09232022

Once run is complete – open the RT-PCR fluidigm software

File open- open BML file

File- export – Table Results with raw data

C:\Users\pbi0\Desktop\CARMEN Analysis\09232022\_2>python 192\_96\_python\_FAM.py --fcount 1691383040 --chip-dim 192 --in-csv 1691383040.csv --in-layout 192\_assignment.xlsx --timepoints 13 --out-dir output

identified crRNAs: 24

identified samples: 192

Traceback (most recent call last):

File "C:\Users\pbi0\Desktop\CARMEN Analysis\09232022\_2\192\_96\_python\_FAM.py", line 253, in <module>

plt\_heatmap(med\_frames, sample\_list, assay\_list, 't18')

File "C:\Users\pbi0\Desktop\CARMEN Analysis\09232022\_2\192\_96\_python\_FAM.py", line 228, in plt\_heatmap

frame = df\_dict[tp][samplelist].reindex(assaylist)

KeyError: 't18'

C:\Users\pbi0\Desktop\CARMEN Analysis\09232022\_2>