

Selection on within-individual variation in flowering time in *Lathyrus vernus*

Data preparation

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Contents

Read data from Excel file	1
Remove empty rows and columns, and standardized columns	1
Rename columns	2
Calculate number of seeds per fruit and proportion of seeds preyed in 1988-89	3
Change column types	3
Standardize traits and relativize fitness	3
Merge data for the 3 years	3

Read data from Excel file

```
data_ids_87 <- read_excel("data/edited/individual_characteristics.xlsx",  
                          sheet = "1987")  
data_ids_88 <- read_excel("data/edited/individual_characteristics.xlsx",  
                          sheet = "1988")  
data_ids_89 <- read_excel("data/edited/individual_characteristics.xlsx",  
                          sheet = "1989")
```

Remove empty rows and columns, and standardized columns

```
data_ids_87 <- data_ids_87 %>% drop_na(ID)%>% # Remove second, empty row
# and 3 rows at the bottom (one empty, two where mean and SD were calculated)
select(-c(...3, ...5, ...19, ...22))%>% # Remove 4 empty columns
select(-c(MeanSTD:IntactREL))
data_ids_88 <- data_ids_88 %>% drop_na(ID)%>% # Remove second, empty row
select(-c(...3, ...5)) # Remove 2 empty columns
data_ids_89 <- data_ids_89 %>% drop_na(ID)%>% # Remove second, empty row
select(-c(...3, ...5)) # Remove 2 empty columns
nrow(data_ids_87)
```

```
## [1] 231
```

```
# 231 rows
nrow(data_ids_88)
```

```
## [1] 169
```

```
# 169 rows
nrow(data_ids_89)
```

```
## [1] 96
```

```
# 96 rows
```

Rename columns

```
data_ids_87 <- data_ids_87 %>%
  rename(id = ID, subplot = Subplot, composite_id = Ind, MFD = `Mean (MFD)`,
    skew = Skew, kurt = Kurtosis, LFD = `Max (LFD)`, FFD = `Min (FFD)`,
    dur = `Range (Duration)`, n_fl = `Flower N`, n_fr = Fruits,
    fr_init = `Fruit init (fr/fl)`, n_seed = `Total seeds`,
    n_preyed_seed = `Preyed seeds`,
    fitness = `Intact seeds (fitness)`,
    n_seed_per_fr = `Seeds per fruit`,
    prop_seed_preyed = `Proportion preyed`,
    imp_seed_preyed = `Imputed values for seed number and predation`)
data_ids_88 <- data_ids_88 %>%
  rename(id = ID, subplot = Subplot, composite_id = Ind, MFD = `Mean (MFD)`,
    skew = Skew, kurt = Kurtosis, LFD = `Max (LFD)`, FFD = `Min (FFD)`,
    dur = `Range (Duration)`, n_fl = `Flower N`, n_fr = Fruits,
    fr_init = `Fruit init (fr/fl)`, n_seed = `Total seeds`,
    n_preyed_seed = `Preyed seeds`,
    fitness = `Intact seeds (fitness)`,
    imp_seed_preyed = `Imputed values for seed number and predation`)
data_ids_89 <- data_ids_89 %>%
  rename(id = ID, subplot = Subplot, composite_id = Ind, MFD = `Mean (MFD)`,
    skew = Skew, kurt = Kurtosis, LFD = `Max (LFD)`, FFD = `Min (FFD)`,
    dur = `Range (Duration)`, n_fl = `Flower N`, n_fr = Fruits,
    fr_init = `Fruit init (fr/fl)`, n_seed = `Total seeds`,
```

```
n_preyed_seed = `Preyed seeds`,
fitness = `Intact seeds (fitness)`,
imp_seed_preyed = Imputed)
```

Calculate number of seeds per fruit and proportion of seeds preyed in 1988-89

```
data_ids_88 <- data_ids_88 %>%
  mutate(n_seed_per_fr = ifelse(n_fr==0, NA, n_seed / n_fr),
         prop_seed_preyed = ifelse(n_seed==0, NA, n_preyed_seed / n_seed))
data_ids_89 <- data_ids_89 %>%
  mutate(n_seed_per_fr = ifelse(n_fr==0, NA, n_seed / n_fr),
         prop_seed_preyed = ifelse(n_seed==0, NA, n_preyed_seed / n_seed))
```

