

How to fill in results Faecal culture (5190)

Quick guide for result reporting

- 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: Answer questions.
- 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

🔥 My Ord	ers My Sch	emes My Regi	stry My	Documents	Administration
_abScala w	elcomes yo	ou !			🝞 Help
 My EQA 				•	Shortcuts & messages
MY REPORTS					SHORTCUTS
 View report View report LATEST 3 NOTIFI You have no notified 	s s < 2015 CATIONS cations.				Choose correct EQA round on the front page or by choosing My Schemes, Fill results
MY ROUNDS					In the view of the closing
Round entry	Closing date	Response Status	Form	Info	
Mycobacterial cu	ture and smear (4)				schemes the closing date
June, 2-2015	21.08.2015	Not sent	LabScala	<u> </u>	and the status of the scheme
Therapeutic drug	s (2)				
August, 3-2015	01.09.2015	Not sent	LabScala		can be seen
General chemistry	y, Daytrol (2)				Open = eForm open for result
August, 8-2015	02.09.2015	Not sent	LabScala	<u> </u>	filling
ANCA and GbmA	b (2)				Net cont - recult filling started
August, 2-2015	03.09.2015	Not sent	LabScala	<u>.</u>	Not sent = result filling started
Urine strip test A,	POC (2)				but results not sent to
August, 3-2015	04.09.2015	Not sent	LabScala	<u>.</u>	Labquality
📾 View all					Accepted = results sent to Labquality

3

Request

- After opening the scheme the request page is opened with the following information
 - Product: Here it can be seen what and how much has been ordered (e.g. Faecal culture (1) means 1 set of samples ordered)
 - Instructions: Scheme spesific instructions can be read and printed
 - **Pre-analytics:** Sample arrival date, quantity received, sample storage conditions etc.
 - Analytics: analyte-specific results can be filled
 - **Postanalytics:** Validate results
 - Exit: Back to the front page
- Also the scheme specific contact information is filled here
 - The contact person receives reminders and important scheme-specific information
 - At least one contact person needs to be named!

Welcome to the round of Faecal culture, October, 3-2015 analysis

🕜 Help

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

MY ORDER	s					MY SCHE	ME SPECIFIC CONTACTS	
Product	Delivery	Opening	Closing	Status	Form	Name	Send E-mail notification to	+
Faecal cult	ture(1)					test	test@test.fi	
	07.10.2015	07.10.2015	30.10.2015	Ordered	LabScala			
➤ Coord	inator conta	tii By p LabS forw	ressing Scala wi vard	the "N Il auto	Next" bu maticall	tton y move y	you	Next Exit

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). If done, if not can be left empty
 - Sample preparation date: If done, if not can be left empty
- Comments can be saved if needed
- Move forward by selecting "Save & next"

Faecal culture, October, 3-2015 pre-analytics

Product	Has eForm	Code	Quantity ordered	Sample arrival date	Quantity received	Sample storage conditions	Sample preparation date
aecal culture, October 3-2015	Yes	5190	1	01.10.2015 📰	1	+2 +8 C	



5

Help

Results – Test selection

- If you have ordered multiple sample sets, the sets can be seen on the top of this view
- If the scheme has multiple samples, the samples are listed on top of the result form (e.g. S001 and S002)
- Result filling is started by answering the question in the "Test selection" part.

	SAMPLE SETS	
First Previous	1	Next Last
Sample S001 Sample S002		
 Test selection 		
Our test selection includes the culture and ident	ification of:	* Choose
Salmonella		- Choose 💌
Shigella		- Choose 💌
Campylobacter		- Choose 💌
Yersinia		- Choose 💌
Aeromonas		- Choose 💌
Plesiomonas		- Choose 💌

Results – Report to the clinician

- Answer to the questions in the report to the clinician part
- From the menu (1) choose the desired microbe name for the finding. The microbe name can also be searched by writing part of the name in the field (2)
- Report also the sending to reference laboratory

▼ Microbe			Add finding 🕂
REPORT TO THE CLINICIAN	2		1
		* Finding	
Report to the clinician		salmo 🗙 🔠	
Finding would normally be sent to a	a reference laboratory?	Aeromonas salmonicida	
		Salmonella Aberdeen	
CULTURE		Salmonella Abony	_
COLTORE		Salmonella Adana	
	Choose	Salmonella Adelaide	
Growth media	Calast	Salmonella Agama	
Growarmedia	- Select -	Salmonella Agona	

Results – Growth media

- From the "Growth media" part you can choose the media/broth used
- You can select multiple options if needed

CULTURE		
	Choose	
Growth media	Chromogenic media, E 💌	
	Check all Uncheck all	
IDENTIFICATION TESTS TO CONFIRM PATHOGEN FIND	Chromogenic media	
	Non-chromogenic media	Choose
Serological method (agglutination test)		- Choose 💌

Results – Identification tests

- Report your results for used identification tests and **save by choosing +**
- You can write comments / specifications on "Comments" field.

