

A critique of irrigation efficiency modeling

R code

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```

# Function to read in all required packages in one go:
loadPackages <- function(x) {
  for(i in x) {
    if(!require(i, character.only = TRUE)) {
      install.packages(i, dependencies = TRUE)
      library(i, character.only = TRUE)
    }
  }
}

# Load the packages
loadPackages(c("data.table", "tidyverse", "sensobol", "wesanderson",
              "cowplot", "parallel", "foreach", "doParallel",
              "countrycode", "ggribes", "scales"))

# Create custom theme
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.key = element_rect(fill = "transparent",
                                     color = NA),
          legend.position = "top",
          strip.background = element_rect(fill = "white"))
}

# Set checkpoint

dir.create(".checkpoint")
library("checkpoint")

checkpoint("2021-08-02",
          R.version = "4.0.3",
          checkpointLocation = getwd())

```

1 Read in data

```
# READ IN DATA -----

# Rohwer data
rohwer <- fread("rohwer_data_all.csv")
rohwer[rohwer == ""] <- NA
rohwer <- rohwer[, Large_fraction := Large_fraction / 100]

# Bos data
bos <- fread("bos_data.csv")
bos <- bos[, Scale := ifelse(Irrigated_area < 10000, "<10.000 ha", ">10.000 ha")]

# Create data set with E_a values as defined by Rohwer
bos.rohwer.ea <- data.table("Irrigation" = c("Surface", "Sprinkler"),
                           "Value" = c(0.6, 0.7),
                           "variable" = "E_a")

# Create data set with E_c values as defined by Rohwer
bos.rohwer.ec <- data.table("Irrigation" = c("Surface", "Sprinkler"),
                           "Value" = c(0.8, 0.95),
                           "variable" = "E_c")

bos.rohwer.all <- rbind(bos.rohwer.ec, bos.rohwer.ea)

# As a function of scale
bos.rohwer.mf.ec <- data.table("Scale" = c("<10.000 ha", ">10.000 ha"),
                              "Value" = c(0.85, 0.59),
                              "variable" = "E_c")

bos.rohwer.mf.ed <- data.table("Scale" = c("<10.000 ha", ">10.000 ha"),
                              "Value" = c(0.81, 0.72),
                              "variable" = "E_d")

bos.rohwer.mf.all <- rbind(bos.rohwer.mf.ec, bos.rohwer.mf.ed)

# PLOT -----

# Field and conveyance efficiency -----

efficiencies_labeller <- c("E_c" = "$E_c$",
                          "E_a" = "$E_a$")

a <- bos %>%
  melt(., measure.vars = c("E_a", "E_c")) %>%
  ggplot(., aes(value, fill = Irrigation, color = Irrigation)) +
  geom_histogram(position = "identity", alpha = 0.4, bins = 15) +
  facet_wrap(~variable, labeller = as_labeller(efficiencies_labeller)) +
```

```

geom_vline(data = bos.rohwer.all, aes(xintercept = Value,
                                     color = Irrigation,
                                     group = variable),
           lty = 2,
           size = 1) +
labs(x = "", y = "Counts") +
theme_AP()

# As a function of scale -----

efficiencies_labeller <- c("E_c" = "$E_c$",
                          "E_a" = "$E_a$",
                          "E_d" = "$E_d$")

b <- melt(bos, measure.vars = c("E_c", "E_a", "E_d")) %>%
na.omit() %>%
ggplot(., aes(value, fill = Scale, color = Scale)) +
geom_histogram(bins = 15, position = "identity", alpha = 0.6) +
labs(x = "Value", y = "Counts") +
scale_fill_manual(values = wes_palette(2, name = "Chevalier1"),
                  name = "Scale") +
facet_wrap(~ variable, labeller = as_labeller(efficiencies_labeller)) +
geom_vline(data = bos.rohwer.mf.all, aes(xintercept = Value,
                                     color = Scale,
                                     group = variable),
           lty = 2,
           size = 1) +
scale_color_manual(values = wes_palette(2, name = "Chevalier1"),
                  name = "Scale",
                  labels = c("<10.000$ ha", ">10.000$ ha")) +
scale_fill_manual(values = wes_palette(2, name = "Chevalier1"),
                  name = "Scale",
                  labels = c("<10.000$ ha", ">10.000$ ha")) +
theme_AP()

