

Global irrigation water demands biased by unreliable irrigation efficiencies

R code

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1 Preliminary steps

```
# 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", "gggridges", "scales", "overlapping",
              "sp", "rworldmap", "ncdf4", "benchmarkme"))

# 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"),
          plot.margin = margin(t = 0, r = 0.3, b = 0, l = 0.3, unit = "cm"))
}

# Set checkpoint

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

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

2 Read in data

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

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

# Jager data
jager <- fread("jager_data.csv")
jager.list <- split(jager, jager$Country)

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

# Solley data (USA)
usa.dt <- fread("usa_efficiency.csv")
usa.dt <- usa.dt[, Efficiency:= consumptive.use / total.withdrawal]

# FAO 1997 data (Irrigation potential in Africa)
fao_dt <- fread("fao_1997.csv")
fao_dt <- fao_dt[, Efficiency:= Efficiency / 100]

# 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 -----

bos2 <- copy(bos)
bos2 <- setnames(bos2, c("E_a", "E_c", "E_d"), c("E[a]", "E[c]", "E[d]"))

# Field and conveyance efficiency -----

a <- bos2 %>%
  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 = label_parsed) +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  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 -----

b <- melt(bos2, 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 = "Irrigation efficiency", y = "Counts") +
  facet_wrap(~ variable, labeller = label_parsed) +
  geom_vline(data = bos.rohwer.mf.all, aes(xintercept = Value,
                                       color = Scale,
                                       group = variable),
            lty = 2,
            size = 1) +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  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()

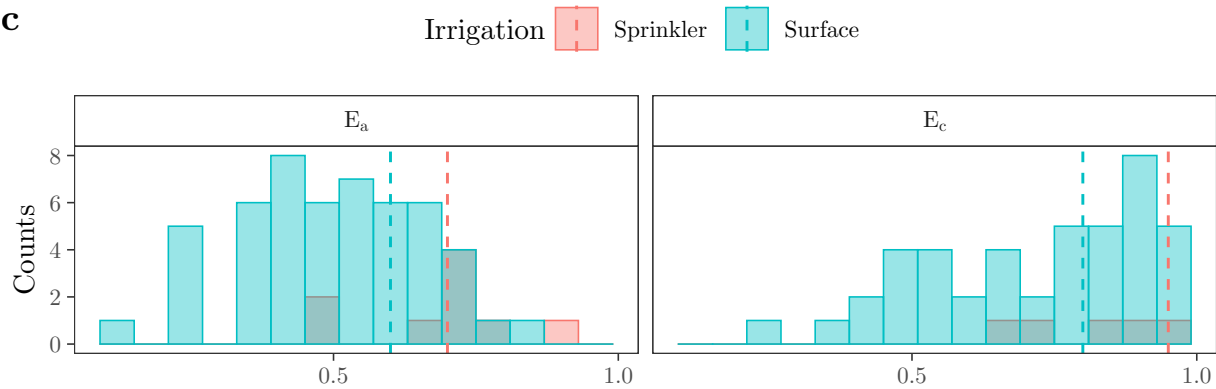
bottom <- plot_grid(a, b, ncol = 1, labels = c("c", "d"))

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

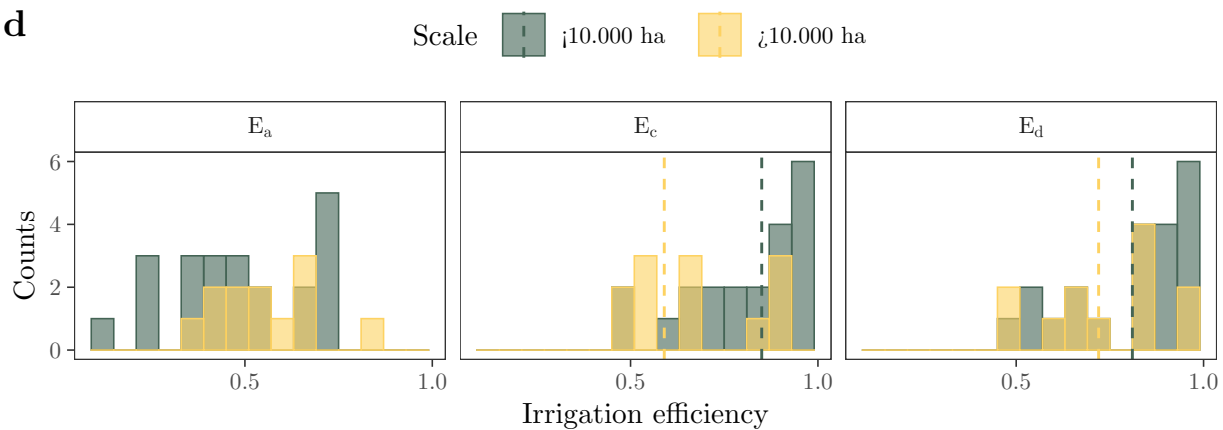
```

bottom

c



d



PLOT USA AND AFRICA -----

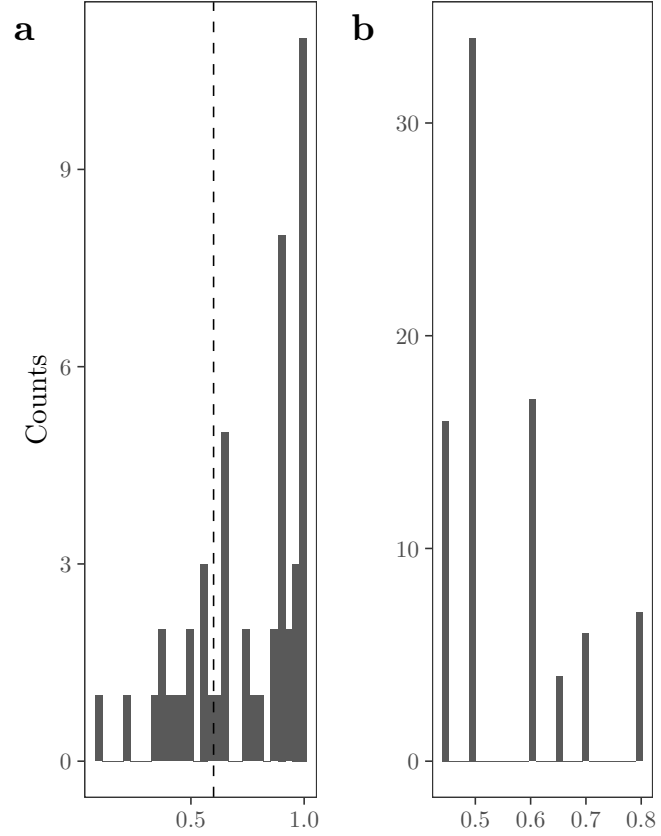
```
c1 <- ggplot(usa.dt, aes(Efficiency)) +
  geom_histogram() +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  geom_vline(xintercept = 0.6, lty = 2) +
  labs(x = "", y = "Counts") +
  theme_AP()

d1 <- ggplot(fao.dt, aes(Efficiency)) +
  geom_histogram() +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  labs(x = "", y = "") +
  theme_AP()

top <- cowplot::plot_grid(c1, d1, ncol = 2, labels = "auto")

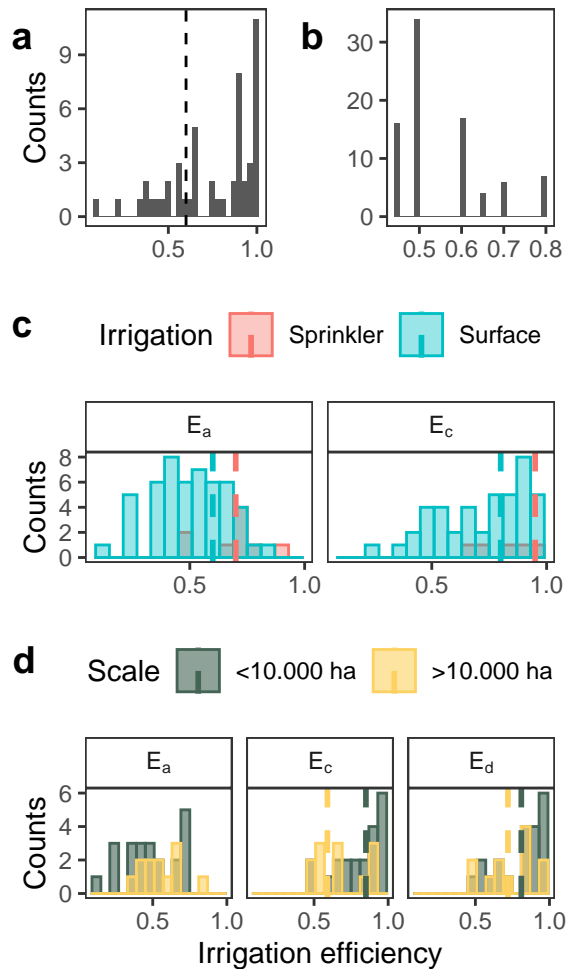
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
## Warning: Removed 3 rows containing non-finite values (stat_bin).
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

top



PLOT MERGED

```
plot_grid(top, bottom, ncol = 1, rel_heights = c(0.3, 0.7))
```



3 The model

3.1 Function to create sample matrix

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

params_algo <- list(
  "Surface" = c("Ea_surf", "Ec_surf", "m", "r_L", "X1", "X2"),
  "Sprinkler" = c("Ea_sprinkler", "Ec_sprinkler", "m"),
  "Micro" = c("Ea_micro", "Ec_micro", "m"),
  "Mixed" = c("Ea_surf", "Ea_sprinkler", "Ec_surf", "Ec_sprinkler", "m", "r_L", "X1", "X2"),
  "Jager" = c("Ea_surf", "Ea_sprinkler", "Ec_surf", "Ec_sprinkler",
              "Ea_micro", "Ec_micro", "m", "r_L", "X1", "X2")
)

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)
}

```

3.2 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, 0.95),
  "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)
},

# REDUCTION IN MANAGEMENT FACTOR DUE TO LARGE-SCALE
# -----
"r_L" = function(x) qunif(x, 0, 0.5),

# IRRIGATED AREA DATASET
# -----

"X1" = function(x) floor(x * (4 - 1 + 1)) + 1,

# THRESHOLD FOR LARGE-SCALE IRRIGATED AREAS
# -----

"X2" = function(x) floor(x * (5 - 1 + 1)) + 1
)

```

3.3 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

n.rows <- nrow(rohwer)
triggers.dt <- rohwer[rep(seq_len(n.rows), 4 * 5)][, .(Country, IFT)] %>%
  .[, X1:= rep(1:4, each = n.rows, times = 5)] %>%
  .[, X2:= rep(1:5, each = n.rows, times = 4)] %>%
  .[, Proportion_large:= runif(n.rows * 4 * 5, min = 0.01, max = 0.7)] %>%
  .[, index:= paste(Country, X1, X2, sep = ".")]

triggers.dt <- setkey(triggers.dt, index)

