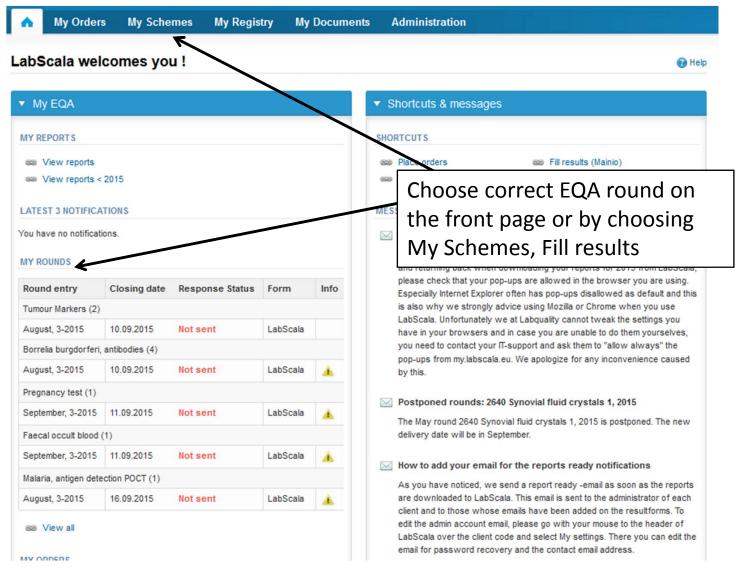


How to fill in results EBV-mononucleosis, spesific antibodies (5641)

Quick guide for result reporting

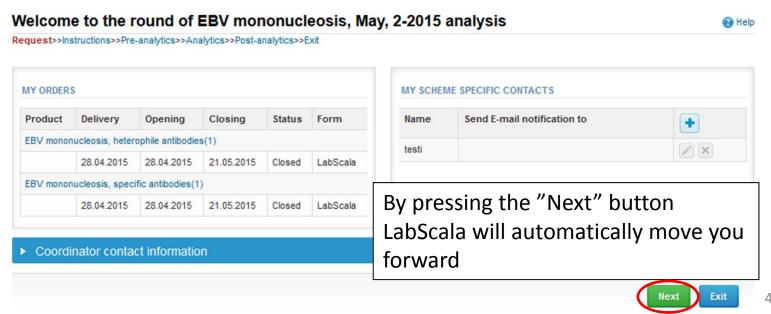
- Start from LabScala home page by choosing My Schemes
 Fill results -> Choose correct round.
- 2. Add Specific Contact, choose Next.
- 3. Read Instructions, choose Next.
- 4. Update Sample registration, choose Save & next
- 5. Reporting results: choose Measurement date, choose Device (method) or if needed, add it. If you can't find your instrumet in the registry, please contact coordinator.
- 6. Save your results as Save as final.
- 7. Validate your results and if needed, update them, finally choose Accept and Send results

Result reporting



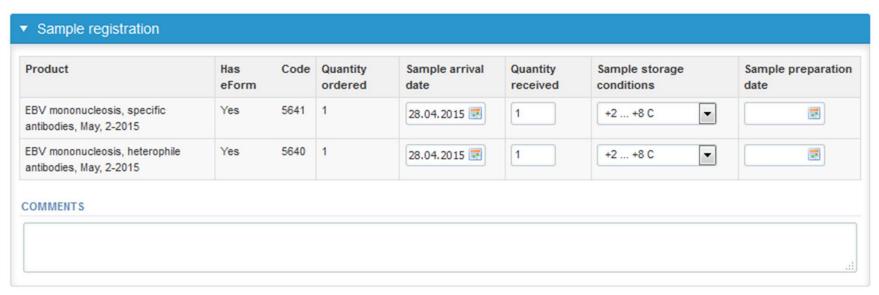
Request

- 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. EBV mononucleosis (1) means 1 set of samples ordered)
 - Instructions: Scheme spesific instructions can be read and printed
 - Pre-analytivs: 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 persons receive reminders and important scheme-specific information
 - At least one contact person needs to be named!



