Supplemental Material for An Exploration of Exploration: Measuring the ability of lexicase selection to find obscure pathways

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Chapter 1

Introduction

This is the supplemental material associated with our 2021 GPTP contribution entitled, An Exploration of Exploration: Measuring the ability of lexicase selection to find obscure pathways. Preprint forthcoming.

1.1 About our supplemental material

This supplemental material is hosted on GitHub using GitHub pages. The source code and configuration files used to generate this supplemental material can be found in [this GitHub repository]. We compiled our data analyses and supplemental documentation into this nifty web-accessible book using bookdown.

Our supplemental material includes the following:

• TODO

1.2 Contributing authors

- Jose Guadalupe Hernandez
- Alexander Lalejini
- Charles Ofria

1.3 Research overview

Abstract:

TODO

Chapter 2

Diagnostic cardinality

2.1 Overview

```
# Relative location of data.
working_directory <-
    "experiments/2021-05-27-cardinality/analysis/"
# working_directory <- "./"

# Settings for visualization
cb_palette <- "Set2"
# Create directory to dump plots
dir.create(paste0(working_directory, "imgs"), showWarnings=FALSE)</pre>
```

2.2 Analysis dependencies

```
library(ggplot2)
library(tidyverse)
library(cowplot)
library(viridis)
library(RColorBrewer)
source("https://gist.githubusercontent.com/benmarwick/2a1bb0133ff568cbe28d/raw/fb53bd97121f7f9ce9
```

These analyses were conducted in the following computing environment:

```
print(version)
...
```

```
## os
                linux-gnu
## system
                x86_64, linux-gnu
## status
## major
               1.0
## minor
## year
               2021
## month
               05
## day
               18
               80317
## svn rev
               R
## language
## version.string R version 4.1.0 (2021-05-18)
## nickname Camp Pontanezen
```

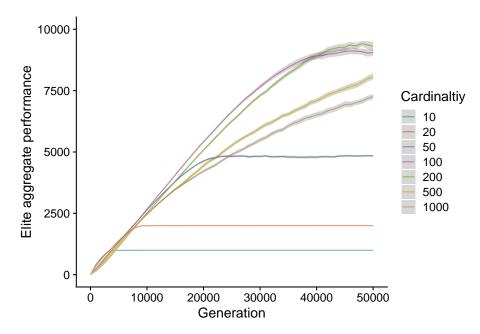
```
data_loc <- paste0(</pre>
  working_directory,
  "data/timeseries-res-1000g.csv"
data <- read.csv(</pre>
  data_loc,
  na.strings="NONE"
data$cardinality <- as.factor(</pre>
  data$OBJECTIVE_CNT
data$selection_name <- as.factor(</pre>
  data$selection_name
data$elite_trait_avg <-</pre>
  data$ele_agg_per / data$OBJECTIVE_CNT
data$unique_start_positions_coverage <-</pre>
  data$uni_str_pos / data$OBJECTIVE_CNT
###### misc ######
# Configure our default graphing theme
theme_set(theme_cowplot())
```

2.4 Performance (max)

Raw aggregate performances. Note that different cardinalities have different score potentials.

```
ggplot(data, aes(x=gen, y=ele_agg_per, color=cardinality)) +
  stat_summary(geom="line", fun=mean) +
  stat_summary(
    geom="ribbon",
    fun.data="mean_cl_boot",
   fun.args=list(conf.int=0.95),
    alpha=0.2,
   linetype=0
  ) +
  scale_y_continuous(
   name="Elite aggregate performance",
   limits=c(0, 10000)
  scale_x_continuous(
   name="Generation"
  ) +
  scale_fill_brewer(
   name="Cardinaltiy",
   palette=cb_palette
  ) +
  scale_color_brewer(
   name="Cardinaltiy",
   palette=cb_palette
  ggsave(
   paste(
      working_directory,
      "imgs/elite_agg_performance_ot.pdf",
      sep=""
   )
 )
```

```
## Saving 6.5 x 4.5 in image
## Warning: Removed 115 rows containing non-finite values (stat_summary).
## Warning: Removed 115 rows containing non-finite values (stat_summary).
## Warning: Removed 115 rows containing non-finite values (stat_summary).
## Warning: Removed 115 rows containing non-finite values (stat_summary).
```

