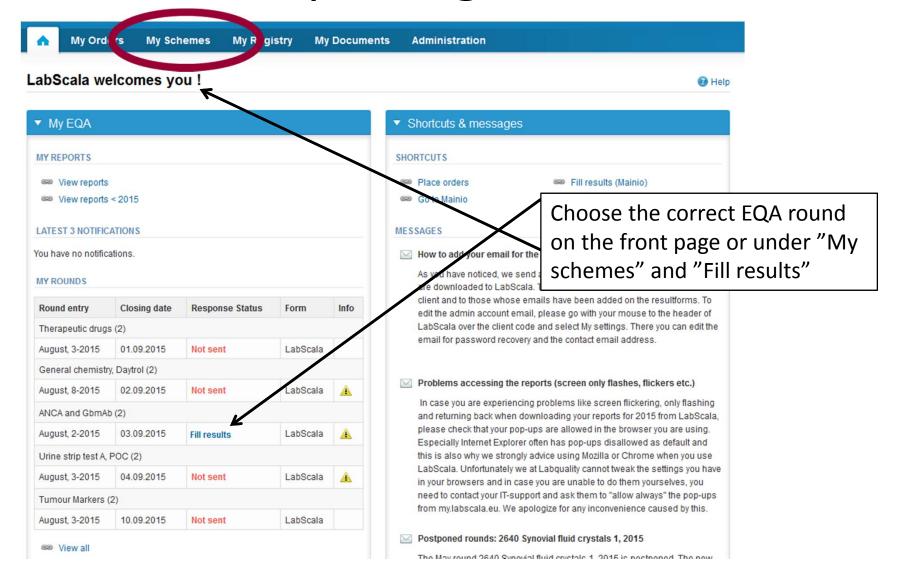


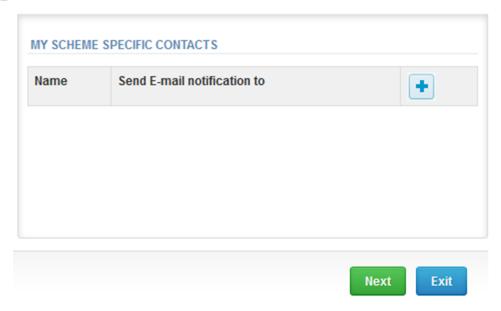
5260
Fungal culture
How to fill in results

Result reporting in LabScala

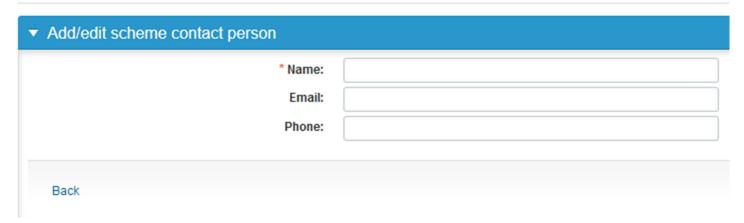


Filling results

- First add your scheme-specific contact info by pressing plus – sign
- Fill in name and email address or phone number
- Save and choose Next



