Simulating Time-Varying Performance

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setup

```
library(ggplot2) # for graphing
library(patchwork) # to add graphs together
library(tibble) # tibbles

devtools::load_all()
```

i Loading impulseResponse

Functions used for analysis

```
params_mat <- function(k_1, tau_1, k_2, tau_2, change_days=NULL, days) {</pre>
  if (length(k_1) == length(tau_1) &&
      length(k_1) == length(tau_2) &&
      length(k_1) == length(k_2) &&
      length(k_1) == length(change_days)+1
    else {stop("check length of parameters")}
  out_matrix <- matrix(0, nrow = days, ncol = 4)</pre>
  colnames(out_matrix) <- c("k_1", "tau_1", "k_2", "tau_2")</pre>
  bound_1 <- 1; bound_2 <- days</pre>
  j \leftarrow 0 # counter for index of k_1, tau_1, etc
  for (elem in c(change_days, days)) {
    j < -j + 1
    bound_2 <- elem
    for (i in bound_1:bound_2) {
      out_matrix[i, ] <- c(k_1[[j]], tau_1[[j]], k_2[[j]], tau_2[[j]])</pre>
    }
    bound_1 <- elem
  return(out_matrix)
```

```
#' Title
#'
#' @param params_mat a n by 4 matrix, where n is the number of days
#' @param training_load an n dimensional vector
#' Oreturn an n? dimensional vector with the performance
#' @export
#'
#' @examples
perf_tv <- function(p_0, params_mat, training_load) {</pre>
  days <- nrow(params_mat)</pre>
  perf_out <- c(rep(NA, days))</pre>
 T_1 \leftarrow 0; T_2 \leftarrow 0
  for (i in 1:days) {
    T_1 <- exp(-1/params_mat[i, "tau_1"])*T_1 + training_load[[i]]
    T_2 <- exp(-1/params_mat[i, "tau_2"])*T_2 + training_load[[i]]
    perf_out[[i]] \leftarrow p_0 + params_mat[i, "k_1"]*T_1 - params_mat[i, "k_2"]*T_2
  return(perf_out)
perf_plot <- function(p_0,</pre>
                       k 1,
                        tau 1,
                       k_2,
                        tau_2,
                       change_days = NULL,
                       days,
                       training_stim) {
  training_load <- c()</pre>
  limit <- 0
  if (training_stim[[1]] == "constant") {
    training_load <- c(rep(training_stim[[2]], days))</pre>
    k <- length(k_1)
```

Computing the limit of the model

I would like to compute the limit of the predicted performance for the time-invariant model Under the assumption of constant training load. We have

$$p(t) = p_0 + k_1 \sum_{i=1}^{t-1} e^{\frac{t-i}{\tau_1}} w(i) + k_2 \sum_{i=1}^{t-1} e^{\frac{t-i}{\tau_2}} w(i)$$

Assume that w(i) = C for all i. Note that

$$\sum_{i=1}^{t-1} e^{\frac{t-i}{\tau_1}} = e^{1/\tau_1} \sum_{i=1}^{t-1} e^{\frac{(t-1)-i}{\tau_1}} = e^{1/\tau_1} \left(-1 + \sum_{i=0}^{s} \left(e^{-1/\tau_1} \right)^i \right)$$

Finding the long-run limit of p(t) then amounts to computing

$$\sum_{i=0}^{\infty} \left(e^{-1/\tau_1} \right)^i = \frac{1}{1 - e^{-1/\tau_1}}$$

Notice that this is a convergent geometric series, $e^{-1/\tau_1} < 1$ when $\tau_1 > 1$ (which we have assumed). Therefore,

$$\sum_{i=1}^{t-1} e^{\frac{t-i}{\tau_1}} = e^{1/\tau_1} \left(-1 + \frac{1}{1 - e^{-1/\tau_1}} \right) = e^{1/\tau_1} \left(\frac{e^{-1/\tau_1}}{1 - e^{-1/\tau_1}} \right) = \frac{1}{1 - e^{-1/\tau_1}}$$

