HIV\_multiomics

Link files

https://doi.org/10.6084/m9.figshare.21120268.v1

https://doi.org/10.6084/m9.figshare.21120271.v1

https://doi.org/10.6084/m9.figshare.21088066.v1

Missing files

MDM

"processing/table\_node\_super\_network\_MDM.csv"

DGE LIMMA

"/home/flomik/Desktop/Code-PHD/3\_layers\_integration/processing/final\_list"

Changes

Data → change “Original\_Scale\_Data\_CLP.xlsx” no IDs → right file : clp\_lipidomics\_data.csv

Data → change “Supplementary Data 1.xlsx” no IDs → right file : cocomo\_norm\_data\_filt.csv

Data → add clp\_um\_new\_data.csv

To clean

Code making network co-expression metabolomics

Code making consensus co-expression network

Check which lipid data used of DGE

Remove one alpha diveristy analysis