News from Scandiatransplant office

June 2020

Introduction

Headlines

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All previous newsletters can be found on the Scandiatransplant web page
http://www.scandiatransplant.org/news/newsletters

Content

Purpose

By this information letter, we wish to communicate to you about status and progress related to the database, collaboration with groups related to Scandiatransplant and on-going working projects.

We hope that you will read it and share the information with whom it might concern.

Do not hesitate to contact us for further information, ideas, problems and help.

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Test date uncertainty on Covid-19 infected patients post transplant

On patients that are Covid-19 positive post transplant the test date is obligatory to register in YASWA. However, we have experienced that in some cases the date is difficult to get hold of.

If this is the case, it is now possible to do the registration and at the same time tick of the new box 'Date uncertainty'.

Change in registration of selected medical values on deceased donors

Entry of data has been changed in the 'medical info' tab on systolic blood pressure, diastolic blood pressure, MAP, pulse rate, diuresis and temperature.

Registration of these parameters have been moved to the bottom of the page and you are now able o register several results (date + time) and follow the development during hospitalization.

When you wish to add a new result, click on 'Add new':
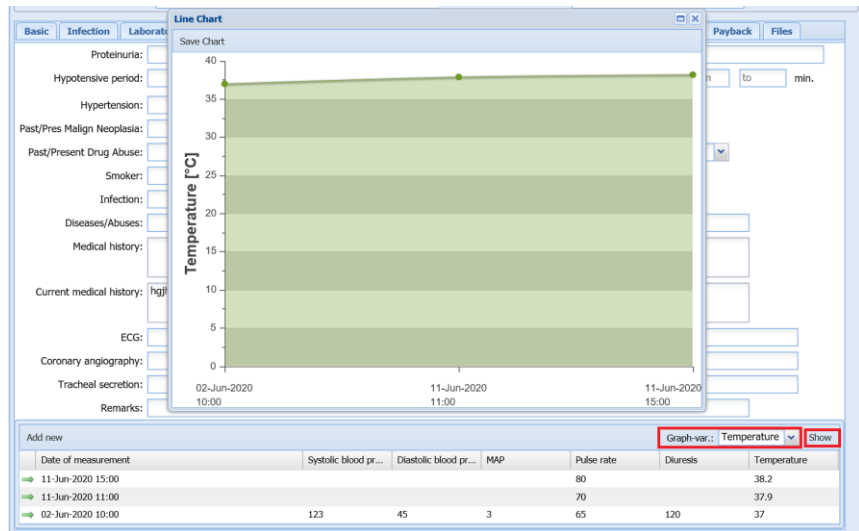
Possibility to add several select medical results

Include label in predefined extractions

In some of the predefined extractions only codes are included, you now have the option to add the label, this is done by ticking of the box in the report execute screen:

If you want to see the development graphical you select a parameter and click on show:

Fusion, transfer previous conclusions

Before a STEP match run the tissue typing labs must import updated Fusion MFI data on their recipients and each time, they need manually to register conclusions that deviate from the default conclusion (everything above MFI of 2000 is positive and everything below negative).
Now a new button/functionality has been added, it will bring forward data from the previous sample where the conclusions are different from default setting 'M: MFI cut-off'. After bringing forward the previous conclusions the user will be able to modify and save the conclusions on the new sample.

Help with transferring previous conclusions

Example…
Several Fusions results have been imported:

On the sample from 12-nov-2019 more antigens have manually been set as unacceptable (the AB list is longer and if you enter the sample you see values where status is different from 'M: MFI cut-off')

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To transfer these results to the newest/next sample you enter the newest/next sample and click on the new 'Add conclusions' button

You will now be able to modify and save data

Next STEP match run

As previously announced the next STEP match run will take place September 1st, 2020.

Deadlines for the upcoming STEP match run:

- Deadline for data entry in YASWA (new pairs and updates) is August 27th, 2020, 10:00 CET
- Scandiatransplant office will do a quality check on data and send a list with pairs to be confirmed by each center August 27th, 2020
- Confirmation of pairs must be done to the Scandiatransplant office no later than August 31st, 2020, 12:00 CET
- STEP match run is made September 1st, 2020.

Summer 2020

You are (almost) on your own 😉

In July user support will be limited, due to summer vacation among the employees at the Scandiatransplant office.

We wish you all a very nice summer 🏖