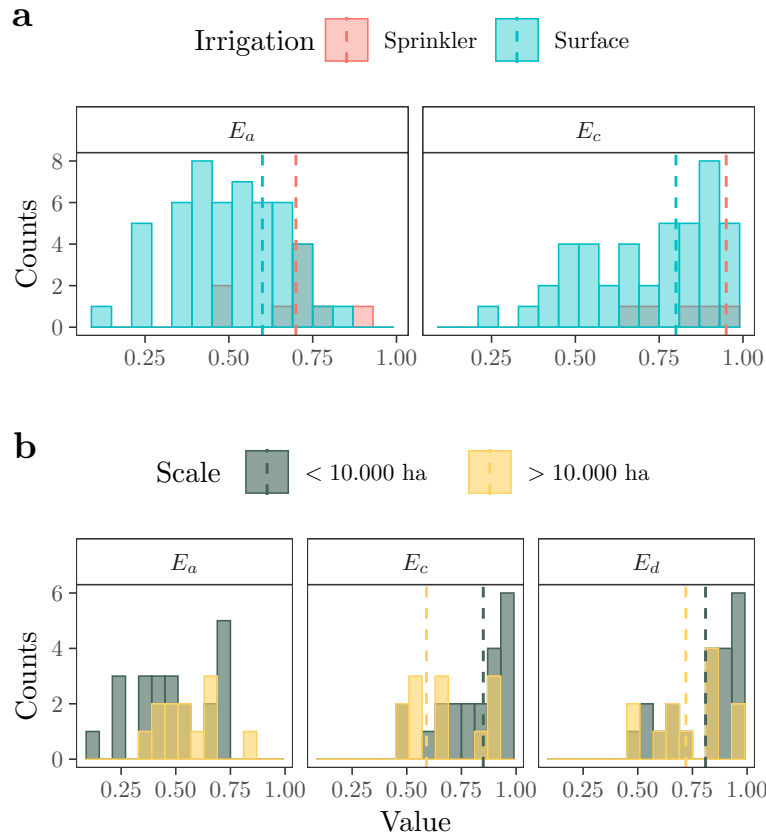
## Scale for 'fill' is already present. Adding another scale for 'fill', which
## will replace the existing scale.

# PLOT MERGED -----

plot_grid(a, b, ncol = 1, labels = "auto")

## Warning: Removed 74 rows containing non-finite values (stat_bin).

```



2 The model

2.1 Function to create sample matrix

```
# CREATE FUNCTION TO DESIGN SAMPLE MATRIX -----

params_algo <- list(
  "Surface" = c("Ea_surf", "Ec_surf", "Proportion_large", "m"),
  "Sprinkler" = c("Ea_sprinkler", "Ec_sprinkler", "Proportion_large", "m"),
  "Micro" = c("Ea_micro", "Ec_micro", "Proportion_large", "m"),
  "Mixed" = c("Ea_surf", "Ea_sprinkler", "Ec_surf", "Ec_sprinkler", "Proportion_large", "m")
)

params_fun <- function(IFT) {
  out <- params_algo[[IFT]]
  return(out)
}

sample_matrix_fun <- function(IFT) {
  params <- params_fun(IFT = IFT)
  mat <- sensobol::sobol_matrices(N = N, params = params)
  out <- list(params, mat)
  names(out) <- c("parameters", "matrix")
}
```

```

    return(out)
}

```

2.2 Define distributions

```

# DEFINE TRUNCATED DISTRIBUTIONS -----

# EA SURFACE -----

Ea.surface <- bos[Irrigation == "Surface"][, .(min = min(E_a, na.rm = TRUE),
                                                    max = max(E_a, na.rm = TRUE))]

shape <- 3.502469
scale <- 0.5444373
minimum <- Ea.surface$min
maximum <- Ea.surface$max
weibull_dist <- sapply(c(minimum, maximum), function(x)
  pweibull(x, shape = shape, scale = scale))

# EC SURFACE -----

Ec.surface <- bos[Irrigation == "Surface"][, .(min = min(E_c, na.rm = TRUE),
                                                    max = max(E_c, na.rm = TRUE))]

shape1 <- 5.759496
shape2 <- 1.403552
minimum.beta <- Ec.surface$min
maximum.beta <- Ec.surface$max
beta_dist <- sapply(c(minimum.beta, maximum.beta), function(x)
  pbeta(x, shape1 = shape1, shape2 = shape2))

