

Minimum Fitness

Shashank Pritam

Table of contents

0.1	Introduction	1
0.1.1	Initial conditions	1
0.2	Materials & Methods	1
0.2.1	Commands for the simulation	1
0.3	Visualization in R	2
0.3.1	Data Loading	3
0.3.2	Plot 1	3
0.3.3	Plot 2	4

0.1 Introduction

What is the impact of insertion bias on the minimum fitness of a population during the invasion of transposable elements (TEs), within the parameter space of bias and cluster size?

0.1.1 Initial conditions

0.2 Materials & Methods

version: invadego0.1.3

0.2.1 Commands for the simulation

The simulations were generated using the code from:

- [sim_storm.py](#)

0.3 Visualization in R

0.3.0.1 Setting the environment

```
library(tidyverse)
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr      1.1.2      v readr      2.1.4
v forcats    1.0.0      v stringr    1.5.0
v ggplot2     3.4.2      v tibble     3.2.1
v lubridate  1.9.2      v tidyr      1.3.0
v purrr       1.0.1
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()     masks stats::lag()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
```

```
library(RColorBrewer)
library(ggplot2)
library(patchwork)
library(dplyr)
theme_set(theme_bw())
```

0.3.0.2 Data loading and parsing

```
# Define and load DataFrame with column names
column_names <- c("rep", "gen", "popstat", "spacer_1", "fwte", "avw", "min_w", "avtes", "avp", "avpopfreq", "fixed", "fwcli", "avcli", "X22")
df <- read_delim('/Users/shashankpritam/github/Insertion-Bias-TE/Simulation-Results_Files/simulations/simulations.csv', column_names)
```

```
Rows: 296 Columns: 22
```

```
-- Column specification -----
```

```
Delimiter: "\t"
```

```
chr  (8): popstat, spacer_1, spacer_2, phase, spacer_3, 3tot, 3cluster, spac...
```

```
dbl (13): rep, gen, fwte, avw, min_w, avtes, avpopfreq, fixed, fwcli, avcli,...
```

```
lgl  (1): X22
```

```
i Use `spec()` to retrieve the full column specification for this data.
```

```
i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

```
# Convert specific columns to numeric
numeric_columns <- c("rep", "gen", "fwte", "avw", "min_w", "avtes", "avpopfreq", "fixed", "f
df[numeric_columns] <- lapply(df[numeric_columns], as.numeric)
```

0.3.1 Data Loading

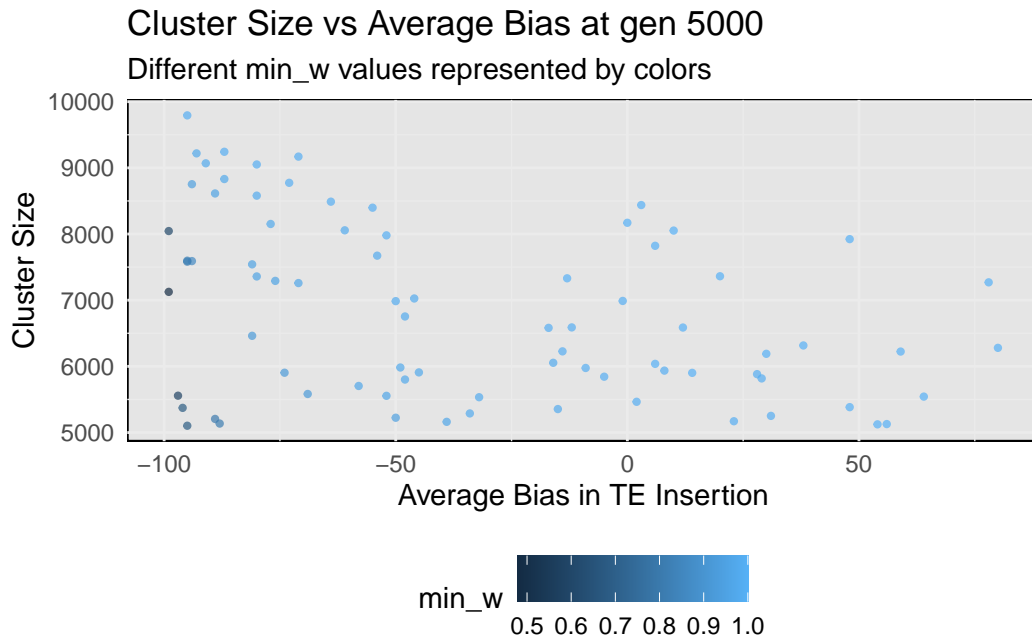
```
# Define color gradient functions
color.gradient <- function(x, colors=c("#D7191C", "#FDAE61", "#A6D96A", "#1A9641"), colsteps=100) {
  # Assign colors based on the 'min_w' column
  df$col <- color.gradient(df$min_w)
  df[df$popstat == "fail-0",]$col <- "grey"
  df$col <- as.factor(df$col)

  # Create and plot the ggplot object
  # Subset the data for gen 5000
  df_gen_5000 <- df[df$gen == 5000,]
```

0.3.2 Plot 1

```
# Plot avbias vs sampleid (Cluster Size) with min_w as color
g_avbias_cluster_size <- ggplot(df_gen_5000, aes(x = avbias, y = sampleid, color = min_w)) +
  geom_point(alpha = 0.7, size = 0.8) +
  ylab("Cluster Size") +
  xlab("Average Bias in TE Insertion") +
  labs(title = "Cluster Size vs Average Bias at gen 5000",
       subtitle = "Different min_w values represented by colors") +
  scale_color_gradient(name = "min_w") +
  theme_minimal() +
  theme(legend.position = "bottom", panel.background = element_rect(fill="grey90"))

# Display the plot
plot(g_avbias_cluster_size)
```



0.3.3 Plot 2

```
library(ggplot2)

# Plot avbias vs sampleid with min_w as color
g_avbias_cluster_size <- ggplot(df_gen_5000, aes(x = avbias, y = sampleid, fill = min_w)) +
  geom_tile() +
  scale_fill_gradient(low = "blue", high = "red", name = "min_w") +
  ylab("Cluster Size") +
  xlab("Average Bias in TE Insertion") +
  labs(title = "Cluster Size vs Average Bias at gen 5000",
       subtitle = "min_w values represented by color gradient") +
  theme_minimal() +
  theme(legend.position = "bottom", panel.background = element_rect(fill="grey90"))

# Display the plot
plot(g_avbias_cluster_size)
```

Cluster Size vs Average Bias at gen 5000

min_w values represented by color gradient