```

3.4 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]]
  out <- list(tmp$parameters, mat)
  names(out) <- c("parameters", "matrix")
  return(out)
}

```

3.5 Define the model

```

# FULL MODEL -----

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

  country.differences <- setdiff(rohwer$Country, jager$Country)
  tmp <- full_sample_matrix(IFT = IFT, Country = Country)
  mat <- tmp$matrix

  if(IFT == "Surface" | IFT == "Mixed" | IFT == "Jager") {
    X1 <- mat[, "X1"]
    X2 <- mat[, "X2"]
    index <- paste(Country, X1, X2, sep = ".")
    Proportion_large <- triggers.dt[index][, Proportion_large]
  }

  if(IFT == "Surface") {

    Mf <- mat[, "m"] - mat[, "r_L"] * 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"] - mat[, "r_L"] * 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 if(IFT == "Micro") {

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

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

    if(Country %in% country.differences == TRUE) {
      next
    }

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

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

    Mf.micro <- mat[, "m"]
    y.micro <- mat[, "Ea_micro"] * mat[, "Ec_micro"] * Mf.micro

    y <- jager.list[[Country]]$Surface_fraction * y.surf +
      jager.list[[Country]]$Sprinkler_fraction * y.spr +
      jager.list[[Country]]$Drip_fraction * y.micro

  }

  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)
}

```

3.6 Define settings

```

# DEFINE SETTINGS -----

N <- 2^14
R <- 10^2
list_continents <- list(c("Africa", "Asia"), c("Americas", "Europe"))

```

3.7 Run the model in parallel

```

# RUN MODEL -----

```

```

new.rohwer <- rohwer[Country %in% jager$Country][, IFT:= "Jager"]
all.dt <- list(rohwer, new.rohwer)

y <- list()
for(j in 1:length(all.dt)) {
  y[[j]] <- mclapply(1:nrow(all.dt[[j]]), function(x)
    full_model(IFT = all.dt[[j]][[x, "IFT"]],
              Country = all.dt[[j]][[x, "Country"]],
              sample.size = N,
              R = R),
    mc.cores = detectCores() * 0.75)
}

```

3.8 Extract model output

```

# EXTRACT MODEL OUTPUT -----

names(y) <- c("Rohwer et al. 2007", "Jägermeyr et al. 2015")

output <- tmp <- list()
for(i in names(y)) {
  output[[i]] <- lapply(y[[i]], function(x) x[["output"]][1:(2 * N)])

  if(i == "Rohwer et al. 2007") {

    names(output[[i]]) <- rohwer$Country

  } else if(i == "Jägermeyr et al. 2015") {

    names(output[[i]]) <- new.rohwer$Country

  }
  tmp[[i]] <- lapply(output[[i]], data.table) %>%
    rbindlist(., idcol = "Country")

  if(i == "Rohwer et al. 2007") {

    tmp[[i]] <- merge(tmp[[i]], rohwer[, .(Country, IFT)], all.x = TRUE) %>%
      .[, IFT:= factor(IFT, levels = c("Surface", "Sprinkler", "Micro", "Mixed"))]

  } else if(i == "Jägermeyr et al. 2015") {

    tmp[[i]] <- tmp[[i]][, IFT:= "Jager"]

  }

  tmp[[i]] <- tmp[[i]][, Continent:= countrycode(tmp[[i]][, Country],

```

```

                                origin = "country.name",
                                destination = "continent"]
}

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

## Warning in countrycode_convert(sourcevar = sourcevar, origin = origin, destination = dest,
uncertainty.dt <- rbindlist(tmp, idcol = "Approach")
uncertainty.dt <- uncertainty.dt[, Study:= ifelse(IFT == "Jager",
                                                "Jägermeyr et al. approach",
                                                "Rohwer et al. approach")]

# EXPORT UNCERTAINTY IN IRRIGATION EFFICIENCY -----

fwrite(uncertainty.dt, "uncertainty.dt.csv")

```

4 Uncertainty analysis

4.1 Coefficient of variation

```

# CALCULATE COEFFICIENT OF VARIATION -----

cv.dt <- uncertainty.dt[, .(sd = sd(V1), mean = mean(V1)),
                          .(Country, Approach, Continent)] %>%
  .[, cv:= sd / mean]

dd <- list()
for (i in 1:length(list_continents)) {
  dd[[i]] <- ggplot(cv.dt[Continent %in% list_continents[[i]]],
                    aes(reorder(Country, cv), cv, color = Approach)) +
    geom_point() +
    scale_color_manual(values = wes_palette("Chevalier1"),
                      labels = c("Jägermeyr et al. approach",
                                "Rohwer et al. approach")) +
    labs(y = "Coefficient of variation",
         x = "") +
    facet_wrap(~Continent, scales = "free_y") +
    scale_y_continuous(limits = c(0, 1),
                      breaks = pretty_breaks(n = 3)) +

    coord_flip() +
    theme_AP() +
    guides(color = guide_legend(nrow = 2, byrow = TRUE))
}

dd

```

[[1]]



##

[[2]]



4.2 Ranges

```
# COMPUTE RANGES -----

calc <- uncertainty.dt[, .(min = min(V1), max = max(V1)), .(Continent, Country)] %>%
  .[, .(range = max - min), .(Continent, Country)] %>%
  .[order(range)]

print(calc, n = Inf)
```

```
##      Continent      Country      range
##  1:      Asia      Cyprus 0.4641096
##  2:      Asia United Arab Emirates 0.4866356
```


##	3:	Asia	Israel	0.4944147
##	4:	Asia	Jordan	0.5057459
##	5:	Europe	Netherlands	0.5673924
##	6:	Americas	Cuba	0.5755095
##	7:	Africa	Tunisia	0.5764236
##	8:	Africa	Mozambique	0.5767331
##	9:	Africa	Zimbabwe	0.5783439
##	10:	Africa	Benin	0.5789929
##	11:	Europe	Italy	0.5801489
##	12:	Europe	France	0.5804556
##	13:	Africa	Namibia	0.5822584
##	14:	Americas	United States	0.5834569
##	15:	Asia	Brunei	0.5857038
##	16:	Asia	Lebanon	0.5861990
##	17:	Asia	Kazakhstan	0.5879825
##	18:	Asia	Saudi Arabia	0.5884782
##	19:	Africa	Zambia	0.5902938
##	20:	Africa	South Africa	0.5905044
##	21:	Americas	Brazil	0.5919833
##	22:	Africa	Swaziland	0.5921023
##	23:	Africa	Ivory Coast	0.5939957
##	24:	Europe	Austria	0.6014893
##	25:	Europe	Belgium	0.6014893
##	26:	<NA>	Byelarus	0.6014893
##	27:	Europe	Denmark	0.6014893
##	28:	Europe	Finland	0.6014893
##	29:	Europe	Germany	0.6014893
##	30:	Europe	Greece	0.6014893
##	31:	Europe	Hungary	0.6014893
##	32:	Europe	Latvia	0.6014893
##	33:	Europe	Lithuania	0.6014893
##	34:	Europe	Luxembourg	0.6014893
##	35:	Africa	Malawi	0.6014893
##	36:	Europe	Slovakia	0.6014893
##	37:	Europe	Sweden	0.6014893
##	38:	Europe	Switzerland	0.6014893
##	39:	Asia	Kuwait	0.6016086
##	40:	Europe	Spain	0.6044159
##	41:	Americas	Canada	0.6058219
##	42:	Africa	Botswana	0.6076223
##	43:	Europe	Ukraine	0.6139551
##	44:	Asia	Qatar	0.6206659
##	45:	Europe	Russia	0.6462099
##	46:	Europe	Czech Republic	0.6472857
##	47:	Europe	United Kingdom	0.6534832
##	48:	Europe	Romania	0.6568256
##	49:	Europe	Croatia	0.6582100
##	50:	Europe	Slovenia	0.6631916

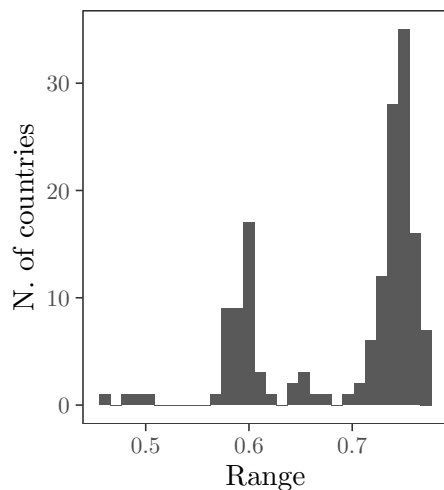
## 51:	Africa	Burkina Faso	0.6718699
## 52:	Africa	Libya	0.6973910
## 53:	Europe	Moldova	0.7086949
## 54:	Africa	Algeria	0.7088434
## 55:	Africa	Chad	0.7142740
## 56:	Asia	Syria	0.7157040
## 57:	Europe	Serbia	0.7175545
## 58:	Oceania	Papua New Guinea	0.7186161
## 59:	Europe	Bulgaria	0.7186650
## 60:	Americas	Jamaica	0.7218540
## 61:	Americas	Mexico	0.7248560
## 62:	Oceania	Australia	0.7267790
## 63:	Europe	Bosnia and Herzegovina	0.7277615
## 64:	Asia	Armenia	0.7278218
## 65:	Africa	Morocco	0.7291735
## 66:	Africa	Ghana	0.7305429
## 67:	Africa	Western Sahara	0.7323095
## 68:	Americas	Argentina	0.7323171
## 69:	Americas	Colombia	0.7324198
## 70:	Americas	Venezuela	0.7328742
## 71:	Asia	Iran	0.7330428
## 72:	Asia	Azerbaijan	0.7333914
## 73:	Asia	China	0.7341298
## 74:	Americas	Ecuador	0.7343571
## 75:	Americas	El Salvador	0.7345590
## 76:	Africa	Gabon	0.7353623
## 77:	Asia	India	0.7353925
## 78:	Americas	Puerto Rico	0.7357588
## 79:	Americas	Costa Rica	0.7360408
## 80:	Americas	Chile	0.7365217
## 81:	Africa	Mauritania	0.7367003
## 82:	Americas	Panama	0.7368114
## 83:	Americas	Bolivia	0.7370244
## 84:	Americas	Suriname	0.7387122
## 85:	Americas	French Guiana	0.7387628
## 86:	Africa	Cameroon	0.7388522
## 87:	Asia	Iraq	0.7389065
## 88:	Oceania	New Zealand	0.7389989
## 89:	Europe	Poland	0.7393614
## 90:	Americas	Paraguay	0.7398433
## 91:	Africa	Guinea	0.7400750
## 92:	Africa	Lesotho	0.7401646
## 93:	Asia	Turkey	0.7402977
## 94:	Africa	Uganda	0.7407207
## 95:	Asia	Malaysia	0.7413572
## 96:	Africa	Kenya	0.7416232
## 97:	Americas	Uruguay	0.7426133
## 98:	Africa	Zaire	0.7427597

## 99:	Africa	Nigeria	0.7428266
## 100:	Asia	Tajikistan	0.7439456
## 101:	Africa	Djibouti	0.7447347
## 102:	Asia	Afghanistan	0.7453029
## 103:	Africa	Ethiopia	0.7453914
## 104:	Asia	Mongolia	0.7458044
## 105:	Asia	Sri Lanka	0.7460123
## 106:	Africa	Senegal	0.7460561
## 107:	Africa	Egypt	0.7465109
## 108:	Americas	Peru	0.7465975
## 109:	Africa	Congo	0.7477196
## 110:	Africa	Mali	0.7481701
## 111:	Asia	Georgia	0.7484232
## 112:	Asia	Japan	0.7485508
## 113:	Asia	Vietnam	0.7488066
## 114:	Americas	Dominican Republic	0.7490021
## 115:	Africa	Angola	0.7494520
## 116:	Africa	Somalia	0.7498927
## 117:	Asia	Thailand	0.7500141
## 118:	Africa	Niger	0.7501017
## 119:	Africa	Eritrea	0.7506740
## 120:	Americas	Guyana	0.7511102
## 121:	Asia	South Korea	0.7513679
## 122:	Asia	Uzbekistan	0.7515984
## 123:	Americas	Guatemala	0.7522521
## 124:	Asia	Indonesia	0.7523546
## 125:	Asia	Bhutan	0.7524556
## 126:	Americas	Nicaragua	0.7524794
## 127:	Asia	Bangladesh	0.7526387
## 128:	Asia	Yemen	0.7533179
## 129:	Americas	Belize	0.7534902
## 130:	Europe	Macedonia	0.7538931
## 131:	Africa	Guinea-Bissau	0.7545221
## 132:	Americas	Haiti	0.7545770
## 133:	Africa	Rwanda	0.7551263
## 134:	Africa	Gambia	0.7551399
## 135:	Asia	Cambodia	0.7552400
## 136:	Asia	Kyrgyzstan	0.7559198
## 137:	Africa	Madagascar	0.7559324
## 138:	Africa	Equatorial Guinea	0.7569197
## 139:	Asia	Nepal	0.7571390
## 140:	Asia	Laos	0.7575573
## 141:	Africa	Tanzania	0.7576033
## 142:	Europe	Albania	0.7584757
## 143:	Asia	Turkmenistan	0.7591862
## 144:	Africa	Sudan	0.7594946
## 145:	Americas	Honduras	0.7600602
## 146:	Africa	Central African Republic	0.7600622