# EA SPRINKLER -----

Ea.sprinkler <- bos[Irrigation == "Sprinkler"][, .(min = min(E_a, na.rm = TRUE),
                                                    max = max(E_a, na.rm = TRUE))]

shape.spr <- 6.9913711
scale.spr <- 0.7451178
minimum.spr <- Ea.sprinkler$min
maximum.spr <- Ea.sprinkler$max
weibull_dist_spr <- sapply(c(minimum.spr, maximum.spr), function(x)
  pweibull(x, shape = shape.spr, scale = scale.spr))

# MANAGEMENT FACTOR (m) -----

shape1.m <- 5.759496
shape2.m <- 1.403552
minimum.m <- 0.65
maximum.m <- 1
beta_dist.m <- sapply(c(minimum.m, maximum.m), function(x)

```

```

pbeta(x, shape1 = shape1.m, shape2 = shape2.m))

# FUNCTION TO TRANSFORM TO APPROPRIATE DISTRIBUTIONS -----

distributions_fun <- list(

  # SURFACE IRRIGATION
  # -----

  "Ea_surf" = function(x) {

    out <- qunif(x, weibull_dist[[1]], weibull_dist[[2]])
    out <- qweibull(out, shape, scale)
  },

  "Ec_surf" = function(x) {

    out <- qunif(x, beta_dist[[1]], beta_dist[[2]])
    out <- qbeta(out, shape1, shape2)
  },

  # SPRINKLER IRRIGATION
  # -----

  "Ea_sprinkler" = function(x) {

    out <- qunif(x, weibull_dist_spr[[1]], weibull_dist_spr[[2]])
    out <- qweibull(out, shape.spr, scale.spr)
  },

  "Ec_sprinkler" = function(x) qunif(x, 0.64, 0.96),

  # MICRO (DRIP) IRRIGATION
  # -----

  "Ea_micro" = function(x) out <- qunif(x, 0.75, 0.9),
  "Ec_micro" = function(x) out <- qunif(x, 0.9, 0.95),

  # PROPORTION LARGE
  # -----

  "Proportion_large" = function(x) x,

  # MANAGEMENT FACTOR
  # -----

  "m" = function(x) {
    out <- qunif(x, beta_dist.m[[1]], beta_dist.m[[2]])

```

```

    out <- qbeta(out, shape1.m, shape2.m)
  }
)

```

2.3 Uncertainty in the proportion of large-scale irrigated areas

```

# DEFINE THE UNCERTAINTY IN THE LARGE FRACTION AT THE COUNTRY LEVEL -----

rohwer.frac <- rohwer[, .(Country, Large_fraction)]
rohwer.frac[, `:=` (min = Large_fraction, max = Large_fraction + 0.1)]

countries.list <- split(rohwer.frac, seq(nrow(rohwer.frac)))
names(countries.list) <- rohwer$Country

```

2.4 Function to create sample matrix and transform to appropriate distributions

```

# FULL ALGORITHM TO CREATE SAMPLE MATRIX -----

full_sample_matrix <- function(IFT, Country) {
  tmp <- sample_matrix_fun(IFT = IFT)
  mat <- tmp[["matrix"]]
  temp <- colnames(mat)
  mat <- sapply(seq_along(temp), function(x) distributions_fun[[temp[x]]](mat[, x]))
  colnames(mat) <- temp
  countries.frac <- countries.list[[Country]]
  mat[, "Proportion_large"] <- qunif(mat[, "Proportion_large"],
                                     countries.frac$min, countries.frac$max)

  out <- list(tmp$parameters, mat)
  names(out) <- c("parameters", "matrix")
  return(out)
}

```

2.5 Run the model

```

# FULL MODEL -----

full_model <- function(IFT, Country, sample.size, R) {

  tmp <- full_sample_matrix(IFT = IFT, Country = Country)
  mat <- tmp$matrix

  if(IFT == "Surface") {

    Mf <- mat[, "m"] - 0.5 * mat[, "Proportion_large"]
    y <- mat[, "Ea_surf"] * mat[, "Ec_surf"] * Mf
  }
}

```



