the bottom, the 'Phenotype Database Studies' dropdown is open, displaying a list of studies.

Selection of Phenotype database data hosted within DASH-IN-enabled Phenotype Database computers.

This screenshot is similar to the previous one but shows the 'Study Assays' dropdown menu open. The 'Diclofenac' study is selected in the 'Phenotype Database Studies' dropdown. The 'Study Assays' dropdown lists various assays, with 'BodyWeight' selected.

Selection of the study Assays associated to Phenotype Database DASH-IN enabled machines.

Setup remote data tables

Data sources were successfully setup on remote servers! Visit the **Federated Analysis** section in the left side-pane.



JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

After having selected all OPAL and Phenotype Database pressing the button “Setup remote data tables” will prepare the DASH-IN technical infrastructure to make federated analysis on different tables hosted on different server but without moving or disclosing any single information out of each hosting server.

Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

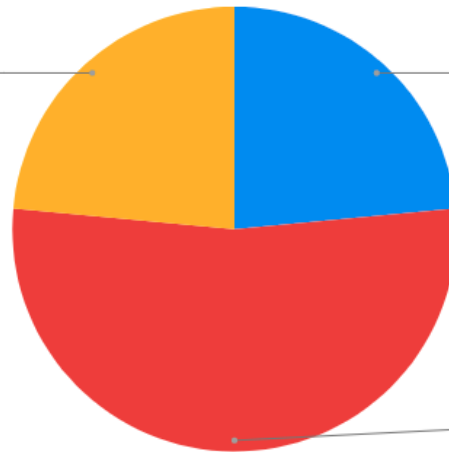
How easy or difficult is it to generate federated analyses?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Dash-in - Setting up data sources - Have you used this functionality

