

## Batch upload of data files

### Samplings Oligo Sequence

Batch upload of data files is implemented on the MicroAqua internal website to ease uploading and parsing large chunks of data. Currently this functionality is available for "Sampling" and "Oligo sequence" data.

1. Click on the button for "sampling batch file upload", under Samplings section.

Upload multiple sampling files in one click

Samplings Batch File Upload

### Samplings

New sampling

List of samplings: you can filter (using the lens icon in the bottom of the grid), sort (clicking on the header column), scroll the data in the grid (using the pagination system)

	Code	Sampling site	Volume (lt)	Partner	Equipment	edit
+	P03-110705-01	CANET / Canet Saint Nazaire lagoon	43.0	3 - UPMC	bucket	
+	P03-110809-01	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	
+	P03-110926-01	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	
+	P03-111019-02	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	
+	P01-111123-02	CANET / Canet Saint Nazaire lagoon	30.0	1 - UniCam	?	
+	P01-110101-01	CANET / Canet Saint Nazaire lagoon	3.0	1 - UniCam	?	
+	P03-111115-01	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	
+	P03-120307-01	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	
+	P03-120403-01	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	
+	P03-120403-02	CANET / Canet Saint Nazaire lagoon	10.0	3 - UPMC	bucket	

2. The data file can be uploaded in formats like .csv(comma separated values) or .tsv(tab separated values). A template for the sample data file is available at the link "Download an example of sampling file". Just download it and fill it with your data.

New Samplings Multiple File Upload (\*.csv or \*.tsv)

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Add Files

Save Files

Download an example of sampling file

Download an example sampling file having a proper sampling file format. Files will only be uploaded in this particular format.

3. To fill data in the example file, some rules have to be taken care of:

a) In the first field, Sampling Site, fill the blank field by Sampling Site Code. Codes are available in the index table of *MicroAqua internal website* > *Sampling Site*.

A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R
Sampling Site	Partner	Sampling date(YYYY-MM-DD)	Volume	Comment	Air Temp.	Moisture	Pressure	Wind Speed	Wind Direction	Water Flow	Light Int.	Rainfall exts.	Depth	Turbidity	Salinity	Tidal Range	Open

**Sampling sites**

Use the Code names to fill the "Sampling Site" field in the Example file

Code	Name	Water use	Water type	Land	Geo pos.
CANET	Canet Saint Nazaire lagoon	Recreative water	Surface water	Grazing land	Canet Saint Nazaire lagoon (Fr
SOLA	SOLA station	Recreative water	Surface water	Grazing land	SOLA station (France)
SCHWENTINE	Schwentine	Recreative water	Surface water	Grazing land	Schwentine (Germany)
MONDONY	Mondony river	Recreative water	Surface water	Grazing land	Mondony river (France)
SAPANCA	Lake of Sapanca	Recreative water	Surface water	Grazing land	Lake of Sapanca (Turkey)
POTENZA	River Potenza (Fonte di Bresc	Resource water	Surface water	Grazing land	River Potenza (Fonte di Bresc
FIASTRA	Lake of Fiastra (100 m far from	Recreative water	Lake	Artificial area	Lake of Fiastra (100 m far from
CHIANTI	River Chienti (Piediripa)	Resource water	Surface water	Open spaces	River Chienti (Piediripa) (Italy)
AMELIE	Mondony river	Recreative water	Surface water	Grazing land	AMELIE (France)
Dodder	River Dodder in Dublin urban ar	Resource water	River	Urban	Dodder Dublin (Ireland)

b) For Partners field, put the Partners code (in upper or lower case). Refer to the following list or check the codes from the MicroAqua website *Partners* page (<http://www.microaqua.eu/beneficiaries>).

University of Camerino	UNICAM
University of Portsmouth	UniPort
Universite Pierre et Marie Curie	UPMC
Veolia Environment Recherche et innovation	VERI
Istituto Superiore Di Sanita	ISS
University of Ireland, Dublin	UCD
National Center of Infectious and Parasitic Diseases	NCIPD
Queens University Belfast	QUB
MariLim Gesellschaft fur Gewasseruntersuchung mbH	Marilim
Istanbul University	Unilst
Universidade de Santiago de Compostela	USC
Scienion AG, Berlin	SCIE

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## List of participants



**Partner : 1**  
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**Principal investigator:**  
 Hayes Paul Kelvin (paul.hayes@port.ac.uk)

**Partner Code**

UniCam Italy [Read more](#)

UniPort United Kingdom [Read more](#)

Save the Example file with the name of your choice and now its ready to be uploaded.

**NOTE1:** The file is already in .CSV format. There is no need to change the format of the file.

**NOTE2:** It is essential to fill in all the field available in the Example file. If the Partner has no data for the respective field, especially for Enviornmental data like pressure, moisture etc, just put "0.0" value in the field. Do not leave any field blank.

