

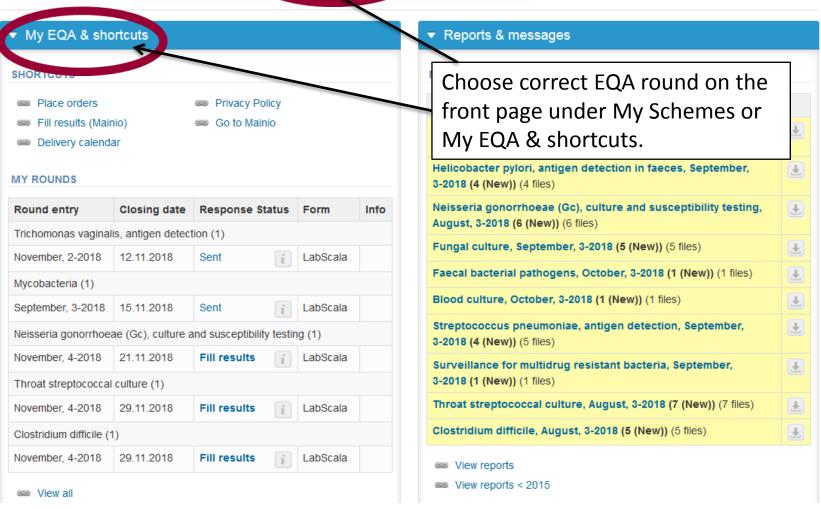
How to fill in results for Streptococcus Group B (GBS) detection (5594)

Quick guide for result reporting

- 1. Start from LabScala home page by choosing My Schemes —> Fill results -> choose correct round.
- 2. Add Specific Contact, choose Next.
- 3. Read Instructions, choose Next.
- 4. Update Sample registration, choose Save & next
- 5. Reporting results: choose the appropriate result/s.
- 6. Save your results as Save as final. Move to next specimen by choosing Next.
- 7. Validate your results and if needed, update them, finally choose Accept and send results.

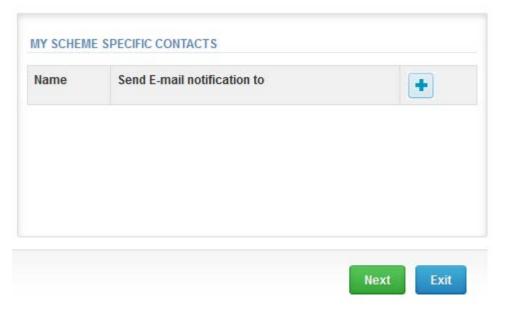
Result reporting



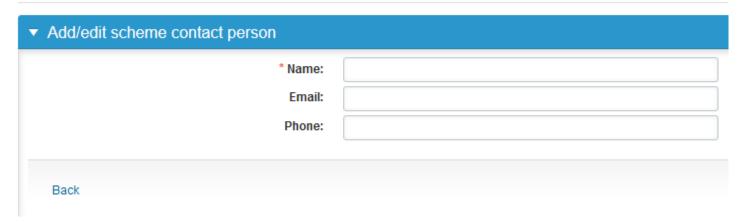


Filling contact info

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



Add scheme contact person



Instructions

 In Instructions the scheme-specific instructions and result forms for internal use can be read and printed by pressing the "Print instructions" and/or "Open result form" button.



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

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

Move forward by selecting "Save & next"

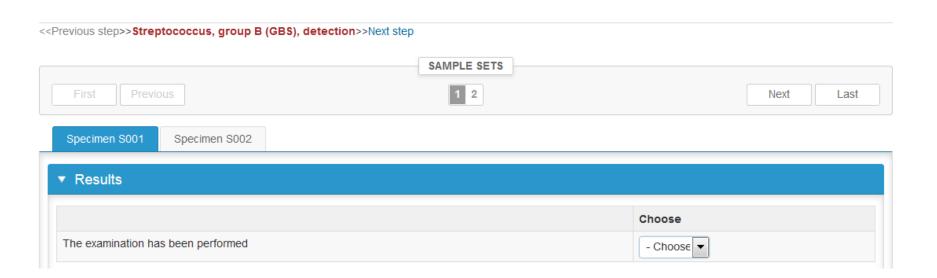
Streptococcus, group B (GBS), detection, November, 4-2018 pre-analytics

LabScala User instructions

Sample registration Sample registration Product Has Code Quantity Sample arrival Quantity Sample storage Sample eForm ordered date received conditions preparation date Streptococcus, group B (GBS), Yes 5594 2 • - Choose detection, November, 4-2018 COMMENTS Save & next

Results – how to start

- If you have ordered multiple sample sets, the sets can be seen on the top of this view
- The samples are listed on top of the result form (e.g. Specimen 001, Specimen 002)
- Results section: choose the appropriate option, if the examination has been performed in a laboratory or on a ward



Results 1/4

• **Report to the clinician**: it is mandatory to give an answer to this section. The Report to the clinician will be scored. Choose the option from the drop-down list.