Yes, but I did not complete the process
23,5%

Yes, I completed the process
23,5%

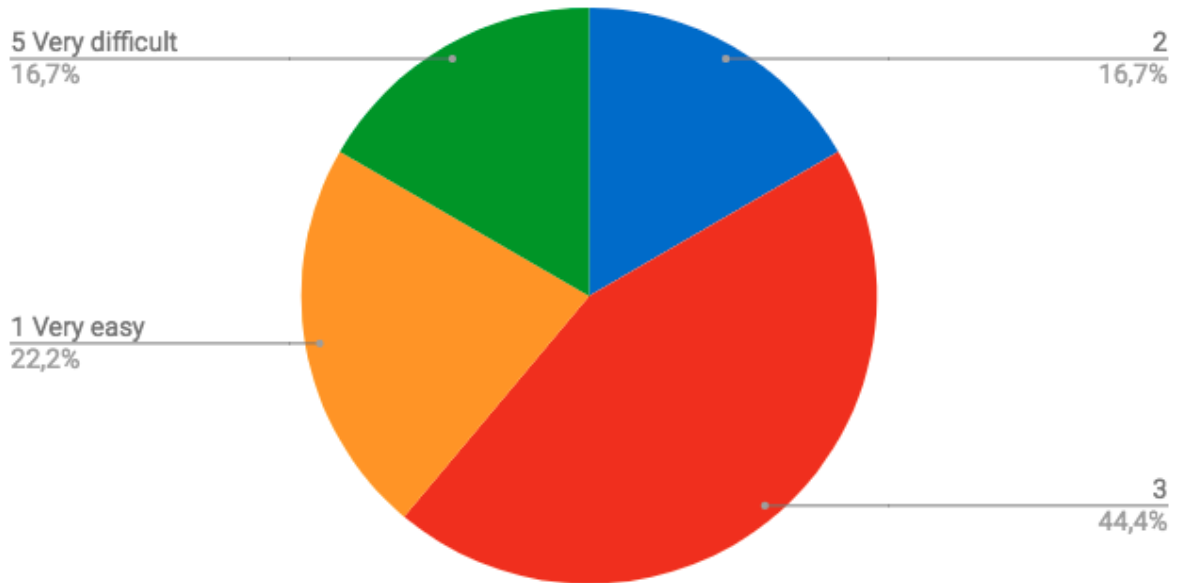


No
52,9%

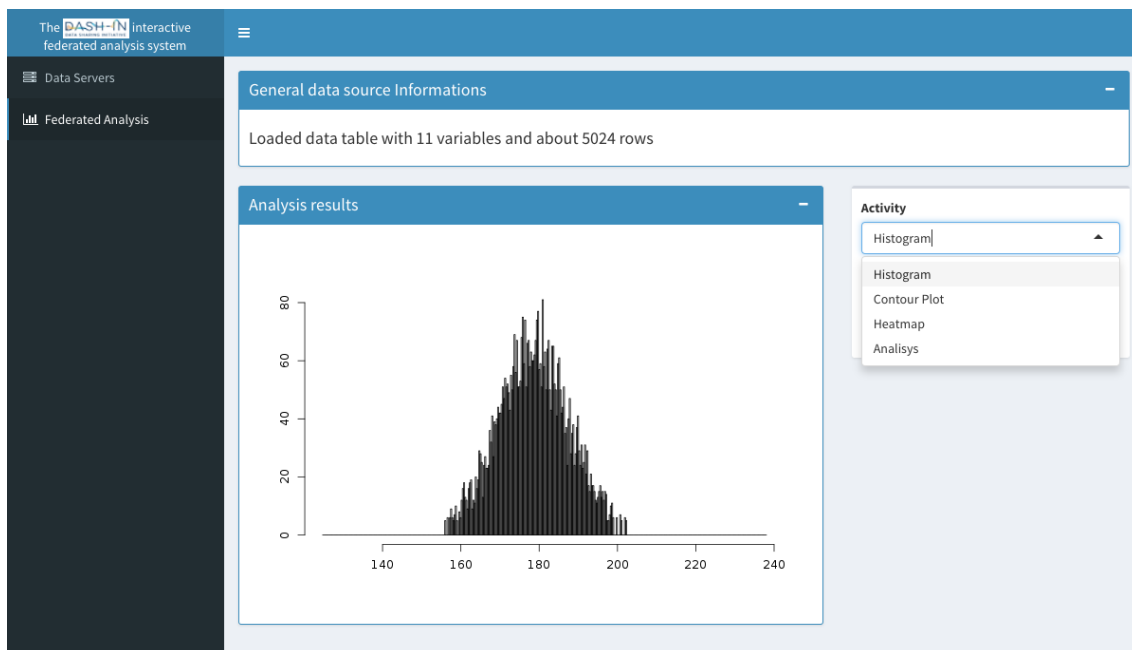


JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

Dash-in - Setting up data sources- How easy or difficult did you find it?



DASH-IN – FEDERATED GRAPHICS

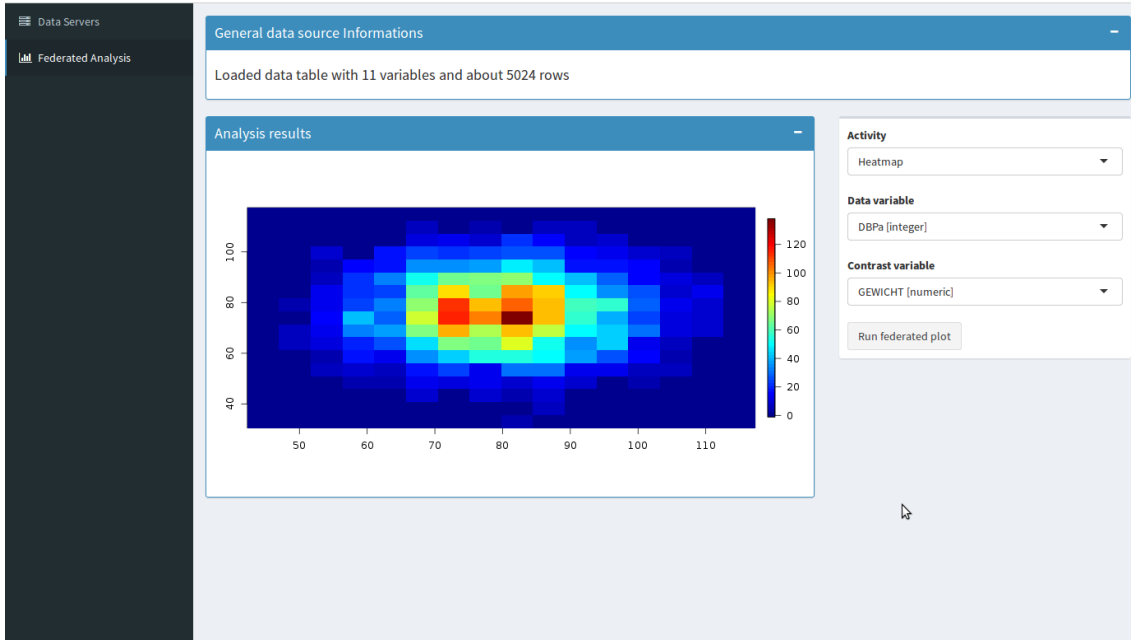


A number of analytical activities from federated data are easily available to the final researcher including **Histograms, Contour Plots, Heatmaps, and Analysis** for phenotype-trait associations (the latter shown in next pages).



