

Should statistics rescue mathematical modelling?

R code

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1 Preliminary functions

```
# PRELIMINARY -----

# Theme for plotting
theme_AP <- function() {
  theme_bw() +
    theme(panel.grid.major = element_blank(),
          panel.grid.minor = element_blank(),
          legend.background = element_rect(fill = "transparent",
                                            color = NA),
          legend.margin=margin(0, 0, 0, 0),
          legend.box.margin=margin(-7,-7,-7,-7),
          legend.key = element_rect(fill = "transparent",
                                     color = NA),
          strip.background = element_rect(fill = "white"))
}

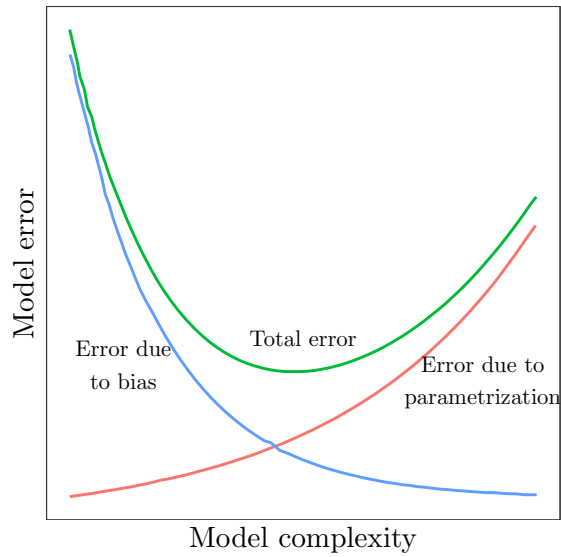
# Load the packages
sensobol::load_packages(c("sensobol", "tidyverse", "data.table", "scales"))
```

2 Figure 1

```
# FIGURE 1 #####

# Read datasets -----
dt <- fread("dataset1.csv")[, type:= "one"]
dt2 <- fread("dataset2.csv")[, type:= "two"]
dt3 <- fread("dataset3.csv")[, type:= "three"]

# Plot -----
rbind(dt, dt2, dt3) %>%
  ggplot(., aes(V1, V2, group = type, color = type)) +
  geom_line(linewidth = 1) +
  labs(x = "Model complexity", y = "Model error") +
  scale_y_continuous(breaks = NULL) +
  scale_x_continuous(breaks = NULL) +
  theme_AP() +
  theme(legend.position = "none") +
  annotate("text", x = 0.15, y = 0.3, label = "Error due \n to bias", size = 3) +
  annotate("text", x = 0.85, y = 0.27, label = "Error due to \n parametrization", size = 3) +
  annotate("text", x = 0.5, y = 0.35, label = "Total error", size = 3)
```



3 Figure 2

```
# FIGURE 2 #####

# Settings -----
N <- 2^8
params <- paste("$x_", 1:2, "$", sep = "")
type <- c("R", "LHS", "QRN")

# Sample matrices -----
mat <- lapply(type, function(x)
  data.table(sobol_matrices(matrices = "A", N = N, params = params, type = x)))
names(mat) <- type

# Plot -----
dt.plot <- rbindlist(mat, idcol = "type")
ggplot(dt.plot, aes(`$x_1$`, `$x_2$`)) +
  geom_point() +
  facet_wrap(~type) +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  scale_y_continuous(breaks = pretty_breaks(n = 3)) +
  theme_AP()
```