```

} else if(IFT == "Sprinkler") {

  Mf <- mat[, "m"]
  y <- mat[, "Ea_sprinkler"] * mat[, "Ec_sprinkler"] * Mf

} else if(IFT == "Mixed") {

  Mf.surf <- mat[, "m"] - 0.5 * mat[, "Proportion_large"]
  y.surf <- mat[, "Ea_surf"] * mat[, "Ec_surf"] * Mf.surf

  Mf.sprink <- mat[, "m"]
  y.sprink <- mat[, "Ea_sprinkler"] * mat[, "Ec_sprinkler"] * Mf.sprink

  y <- 0.5 * y.surf + 0.5 * y.sprink

} else {
  Mf <- mat[, "m"]
  y <- mat[, "Ea_micro"] * mat[, "Ec_micro"] * Mf
}

ind <- sobol_indices(N = sample.size, Y = y, params = tmp$parameters,
                    boot = TRUE, R = R)
out <- list(y, ind)
names(out) <- c("output", "indices")
return(out)
}

```

2.6 Define settings

```
# DEFINE SETTINGS -----
```

```

N <- 2^13
R <- 10^2

```

2.7 Run model

```
# RUN MODEL -----
```

```

y <- mclapply(1:nrow(rohwer), function(x)
  full_model(IFT = rohwer[[x, "IFT"]],
             Country = rohwer[[x, "Country"]],
             sample.size = N,
             R = R),
  mc.cores = detectCores() * 0.75)

```

2.8 Extract model output

```
# EXTRACT MODEL OUTPUT -----  
  
output <- lapply(y, function(x) x[[1]])  
names(output) <- rohwer$Country  
tmp <- lapply(output, data.table) %>%  
  rbindlist(., idcol = "Country") %>%  
  merge(., rohwer[, .(Country, IFT)], all.x = TRUE)  
  
tmp <- tmp[, Continent:= countrycode(tmp[, Country], origin = "country.name",  
                                     destination = "continent")] %>%  
  .[, IFT:= factor(IFT, levels = c("Surface", "Sprinkler", "Micro", "Mixed"))]
```

```
## Warning in countrycode_convert(sourcevar = sourcevar, origin = origin, destination = dest,
```

2.9 Uncertainty analysis

```
# PLOT UNCERTAINTY -----  
  
plot_ggridges <- function(dt, Cont) {  
  pp <- ggplot(dt[Continent == Cont], aes(x = V1, y = fct_reorder(Country, V1),  
                                           fill = IFT), alpha = 0.5) +  
    geom_density_ridges(scale = 2) +  
    labs(x = "Irrigation efficiency", y = "") +  
    facet_wrap(~Continent) +  
    theme_AP() +  
    theme(legend.position = "none")  
  return(pp)  
}  
  
a <- plot_ggridges(dt = tmp, Cont = "Africa") +  
  scale_fill_manual(labels = c("Surface", "Sprinkler", "Mixed"),  
                    values = c("#D8B70A", "#02401B", "#81A88D"),  
                    name = "Irrigation")  
  
b <- plot_ggridges(dt = tmp, Cont = "Americas") +  
  scale_fill_manual(labels = c("Surface", "Mixed"),  
                    values = c("#D8B70A", "#81A88D"),  
                    name = "Irrigation")  
  
c <- plot_ggridges(dt = tmp, Cont = "Asia") +  
  scale_fill_manual(labels = c("Surface", "Micro", "Mixed"),  
                    values = c("#D8B70A", "#A2A475", "#81A88D"),  
                    name = "Irrigation")  
  
d <- plot_ggridges(dt = tmp, Cont = "Europe") +  
  scale_fill_manual(labels = c("Surface", "Sprinkler", "Mixed"),
```

```

        values = c("#D8B70A", "#02401B", "#81A88D"),
        name = "Irrigation")

legend.africa <- get_legend(a + theme(legend.position = "top"))