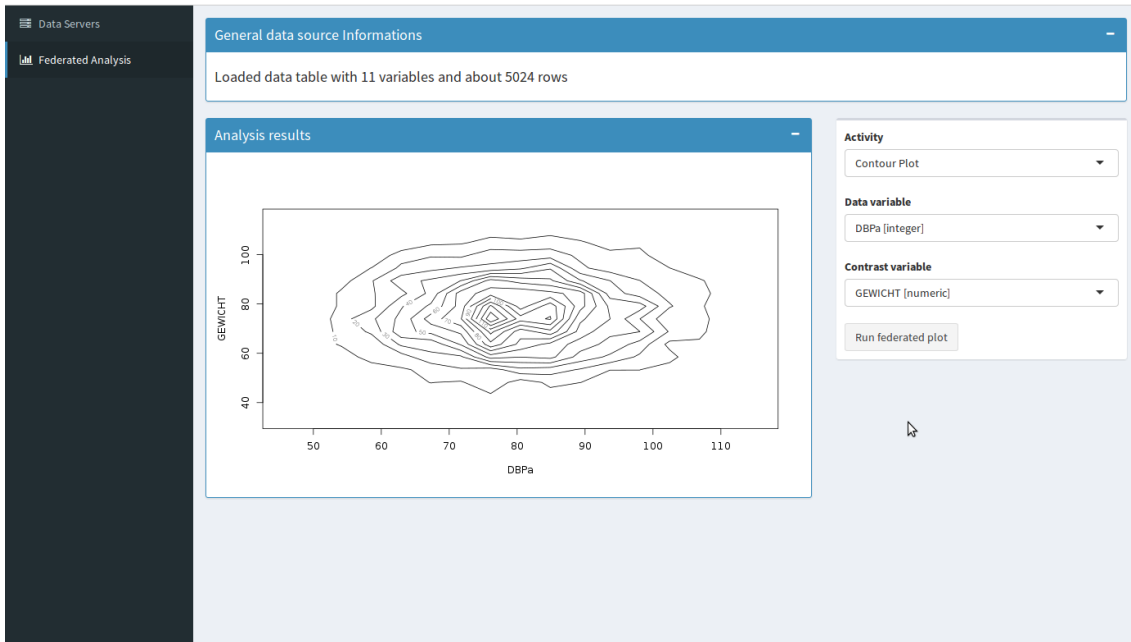
JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

≡ The **DASH-IN** interactive federated analysis system
DATA SHARING INITIATIVE



Federated heatmaps

≡ The **DASH-IN** interactive federated analysis system
DATA SHARING INITIATIVE



Federated contour plots



JOINT PROGRAMMING INITIATIVE – A HEALTHY DIET FOR A HEALTHY LIFE EUROPEAN NUTRITION PHENOTYPE ASSESSMENT AND DATA SHARING INITIATIVE

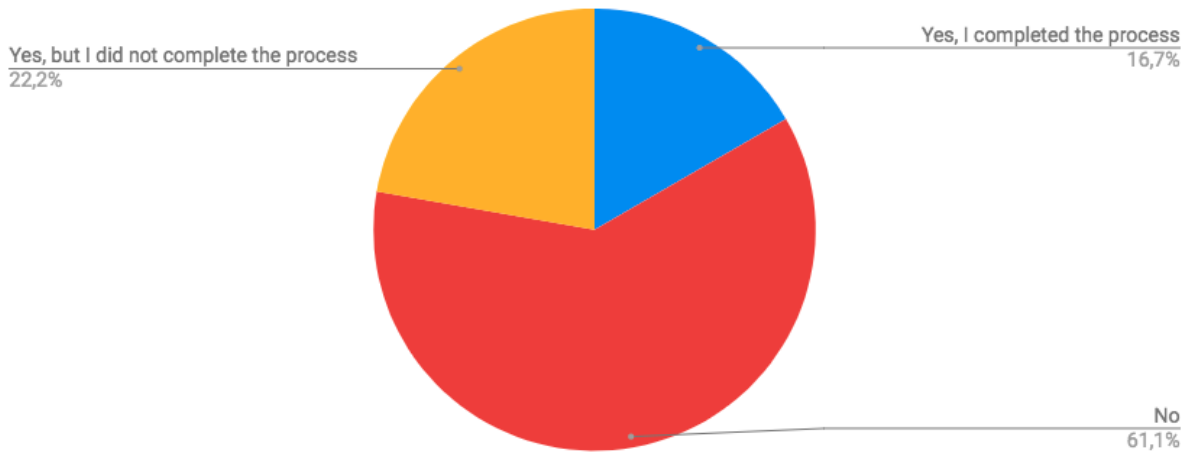
Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