4. Once the data files are ready, simply click on "Add files" button *multiple times* to get the file upload tag. Maximum of 10 files can be uploaded together with one click. Just browse your files and provide the location. Then click on "Save Files". A unique sampling code will be assigned to individual rows with in the data file. Once the data is parsed sucessfully, filter samples can be associated manually with each sampling entry.

New Samplings Multiple File Upload (\*.csv or \*.tsv) [Download an example of sampling file](#)

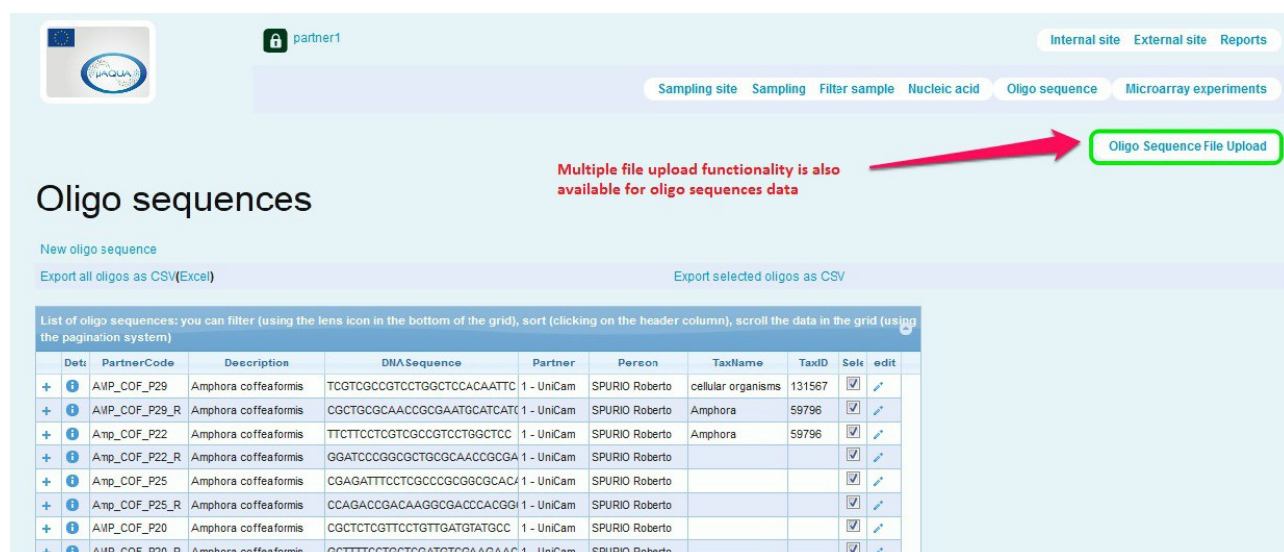
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**Add Files** **Save Files**

**Browse your local computer filesystem to select sampling files(s)**



5. The same procedure has to be followed with "Oligo Sequence" batch files.



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Sampling site Sampling Filter sample Nucleic acid Oligo sequence Microarray experiments