```
## 147:    Africa                Liberia 0.7619743
## 148:    Africa                Sierra Leone 0.7631982
## 149:     Asia                Burma 0.7636254
## 150:    Africa                Togo 0.7637762
## 151:    Europe                Norway 0.7643408
## 152:     Asia                North Korea 0.7660183
## 153:    Africa                Burundi 0.7663720
## 154:    Europe                Portugal 0.7670973
## 155: Americas                Trinidad 0.7678994
## 156:     Asia                Philippines 0.7739224
## 157:     Asia                Pakistan 0.7744971
## 158:     Asia                Oman 0.7747888
##      Continent                Country      range
```

```
ggplot(calc, aes(range)) +
  geom_histogram() +
  labs(x = "Range", y = "N. of countries") +
  theme_AP()
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



```
# COMPARE RANGES -----
ranges_empirical <- uncertainty.dt[, .(higher = max(V1), lower = min(V1)), IFT] %>%
  .[, Study:= "This study"]%>%
  .[!IFT == "Jager"]

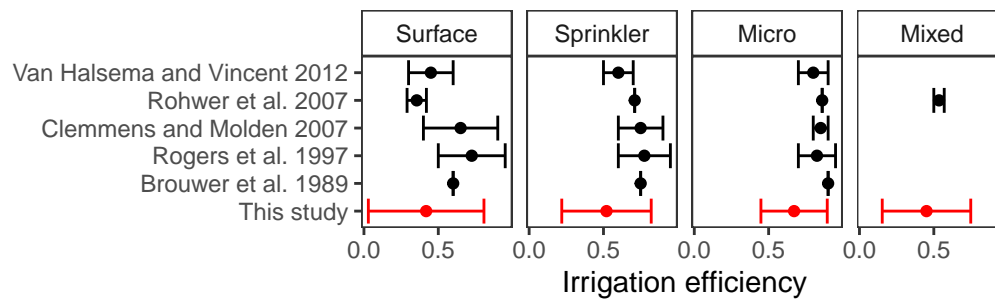
ranges_efficiencies <- fread("ranges_efficiencies.csv")

rbind(ranges_empirical, ranges_efficiencies)[, mean.value:= (higher + lower) / 2] %>%
  .[, Study:= factor(Study, levels = c("This study",
    "Brouwer et al. 1989",
    "Rogers et al. 1997",
    "Clemmens and Molden 2007",
    "Rohwer et al. 2007",
```

```

"Van Halsema and Vincent 2012"))] %>%
na.omit() %>%
ggplot(., aes(mean.value, Study, color = ifelse(Study == "This study", "red", "black"))) +
geom_point() +
scale_x_continuous(breaks = pretty_breaks(n = 3)) +
geom_errorbar(aes(xmin = lower, xmax = higher)) +
scale_color_identity() +
facet_wrap(~IFT, ncol = 4) +
labs(x = "Irrigation efficiency", y = "") +
theme_AP()

```



4.3 Overlap between irrigation efficiencies

```

# CHECK OVERLAP -----

dd <- uncertainty.dt[!Continent == "Oceania"][Study == "Rohwer et al. approach"] %>%
  split(., .$Continent, drop = TRUE)

overlap.dt <- lapply(dd, function(x) split(x, x$IFT, drop = TRUE)) %>%
  lapply(., function(x) lapply(x, function(y) y[, V1])) %>%
  lapply(., function(x) overlap(x)$OV)

overlap.dt

## $Africa
## Surface-Sprinkler      Surface-Mixed      Sprinkler-Mixed
##           0.2642237           0.4604639           0.4606540
##
## $Americas
## Surface-Mixed
##           0.4477437
##
## $Asia
## Surface-Micro Surface-Mixed      Micro-Mixed
##           0.05075313      0.44599065      0.06876229
##
## $Europe
## Surface-Sprinkler      Surface-Mixed      Sprinkler-Mixed
##           0.2616758           0.4542842           0.4650725

```

```

ff <- uncertainty.dt[!Continent == "Oceania"] %>%
  .[Country %in% intersect(rohwer[, Country], jager[, Country])] %>%
  split(., .$Country, drop = TRUE) %>%
  lapply(., function(x) split(x, x$Approach, drop = TRUE)) %>%
  lapply(., function(x) lapply(x, function(y) y[, V1])) %>%
  lapply(., function(x) overlap(x)$OV) %>%
  lapply(., data.table) %>%
  rbindlist(., idcol = "Country") %>%
  .[, Continent:= countrycode(., Country,
                             origin = "country.name",
                             destination = "continent")]

list_continents <- list(c("Africa", "Asia"), c("Americas", "Europe"))

# PLOT OVERLAP -----

dd <- list()
for(i in 1:length(list_continents)) {
  dd[[i]] <- ff[Continent %in% list_continents[[i]]] %>%
    ggplot(., aes(reorder(Country, V1), V1)) +
    geom_point() +
    scale_color_discrete(name = "GM") +
    labs(y = "Overlap", x = "") +
    facet_wrap(~Continent, scales = "free_y") +
    coord_flip() +
    theme_AP()
}

dd

## [[1]]

```

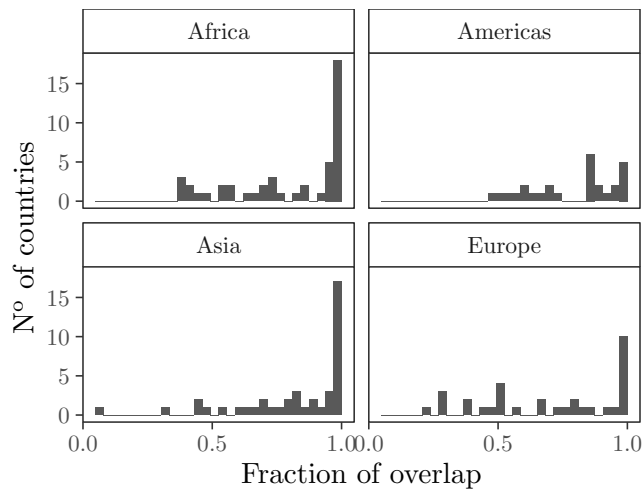


[[2]]

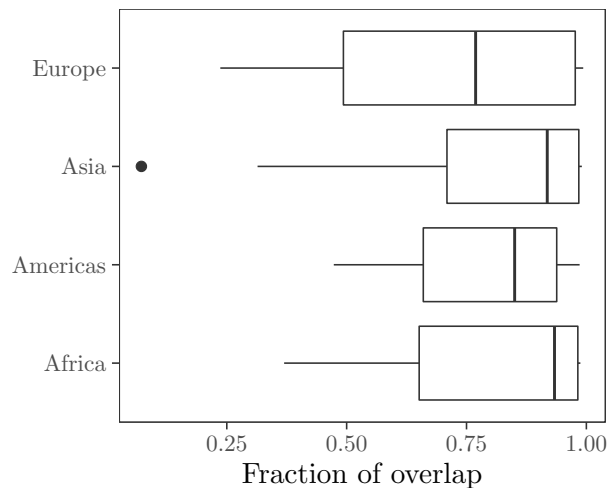


PLOT OVERLAP AS HISTOGRAMS AND BOXPLOTS

```
ggplot(ff, aes(V1)) +
  geom_histogram() +
  facet_wrap(~Continent) +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  theme_AP() +
  labs(x = "Fraction of overlap", y = "Nº of countries")
```

```
ggplot(ff, aes(Continent, V1)) +
  geom_boxplot() +
  coord_flip() +
  theme_AP() +
  labs(y = "Fraction of overlap", x = "")
```



