Raw Counts Data Frame

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```
library(ProjectTemplate)
cwd <- getwd()
setwd("../")
load.project()

## Warning in .load.config(override.config): Your configuration file
## is missing the following entries: data_loading_header, data_ignore,
## logging_level, cache_loaded_data, sticky_variables. Defaults will be used.

## Warning in .check.version(config): Your configuration is compatible with version 0.7 of the ProjectT
## Please run ProjectTemplate::migrate.project() to migrate to the installed version 0.8.
setwd(cwd)
pipeline_dir <- "../../mgtst_pipelines"
mrexp <- get_mrexp(pipeline_dir)</pre>
```

Objective

Generate a tidy dataframe with count data.

```
get_count_df <- function(mrobj, agg_genus = FALSE, css = TRUE){</pre>
      if(agg_genus){
            mrobj <- aggregateByTaxonomy(mrobj, lvl = "Rank6",</pre>
                                           norm = FALSE, log = FALSE, sl = 1)
      }
      if(css == TRUE){
            mrobj <- cumNorm(mrobj, p = 0.75)</pre>
            count_mat <- MRcounts(mrobj, norm = TRUE, log = FALSE, sl = 1000)</pre>
      }else{
            count_mat <- MRcounts(mrobj, norm = FALSE, log = FALSE, sl = 1)</pre>
       count_mat %>%
            as.data.frame() %>%
            rownames_to_column(var = "feature_id") %>%
            gather("id","count", -feature_id)
}
## Converting count matrix to data frame
counts_df <- mrexp %>%
      map_df(get_count_df, css = FALSE, .id = "pipe") %>%
      ## Adding sample metadata
      left_join(pData(mrexp$dada2))
```

```
## Joining, by = "id"
saveRDS(counts_df, "~/Desktop/raw_counts.RDS")
```