Pre-analytics

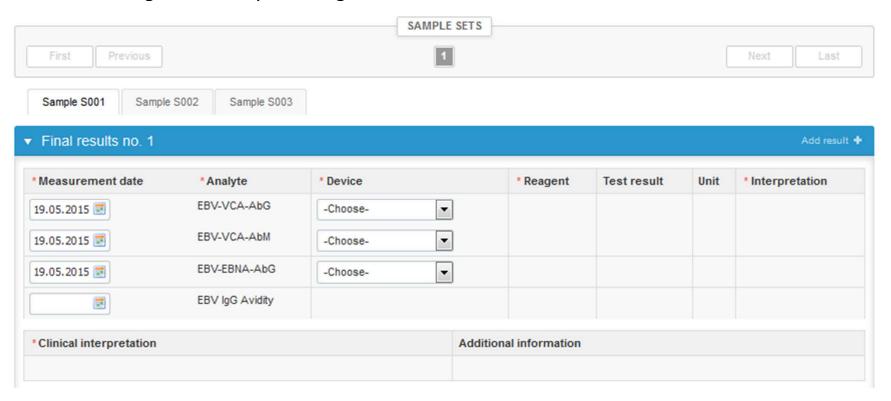
- In Pre-analytics, the following is asked
 - Sample arrival date
 - Quantity received: How many sample sets were received
 - Sample storage condition: How have the samples been stored before analysis (refrigerator, room temperature, freezer, other)
 - Sample preparation date: if done, if not, can be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





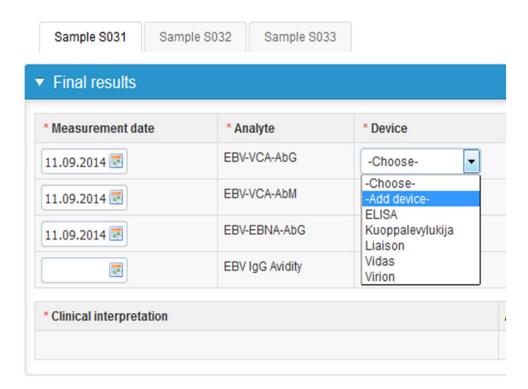
Results – how to start

- In Analytics the analyte-specific results can be filled
- If you have ordered multiple sample sets, the sets can be seen on the top of this view
- If the scheme has multiple sample, the samples are listed on top of the result form (e.g. S001, S002 and S003)
- Result filling is started by selecting "Measurement date"



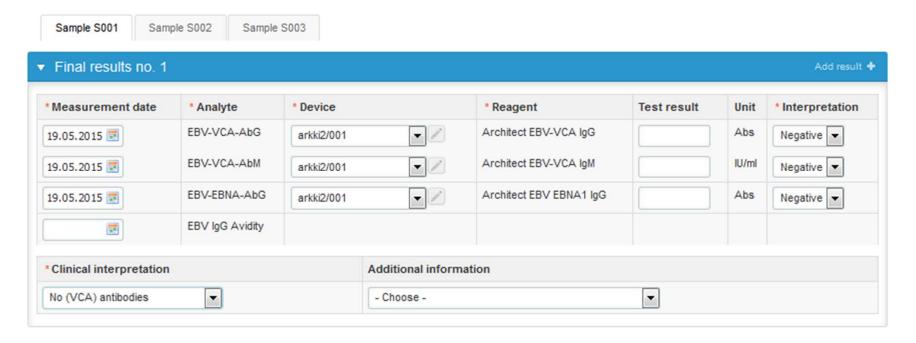
Results – selecting the device

- Analyte-specific results are filled by selecting a previously filled device from the dropdown list or by selecting "-Add device-"
 - Devices can be also added beforehand from My Registry – My devices
- See instructions starting from slide 13.