CHECK CORRESPONDENCE BETWEEN SHARES OF IFT AND PREDOMINANT TECHNOLOGY -----

Retrieve countries where overlap is <0.3

```
merge(jager, rohwer, by = c("Country")) %>%
  .[Country %in% ff[V1 < 0.3][, Country]] %>%
  .[, .(Country, Surface_fraction, Sprinkler_fraction, Drip_fraction, IFT)]
```

##	Country	Surface_fraction	Sprinkler_fraction	Drip_fraction	IFT
## 1:	Macedonia	0.891	0.091	0.018	Sprinkler
## 2:	Moldova	0.298	0.636	0.066	Surface
## 3:	Oman	0.793	0.113	0.094	Micro
## 4:	Slovenia	0.000	0.693	0.307	Mixed
## 5:	Spain	0.297	0.226	0.478	Mixed

```

# PLOT UNCERTAINTY -----
gg <- list()
for (i in 1:length(list_continents)) {
  gg[[i]] <- ggplot(uncertainty.dt[Continent %in% list_continents[[i]]],
    aes(x = V1, y = fct_reorder(Country, V1), fill = Study)) +
    geom_density_ridges(scale = 2, alpha = 0.3) +
    labs(x = "Irrigation efficiency", y = "") +
    facet_wrap(~Continent, scales = "free") +
    scale_x_continuous(breaks = pretty_breaks(n = 3),
      limits = c(0, 1)) +
    scale_fill_manual(values = wes_palette("Chevalier1")) +
    theme_AP() +
    theme(legend.position = "top") +
    guides(fill = guide_legend(nrow = 2, byrow = TRUE))
}

```

```

# MERGE PLOTS -----

```

```

gg

```

```

## [[1]]

```

```

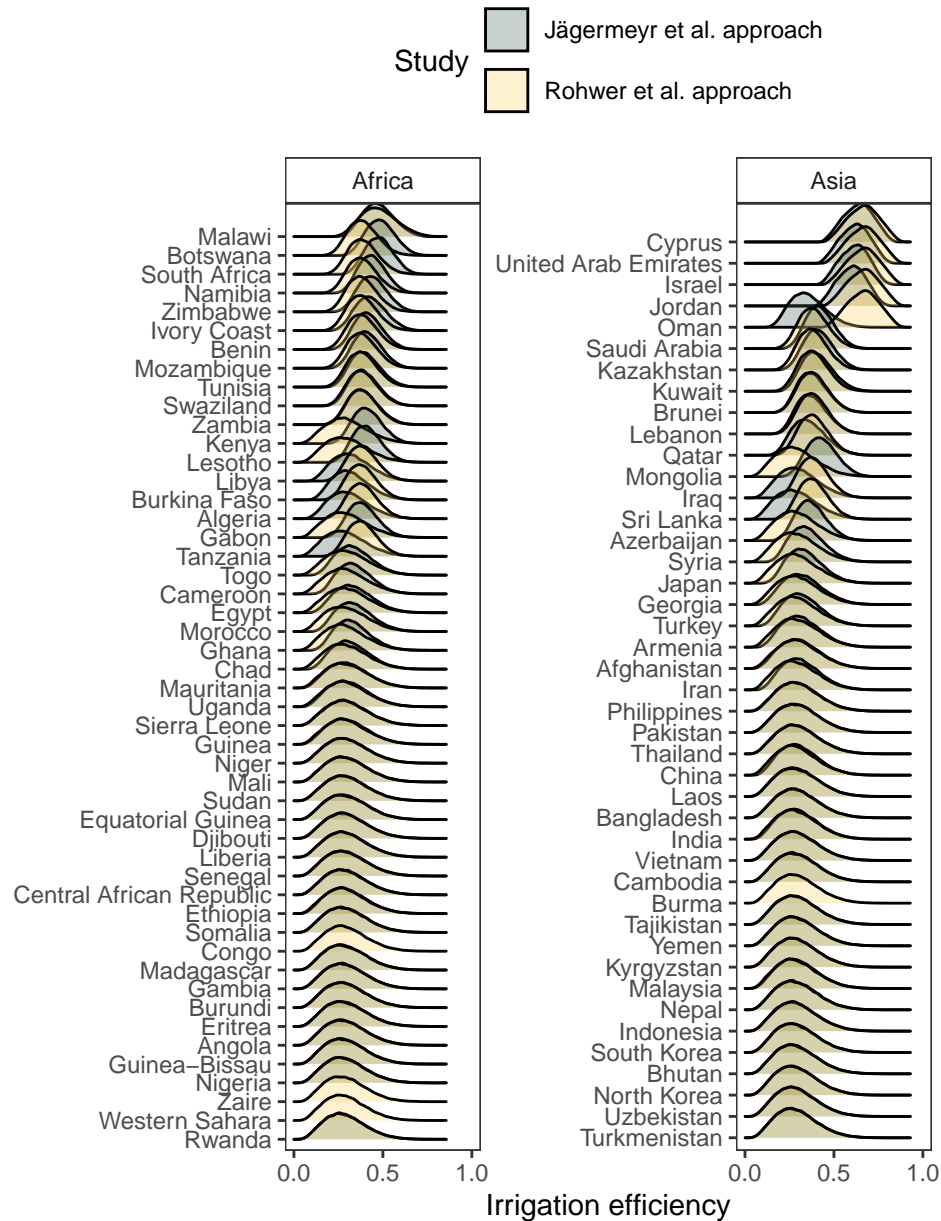
## Picking joint bandwidth of 0.0116

```

```

## Picking joint bandwidth of 0.0117

```





```
##
```

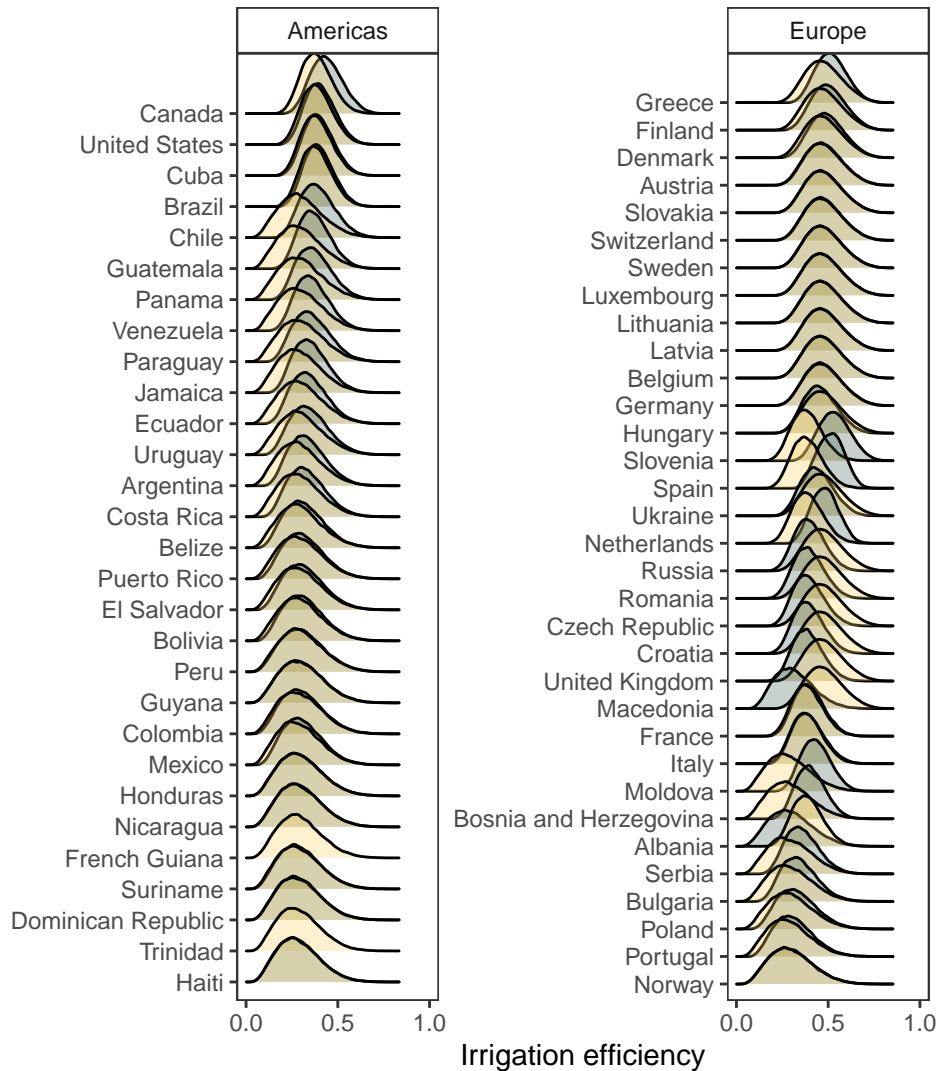
```
## [[2]]
```

```
## Picking joint bandwidth of 0.0119
```

```
## Picking joint bandwidth of 0.0107
```

Study

 Jägermeyr et al. approach
 Rohwer et al. approach



PLOT UNCERTAINTY IN EACH IRRIGATION TECHNOLOGY -----

```

gg <- list()

for(i in 1:length(list_continents)) {
  gg[[i]] <- uncertainty.dt[Approach == "Rohwer et al. 2007"][Continent %in% list_continents[i]]
  ggplot(., aes(x = V1, y = fct_reorder(Country, V1), fill = IFT)) +
    geom_density_ridges(scale = 2, alpha = 0.3) +
    labs(x = "Irrigation efficiency", y = "") +
    facet_wrap(~Continent, scales = "free") +
    scale_x_continuous(breaks = pretty_breaks(n = 3),
                      limits = c(0, 1)) +
    theme_AP() +

