

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

1. Connect to <https://smrt-igt.unil.ch>
2. Go to "User data"
3. Select either "Samples" or "libraries"
4. Select all table lanes and copy
5. Paste into Excel (2 tabs, 1 for input samples, 1 for libraries)
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_PacBio" as subject

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Plateforme GTF

PACIFIC BIOSCIENCES

Home **User data** Tracking Account

Samples submitted from the Lausanne Genomic Technologies Facility lab

Search:

To match the beginning of a word, precede it with '^'. To match the end of a word, follow it with '\$'.

Samples **Libraries**

ID	Name	Type	Date	Organism	Project	By
1507	1_101	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1576	10_110	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1661	100_1104	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1662	101_1105	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1663	102_1106	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1664	103_1107	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1665	104_1108	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1666	105_1110	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1667	106_1201	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1668	107_1204	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1669	108_1207	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne
1670	109_1208	Other	2021-09-15	Other	Amplicons sequencing GTF	acharpagne