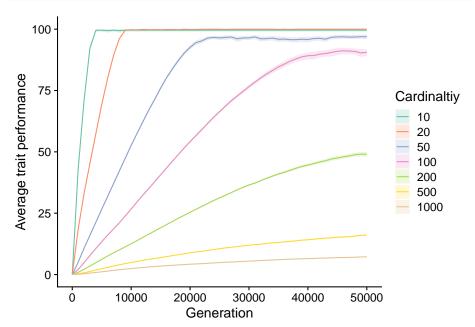


```
elite_trait_ave_fit <- ggplot(</pre>
    data,
    aes(
      x=gen,
      y=elite_trait_avg,
      color=cardinality,
      fill=cardinality
    )
  ) +
  stat_summary(geom="line", fun=mean) +
 stat_summary(
    geom="ribbon",
   fun.data="mean_cl_boot",
   fun.args=list(conf.int=0.95),
    alpha=0.2,
   linetype=0
  ) +
  scale_y_continuous(
    name="Average trait performance",
   limits=c(0, 100)
 ) +
  scale_x_continuous(
    name="Generation"
 scale_fill_brewer(
```

```
name="Cardinaltiy",
   palette=cb_palette
) +
scale_color_brewer(
   name="Cardinaltiy",
   palette=cb_palette
) +
ggsave(
   paste(working_directory, "imgs/elite_trait_average_ot.pdf", sep="")
)
```

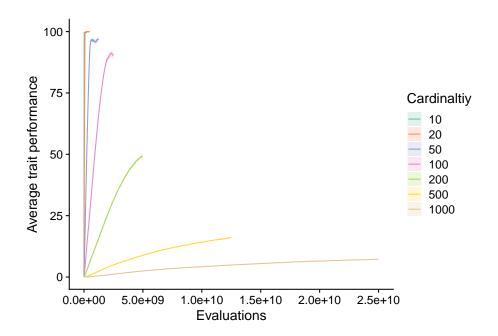
```
## Saving 6.5 \times 4.5 in image
```

```
elite_trait_ave_fit
```



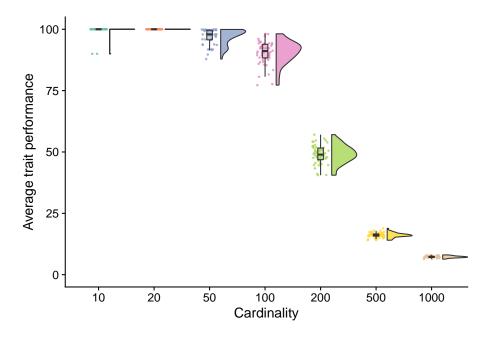
```
ggplot(
   data,
   aes(
      x=evaluations,
      y=elite_trait_avg,
      color=cardinality,
      fill=cardinality
)
) +
stat_summary(geom="line", fun=mean) +
stat_summary(
```

```
geom="ribbon",
  fun.data="mean_cl_boot",
  fun.args=list(conf.int=0.95),
  alpha=0.2,
  linetype=0
) +
scale_y_continuous(
  name="Average trait performance",
  limits=c(0, 100)
) +
scale_x_continuous(
  name="Evaluations"
) +
scale_fill_brewer(
  name="Cardinaltiy",
  palette=cb_palette
) +
scale_color_brewer(
  name="Cardinaltiy",
  palette=cb_palette
)
```



2.4.1 Final performance

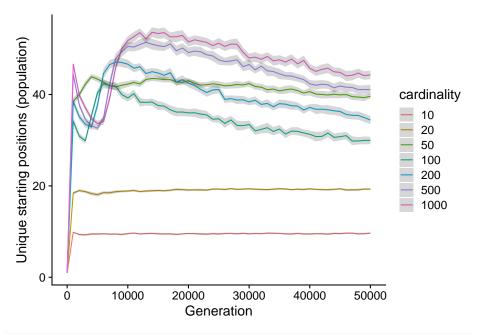
```
final_data <- filter(data, gen==max(data$gen))</pre>
elite_trait_ave_fit_final <- ggplot(</pre>
    final data,
    aes(x=cardinality, y=elite_trait_avg, fill=cardinality)
  geom_flat_violin(
    position = position_nudge(x = .2, y = 0),
    alpha = .8,
    scale="width"
  ) +
  geom_point(
    mapping=aes(color=cardinality),
    position = position_jitter(width = .15),
    size = .5,
    alpha = 0.8
  ) +
  geom_boxplot(
    width = .1,
    outlier.shape = NA,
    alpha = 0.5
  ) +
  scale_y_continuous(
    name="Average trait performance",
    limits=c(0, 100)
  ) +
  scale_x_discrete(
   name="Cardinality"
  scale_fill_brewer(
   name="Cardinaltiy",
   palette=cb_palette
 ) +
  scale_color_brewer(
   name="Cardinaltiy",
   palette=cb_palette
 ) +
    legend.position="none"
elite_trait_ave_fit_final
```