```

```

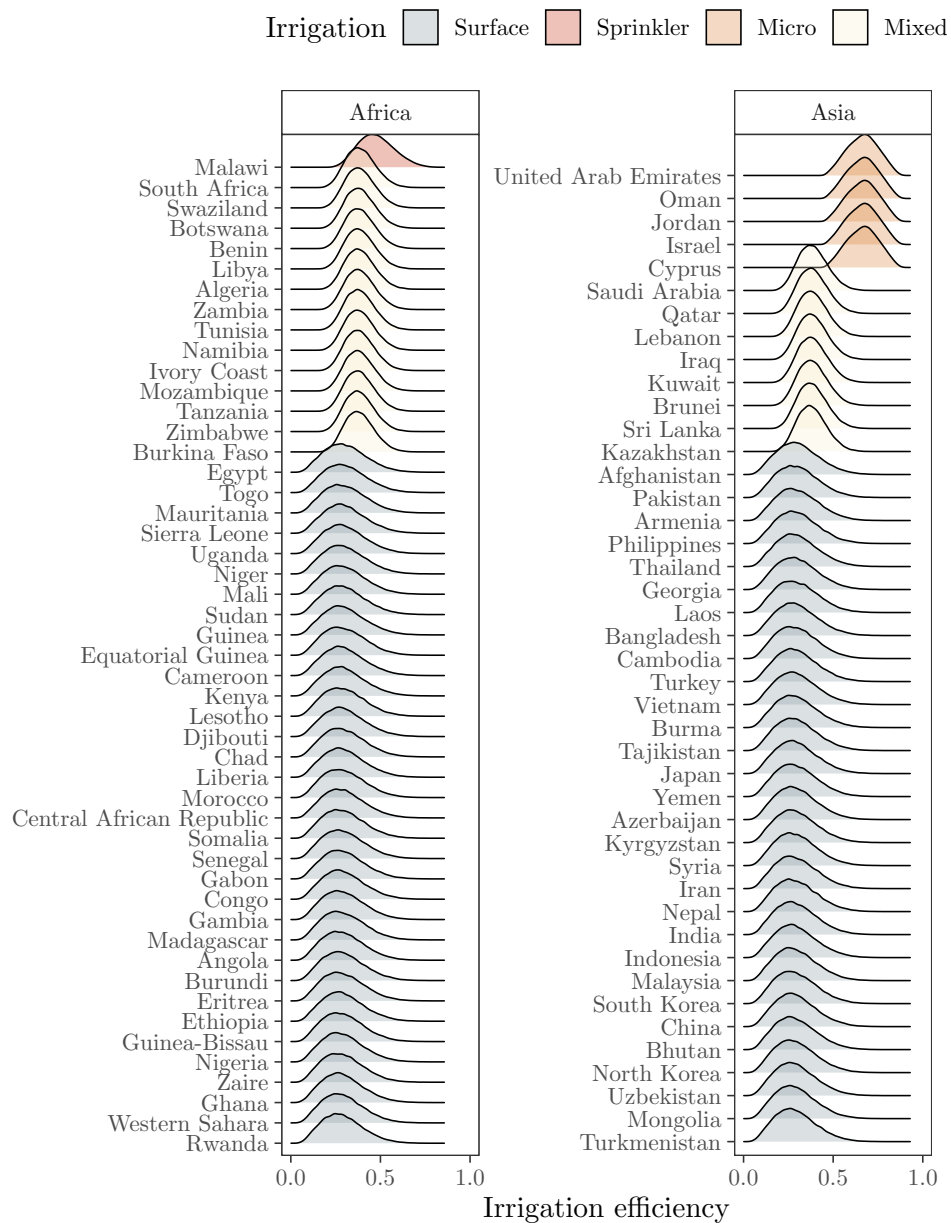
    theme(legend.position = "top")
  if(i == 1) {
    gg[[i]] <- gg[[i]] +
      scale_fill_manual(values = c("#899DA4", "#C93312", "#DC863B", "#FAEFD1"),
                        labels = c("Surface", "Sprinkler", "Micro", "Mixed"),
                        name = "Irrigation")
  } else if(i == 2) {
    gg[[i]] <- gg[[i]] +
      scale_fill_manual(values = c("#899DA4", "#C93312", "#FAEFD1"),
                        labels = c("Surface", "Sprinkler", "Mixed"),
                        name = "Irrigation")
  }
}
gg

```

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

```
## Picking joint bandwidth of 0.0118
```

```
## Picking joint bandwidth of 0.0119
```

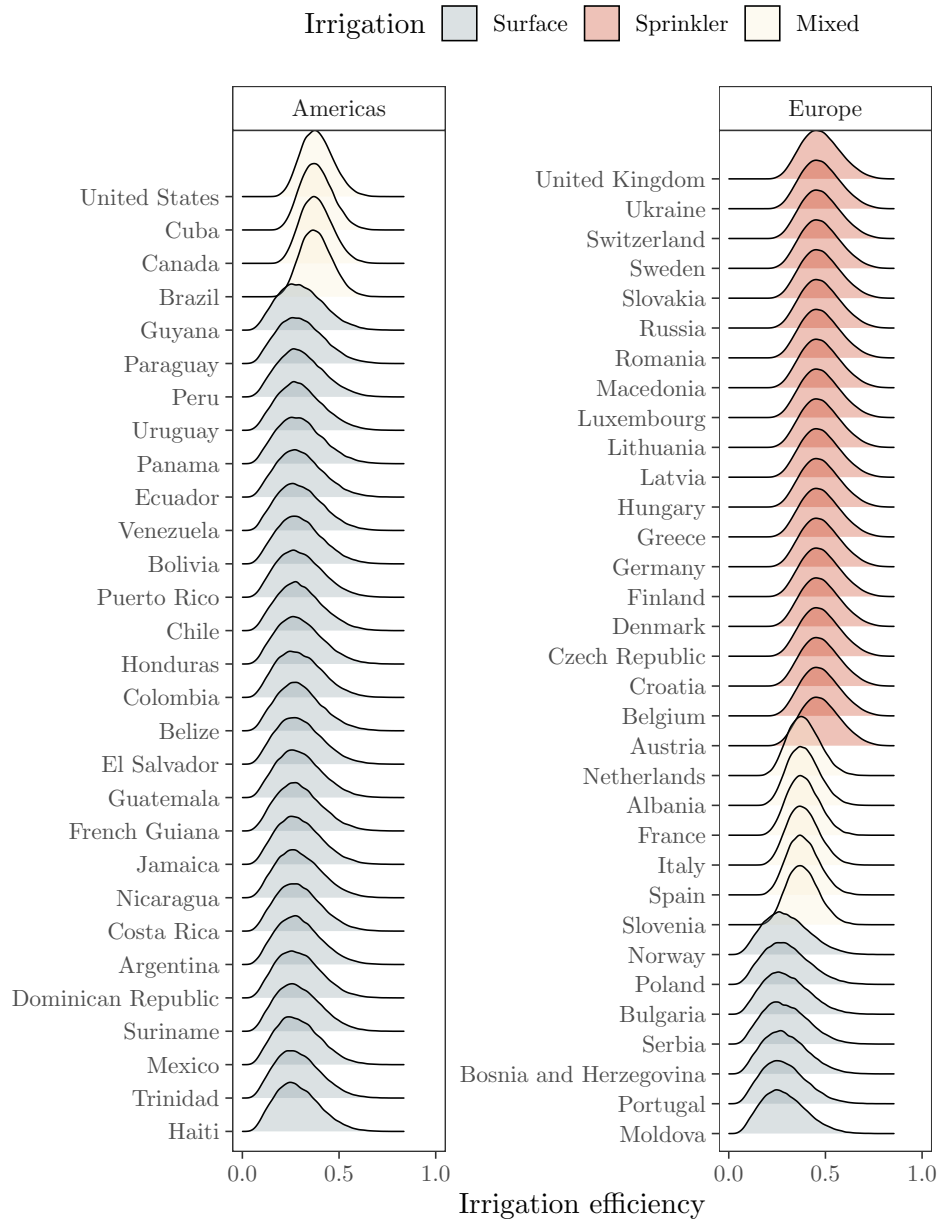


##

[[2]]

Picking joint bandwidth of 0.0124

Picking joint bandwidth of 0.0112



PLOT ROHWER ET AL.'S IRRIGATION EFFICIENCY VALUES -----

```
rohwer[, Continent:= countrycode(rohwer[, Country], origin = "country.name",
                                destination = "continent")]
```

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

```
dd <- list()
for (i in 1:length(list_continents)) {
  dd[[i]] <- ggplot(rohwer[Continent %in% list_continents[[i]]],
                    aes(x = Ep, y = fct_reorder(Country, Ep), color = IFT)) +
  geom_point() +
  labs(x = "Irrigation efficiency", y = "") +
  scale_x_continuous(breaks = pretty_breaks(n = 3),
```

```

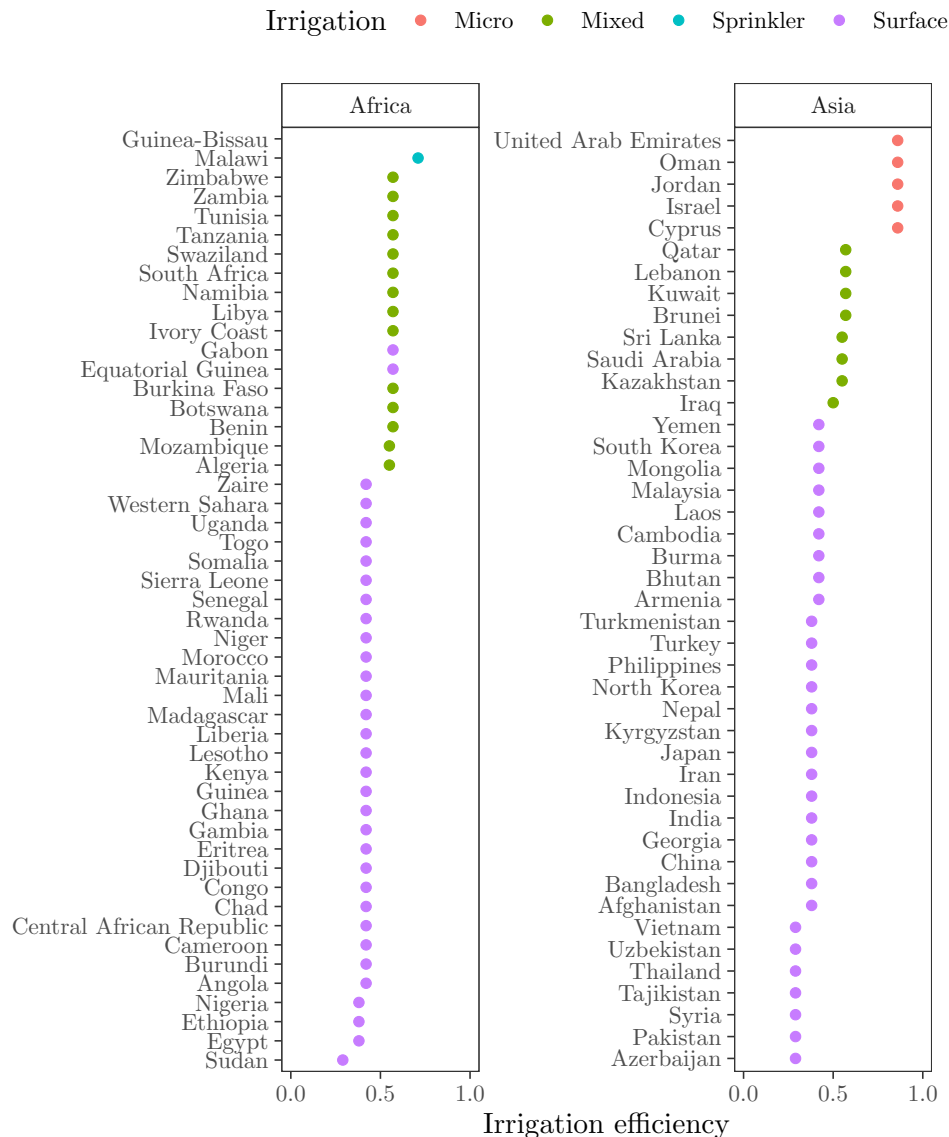
    limits = c(0, 1)) +
  facet_wrap(~Continent, scales = "free") +
  scale_color_discrete(name = "Irrigation") +
  theme_AP()
}