Therefore

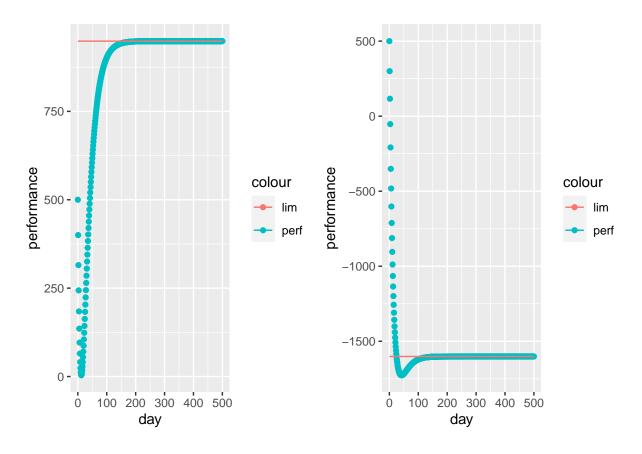
$$\lim_{t \to \infty} p(t) = p_0 + C \left(\frac{k_1}{1 - e^{-1/\tau_1}} - \frac{k_2}{1 - e^{-1/\tau_2}} \right)$$

This is how we compute the red line in perf_plot.

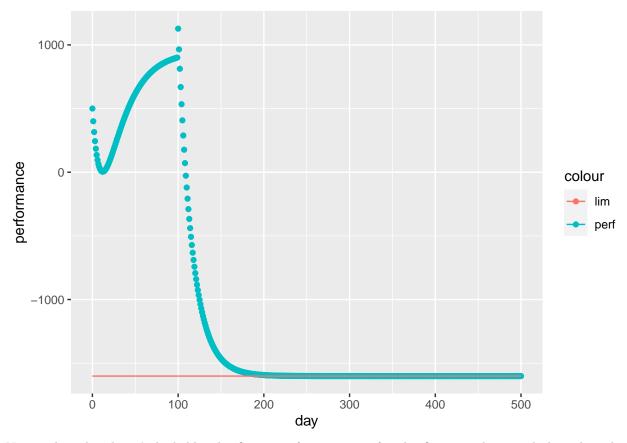
Exploring the time, varying model

This is a time-invariant plot with our new function, to check that things are working correctly.

```
k_1 = 2,
tau_1 = 20,
k_2 = 4,
tau_2 = 15,
days = 500,
training_stim = list("constant", 100))
```

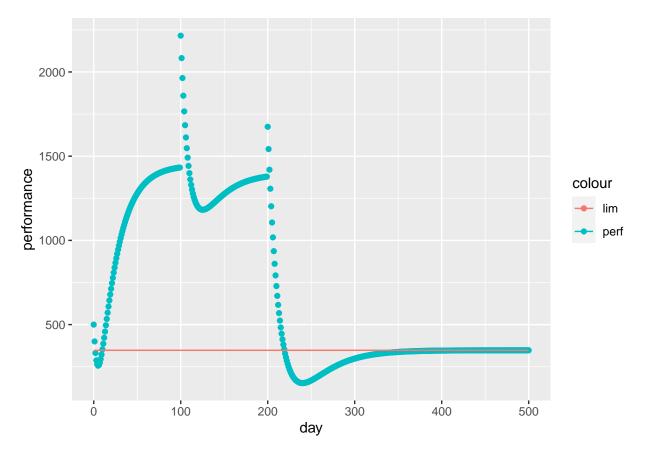


Adding one change date



Notice that this doesn't look like the first set of parameters for the first 100 days, and then the other distribution for the rest of the days. Even though it looks like a new curve at the change day, it is reflecting a change of the parameters.

Adding an additional change dage



To have the effects of an initial negative effect, and a long-run positive effect to a constant stimulus, it seems to be necessary that $k_1 < k_2$ and τ_1 has to be much bigger than

Lets try to use $\frac{k_1}{1-e^{-1/\tau_1}} - \frac{k_2}{1-e^{-1/\tau_2}}$ in the "metric" to evaluate the distance between parameter sets We can write

$$\operatorname{dist}_{\operatorname{perf}}(z,z') = |f(z) - f(z')|$$

where $f:(\mathbb{R}^2 \times \mathbb{R}_{>0})^2 \to \mathbb{R}$ is given by

$$z = (k_1, \tau_1, k_2, \tau_2) \mapsto \frac{k_1}{1 - e^{-1/\tau_1}} - \frac{k_2}{1 - e^{-1/\tau_2}}$$

 $\operatorname{dist}_{\operatorname{perf}}$ is a metric, from the properties of the absolute value, except for the fact that the distance between two points has to be positive, since f is not injective.