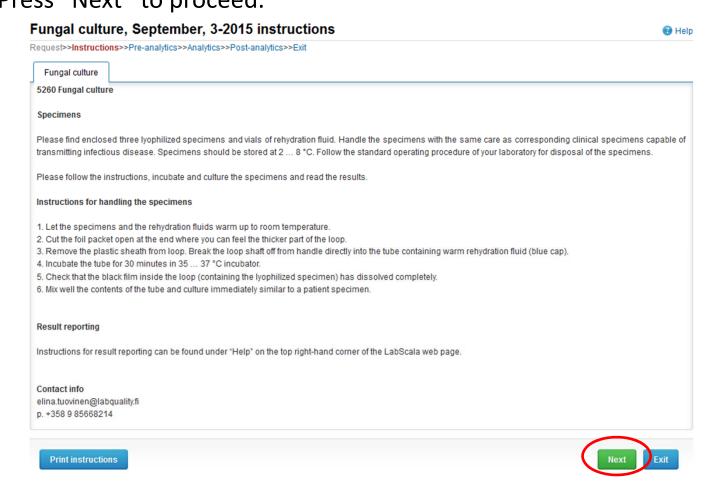
Add scheme contact person



Instructions

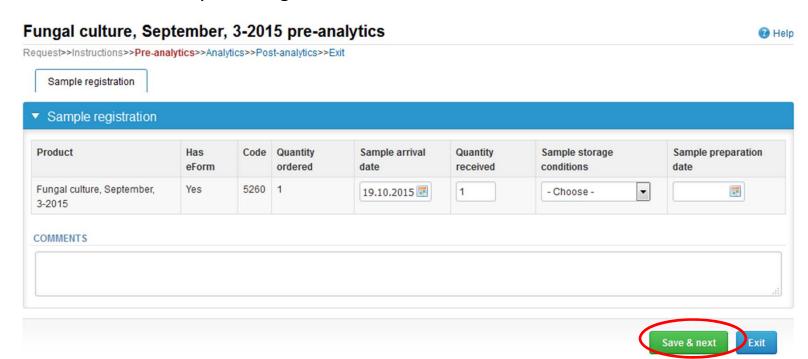
In Instructions the scheme-specific instructions can be read and printed by pressing the "Print instructions" button on the bottom of the page.

Press "Next" to proceed.

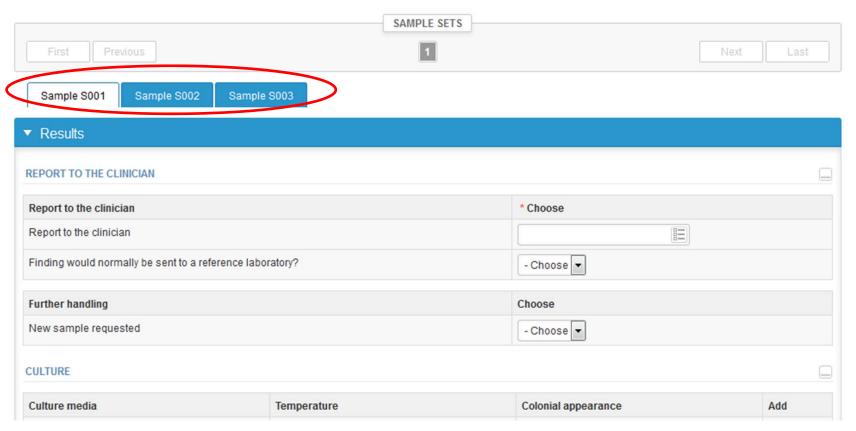


Pre-analytics

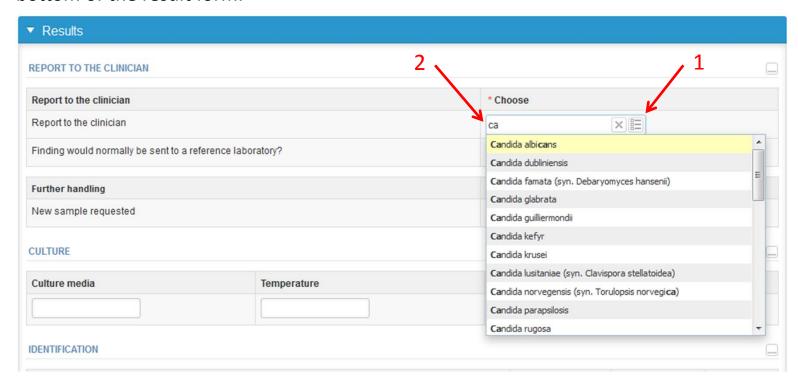
- In Pre-analytics, the following is asked
 - Sample arrival date
 - Quantity received: How many sample sets were received
 - Sample storage condition: How have the samples been stored before analysis (refrigerator, room temperature, freezer, other)
 - Sample preparation date: if done, if not, can be left empty
- Comments can be saved if needed
- Move forward by selecting "Save & next"



- The sample-specific results can be filled on Analytics view.
- If you have ordered multiple sample sets, the sets can be seen on the top of this view.
- There are own result sheets for each sample. The samples are listed on top of the result form (e.g. S001, S002, S003).