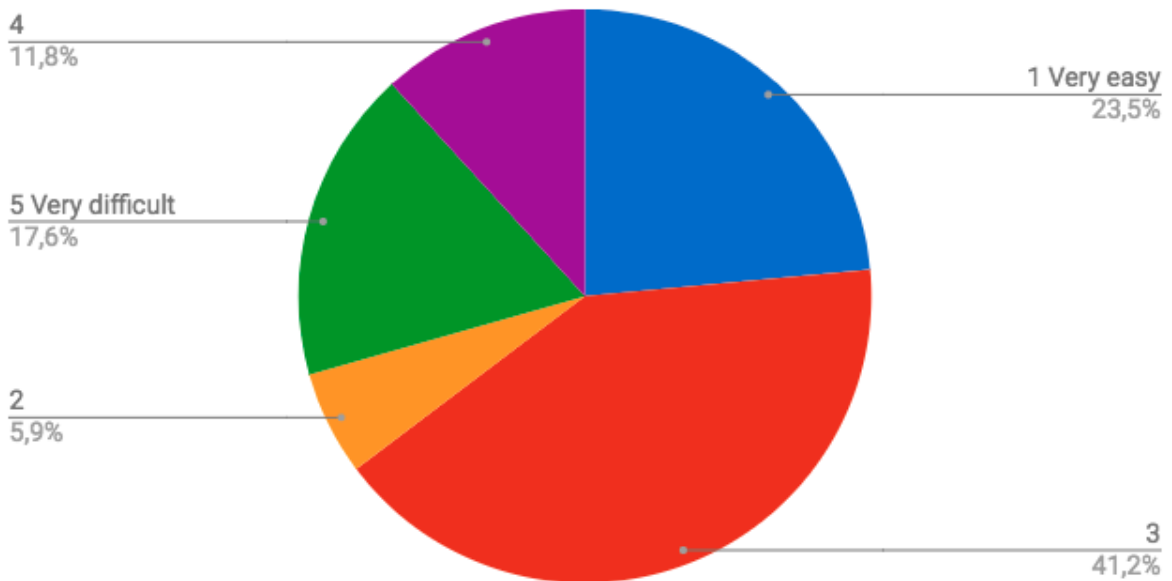
How easy or difficult is it to generate federated graphics?

- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Dash-in - Federated graphics - Have you used this functionality



Dash-in - Setting up data sources- How easy or difficult did you find it?





DASH-IN – FEDERATED ANALYSES

The screenshot shows the DASH-IN interface. The left sidebar contains 'Data Servers' and 'Federated Analysis'. The main area is divided into two panels: 'General data source Informations' and 'Analysis results'. The 'General data source Informations' panel shows 'Loaded data table with 11 variables and about 5024 rows'. The 'Analysis results' panel contains instructions: 'Specify model parameters and click 'Run federated analysis''. Below this, there are dropdown menus for 'Activity' (set to 'Analysis'), 'Analysis type' (set to 'Generalized linear model (GLM)'), and 'Output distribution' (set to 'Poisson (link=log)'). There is a text input field for 'Explanatory variables' containing 'SMK4A1' and a 'Run federated analysis' button.

The Analysis toolbox offers linear regressions and generalized linear models for the binomial and Poisson distributions. The analytical results are the aggregation of statistics computed at each hosting institution increasing the power while at the same time ensuring complete data privacy and confidentiality as the data never leaves the hosting data servers.