• **Culture Plate**: you may report results for selective and chromogenic medium. Always press the plus (+) button at the end of the row to add the result.



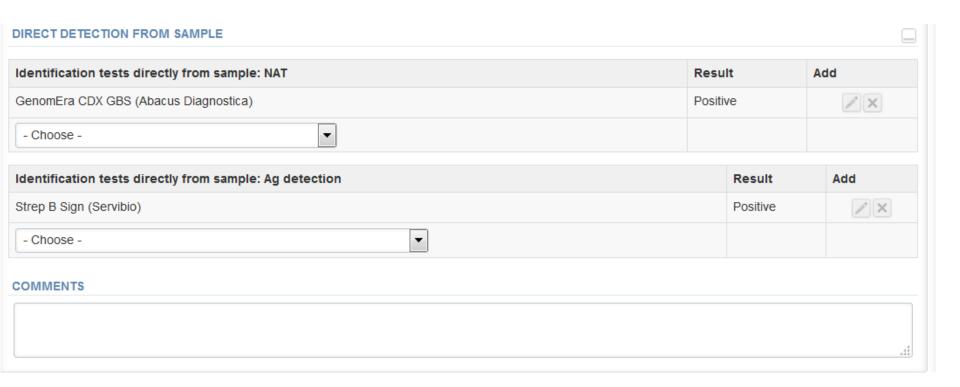
Results 2/4

• **Identification test performed on colony**: Note that this section is intended for tests that are performed on colony. You may report multiple test results.

DENTIFICATION TEST PERFORMED ON COLONY								
Agglutination tests					Result		Add	
Microgen Strep Rapid Test (Microgen Bioproducts)					Positive		X	
BBL Streptocard Acid Latex Test (Becton Dickinson)					Positive		X	
- Choose -	•							
Commercial identification kits and analyzers			F	Result			Add	
apid ID 32 Strep (bioMerieux)			5	Streptococcus agalactiae			/ X	
- Choose -								
dentification tests: MALDI-TOF Result							Add	
Microflex LT (Bruker)		Streptococcus agalactiae					X	
- Choose -								
Identification tests on colony: NAT					Result	Add	d	
HG Group B Streptococcus (HiberGene Diagnostics)					Positive		X	
- Choose - ▼								
			Result					
Chromogenic medium			Positive	• ▼				

Results 3/4

- Identification tests performed direct from sample: Note that this section is intended for tests that are performed directly on sample. You may report multiple test results.
- The Comments section is intended for any comments you want to make on the test results, samples or the round.



Results 4/4

- Finally select
 - "Save as draft" if you do not yet want to send in the results. Results are not proccessed in Labquality
 - "Save as final" if you see the results as being final
- From this page you can return to the front page by selecting "Back to list" or clear the whole page by selecting "Clear all page data"



Results, edit data

- If "Save as final" was selected the results are saved and the table is closed. The
 data can be edited by selecting "Edit data" from the bottom of the page
- By selecting "Next" LabScala takes you automatically to the next sample or sample set to be filled



Post-analytics 1/2

- 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 analyte-specific validation limit errors 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)

Validate results

Product	Sample set	Sample	Errors	Last saved	Date	Status
		•				
Streptococcus, group B (GBS), detection	1	Specimen S001	OK	Yvonne Björkman	12.11.2018 09:44	Accepted
Streptococcus, group B (GBS), detection	1	Specimen S002	OK	Teija Häkkinen	17.10.2018 13:19	Draft
Streptococcus, group B (GBS), detection	2	Specimen S001	OK	Yvonne Björkman	12.11.2018 09:46	Open
Streptococcus, group B (GBS), detection	2	Specimen S002	ОК	Yvonne Björkman	12.11.2018 09:46	Open

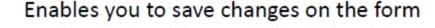
Accept and send results

Post-analytics 2/2

- The results can be edited by selecting the sample
- When you evaluate that your results are finalized and are ready to be sent to Labquality, select "Accept and send results"
- If you wish not to send the results yet, select "Exit"
- The results can be edited as long as the round is open, even if "Accept and send results" has been selected

LabScala buttons







Takes you back to the previous view



Enables you to add some information. In tables it adds a row.



Edit button enables you to edit texts and information





Delete button enables you to delete texts and information



Accept button marks something as being accepted or valid



Lookup button marks a search field where you can enter text to be searched for



List button marks a field where you can search from the background register



Questions?

- In case you have questions, please contact:
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