```

```
dd
```

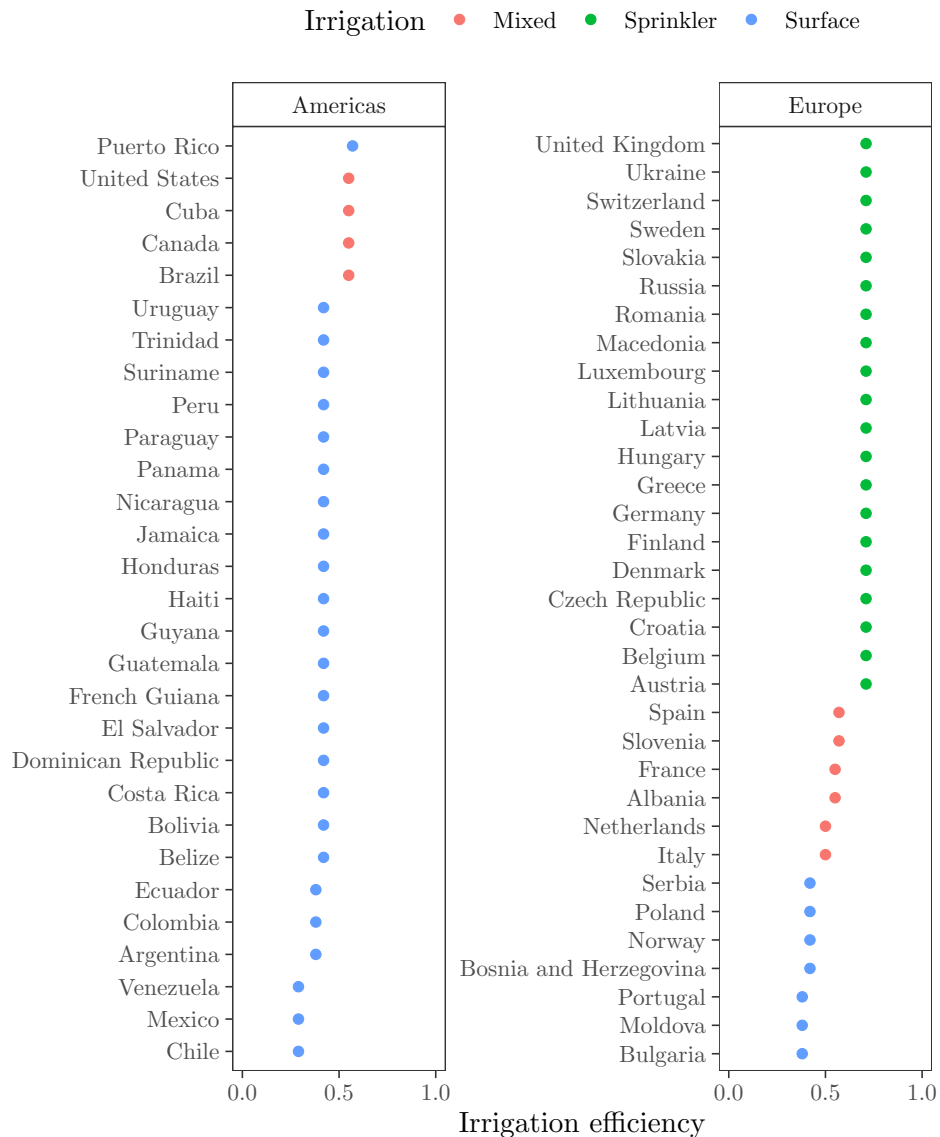
```
## [[1]]
```

```
## Warning: Removed 1 rows containing missing values (geom_point).
```



```
##
```

```
## [[2]]
```

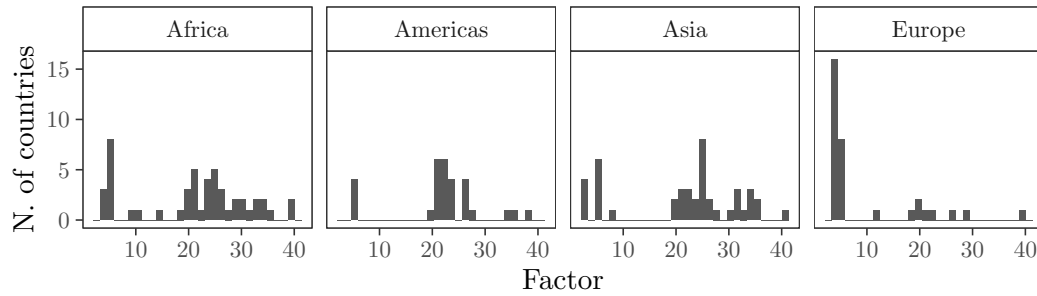
```
# CALCULATE THE UNCERTAINTY IN THE RANGES -----

selection_continents <- c("Africa", "Asia", "Americas", "Europe")

factor_unc <- uncertainty.dt[, .(min = min(V1), max = max(V1)), .(Continent, Country)] %>%
  .[Continent %in% selection_continents] %>%
  .[, factor:= max / min]

ggplot(factor_unc, aes(factor)) +
  geom_histogram() +
  facet_wrap(~Continent, ncol = 4) +
  labs(x = "Factor", y = "N. of countries") +
  theme_AP()

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```



*# Number of countries whose irrigation water withdrawals fluctuate a factor of x
due to uncertainty in irrigation efficiency*

```
factor_unc %>%
  .[, factor:= floor(max / min)] %>%
  .[, .(number.countries = .N), factor] %>%
  .[order(factor)] %>%
  print()
```

```
##      factor number.countries
## 1:      2              4
## 2:      3             15
## 3:      4             27
## 4:      5              3
## 5:      7              1
## 6:      9              1
## 7:     10              1
## 8:     12              1
## 9:     14              1
## 10:    18              1
## 11:    19              6
## 12:    20             11
## 13:    21             11
## 14:    22              7
## 15:    23              9
## 16:    24             11
## 17:    25              6
## 18:    26              7
## 19:    27              1
## 20:    28              3
## 21:    29              3
## 22:    30              2
## 23:    31              4
## 24:    32              1
## 25:    33              4
## 26:    34              5
## 27:    35              3
## 28:    37              1
## 29:    38              1
## 30:    39              2
```

```
## 31:      40      1
##      factor number.countries
```

4.4 Retrieve data from ISIMIP

```
# FUNCTIONS TO EXTRACT DATA FROM .NC FILES -----

coords2country = function(points) {
  countriesSP <- rworldmap::getMap(resolution = 'low')
  pointsSP = sp::SpatialPoints(points, proj4string=CRS(proj4string(countriesSP)))
  indices = sp::over(pointsSP, countriesSP)
  indices$ADMIN
}

# Function to load and extract data from .nc files from ISIMIP
open_nc_files <- function(file, dname, selected.years, vec) {
  ncin <- nc_open(file)
  # get longitude, latitude, time
  lon <- ncvar_get(ncin, "lon")
  lat <- ncvar_get(ncin, "lat")
  # Get variable
  tmp_array <- ncvar_get(ncin, dname)
  m <- lapply(selected.years, function(x) vec[[x]])

  out <- lapply(m, function(x) {
    tmp_slice <- lapply(x, function(y) tmp_array[, , y])
    # create dataframe -- reshape data
    # matrix (nlon*nlat rows by 2 cols) of lons and lats
    lonlat <- as.matrix(expand.grid(lon,lat))
    # vector of `tmp` values
    tmp_vec <- lapply(tmp_slice, function(x) as.vector(x))
    # create dataframe and add names
    tmp_df01 <- lapply(tmp_vec, function(x) data.frame(cbind(lonlat, x)))
    names(tmp_df01) <- x
    da <- lapply(tmp_df01, data.table) %>%
      rbindlist(., idcol = "month") %>%
      na.omit()
    # Convert coordinates to country
    Country <- coords2country(da[1:nrow(da), 2:3])
    df <- cbind(Country, da)
    setDT(df)
    out <- na.omit(df)[, .(Water.Withdrawn = sum(x)), Country]
    out[, Water.Withdrawn:= Water.Withdrawn * 10000]
    out[, Continent:= countrycode(out[, Country],
                                   origin = "country.name",
                                   destination = "continent")] %>%
    .[, Code:= countrycode(out[, Country],
```

```

        origin = "country.name",
        destination = "un")] %>%
    .[, Country:= countrycode(out[, Code],
        origin = "un",
        destination = "country.name")] %>%
    .[!Continent == "Oceania"]
    setcolorder(out, c("Country", "Continent", "Code", "Water.Withdrawn"))
  })
  return(out)
}

# READ IN NC FILES -----

# Define settings
vecs <- 1:((2010 - 1970) * 12)
vec <- split(vecs, ceiling(seq_along(vecs) / 12))
names(vec) <- 1971:2010
selected.years <- "2010"
dname <- "pirrww"

files <- list("h08_wfdei_nobc_hist_varsoc_co2_pirrww_global_monthly_1971_2010.nc",
             "pcr-globwb_wfdei_nobc_hist_varsoc_co2_pirrww_global_monthly_1971_2010.nc",
             "lpjml_wfdei_nobc_hist_varsoc_co2_pirrww_global_monthly_1971_2010.nc",
             "watergap2_wfdei_nobc_hist_varsoc_co2_pirrww_global_monthly_1971_2010.nc")

names.isimip <- c("H08", "PCR-GLOBWB", "LPJmL", "WaterGap")

isimip.dt <- mclapply(files, function(x)
  open_nc_files(file = x, dname = dname, selected.years = selected.years, vec = vec),
  mc.cores = detectCores() * 0.75)

# EXTRACT CORRECTIVE COEFFICIENTS FOR IRRIGATION EFFICIENCY FOR LPJML -----

ncin <- nc_open("irrigation_project_efficiencies.nc")
lon <- ncvar_get(ncin, "lon")
lat <- ncvar_get(ncin, "lat")
tmp_array <- ncvar_get(ncin)
lonlat <- as.matrix(expand.grid(lon,lat))
da <- na.omit(cbind(lonlat, as.vector(tmp_array))) %>%
  data.frame() %>%
  na.omit()
Country <- coords2country(da[1:nrow(da), 1:2])
lpjml_efficiencies <- cbind(Country, da) %>%
  na.omit() %>%
  data.table() %>%
  .[, .(Ep = mean(V3)), Country]

```