2.5 Unique starting positions (population)

```
ggplot(data, aes(x=gen, y=uni_str_pos, color=cardinality)) +
  stat_summary(geom="line", fun=mean) +
  stat_summary(
    geom="ribbon",
    fun.data="mean_cl_boot",
   fun.args=list(conf.int=0.95),
    alpha=0.2,
   linetype=0
  ) +
  scale_y_continuous(
    name="Unique starting positions (population)",
  scale_x_continuous(
   name="Generation"
  ) +
  ggsave(
    paste(working_directory, "imgs/pop_unique_starting_positions_ot.pdf", sep="")
  )
```

Saving 6.5×4.5 in image



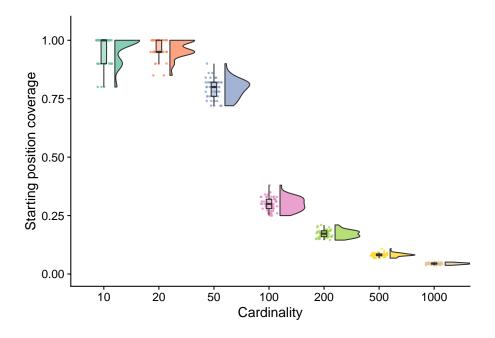
```
unique_start_positions_coverage_fig <- ggplot(data, aes(x=gen, y=unique_start_positions_coverage)
  stat_summary(geom="line", fun=mean) +
  stat_summary(
    geom="ribbon",
    fun.data="mean_cl_boot",
   fun.args=list(conf.int=0.95),
    alpha=0.2,
   linetype=0
  ) +
  scale_y_continuous(
   name="Starting position coverage",
   limits=c(0.0, 1.05)
  ) +
  scale_x_continuous(
   name="Generation"
  ) +
  scale_fill_brewer(
   name="Cardinaltiy",
   palette=cb_palette
  ) +
  scale_color_brewer(
   name="Cardinaltiy",
   palette=cb_palette
  ) +
  ggsave(
```

```
paste(working_directory, "imgs/pop_unique_starting_position_coverage_ot.pdf", sep=
## Saving 6.5 \times 4.5 in image
unique_start_positions_coverage_fig
    1.00
Starting position coverage 0.75 0.50 0.25
                                                                     Cardinaltiy
                                                                        10
                                                                        20
                                                                        50
                                                                        100
                                                                        200
                                                                        500
                                                                        1000
    0.00
                   10000
                             20000
                                        30000
                                                  40000
           Ò
                                                            50000
                               Generation
```

2.5.1 Final coverage

```
final_unique_start_positions_coverage_fig <- ggplot(final_data, aes(x=cardinality, y=)) +
  geom_flat_violin(
    position = position_nudge(x = .2, y = 0),
    alpha = .8,
    scale="width"
) +
  geom_point(
    mapping=aes(color=cardinality),
    position = position_jitter(width = .15),
    size = .5,
    alpha = 0.8
) +
  geom_boxplot(
    width = .1,
    outlier.shape = NA,</pre>
```

```
alpha = 0.5
 ) +
  scale_y_continuous(
    name="Starting position coverage",
    limits=c(0, 1.05)
 ) +
  scale_x_discrete(
    name="Cardinality"
  scale_fill_brewer(
    name="Cardinaltiy",
    palette=cb_palette
 ) +
  scale_color_brewer(
    name="Cardinaltiy",
    palette=cb_palette
 ) +
  theme(
    legend.position="none"
 )
{\tt final\_unique\_start\_positions\_coverage\_fig}
```



2.6 Manuscript figures

```
grid <- plot_grid(</pre>
  elite_trait_ave_fit +
    ggtitle("Performance over time") +
    theme(legend.position="none"),
  elite_trait_ave_fit_final +
    ggtitle("Final performance") +
    theme(),
  unique_start_positions_coverage_fig +
    ggtitle("Start position coverage over time") +
    guides(color = guide_legend(nrow = 1), fill=guide_legend(nrow = 1)) +
    theme(
      legend.position="bottom",
      legend.box="horizontal"
  final_unique_start_positions_coverage_fig +
    ggtitle("Final start position coverage") +
    theme(),
  nrow=2,
  ncol=2,
  rel_widths=c(2,1),
  labels="auto"
save_plot(
 paste(working_directory, "imgs/cardinality-panel.pdf", sep=""),
 base_width=12,
 base_height=10
grid
```

