HIV\_multiomics

Files verified =

SNF\_cross\_validation.Rmd

Produced files

"processing/cluster\_SNF\_3\_omics\_3\_clusters.csv"

Missing files

"processing/table\_clinical\_data.csv"

"data/clp\_lipidomics\_data.csv"

"data/cocomo\_norm\_data\_filt.csv"

"processing/relative\_abundance\_table.csv"