# Simulating Time-Varying Performance

Andrew Glover

2024-01-23

#### setup

```
library(ggplot2) # for graphing
library(patchwork) # to add graphs together
library(tibble) # tibbles
devtools::load_all()
## i Loading impulseResponse
## 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
```

## Functions used for analysis

#### Generating the parameters matri

```
params_mat <- function(k_1, tau_1, k_2, tau_2, change_days=NULL, days) {
  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")
bound_1 <- 1; bound_2 <- days
j <- 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]])
    }
    bound_1 <- elem
}
return(out_matrix)
}</pre>
```

#### A note on recursive equations:

All of the time-invariant models here rely on the following recursion equations:

$$p(t) = p_0 + k_1 \cdot g(t) - k_2 \cdot h(t)$$

where g(0) = h(0) = 0 and

$$g(i) = e^{-1/\tau_1} (g(i-1) + w(i)), \quad h(i) = e^{-1/\tau_2} (g(i-1) + w(i))$$

We want to have recursive equations that allow the parameters to change over time. Breaking down either the fitness or the fatigue term in the model, we get that

Fitness(t) = 
$$k_1 \sum_{i=0}^{t-1} e^{t-i} w(i) = k_1 e^{-1/\tau_1} w(t) + k_1 e^{-2/\tau_1} w(t-1) + \dots + k_1 e^{t/\tau_1} w(0)$$

Lets say that at time  $t_{\text{new}}$ ,  $k_1$  changes to  $k_{\text{new}}$  and  $t_1$  changes to  $t_{\text{new}}$ . We would expect that the new model looks something like

Fitness(t) = 
$$k_{new}e^{-1/\tau_{new}}w(t) + \dots + k_{new}e^{-(t-t_{new})/\tau_{new}}w(t_{new}) + k_1e^{-(t-t_{new})/\tau_{new}-1/\tau_1}w(t_{new}-1) + \dots + k_1e^{-(t-t_{new})/\tau_{new}-t_{new}/\tau_{new}}$$

So, I propose a new model for performance:

$$p(t) = p_0 + \sum_{i=0}^{t-1} k_1^i \exp\left(-\sum_{j=i}^t (\tau_1^j)^{-1}\right) w(i) + \sum_{i=1}^{t-1} k_2^i \exp\left(-\sum_{j=i}^t (\tau_2^j)^{-1}\right)$$

where the superscript in the parameters indicates the parameter to be used for that day, i.e.  $k_1^i$  indicates the  $k_1$  parameter on the *i*th day.

I made a naieve implementation of the time varying model, before I wrote the above model for performance down. The code below

```
mat_to_perf_old <- function(p_0, params_mat, training_load) {
  days <- nrow(params_mat)
  perf_out <- c(rep(NA, days))
  T_1 <- 0; T_2 <- 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]])</pre>
```

```
perf_out[[i]] <- p_0 + params_mat[i, "k_1"]*T_1 - params_mat[i, "k_2"]*T_2
}
return(perf_out)
}</pre>
```

fails because it would instead calculates this

$$Fitness(t) = k_{new}e^{-1/\tau_{new}}w(t) + \dots + k_{new}e^{-(t-t_{new})/\tau_{new}}w(t_{new}) + k_{new}e^{-(t-t_{new})/\tau_{new}-1/\tau_1}w(t_{new}-1) + \dots + k_{new}e^{-(t-t_{new})/\tau_1}w(t_{new}-1) + \dots + k_{new}e^{-(t-t_{new})/\tau_1}w($$

Insead of what we want. In works it goes back and changes the  $k_1$  to  $k_{\text{new}}$  on the previous days before the model switched to  $k_n ew$ .

The old implementation relies on this recursive form of the model:

$$p(t) = p_0 + k_1^t g(t) - k_2^t h(t)$$

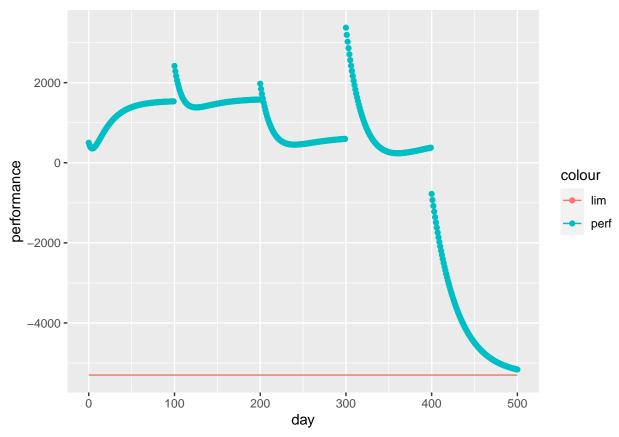
the recursive equations of g(0) = h(0) = 0 and

$$g(i) = e^{-1/\tau_1}(g(i-1) + w(i)) \quad h(i) = e^{-1/\tau_2}(h(i-1) + w(i))$$

What happens in practice with this old recursive equation is that the model jump very rapidly when the parameteters are changed.