IDENTIFICATION TESTS TO CONFIRM PATHOGEN FINDING						
Serological method (agglutination test)				Choose Yes 💌		
Commercial identification test kits and analyzers	Profile number		* Result		Proba	bility % Add
Identification by MALDI-TOF MS	123456789	Score	Salmonella sp. Probability	× =	*Result	Add
- Choose -						
Nucleic acid detection - Choose -			*Result			Add

COMMENTS



Results

- If needed add the next finding by choosing "Add finding +"
- Finally choose "Save as final" to save the reported finding/findings

							Add finding
REPORT TO THE CLINICIAN							E
				* Finding			
Report to the clinician						E	
Finding would normally be sent to	a reference laboratory?			- Choose 💌			
ULTURE							(
	Choose						
Growth media	- Select	- Select -					
DENTIFICATION TESTS TO CON	FIRM PATHOGEN FINDING						
DENTIFICATION TESTS TO CON Serological method (agglutination Commercial identification tes	FIRM PATHOGEN FINDING test)	Profile numb	er * Re	sult	Choose Yes	Probability %	Add
DENTIFICATION TESTS TO CON Serological method (agglutination Commercial identification tes VITEK 2	FIRM PATHOGEN FINDING test) at kits and analyzers	Profile number 123456789	er * Re Salm	sult ionella sp.	Choose Yes	Probability % 99 %	Add
DENTIFICATION TESTS TO CON Serological method (agglutination Commercial identification tes VITEK 2	FIRM PATHOGEN FINDING test) st kits and analyzers	Profile number 123456789	er Re Salm	sult ionella sp.	Choose Yes •	Probability % 99 %	Add
Serological method (agglutination Commercial identification tes VITEK 2	test) st kits and analyzers	Profile number 123456789	er • Re Salm Score	sult ionella sp. Probability	Choose Yes	Probability % 99 % Result	Add
Serological method (agglutination Commercial identification tes VITEK 2	FIRM PATHOGEN FINDING test) at kits and analyzers	Profile number 123456789	er * Re Salm Score	esult nonella sp. Probability	Choose Yes v	Probability % 99 % Result	Add
DENTIFICATION TESTS TO CON Serological method (agglutination Commercial identification tes VITEK 2	FIRM PATHOGEN FINDING test) st kits and analyzers	Profile number 123456789	er Re Salm Score	sult ionella sp. Probability	Choose Yes	Probability % 99 % *Result Add	Add

COMMENTS

Back to list Clear all page data



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 is being validated
 - **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

Validate results

- Status: Status of the results (Accepted, Draft, Error)

Faecal culture, October, 3-2015 postanalytics

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

 Validation results 						
Product	Sample set	Sample	Errors	Last saved	Date	Status
Faecal culture	1	Sample S001	ок	Paakkanen, Jaana	02.10.2015 15:36	Accepted
Faecal culture	1	Sample S002	ок	Paakkanen, Jaana	02.10.2015 14:51	Accepted

Exit

11

Accept and send results

Help

Post-analytics

- The results can be edited by selecting the correct sample
- When all of the states 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"
- The results can be edited as long as the round is open, even if "Accept and send results" has been selected

Faecal culture, October, 3-2015 postanalytics

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

Va	lida	te	re	su	Its
----	------	----	----	----	-----

▼ Validation results							
Product	Sample set	Sample	Errors	Last saved	Date	Status	
Faecal culture	1	Sample S001	ОК	Paakkanen, Jaana	02.10.2015 15:36	Accepted	
Faecal culture	1	Sample S002	ок	Paakkanen, Jaana	02.10.2015 14:51	Accepted	



Help

LabScala buttons

Save	Enables you to save changes on the form
Back	Takes you back to the previous view
•	Enables you to add some information. In tables it adds a row.
2	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
Q	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
•	To the Home page

Questions?

- In case you have questions, please contact:
 - Jaana Paakkanen
 jaana.paakkanen@labquality.fi
 +358 9 8566 8236
 - Jonna Pelanti
 jonna.pelanti@labquality.fi
 +358 9 8566 8211