Since

$$e^{n/x} \approx 1 - \frac{n}{x}$$

we have that

$$1 - e^{-1/x} \approx 1/x$$

Therefore,

$$f(z) = \frac{k_1}{1 - e^{-1/\tau_1}} - \frac{k_2}{1 - e^{-1/\tau_2}} \approx k_1 \tau_1 - k_2 \tau_2$$

I don't believe that this approximation is very good, but it provides a good intuition about this function.

```
lim_func <- function(k_1, tau_1, k_2, tau_2) {
    k_1/(1-exp(-1/tau_1))-k_2/(1-exp(-1/tau_2))
}

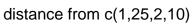
params_dist <- function(params_1, params_2) {
    abs(lim_func(params_1[[1]], params_1[[2]], params_1[[3]], params_1[[4]])-
        lim_func(params_2[[1]], params_2[[2]], params_2[[3]], params_2[[4]])
    )
}</pre>
```

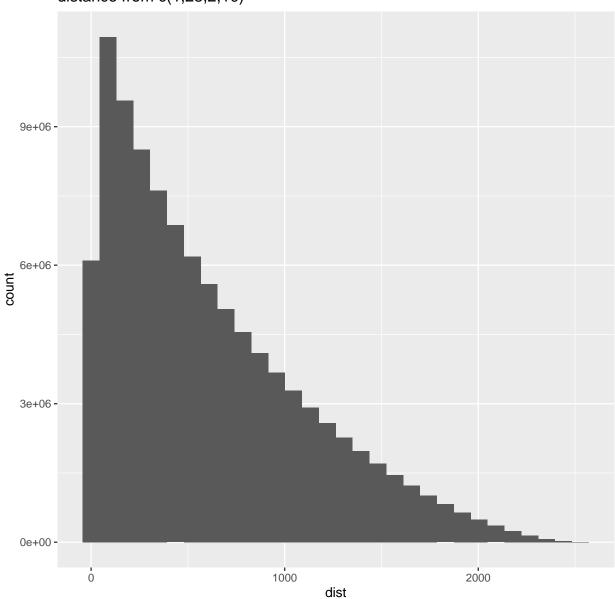
Applying the "norm" to figure out "distance" from a point

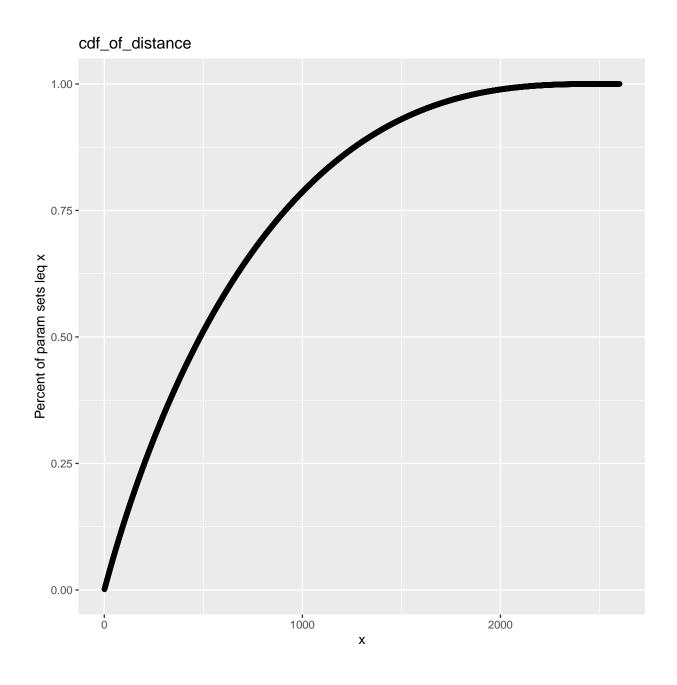
Don't run this chunk locally

```
params_grid <- expand.grid(k_1 = seq(1,50, length.out=100),</pre>
            tau 1 = seq(1,50, length.out=100),
            k_2 = seq(1,50, length.out=100),
            tau_2 = seq(1,50, length.out=100)
params matrix <- as.matrix(params grid)</pre>
dist_vec <- c(rep(0, 100000000))
iter_fn <- function(i, params_matrix) {</pre>
  params_dist(params_matrix[i, ], c(1,25,2,10))
# parallel computing is a factor for this computation, took a few minuites
dist_vec <- parallel::mcmapply(</pre>
  iter_fn,
  i = c(1:100000000),
 MoreArgs = list(params_matrix = params_matrix),
  mc.cores = floor(.9 * parallel::detectCores())
# a 100,000,000 element, 100 MB vector
save(dist_vec, file = stringr::str_c(rprojroot::find_rstudio_root_file(),
                                      "/generated_data/dist_vec.RData"))
# so we don't have to do the computation again
dist_tib <- tibble::tibble(</pre>
  "dist" = dist_vec
plot_hist <- ggplot(dist_tib, aes(x=dist)) +</pre>
  geom_histogram() +
  labs(title = "distance from c(1,25,2,10)")
plot_hist
# so we can call the outputed plot in this markdownfile, to save time.
ggsave(filename = "dist from single set.pdf",
       plot_hist,
```

```
path = stringr::str_c(rprojroot::find_rstudio_root_file(),"/plots"),
       device = "pdf")
# this took awhile
cdf_vec <- c(rep(0, 2601))
for (i in 1:2601) {
  cdf[[i]] <- length(which(dist_vec<=i))</pre>
}
cdf_data <- tibble(</pre>
 x'' = c(1:2601),
  "percent_leq_x" = cdf/100000000
# to save computation time later
save(cdf_vec, file = stringr::str_c(rprojroot::find_rstudio_root_file(),
                                     "/generated_data/cdf_vec.RData"))
cdf_plot <- ggplot(cdf_data, aes(x=x, y= percent_leq_x)) +</pre>
  geom_point() +
  labs(x = "x",
       y = "Percent of param sets leq x",
       title = "cdf_of_distance ")
cdf_plot
ggsave(filename = "cdf_plot.pdf",
       cdf_plot,
       path = stringr::str_c(rprojroot::find_rstudio_root_file(),"/plots"),
       device = "pdf")
```