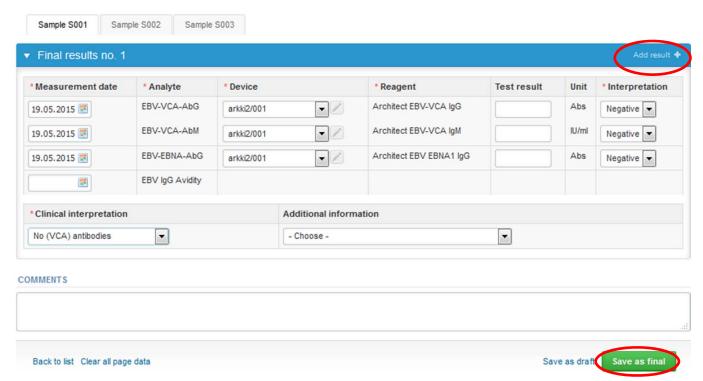
Results

- Saved device can be chosen from the dropdown list in the result form
- "Test result": absorbance or other numerical value if available (not mandatory)
- Choose the correct option from the "Interpretation" (mandatory)
- Choose the correct option from "Clinical interpretation" (mandatory)
- Choose the correct option from "Additional information" (not mandatory)



Results

- If you want to report additional results from the same specimen, please select the "Add result +" on the result form and a new result field will be opened.
- Finally, select
 - "Save as final" if you see the results as being final
 - "Save as draft" if you do not yet want to send in the results. Results are not proccessed in Labquality
- 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"



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

Validate results

▼ \	Vali	dat	ion	resu	its

Product	Sample set	Sample	Errors	Last saved	Date	Status
Preanalytics		Registration	ОК	Paakkanen, Jaana	10.09.2015 10:37	Accepted
EBV mononucleosis, specific antibodies	1	Sample S001	Field "Clinical interpretation" is mandatory in form "Final results no. 1" of sample set no. "1".	Paakkanen, Jaana	10.09.2015 15:02	Error
EBV mononucleosis, specific antibodies	1	Sample S002	ок	Paakkanen, Jaana	10.09.2015 13:19	Accepted
EBV mononucleosis, specific antibodies	1	Sample S003	ок	Paakkanen, Jaana	10.09.2015 13:19	Accepted

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

Validate results

Product	Sample set	Sample	Errors	Last saved	Date	Status
Preanalytics		Registration	ок	Paakkanen, Jaana	10.09.2015 10:37	Accepted
EBV mononucleosis, specific antibodies	1	Sample S001	ок	Paakkanen, Jaana	10.09.2015 15:03	Accepted
EBV mononucleosis, specific antibodies	1	Sample S002	ок	Paakkanen, Jaana	10.09.2015 13:19	Accepted
EBV mononucleosis, specific antibodies	1	Sample S003	OK	Paakkanen, Jaana	10.09.2015 13:19	Accepted

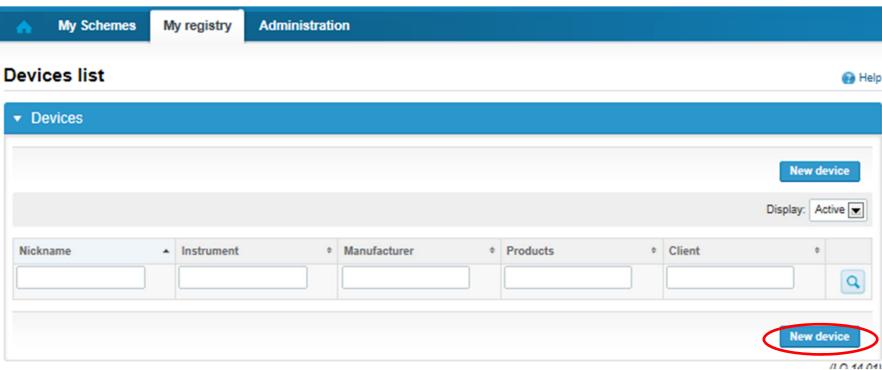
Adding your own devices

- There are two ways to add devices in LabScala:
- 1) By selecting "-Add device-" in the "Final results" page
- 2) By choosing "My devices" under "My registry" in the top header you can fill in your device information even before a round is opened for result filling