If we plot the performance using this old way with the parameter sets

	1-100	101-200	201-300	301-400	401-500
k_1	1	2	3	5	6
k_2	20	25	30	40	40
$tau_1$	2	4	6	8	10
$tau\_2$	5	10	15	25	30

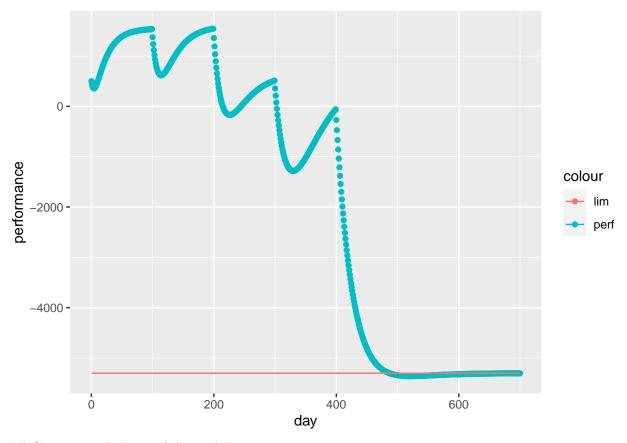


we will get

This method plot the performance in a discontinuous and unnatural way.

#### Improved model

```
mat_to_perf <- function(p_0, params_mat, training_load) {
   days <- nrow(params_mat)
   perf_out <- c(rep(NA, days))
   T_1 <- 0; T_2 <- 0
   for (i in 1:days) {
      T_1 <- exp(-1/params_mat[i, "tau_1"])*(T_1 + params_mat[i, "k_1"]*training_load[[i]])
      T_2 <- exp(-1/params_mat[i, "tau_2"])*(T_2 + params_mat[i, "k_2"]*training_load[[i]])
      perf_out[[i]] <- p_0 + T_1 - T_2
   }
   return(perf_out)
}</pre>
```



## 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

$$p(t) = p_0 + k_1 \sum_{i=0}^{t-1} e^{\frac{t-i}{\tau_1}} w(i) + k_2 \sum_{i=0}^{t-1} e^{\frac{t-i}{\tau_2}} w(i)$$

Assume that w(i) = C for all i. This takes the convention that  $p_0$  happens on day 0. Note that

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

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,

$$\sum_{i=0}^{t-1} e^{\frac{t-i}{\tau_1}} = -1 + \frac{1}{1 - e^{-1/\tau_1}} = \frac{e^{-1/\tau_1}}{1 - e^{-1/\tau_1}}$$

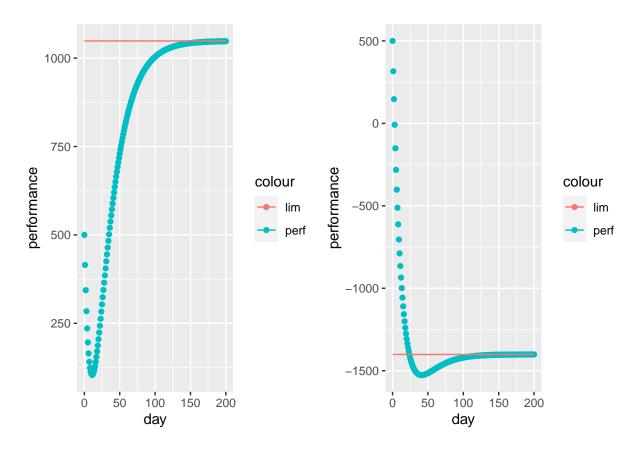
Therefore

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

This is how we compute the red line in perf\_plot.

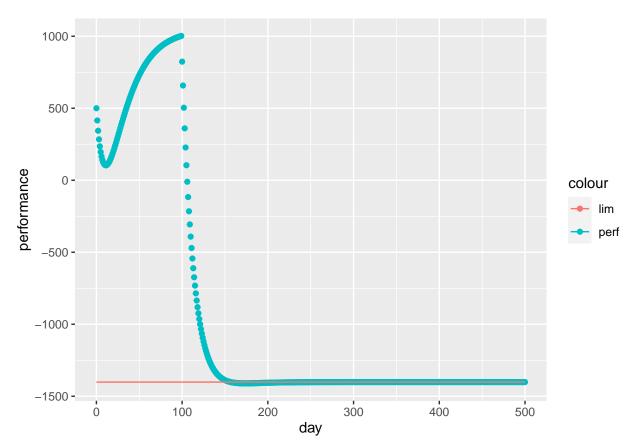
## Exploring the time, varying model

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



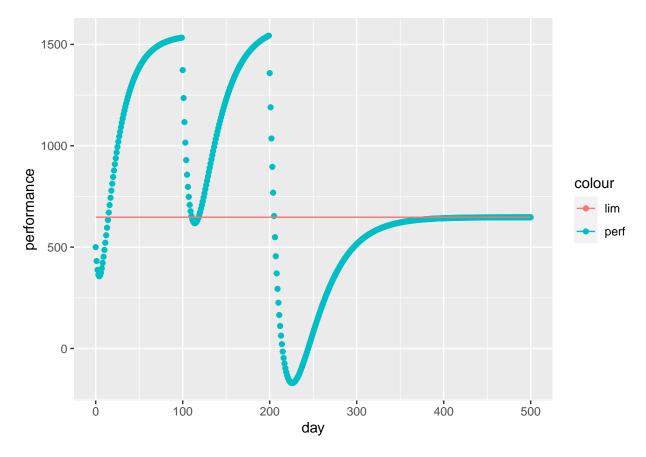
Adding one change date

```
change_days = c(100),
days = 500,
training_stim = list("constant", 100))$plot
```