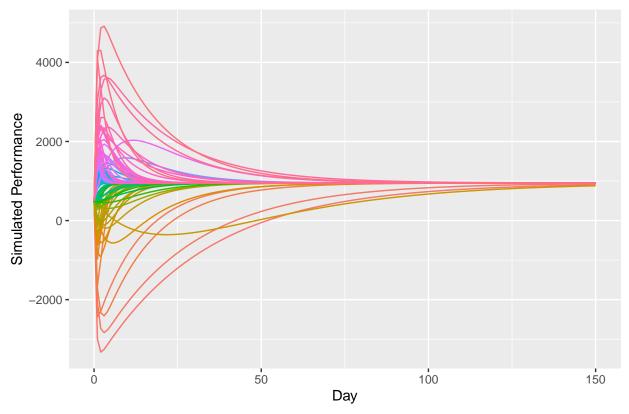
Generating many performance curves with the same limit and plotting $% \left(1\right) =\left(1\right) +\left(1\right)$

```
set.seed(443)
C_1 <- round(lim_func(1,25,2,10), digits = 1)
n <- 100
vec_1 <- runif(n, 5, 50) + C_1
k_1_same <- mapply(function(i) {runif(1, 1, vec_1[[i]])}, c(1:n))
tau_1_same <- -1/log(1-k_1_same/vec_1)
vec_2 <- vec_1-C_1</pre>
```

```
k_2_same <- mapply(function(i) {runif(1, 1, vec_2[[i]])}, c(1:n))</pre>
tau_2_same \leftarrow -1/log(1-k_2_same/vec_2)
day <- 150
p_0 <- 500
training_load <- c(rep(100, day))
same_test_tib <- tibble(</pre>
 "day" = c(0:day)
for (i in 1:n) {
  same_test_tib[, stringr::str_c("p", i)] <- c(p_0,</pre>
                                                 perf_tv(
                                                  p_0 = p_0,
                                                  params_mat = params_mat(
                                                    k_1 = k_1_{same}[[i]],
                                                     tau_1 = tau_1_same[[i]],
                                                    k_2 = k_2 = [[i]],
                                                     tau_2 = tau_2_same[[i]],
                                                    days = day
                                                  ),
                                                  training_load = training_load
                                                 ))
}
pal_color <- scales::hue_pal()(n)</pre>
names(pal_color) <- names(same_test_tib[2:n+1])</pre>
cols <- as.list(names(same_test_tib[2:n+1]))</pre>
min_or_max_vec <- c(rep(0,n))</pre>
for (i in 1:n) {
  if (round(max(same_test_tib[, i+1]), digits = 1) \le 950){
    min_or_max_vec[[i]] <- min(same_test_tib[, i+1])</pre>
  else {
    min_or_max_vec[[i]] <- max(same_test_tib[, i+1])</pre>
  }
}
# reordering the columns so the plot looks nice
ord_min_or_max_vec <- sort(min_or_max_vec)</pre>
ord_min_or_max_vec
##
     [1] -3602.189461 -3327.611469 -2832.581326 -2432.440204 -2408.966002
##
     [6] -1648.104660 -975.945776 -918.660903 -626.968981 -572.658665
## [11] -565.992065 -565.048350 -392.771165 -389.042347 -356.617522
   [16] -335.648616 -188.585342 -170.469789 -113.114856
                                                                -81.874936
##
## [21]
          -28.535047
                         -3.657609
                                     277.377738
                                                   309.045056 332.437935
## [26]
           332.500656 340.168229
                                      365.358637
                                                   398.443765 403.399607
## [31]
           434.195087
                                                  480.863142
                        446.938957
                                      471.477564
                                                                 500.000000
## [36]
           500.000000 500.000000
                                      500.000000 500.000000 500.000000
```

```
##
    [41]
           500.000000
                        500.000000
                                     500.000000
                                                  950.139202
                                                               950.904572