Own devices

- Both choices takes you to your "Devices list" where you are able to see all your devices that you have saved in LabScala
- If you have saved no devices, the list is empty
- Adding devices is done by selecting "New device"

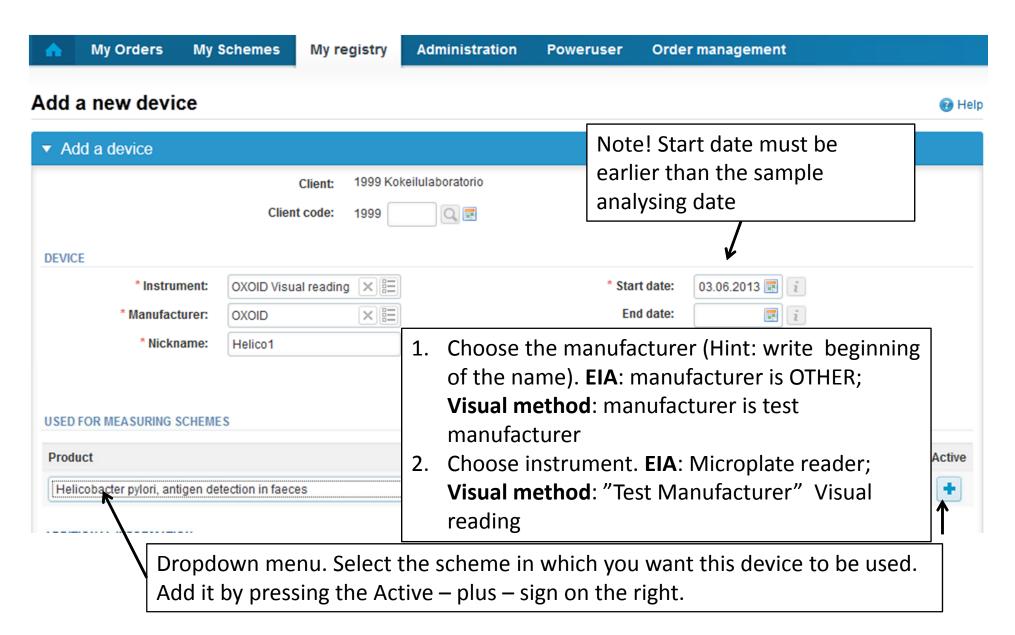


Add a new device Help Add a device Client Client code Access to the background register DEVICE Start date: * Instrument: End date: Manufacturer: Location: * Nickname: lient system ID: Free text – fields. If something is written, LabScala searches **USED FOR MEASURING SCHEMES** the background register for matches. Hint! If you write a part of your instrument name you do not Product Active need to fill in the manufacturer, LabScala does it for you! Hormones A -ADDITIONAL INFORMATION Dropdown menu. Select the scheme in which you want this device to be used. Add it by pressing the Active – plus – sign on the right.

NB! * - marked fields are mandatory!

