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

- 1. Connect to <a href="https://uhts-lgtf.unil.ch">https://uhts-lgtf.unil.ch</a>
- 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 <u>contactGTF@unil.ch</u>, using "GTF\_Recovery\_2024\_Illumina" as subject

0000	Laboratory Submissions (Show All) Submit New Samples   Submit New Libraries						X 🕹			
	ID	Name	Submission	Туре	Date	Organism	Project	Laboratory	Ву	
	65896	Rad03	library	Custom	2021-12-20	Other	test_RAD_UDI	Genomic Technologies Fa	rsermier	
me	65897	Rod04	library	Custom	2021-12-20	Other	test_RAD_UDI	Genomic Technologies Fa	rsermier	
r data	65898	Rad05	library	Custom	2021-12-20	Other	test_RAD_UDI	Genomic Technologies Fa	rsermier	
	65899	Rad06	library	Custom	2021-12-20	Other	test_RAD_UDI	Genomic Technologies Fa	rsermier	
submissions	65900	Rad07	library	Custom	2021-12-20	Other	test_RAD_UDI	Genomic Technologies Fa	raermier	
	65901	Rad08	Ibrary	Custom	2021-12-20	Other	test_RAD_UDI	Genomic Technologies Fa	rsermier	
laboration	66710	Sch_glob1	sample	total RNA	2021-11-29	Homo saplens	Globin test	Genomic Technologies Fa	rsermier	
missions	66709	Sch_glob2	sample	total RNA	2021-11-29	Homo sapiens	Globin test	Genomic Technologies Fa	rsermier	
data	66708	Sch_glob3	sample	total RNA	2021-11-29	Homo sapiens	Globin test	Genomic Technologies Fa	rsermier	
	62235	Last4000	library	Custom	2021-11-16	Other	WGA sequenciung	Genomic Technologies Fa	acharpagne	
laboration data	61630	PhDINOVA_v3_28	library	Custom	2021-11-10	Enterobacteria phage phi	PhiX-Sequencing-Control	Genomic Technologies Fa	ebeaudoing	
34	61217	PhiXUDP88_2I8_4KPE	Ibrary	Custom	2021-10-26	Enterobacteria phage phi	PhiX-Sequencing-Control	Genomic Technologies Fa	ebeaudoing	
ount	60878	PhbxUDP88_I6	library	Custom	2021-10-18	Enterobacteria phage phi	PhiX-Sequencing-Control	Genomic Technologies Fa	ebeaudoing	
L017 - 041 ( = 1.1.0	60877	PhixUDP88_I68	library	Custom	2021-10-18	Enterobacteria phage phi	PhiX-Sequencing-Control	Genomic Technologies Fa	ebeaudoing	
	60804	PhiXUDP88_218	library	Custom	2021-10-11	Enterobacteria phage phi	PhiX-Sequencing-Control	Genomic Technologies Fa	ebeaudoing	
r(customer)	59171	PhiXUDP37_28_4KPE	library:	Custom	2021-08-23	Enterobacteria phage phi	PhiX-Sequencing-Control	Genomic Technologies Fa	ebeaudoing	
Locout	59523	Ahu_HiC	sample	genomic DNA	2021-05-25	Other	ERGA	Genomic Technologies Fa	mdupasquier	
Constant of the second s	58561	A118_Tnpos	sample	genomic DNA	2021-05-12	Other	Tnseq_IDT_Biotin	Genomic Technologies Fa	mdupasquier	
	55639	CT16_pos	sample	PCR clone/product	2021-03-04	Other	COV sequencing EDEN p	Genomic Technologies Fa	mdupasquier	
	55840	CT18_8	sample	PCR clone/product	2021-03-04	Other	COV sequencing EDEN p	Genomic Technologies Fa	mdupasquier	
ואַט	55641	CT24_4	sample	PCR clone/product	2021-03-04	Other	COV sequencing EDEN p	Genomic Technologies Fa	mdupasquier	
	55642	CTLneg	sample	PCR clone/product	2021-03-04	Other	COV sequencing EDEN p	Genomic Technologies Fa	mdupasquier	
N YORK IN CAYPING	48966	05b	sample	total RNA	2021-02-16	Other	GTF-Timema Schwander	Genomic Technologies Fa	kbojkowska	

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.