Results Section: Pipeline Design and Processing 43,000+ genomes

In this notebook will be generating statistics and plots related to processing 43,000+ genomes on Seven Bridges Cancer Genomics Cloud (CGC) platform.

Load Up Packages

```
library(staphopia)
library(ggplot2)
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
## filter, lag

## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
```

Read In The Data

```
results <- read.table("../data/cgc-runs.txt", header = TRUE, sep = "\t")
colnames(results)
## [1] "name"
                      "status"
                                     "project"
                                                   "app"
                                                                  "created_by"
## [6] "total_time" "run_time"
                                     "queue_time" "price"
This leaves use with 9 columns:
  1. name: Name of the job
  2. status: Job's status
  3. project: CGC project job was executed from.
  4. app: CGC app used to execute the job.
  5. created_by: User who submitted the job.
  6. total time: Total amount of time (in minutes) a job was queued and run
  7. run_time: Total amount of time (in minutes) a job took to complete
  8. queue time: Total amount of time (in minutes) a job was queued
  9. price: Total cost of the run
```

Clean Up The Data

Before we generate statistics and plots, we need to clean the data. There are jobs where the *run_time* and *price* were not properly reported from CGC. We will filter samples where the *run_time* is 0.

```
results_clean <- results[results$run_time > 0, ]
nrow(results) - nrow(results_clean)
```

```
## [1] 11424
```

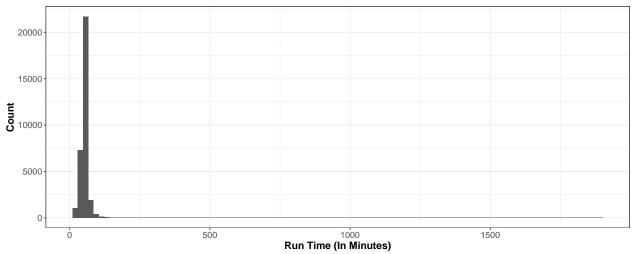
Job Summary

Run Time Summary

```
summary(results_clean$run_time)
##
     Min. 1st Qu. Median
                             Mean 3rd Qu.
##
      9.75
            47.26
                    51.23
                            52.39
                                    56.26 1883.70
Number of Jobs With > 120 Minute Runtime
nrow(results_clean[results_clean$run_time > 120, ])
## [1] 160
Summary of Jobs With Run Time Between 10 and 120 Minutes
summary(results_clean[between(results_clean$run_time, 10, 120), ]$run_time)
##
     Min. 1st Qu. Median
                             Mean 3rd Qu.
                                             Max.
##
     10.83
           47.24
                    51.19
                            51.76
                                    56.11 119.79
```

Plots

Run Time (Complete)



Pipeline Run Time (Between 10-120 Minutes)

```
p <- ggplot(data=results_clean[between(results_clean$run_time, 10, 120),], aes(run_time)) +
    xlab("Run Time (In Minutes)") +
    ylab("Count") +
    geom_histogram(bins=100) +
    theme_bw() +
    theme(axis.text=element_text(size=12),
          axis.title=element_text(size=14,face="bold"))
p
 2500
 2000
 1500
 1000
  500
                                                                          100
                                                                                             125
                                         Run Time (In Minutes)
# Output plot to PDF and PNG
staphopia::write_plot(p, paste0(getwd(), '/images/figure-x-pipeline-run-time'))
```

Session Info

```
sessionInfo()
```

```
## R version 3.4.3 (2017-11-30)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Ubuntu 16.04.2 LTS
##
## Matrix products: default
## BLAS: /usr/lib/libblas/libblas.so.3.6.0
## LAPACK: /usr/lib/lapack/liblapack.so.3.6.0
##
## locale:
## [1] LC_CTYPE=en_US.UTF-8
                                  LC_NUMERIC=C
## [3] LC_TIME=en_US.UTF-8
                                  LC_COLLATE=en_US.UTF-8
## [5] LC_MONETARY=en_US.UTF-8
                                  LC_MESSAGES=en_US.UTF-8
## [7] LC_PAPER=en_US.UTF-8
                                  LC_NAME=C
## [9] LC_ADDRESS=C
                                  LC_TELEPHONE=C
## [11] LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
## attached base packages:
## [1] stats
                graphics grDevices utils
                                              datasets methods
                                                                  base
##
## other attached packages:
## [1] dplyr_0.7.4
                   ggplot2_2.2.1 staphopia_0.1.9
```

```
##
## loaded via a namespace (and not attached):
## [1] Rcpp_0.12.15
                        bindr_0.1.1
                                                          magrittr_1.5
                                         knitr_1.20
## [5] munsell_0.4.3
                        colorspace_1.3-2 R6_2.2.2
                                                          rlang_0.1.6
## [9] stringr_1.2.0
                                                          grid_3.4.3
                        plyr_1.8.4
                                         tools_3.4.3
## [13] gtable_0.2.0
                        htmltools_0.3.6 assertthat_0.2.0 yaml_2.1.18
## [17] lazyeval_0.2.1
                        rprojroot_1.3-2 digest_0.6.15
                                                          tibble_1.4.2
## [21] bindrcpp_0.2
                                         evaluate_0.10.1 rmarkdown_1.9
                        glue_1.2.0
## [25] labeling_0.3
                                         compiler_3.4.3
                                                          pillar_1.1.0
                        stringi_1.1.6
## [29] scales_0.5.0
                        backports_1.1.2 pkgconfig_2.0.1
```