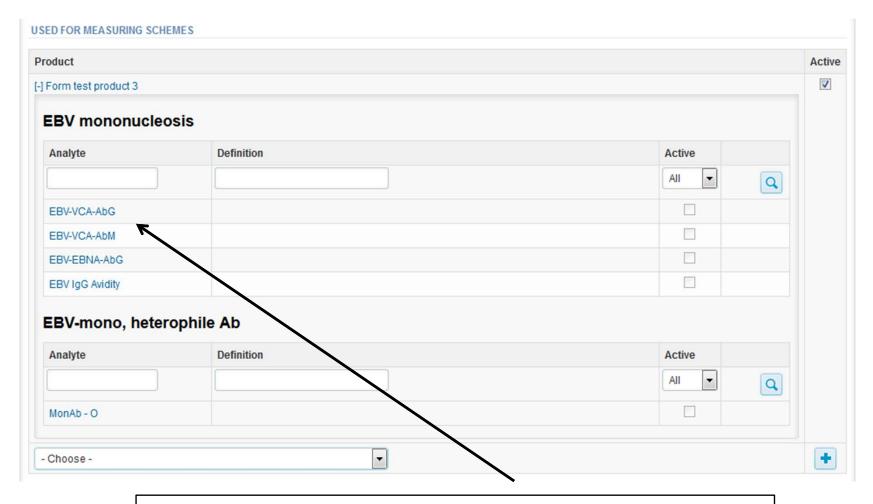
Back

Save



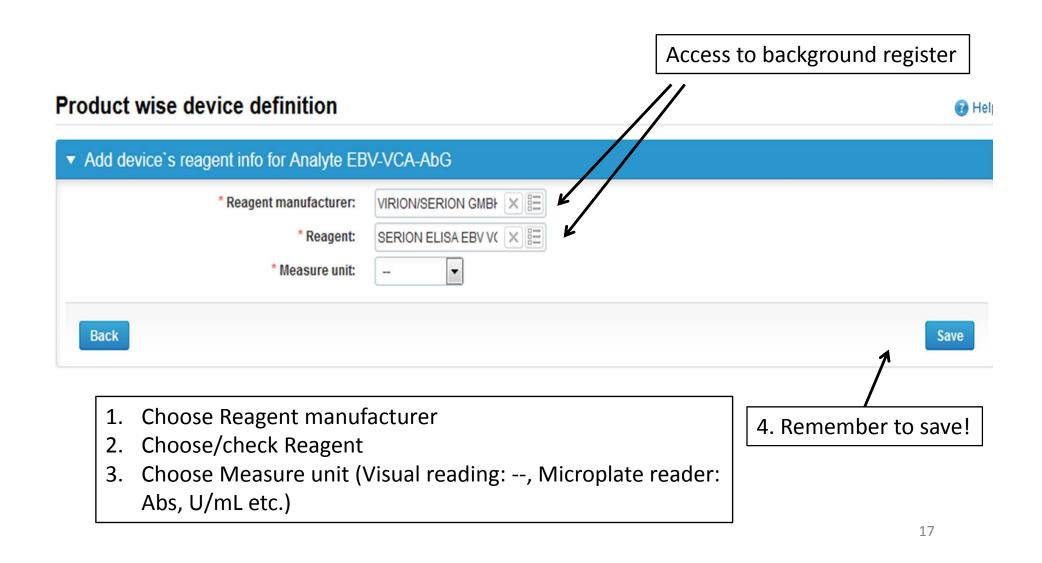
NB! * - marked fields are mandatory!

