Merge featureCounts read count output files into one

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General directory setting

```
wd <- here::here()
shared <- fs::path_dir(wd), "shared")</pre>
```

Loading packages

```
library(magrittr)
library(ggplot2)
```

Load common R scripts

```
#source(fs::path(wd, "script_r", "MISC.R"))
#source(fs::path(here::here(), "script_r", "MISC_PALETTE.R"))
```

Merge featureCounts outputs

```
dirs_output <-
    c(
        fs::path("data_preproc", "readcount", "count_ribo_uorf_psite")
)

for(dir_output in dirs_output) {
    count_files <- fs::dir_ls(fs::path(wd, dir_output), regexp = "counts.txt$")

    count_df <-
        count_files %>%
        purrr::map(ngsmisc::fC_read_count) %>%
        purrr::map(ngsmisc::rename_fpath_bam) %>%
        ngsmisc::fC_merge_count() %>%
        ngsmisc::fC_calc_tpm() %>%
        dplyr::select(-(Chr:Strand))
    readr::write_csv(count_df, fs::path(wd, dir_output, "count_by_gene.csv"))
}
```

Sessioninfo

```
sessionInfo()
```

```
R version 4.2.1 (2022-06-23)
Platform: aarch64-apple-darwin20 (64-bit)
Running under: macOS Ventura 13.1

Matrix products: default
BLAS: /Library/Frameworks/R.framework/Versions/4.2-arm64/Resources/lib/libRblas.0.dylib
LAPACK: /Library/Frameworks/R.framework/Versions/4.2-arm64/Resources/lib/libRlapack.dylib

locale:
[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
```

```
attached base packages:
                                                       methods
[1] stats
              graphics grDevices datasets utils
                                                                 base
other attached packages:
[1] ggplot2_3.4.2 magrittr_2.0.3
loaded via a namespace (and not attached):
 [1] compiler_4.2.1
                         pillar_1.9.0
                                              BiocManager_1.30.18
 [4] tools_4.2.1
                         bit_4.0.5
                                              digest_0.6.31
 [7] jsonlite_1.8.4
                         evaluate_0.20
                                              lifecycle_1.0.3
[10] tibble_3.2.1
                         gtable_0.3.1
                                              pkgconfig_2.0.3
[13] rlang_1.1.0
                         cli_3.6.0
                                              rstudioapi_0.14
                         yaml_2.3.6
[16] parallel_4.2.1
                                              xfun_0.40
[19] fastmap_1.1.0
                         withr_2.5.0
                                              dplyr_1.1.1
[22] stringr_1.5.0
                         knitr_1.42
                                              hms_1.1.3
[25] generics_0.1.3
                         fs_1.5.2
                                              vctrs_0.6.1
[28] ngsmisc_0.4.0
                         bit64_4.0.5
                                              rprojroot_2.0.3
[31] grid_4.2.1
                         tidyselect_1.2.0
                                              glue_1.6.2
[34] here_1.0.1
                         R6_2.5.1
                                              fansi_1.0.3
[37] vroom_1.6.0
                         rmarkdown_2.24
                                              tzdb_0.3.0
[40] readr_2.1.4
                         purrr_1.0.1
                                              scales_1.2.1
                         colorspace_2.0-3
[43] htmltools_0.5.3
                                              renv_1.0.3
[46] utf8_1.2.2
                         stringi_1.7.12
                                              munsell_0.5.0
[49] crayon_1.5.2
```