**Oligo sequences**

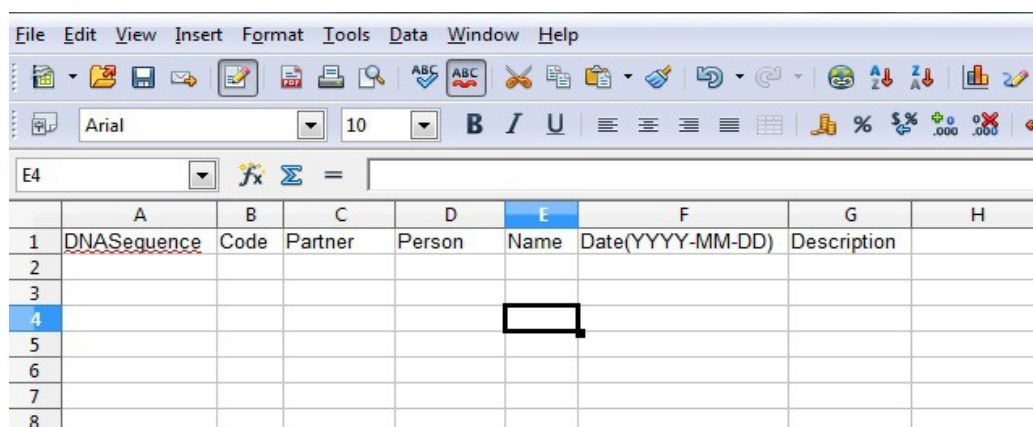
New oligo sequence

Export all oligos as CSV(Excel) Export selected oligos as CSV

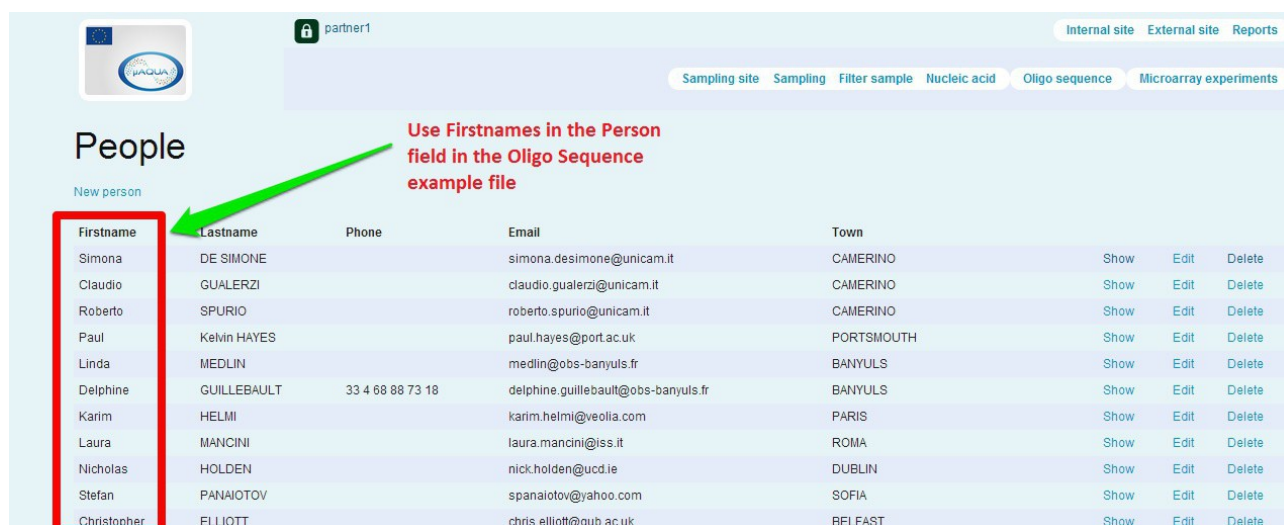
List of oligo sequences: you can filter (using the lens icon in the bottom of the grid), sort (clicking on the header column), scroll the data in the grid (using the pagination system)

Deti	PartnerCode	Description	DNASequence	Partner	Person	TaxName	TaxID	Sele	edit
+ i	AMP_COF_P29	Amphora coffeiformis	TCGTGCGCGTCTGGCTCCACAATTC	1 - UniCam	SPURIO Roberto	cellular organisms	131567	<input checked="" type="checkbox"/>	
+ i	AMP_COF_P29_R	Amphora coffeiformis	CGCTGCGCAACCGCGAATGCATCATC	1 - UniCam	SPURIO Roberto	Amphora	59796	<input checked="" type="checkbox"/>	
+ i	Atmp_COF_P22	Amphora coffeiformis	TTCTTCTCTGTCGCGCTCTGCTCC	1 - UniCam	SPURIO Roberto	Amphora	59796	<input checked="" type="checkbox"/>	
+ i	Atmp_COF_P22_R	Amphora coffeiformis	GGATCCCGGCGCTGCGCAACCGCGA	1 - UniCam	SPURIO Roberto			<input checked="" type="checkbox"/>	
+ i	Atmp_COF_P25	Amphora coffeiformis	CGAGATTCTCTGCGCGCGCGCAC	1 - UniCam	SPURIO Roberto			<input checked="" type="checkbox"/>	
+ i	Atmp_COF_P25_R	Amphora coffeiformis	CCAGACGACAAAGCGGACCGG	1 - UniCam	SPURIO Roberto			<input checked="" type="checkbox"/>	
+ i	AMP_COF_P20	Amphora coffeiformis	CGCTCTGTTCTGTTGATGTATGCC	1 - UniCam	SPURIO Roberto			<input checked="" type="checkbox"/>	
+ i	AMP_COF_P20_R	Amphora coffeiformis	GCTTTTCTGCTCGATGTCGAAGAAC	1 - UniCam	SPURIO Roberto			<input checked="" type="checkbox"/>	

6. In the example file, fill the *Partner* field with partner code and *Person* field with the Firstnames of the person uploading the file. For partner please refer to the section 3.b above and for Persons, please refer the following image. Firstnames of all the persons involved in MicroAqua are available here: <http://bioaqua1.microaqua.eu/people>



	A	B	C	D	E	F	G	H
1	DNASquence	Code	Partner	Person	Name	Date(YYYY-MM-DD)	Description	
2								
3								
4								
5								
6								
7								
8								



Internal site External site Reports

Sampling site Sampling Filter sample Nucleic acid Oligo sequence Microarray experiments

**People**

New person

Use Firstnames in the Person field in the Oligo Sequence example file

Firstname	Lastname	Phone	Email	Town	Show	Edit	Delete
Simona	DE SIMONE		simona.desimone@unicam.it	CAMERINO			
Claudio	GUALERZI		claudio.gualerzi@unicam.it	CAMERINO			
Roberto	SPURIO		roberto.spurio@unicam.it	CAMERINO			
Paul	Kelvin HAYES		paul.hayes@port.ac.uk	PORTSMOUTH			
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Christopher	ELLIOTT		chris.elliott@qub.ac.uk	BELFAST			