≡ The **DASH-IN** interactive federated analysis system
DATA SHARING INITIATIVE

This screenshot shows the 'Analysis results' section of the DASH-IN interface. It displays the evaluated model: 'GESLACHT ~ LENGTE + SMK11#num + DBPa * GEWICHT' family 'gaussian' link 'identity'. Below this is a table of model coefficients:

Variables	Estimate	Std. Error	z-value	p-value	low0.95CI	high0.95CI
(Intercept)	1.699	0.284	5.991	0	1.143	2.255
LENGTE	0	0.001	-0.1	0.92	-0.002	0.001
SMK11_conv	-0.04	0.018	-2.249	0.025	-0.074	-0.005
DBPa	-0.002	0.003	-0.687	0.492	-0.008	0.004
GEWICHT	-0.002	0.003	-0.664	0.507	-0.009	0.004
DBPa:GEWICHT	0	0	0.674	0.382	0	0

Below the table, statistical metrics are provided: Null deviance: 1255.7 on 5023 degrees of freedom; Residual deviance: 1253.56 on 5018 degrees of freedom; Multiple R-squared: 0.0017 Adjusted R-squared: 5e-04; F-statistic: 1.7126 on 5 and 5018 DF, p-value: 0.128.

The 'Help' section at the bottom explains the dependent variable and explanatory variables. The dependent variable is 'GESLACHT (factor)'. The explanatory variables are 'LENGTE + SMK11#num + DBPa * GEWICHT'. A note states: 'These are variable names we want to test for association with the dependent one. More than one explanation variables can be separated by the '+' operator. The interaction between two variables can be tested as well (example: ...)

Federated linear regressions



≡ The **DASH-IN** interactive federated analysis system
DATA SHARING INITIATIVE

Data Servers

Federated Analysis

General data source Informations

Loaded data table with 11 variables and about 5024 rows

Analysis results

Evaluated model 'GESLACHT ~ LENGTE + SMK11#num + DBPa * GEWICHT' family 'binomial' link 'logistic'

Variables	Estimate	Std. Error	z-value	p-value	low0.95CLLP	high0.95CLLP	P_OR	low0.95CLP_OR	high0.95CLP_OR
(Intercept)	0.799	1.137	0.703	0.482	-1.429	3.028	0.69	0.193	0.954
LENGTE	0	0.003	-0.1	0.92	-0.006	0.006	1	0.994	1.006
SMK11_conv	-0.159	0.071	-2.248	0.025	-0.298	-0.02	0.853	0.742	0.98
DBPa	-0.009	0.013	-0.687	0.492	-0.033	0.016	0.991	0.967	1.016
GEWICHT	-0.009	0.013	-0.664	0.507	-0.034	0.017	0.991	0.966	1.017
DBPa:GEWICHT	0	0	0.874	0.382	0	0	1	1	1

Residual deviance: 6954.97 on 5018 degrees of freedom

Activity

Analysis

Analysis type:
Generalized linear model (GLM)

Output distribution:
Binomial (link-logistic)

Dependent variable:
GESLACHT (factor)

Explanatory variables:
LENGTE + SMK11#num + DBPa * GEWICHT

Run federated analysis

Help

Dependent variable:
The variable we would like to predict using the explanatory variables. If the selected dependent variable is categorical and the output distribution is continuous (as in linear regression or GLM-Poisson case) the variable will be automatically converted to numeric.

Explanatory variables:
These are variables we use to test for association with the dependent one. More than one explanatory variable can be selected by the '+' operator. The interaction between two variables can be tested as well (see help).

Federated GLM modeling the output as a binomial distribution (logit link function)

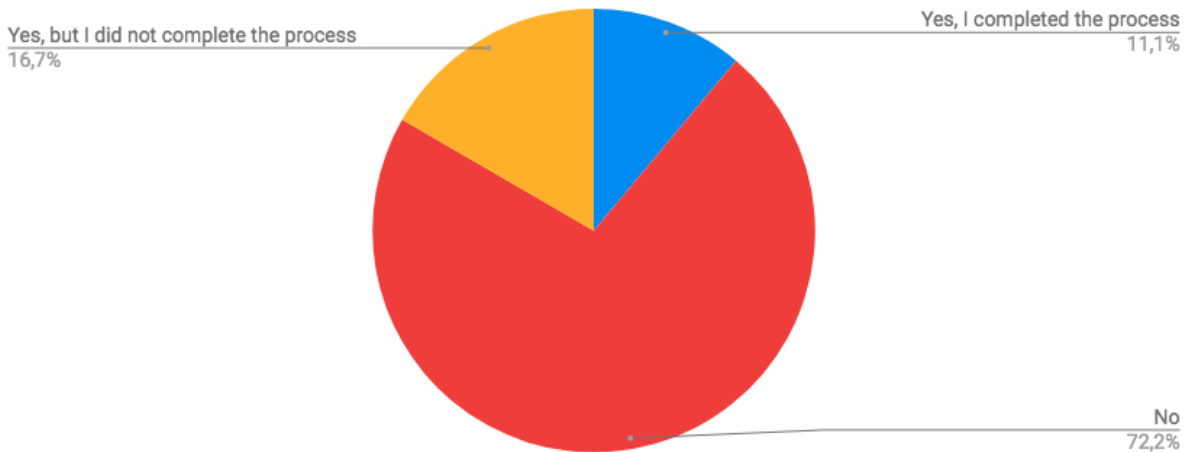
Have you used this functionality?

