

Normalization

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Load libraries and other scripts

Taxonomy agglomeration

```
## $ASV_tissue_V3_B2.csv
## [1] 368 95
##
## $ASV_tissue_V3_B1.csv
## [1] 368 1
##
## $ASV_CVL_V3_B1.csv
## [1] 368 111
```

Merging and renaming

```
## [1] "ASV_tissue_V3_B2.csv" "ASV_tissue_V3_B1.csv" "ASV_CVL_V3_B1.csv"
## $ASV_CVL_V3_B1.csv
## [1] 368 111
##
## $ASV_tissue_V3
## [1] 368 96
```

normalize and save microbiome datasets

```
## $ASV_CVL_V3_B1.csv
## NULL
##
## $ASV_tissue_V3
## NULL
```

normalize and save trx datasets