##
   Γ46]
          954.043203 979.344754 1010.548716 1011.162789 1065.364592
##
   [51] 1091.073545 1104.515909
                                    1126.229670 1129.689630 1160.708664
   [56] 1174.448884 1175.673531
                                    1257.279924 1257.542711 1265.364346
##
##
   [61]
         1301.476567 1326.798975
                                    1339.303557
                                                 1368.451567
                                                              1371.699938
##
  [66]
         1454.878860 1501.289621
                                    1528.227053 1542.571439 1586.709490
  [71] 1587.609994 1593.589464
                                    1609.753998 1652.418754 1661.571624
##
## [76] 1677.399624 1682.020849
                                    1684.326918 1759.961127 1781.524675
##
   Г817
         1848.557754 1934.673381
                                    2031.722969
                                                 2049.825880
                                                              2076.976777
##
  [86] 2136.624471 2146.203990
                                    2328.957673
                                                 2358.956663
                                                             2362.264050
##
  [91] 2371.730776 2396.775787
                                    2408.549590
                                                 2607.199206 3102.314686
##
   [96] 3617.734852 3675.639829
                                    4100.450575 4302.409166 4916.749802
permutation_index <- c(rep(NA, n))</pre>
for (i in 1:n) {
  value <- min_or_max_vec[[i]]</pre>
  output_index <- which(ord_min_or_max_vec == value)</pre>
  while(is.na(permutation_index[[i]])) {
    if (output_index[[1]] %in% permutation_index){
      output_index <- output_index[-1]</pre>
   }
    else {
      permutation_index[[i]] <- output_index[[1]]</pre>
   }
 }
}
same_test_tib_new <- same_test_tib</pre>
for (i in 1:n) {
  same_test_tib_new[, permutation_index[[i]] + 1] <- same_test_tib[,i + 1]</pre>
}
plot_same <- ggplot(data = same_test_tib_new, aes(x = day)) +</pre>
    lapply(cols, function(x) {
      geom_line(aes(y = .data[[x]], color = x))
   }) +
    scale_color_manual(values = pal_color) +
 labs(x = "Day",
       y = "Simulated Performance",
       title = "Simulated Performacne of Parameter Sets with Same Limit") +
  theme(legend.position = "none")
plot same
```

Simulated Performacne of Parameter Sets with Same Limit



The colors were sorted by the maximum value. It seems that the maximum value does not determine a convergence rate; there are some curves that peak late, but this generally doesn't happen.

Even though our metric is not truly a metric, it will probably become one when also restrict our curves with minimizing the sum of squared error.

```
p_0=500
k_1 <- seq(1, 50, length.out = 50)
k_2 <- seq(1, 50, length.out = 50)
tau_1 <- seq(1, 50, length.out = 50)
tau_2 <- seq(1, 50, length.out = 50)
days <- 100
training_load_1 <- c(rep(100, days))
params_grid_1 <- as_tibble(expand.grid(k_1, tau_1, k_2, tau_2))
output_matrix <- params_grid_1
output_matrix$performance</pre>
```

Warning: Unknown or uninitialised column: 'performance'.

NULL