Adding reagent information

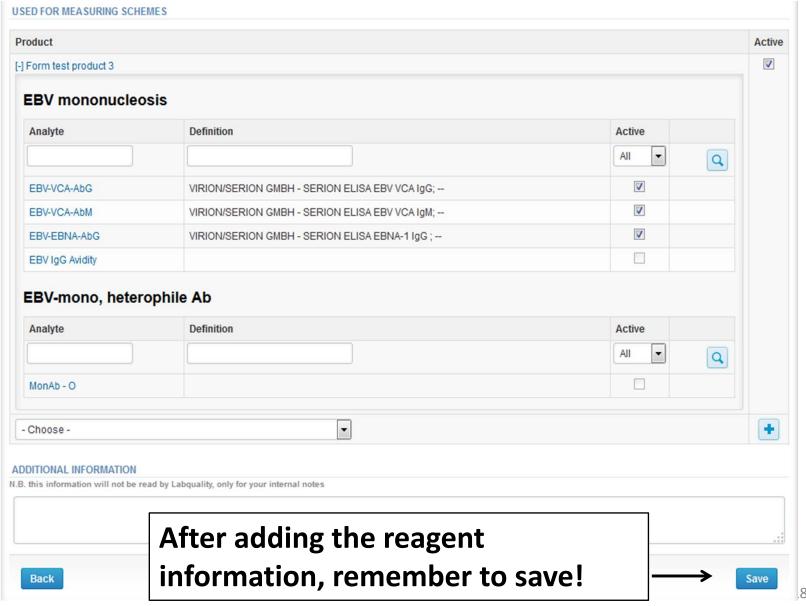


1. Add the analyte-specific reagent info by clicking the analyte

Adding reagent information

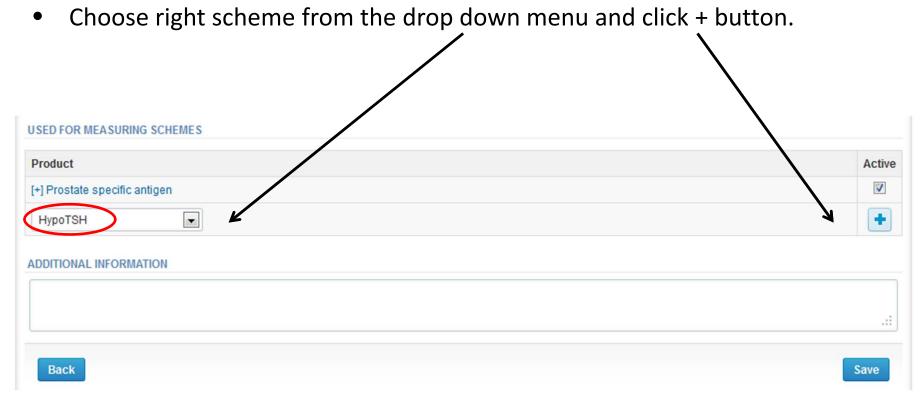


Saving device information

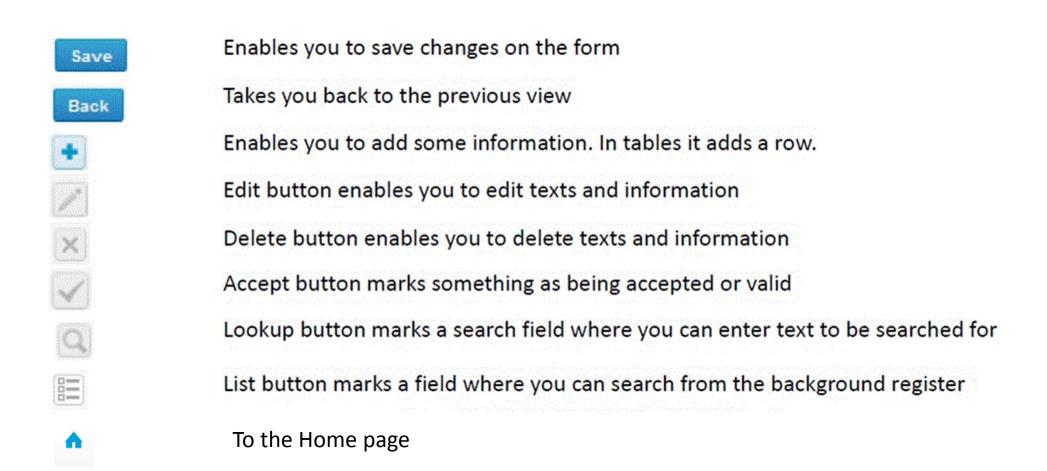


Own devices

- If you have used the same device already on other schemes, just add new scheme and analytes for that device from the dropdown list.
- Open device from the edit button on the right (pencil)



LabScala buttons



Questions?

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