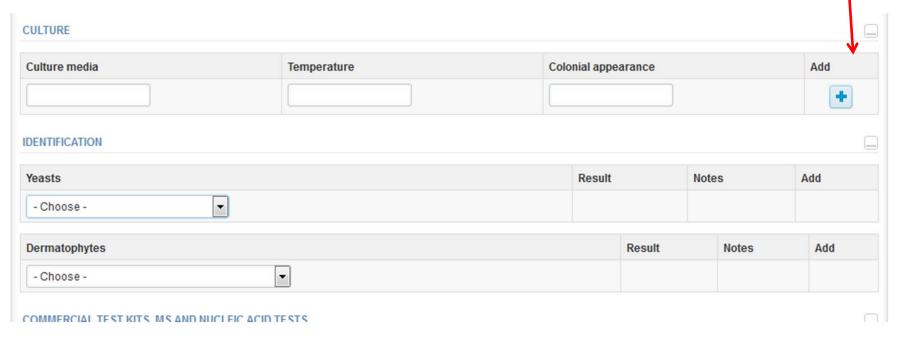


- The first two questions on the form are mandatory.
- Choose from the dropdown menu (1) your report to the clinician. You can also seach for a
 microbe by writing a part of its name to the field (2). If you cannot find the desired microbe,
 you may also choose "Other finding" and specify it in the "Comments" field.
- Answer to the question of sending the sample to a reference laboratory.
- The question of further handling is not mandatory.
- Comments and additional information can be recorded in the "Comments" field in the bottom of the result form.

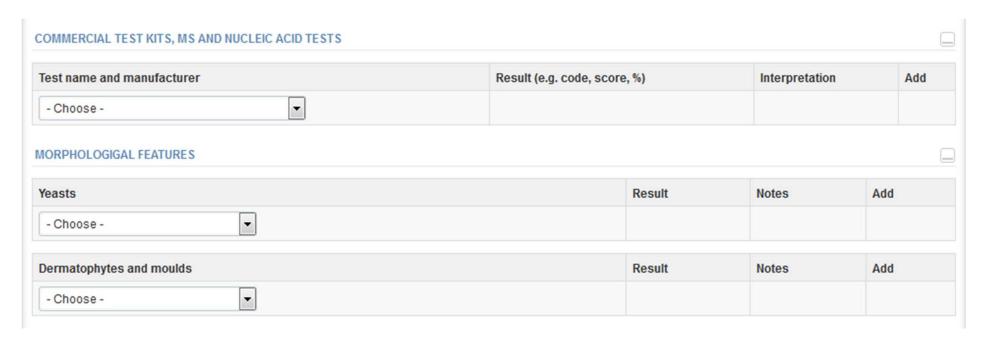


- In the "Culture" section you can report the culture media used, culture conditions and colonial appearance. You can report multiple rows. NB! You must save your data by choosing +.

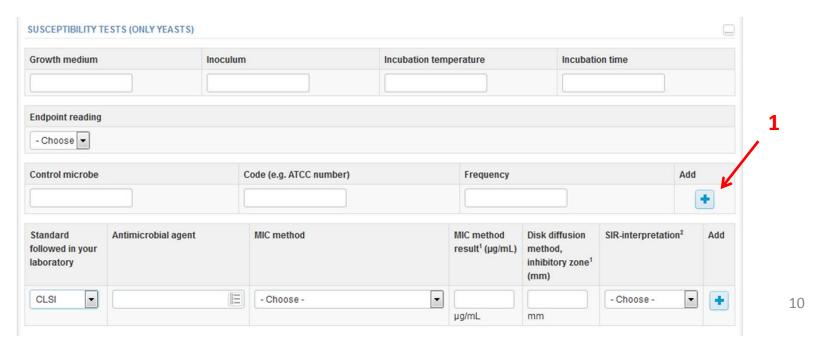
 Otherwise the data will be lost.
- You can fill in the rest of the form so far you have made findings. Please report only the items, which are important for the identification.
- In the section "Identification" you can report results of identification tests. The tests are chosen from the drop down menu. Remember to save the row by choosing +



- In the section "Commercial test kits, MS and nucleic acid tests" you can report the results of identification tests. If you cannot find the desired option from the dropdown menu, you may also write "Other finding" and specify it in the "Comments" field. You can report multiple result rows. Save your results by choosing +
- In the section "Morphological features" you can report the morphological findings observed. You can report many result rows. <u>Save your result by choosing +</u>



- <u>The section "Susceptibility tests" is intended only for yeasts.</u> It is not mandatory to report susceptibility tests.
- Answer the questions regarding culture conditions, endpoint reading and control microbe.
 Remember to save by choosing + (1)
- Choose the standard that you follow and the antimicrobial agent.
- Report either MIC or disk diffusion method and SIR interpretation.
- Choose + to save the result and to add a new antimicrobial agent.
- Continue reporting the results of other antimicrobial agents and answer to possible additional questions.