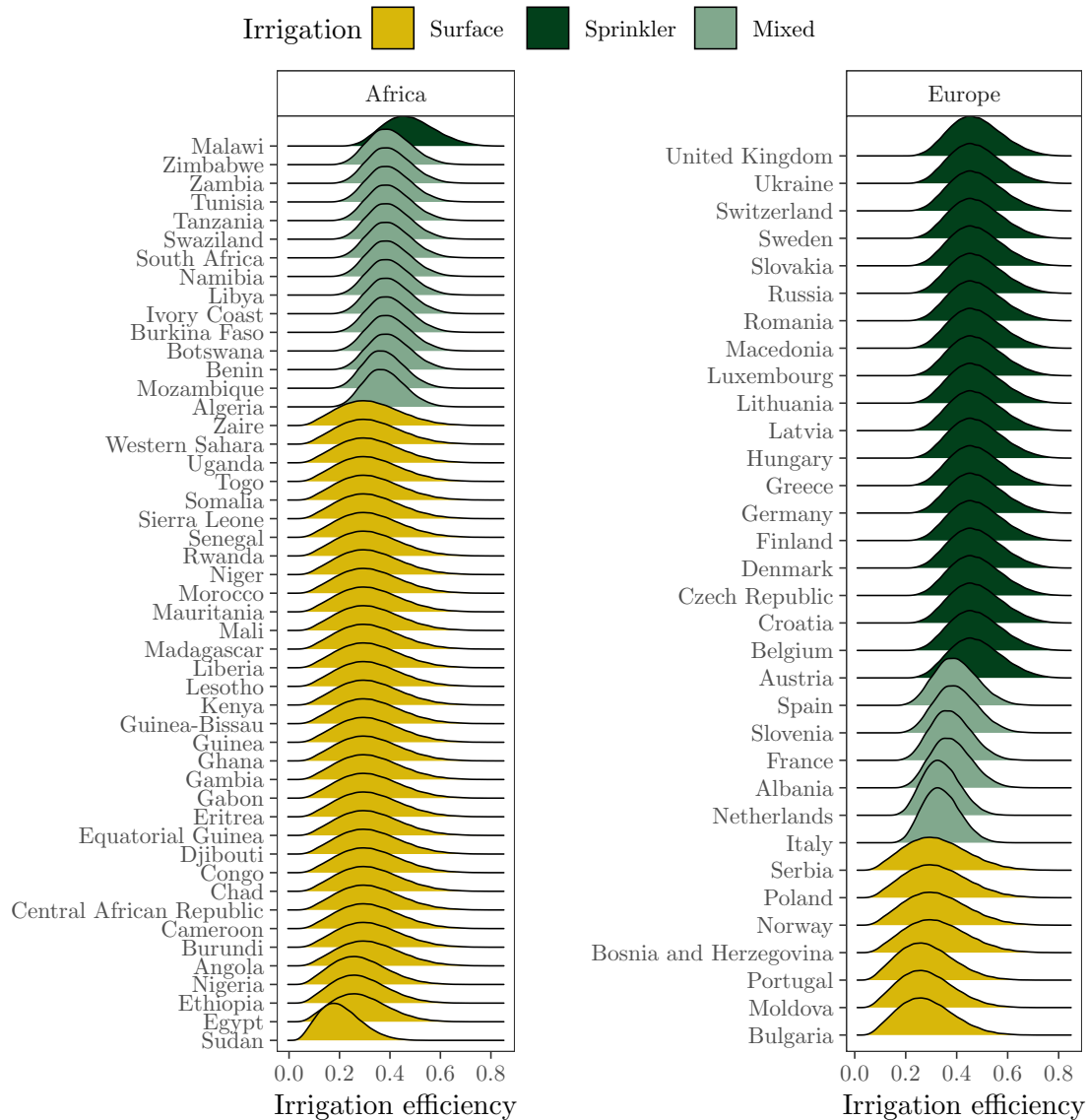
## Picking joint bandwidth of 0.0109
legend.asia <- get_legend(c + theme(legend.position = "top"))

## Picking joint bandwidth of 0.00995
# MERGE PLOTS -----

bottom <- plot_grid(a, d, ncol = 2)

## Picking joint bandwidth of 0.0109
## Picking joint bandwidth of 0.0101
plot_grid(legend.africa, bottom, ncol = 1, rel_heights = c(0.05, 0.95))

```



```
bottom <- plot_grid(b, c, ncol = 2)
```

```
## Picking joint bandwidth of 0.0112
```

```
## Picking joint bandwidth of 0.00995
```

```
plot_grid(legend.asia, bottom, ncol = 1, rel_heights = c(0.05, 0.95))
```