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  $\tau_1$  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))
}

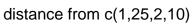
params_dist <- function(params_1, params_2) {
   abs(lim_func(params_1[[1]], params_1[[2]], params_1[[3]], params_1[[4]])-
        lim_func(params_2[[1]], params_2[[2]], params_2[[3]], params_2[[4]])
   )
}</pre>
```

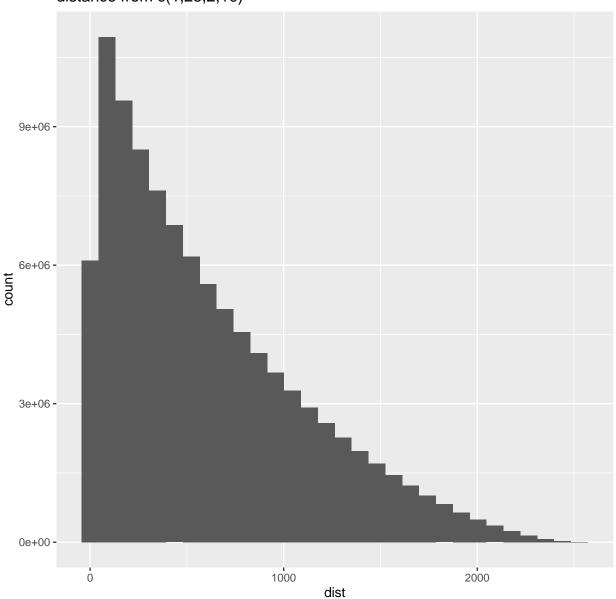
### Applying the "norm" to figure out "distance" from a point

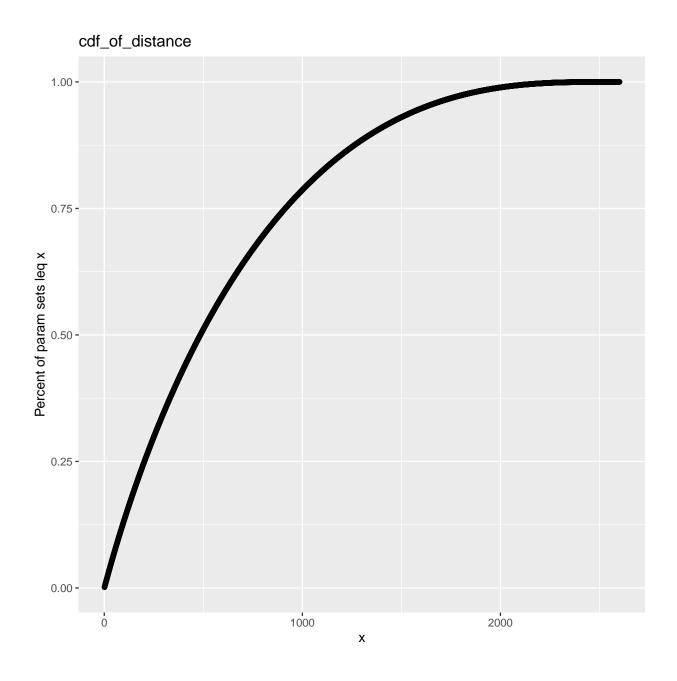
Don't run this chunk locally

```
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 \leftarrow 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))</pre>
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,
                                                        k_1 = k_1_{same}[[i]],
                                                        tau_1 = tau_1_same[[i]],
                                                        k_2 = k_2_{same}[[i]],
                                                        tau_2 = tau_2_same[[i]],
                                                        days = day,
                                                      training_stim = list("constant", 100)
                                                    ))
}
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) <= 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
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
for (i in 1:n) {
    same_test_tib_new[, permutation_index[[i]] + 1] <- same_test_tib[,i + 1]
}

plot_same <- ggplot(data = same_test_tib_new, aes(x = day)) +
    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 Performance of Parameter Sets with Same Limit") +
    theme(legend.position = "none")

plot_same</pre>
```

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 \leftarrow seq(1, 50, length.out = 50)
k_2 \leftarrow seq(1, 50, length.out = 50)
tau_1 \leftarrow seq(1, 50, length.out = 50)
tau_2 \leftarrow seq(1, 50, length.out = 50)
days <- 100
training_load_1 <- c(rep(100, days))</pre>
params_grid_1 <- as_tibble(expand.grid(k_1, tau_1, k_2, tau_2))</pre>
output_matrix <- params_grid_1</pre>
output_matrix$performance
## Warning: Unknown or uninitialised column: 'performance'.
## NULL
for (i in nrow(params_grid_1)){
  invariant_perf(params = c(p_0, params_grid_1[i, ]),
                   training_load = training_load_1)
}
```