- You can write additional information and other comments of the samples, tests or result form in the "Comments" field at the end of the page.
- When all the data has been reported, choose "Save as final".
- It is possible to edit data by choosing "Edit data".
- "Next" will take you to the next sample.
- Report the results for the samples S002 and S003 and choose "Save as final" for each sample.



Post-analytics

- Next the user is taken to the result validation
 - Product: what part of the process is being validated
 - Sample set: which sample set is being validated
 - Sample: which sample
 - Errors: if there are reportingerrors these are shown here
 - Last saved: The user who has saved the results
 - Date: date of last saving
 - Status: status of the results (Accepted, Draft, Error, Open)

Fungal culture, September, 3-2015 postanalytics



Request>>Instructions>>Pre-analytics>>Analytics>>Post-analytics>>Exit

Validate results

Product	Sample set	Sample	Errors	Last saved	Date	Status
Preanalytics		Registration	OK	Teija Häkkinen	19.10.2015 16:07	Accepted
Fungal culture	1	Sample S001	OK	Tanja Elo	29.10.2015 14:03	Draft
Fungal culture	1	Sample S002	OK	Elina Tuovinen	27.10.2015 14:06	Accepted
Fungal culture	1	Sample S003	OK	Elina Tuovinen	27.10.2015 15:14	Accepted

Accept and send results

Exit

Post-analytics

- The results can be edited by selecting the correct sample.
- When all of the statuses are such that the results can be sent to Labquality, select "Accept and send results".
- If you wish not to send the results yet, select "Exit".
- Note: The results can be edited as long as the round is open, even if "Accept and send results" has been selected.

Fungal culture, September, 3-2015 postanalytics



Request>>Instructions>>Pre-analytics>>Analytics>>Post-analytics>>Exit

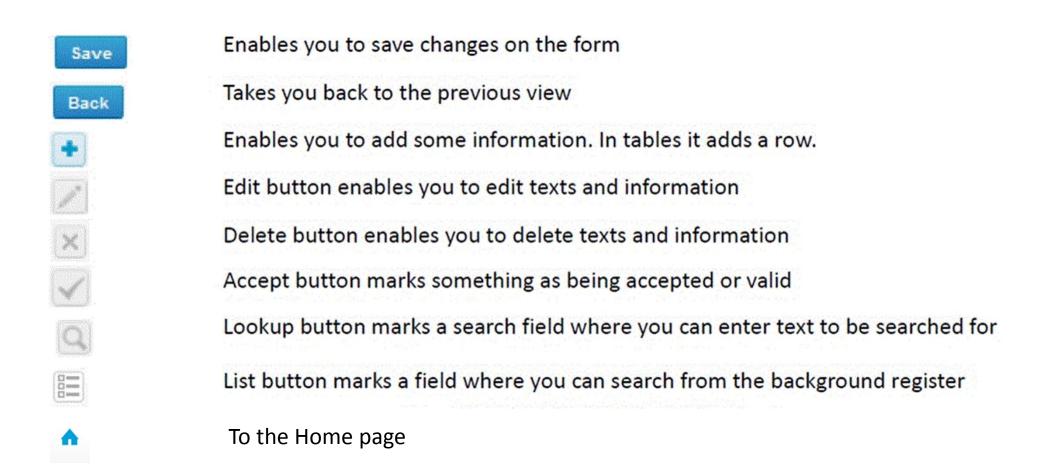
Validate results

Product	Sample set	Sample	Errors	Last saved	Date	Status
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Fungal culture	1	Sample S002	OK	Elina Tuovinen	27.10.2015 14:06	Accepted
Fungal culture	1	Sample S003	OK	Elina Tuovinen	27.10.2015 15:14	Accepted

Accept and send results

Exit

LabScala buttons



Questions?

- In case you have questions, please contact:
 - Elina Tuovinenelina.tuovinen@labquality.fi+358 9 8566 8214
 - Jonna Pelantijonna.pelanti@labquality.fi+358 9 8566 8211