Change column types

```
data_ids_87 <- data_ids_87 %>%
  mutate(imp_seed_preyed = as.factor(imp_seed_preyed))
data_ids_88 <- data_ids_88 %>%
  mutate(imp_seed_preyed = as.factor(imp_seed_preyed))
data_ids_89 <- data_ids_89 %>%
  mutate(imp_seed_preyed = as.factor(imp_seed_preyed))
# See if I keep integer values as "double"!
```

Standardize traits and relativize fitness

```
data_ids_87 <- data_ids_87 %>%
  mutate(across(c(MFD:n_fl), scale, .names = "{col}_std"))%>%
  mutate(fitness_rel = fitness / mean(fitness))
data_ids_88 <- data_ids_88 %>%
  mutate(across(c(MFD:n_fl), scale, .names = "{col}_std"))%>%
  mutate(fitness_rel = fitness / mean(fitness))
data_ids_89 <- data_ids_89 %>%
  mutate(across(c(MFD:n_fl), scale, .names = "{col}_std"))%>%
  mutate(fitness_rel = fitness / mean(fitness))
```

Merge data for the 3 years

```
data_ids_87 <- data_ids_87 %>%
  mutate(year = as.integer(1987))
data_ids_88 <- data_ids_88 %>%
```

```

mutate(year = as.integer(1988))
data_ids_89 <- data_ids_89 %>%
  mutate(year = as.integer(1989))
data_ids <- full_join(full_join(data_ids_87,data_ids_88),data_ids_89)
data_ids <- data_ids %>% mutate(year = as.factor(year))

```

```
sessionInfo()
```

```

## R version 4.2.2 (2022-10-31 ucrt)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 22621)
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=English_United States.utf8
## [2] LC_CTYPE=English_United States.utf8
## [3] LC_MONETARY=English_United States.utf8
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United States.utf8
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] moments_0.14.1   lubridate_1.9.0  timechange_0.1.1 readxl_1.4.1
## [5] forcats_0.5.2    stringr_1.5.0    dplyr_1.0.10    purrr_1.0.0
## [9] readr_2.1.3      tidyr_1.2.1      tibble_3.1.8    ggplot2_3.4.0
## [13] tidyverse_1.3.2
##
## loaded via a namespace (and not attached):
## [1] tidyselect_1.2.0   xfun_0.36        haven_2.5.1
## [4] gargle_1.2.1       colorspace_2.0-3 vctrs_0.5.1
## [7] generics_0.1.3     htmltools_0.5.4  yaml_2.3.6
## [10] utf8_1.2.2         rlang_1.0.6      pillar_1.8.1
## [13] withr_2.5.0        glue_1.6.2       DBI_1.1.3
## [16] dbplyr_2.2.1       modelr_0.1.10    lifecycle_1.0.3
## [19] munsell_0.5.0      gtable_0.3.1     cellranger_1.1.0
## [22] rvest_1.0.3        evaluate_0.19    knitr_1.41
## [25] tzdb_0.3.0         fastmap_1.1.0    fansi_1.0.3
## [28] broom_1.0.2        scales_1.2.1     backports_1.4.1
## [31] googlesheets4_1.0.1 jsonlite_1.8.4   fs_1.5.2
## [34] hms_1.1.2          digest_0.6.31    stringi_1.7.8
## [37] grid_4.2.2         cli_3.5.0        tools_4.2.2
## [40] magrittr_2.0.3     crayon_1.5.2     pkgconfig_2.0.3
## [43] ellipsis_0.3.2     xml2_1.3.3       reprex_2.0.2
## [46] googledrive_2.0.0  assertthat_0.2.1 rmarkdown_2.19
## [49] httr_1.4.4         rstudioapi_0.14  R6_2.5.1
## [52] compiler_4.2.2

```