```

# ARRANGE NC FILES -----

names(isimip.dt) <- names.isimip

isimip.dt <- lapply(isimip.dt, function(x) rbindlist(x)) %>%
  rbindlist(., idcol = "Model") %>%
  na.omit() %>%
  # To correct for duplicate country in Cyprus
  .[, .(Water.Withdrawn = mean(Water.Withdrawn)), .(Model, Country, Continent, Code)]

lpjml_harmonized <- merge(isimip.dt[Model == "LPJmL"], lpjml_efficiencies, all.x = TRUE) %>%
  .[, Water.Withdrawn:= Water.Withdrawn * Ep] %>%
  .[, Ep:= NULL]

isimip.dt <- rbind(isimip.dt[!Model == "LPJmL"], lpjml_harmonized)

fwrite(isimip.dt, "isimip.dt")

# MERGE UNCERTAINTY IN EP WITH ISIMIP DATA -----

efficiency.dt <- copy(uncertainty.dt) %>%
  setnames(., "V1", "Ep")

ghm.dt <- dcast(isimip.dt, Country + Continent + Code ~ Model, value.var = "Water.Withdrawn")
full.dt <- merge(efficiency.dt, ghm.dt, by = c("Country", "Continent"), all.x = TRUE) %>%
  .[, (names.isimip):= lapply(.SD, function(x) x / Ep), .SDcols = names.isimip]
tmp.dt <- melt(full.dt, measure.vars = names.isimip, variable.name = "Model",
  value.name = "IWW_corrected")
ghm.large <- melt(ghm.dt, measure.vars = names.isimip, variable.name = "Model",
  value.name = "IWW")
gm.uncertainty <- tmp.dt[, .(min = min(IWW_corrected), max = max(IWW_corrected)),
  .(Country, Continent, Model)]
gm.dt <- merge(ghm.large, gm.uncertainty)

```

4.5 Retrieve data from ISIMIP (climate change in 2050)

```

# READ IN FILES ON CLIMATE CHANGE UNCERTAINTY (2050) -----

files <- list(
  "watergap2_miroc5_ewembi_rcp85_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "watergap2_miroc5_ewembi_rcp60_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "watergap2_miroc5_ewembi_rcp45_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "watergap2_miroc5_ewembi_rcp26_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "lpjml_miroc5_ewembi_rcp85_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "lpjml_miroc5_ewembi_rcp60_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "lpjml_miroc5_ewembi_rcp26_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
  "pcr-globwb_miroc5_ewembi_rcp60_2005soc_co2_pirrww_global_monthly_2006_2099.nc",

```

```

"pcr-globwb_miroc5_ewembi_rcp26_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
"h08_miroc5_ewembi_rcp85_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
"h08_miroc5_ewembi_rcp60_2005soc_co2_pirrww_global_monthly_2006_2099.nc",
"h08_miroc5_ewembi_rcp26_2005soc_co2_pirrww_global_monthly_2006_2099.nc"
)

vecs <- 1:((2099 - 2005) * 12)
vec <- split(vecs, ceiling(seq_along(vecs) / 12))
names(vec) <- 2006:2099
dname <- "pirrww"
selected.years <- as.character(seq(2030, 2050, 10))

# Read in datasets
isimip.climate <- mclapply(
  files, function(x)
    open_nc_files(file = x, dname = dname, selected.years = selected.years, vec = vec),
  mc.cores = detectCores() * 0.75
)

# ARRANGE DATASETS -----

ghms <- c(rep("WaterGap", times = 4),
  rep("LPJmL", times = 3),
  rep("PCR-GLOBWB", times = 2),
  rep("H08", times = 3))

climate_scenario <- c(85, 60, 45, 26, 85, 60, 26, 60, 26, 85, 60, 26)
names.isimip <- paste(ghms, climate_scenario, sep = "/")

# Name the slots
names(isimip.climate) <- names.isimip

for(i in names(isimip.climate)) {
  names(isimip.climate[[i]]) <- selected.years
}

# Arrange data
isimip.climate.dt <- lapply(isimip.climate, function(x) rbindlist(x, idcol = "Year")) %>%
  rbindlist(., idcol = "Model") %>%
  .[!Continent == "Oceania"] %>%
  separate(., "Model", c("Model", "Climate scenario"), "/") %>%
  na.omit() %>%
  .[Year == 2050]

# Export
fwrite(isimip.climate.dt, "isimip.climate.dt.csv")

```

```

# PLOT RANGES OF STRUCTURAL UNCERTAINTY AND RANGES OF
# STRUCTURAL UNCERTAINTY + UNCERTAINTY IN IRRIGATION EFFICIENCY +
# UNCERTAINTY IN CLIMATE CHANGE -----

countries_list <- c("Egypt", "Sudan", "South Africa", "Morocco", "Madagascar",
  "United States", "Mexico", "Brazil", "Chile", "Peru",
  "India", "China", "Pakistan", "Iran", "Indonesia",
  "Italy", "Spain", "France", "Ukraine", "Romania")

range.gm <- gm.dt %>%
  .[, .(min = min(IWW, na.rm = TRUE), max = max(IWW, na.rm = TRUE)),
    .(Country, Continent)] %>%
  .[, Approach:= "GM"]

range.study <- gm.dt %>%
  .[, .(min = min(min, na.rm = TRUE), max = max(max, na.rm = TRUE)),
    .(Country, Continent)] %>%
  .[, Approach:= "GM + uncertainty in irrigation efficiency"]

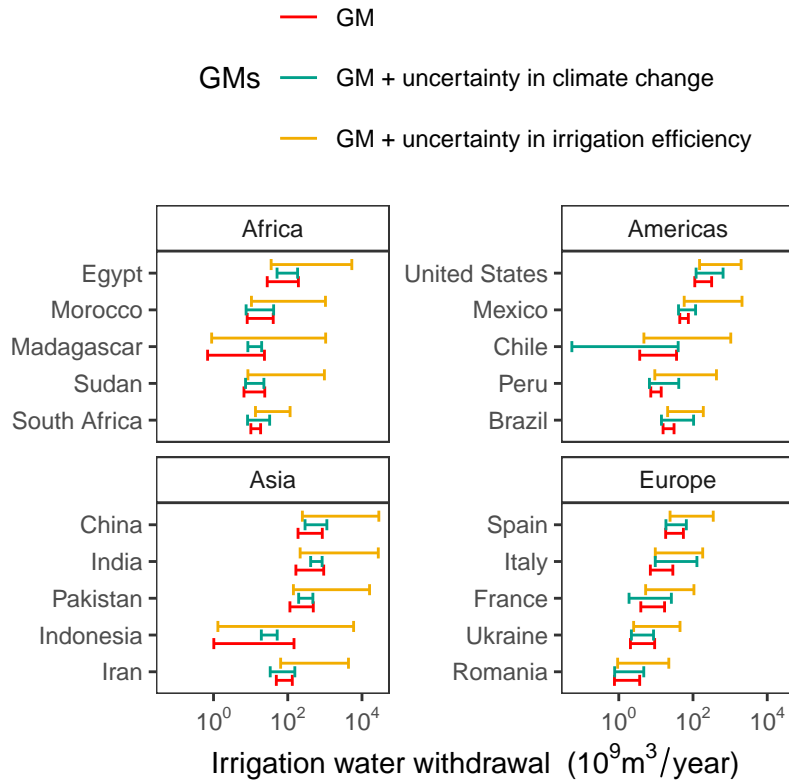
range.climate <- isimip.climate.dt %>%
  .[, .(min = min(Water.Withdrawn), max = max(Water.Withdrawn)),
    .(Country, Continent)] %>%
  .[, Approach:= "GM + uncertainty in climate change"]

all.uncertainties <- rbind(range.gm, range.study, range.climate) %>%
  .[, mean:= (min + max) / 2]

# Substitute 0 by NA -----
all.uncertainties[all.uncertainties == 0] <- NA

all.uncertainties %>%
  .[Country %in% countries_list] %>%
  ggplot(. , aes(reorder(Country, mean), mean, color = Approach)) +
  geom_errorbar(aes(ymin = min,
    ymax = max),
    position = position_dodge(0.7)) +
  scale_y_log10(breaks = trans_breaks("log10", function(x) 10 ^ (2 * x)),
    labels = trans_format("log10", math_format(10 ^ .x))) +
  scale_color_manual(name = "GMs", values = wes_palette("Darjeeling1")) +
  labs(y = expression(paste("Irrigation water withdrawal ", " ", "(", 10^9, m^3/year, " ", " ")),
    x = "") +
  facet_wrap(~Continent, scales = "free_y") +
  coord_flip() +
  theme_AP() +
  guides(color = guide_legend(nrow = 3, byrow = TRUE))

```



```
# EXPORT -----
fwrite(all.uncertainties, "all.uncertainties.csv")

# PLOT RANGES OF STRUCTURAL UNCERTAINTY AND RANGES OF
# STRUCTURAL UNCERTAINTY + UNCERTAINTY IN IRRIGATION EFFICIENCY (COMPLETE) -----

vec1 <- all.uncertainties[Approach == "GM", Country]
vec2 <- all.uncertainties[Approach == "GM + uncertainty in climate change", Country]
vec3 <- all.uncertainties[Approach == "GM + uncertainty in irrigation efficiency", Country]
common_countries <- Reduce(intersect, list(vec1, vec2, vec3))

