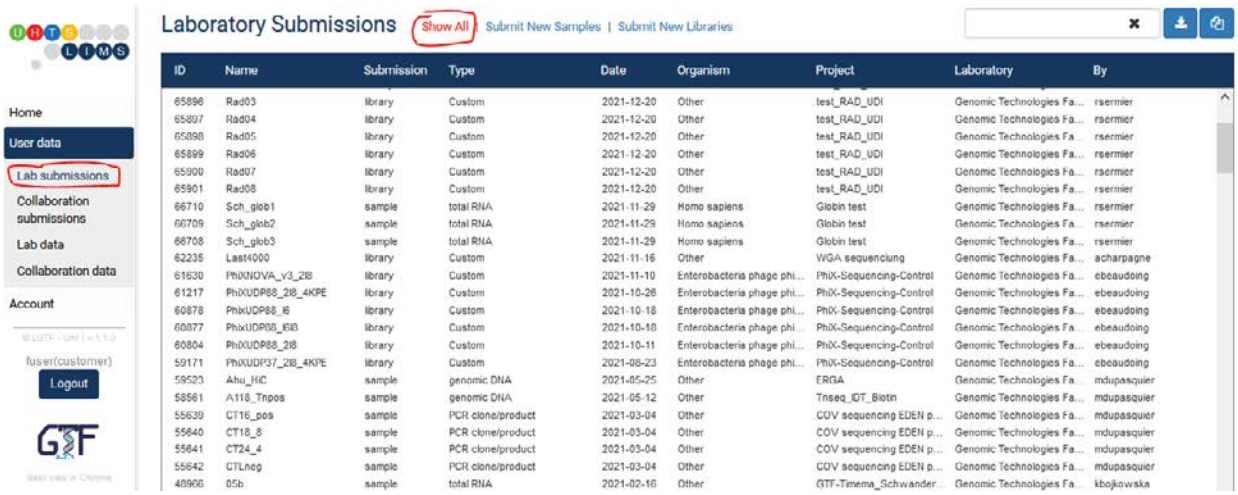
Procedure to request samples/libraries recovery (for Illumina sequencing)

1. Connect to https://uhts-lgtf.unil.ch
2. Go to "Lab Submissions"
3. Click on "Show All"
4. Copy all data with the button
5. Paste into Excel
6. Use standard sorting tools to select the samples/libraries you would like to recover
7. Copy/Paste full corresponding lines into a new xls file
8. Send this file to contactGTF@unil.ch, using "GTF_Recovery_2024_Illumina" as subject



Please take note: if you have submitted samples (RNA, DNA) to the GTF, you will automatically get the resulting libraries back if you put the submitted samples into the excel file for sample recovery.