- No
- Yes, but I did not complete the process
- Yes, I completed the process

How easy or difficult is it to generate federated analyses?

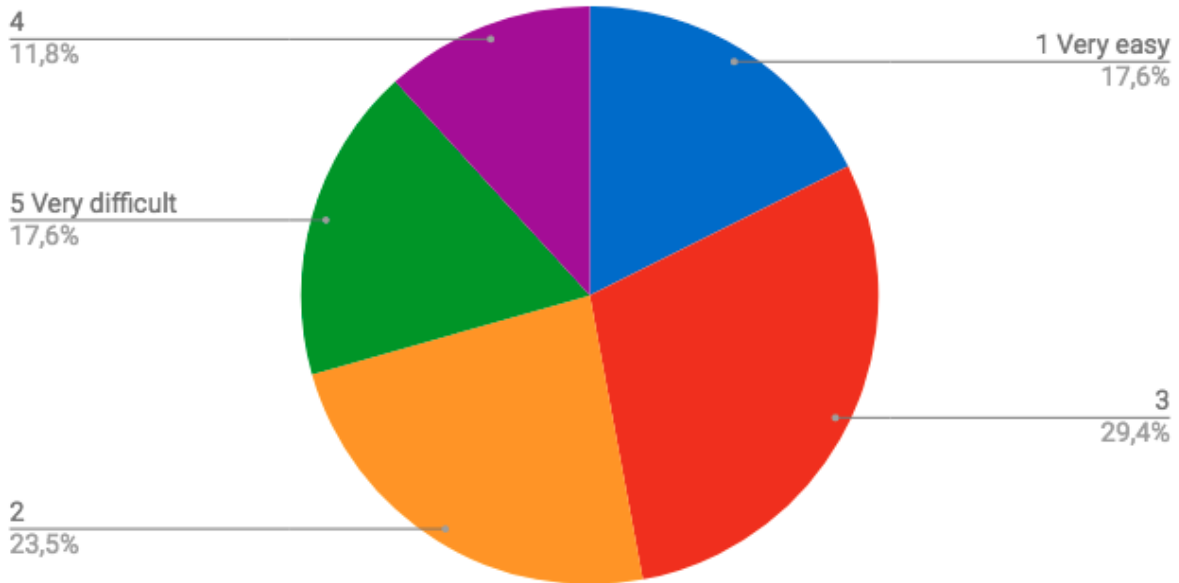
- 1 Very easy
- 2
- 3
- 4
- 5 Very difficult

Dash-in - Federated analyses - Have you used this functionality





Dash-in - Federated analyses - How easy or difficult did you find it?



DASH-IN – DO YOU HAVE ANY COMMENTS ON THE WEB BASED DASH-IN FEDERATED ANALYSES?

Federated Analyses section in the left side pane remained always empty.
Does not work so far. Just after connection (Analysis) : Error: An error has occurred. Check your logs or contact the app author for clarification. There is No DbNP available. Where could I found the list of studies of all partners ?
not used
I received error messages: "An error has occurred. Check your logs or contact the app author for clarification." 2. I wasn't able to find (and analyze) any DBNP studies 3. I still find it weird to be forced to assess potential difficulties related to a "functionality" that I could not use at all!
no
Some tasks are not immediate.
1) in the "Data Server" section, when selecting the data sever source and table, after clicking "Setup remote data tables" a progress bar and a progress status appear at the top of the page. It could be useful to have it just next to the button to monitor the upload process, and to have feedback that something is indeed ongoing 2) Got an error on all data servers "Error: An error has occurred. Check your logs or contact the app author for clarification.". This error did not prevent me from doing the analysis
Need Tutorials
it would be better if all the analysis steps and result could be recorded and downloaded.
A local explanation is suggested.