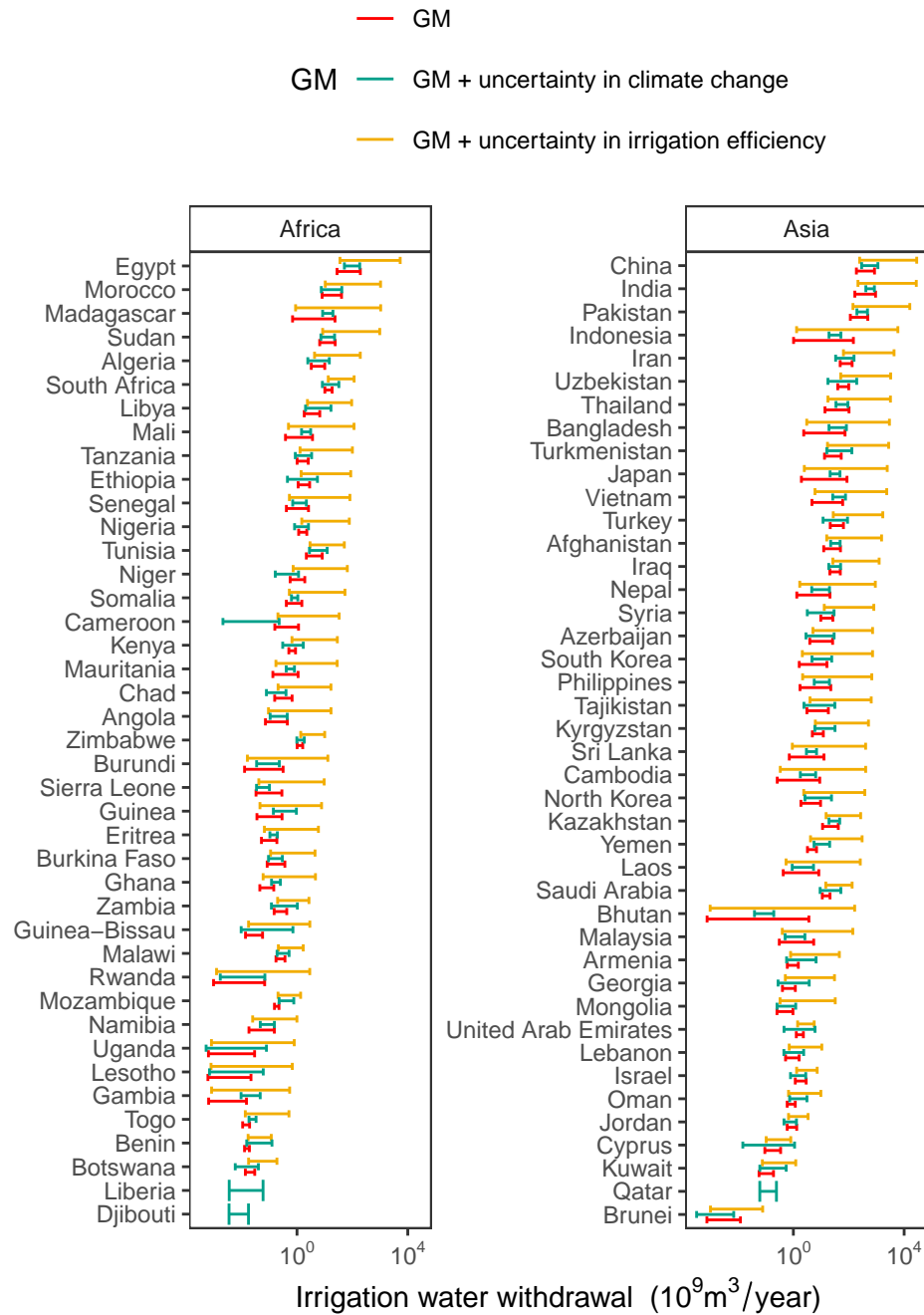
dd <- list()
for (i in 1:length(list_continents)) {
  dd[[i]] <- all.uncertainties %>%
    .[Country %in% common_countries] %>%
    na.omit() %>%
    .[Continent %in% list_continents[[i]]] %>%
    ggplot(., aes(reorder(Country, mean), mean, color = Approach)) +
    geom_errorbar(aes(ymin = min,
                      ymax = max),
                  position = position_dodge(0.7)) +
    scale_y_log10(breaks = trans_breaks("log10", function(x) 10 ^ (2 * x)),
                  labels = trans_format("log10", math_format(10 ^ .x))) +
    scale_color_manual(name = "GM", values = wes_palette("Darjeeling1")) +
    labs(y = expression(paste("Irrigation water withdrawal ", " ", "(", 10^9, " m^3/year, " ", " ")),
         x = "") +
}
```



```
facet_wrap(~Continent, scales = "free_y") +
coord_flip() +
theme_AP() +
guides(color = guide_legend(nrow = 3, byrow = TRUE))
}
```

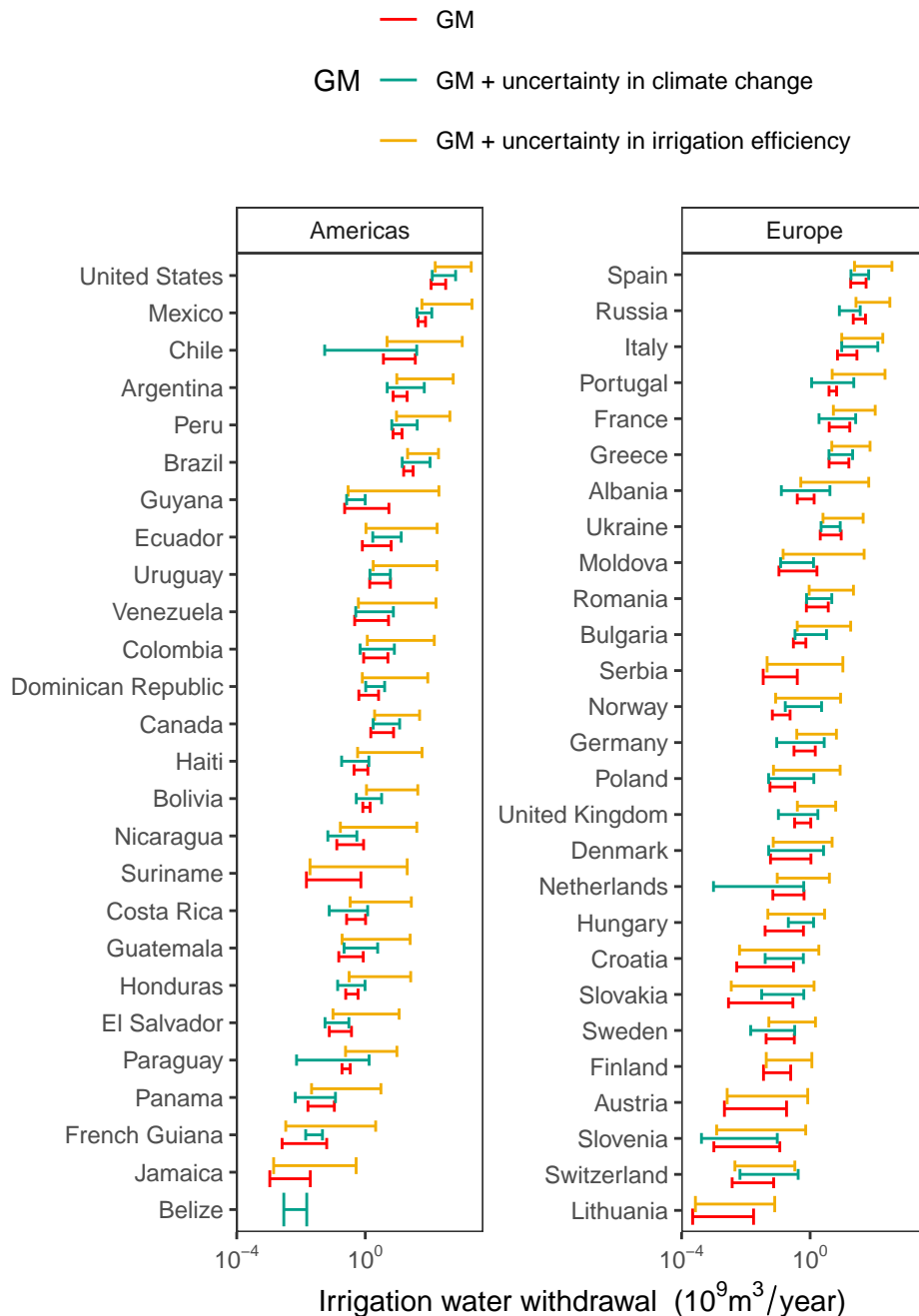
```
dd
```

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



```
##
```

```
## [[2]]
```



```
# COMPARE RANGES -----

all.uncertainties[, range:= max - min]
dd <- dcast(all.uncertainties, Country + Continent ~ Approach, value.var = "range") %>%
  na.omit() %>%
  .[, maxCol:= max.col(., 3:5, ties.method = "first")]

# check which countries show the largest ranges in climate uncertainty
lapply(1:3, function(x) dd[maxCol == x])
```

```
## [[1]]
## Empty data.table (0 rows and 6 cols): Country,Continent,GM,GM + uncertainty in climate change
##
## [[2]]
##           Country Continent      GM GM + uncertainty in climate change
## 1:      Benin      Africa 0.00634165      0.1095379
## 2:      Cyprus      Asia 0.25356972      1.0901057
## 3: Switzerland      Europe 0.06956323      0.4174803
## 4: United Arab Emirates      Asia 1.02446010      5.5930987
##      GM + uncertainty in irrigation efficiency maxCol
## 1:      0.1006289      2
## 2:      0.7022605      2
## 3:      0.3293323      2
## 4:      4.2191107      2
##
## [[3]]
##           Country Continent      GM GM + uncertainty in climate change
## 1: Afghanistan      Asia 37.8147475      26.5443113
## 2:  Albania      Europe 0.9324792      4.0007740
## 3:  Algeria      Africa 6.7659506      12.2469498
## 4:  Angola      Africa 0.3768065      0.3377945
## 5: Argentina      Americas 12.6584940      63.6812769
## ---
## 118: Venezuela      Americas 4.8093549      6.9755145
## 119:  Vietnam      Asia 55.5815237      49.9370651
## 120:  Yemen      Asia 3.6273127      14.8781794
## 121:  Zambia      Africa 0.2776971      0.9126939
## 122: Zimbabwe      Africa 0.5964828      0.8408030
##      GM + uncertainty in irrigation efficiency maxCol
## 1:      1515.858339      3
## 2:      66.132774      3
## 3:      188.296820      3
## 4:      16.773342      3
## 5:      535.053389      3
## ---
## 118:      158.353988      3
## 119:      2342.467615      3
## 120:      299.368661      3
## 121:      2.500587      3
## 122:      8.635619      3
```

5 Sensitivity analysis

```
# SAMPLE MATRIX DISTRIBUTIONS -----
# Define labels
label_facets <- c("Ea_surf" = "$E_{a_{su}}$",
```

```

"Ec_surf" = "$E_{c_{su}}$",
"Ea_sprinkler" = "$E_{a_{sp}}$",
"Ec_sprinkler" = "$E_{c_{sp}}$",
"Ea_micro" = "$E_{a_{mi}}$",
"Ec_micro" = "$E_{c_{mi}}$",
"Proportion_large" = "$f_L$",
"m" = "$m$",
"r_L" = "$r_L$"

```

```

mat <- data.table(full_sample_matrix(IFT = "Jager", Country = "Spain")$matrix)
mat <- mat[, Proportion_large:= NULL]

```

```

## Warning in `[.data.table`(mat, , `:=`(Proportion_large, NULL)): Column
## 'Proportion_large' does not exist to remove

```

```

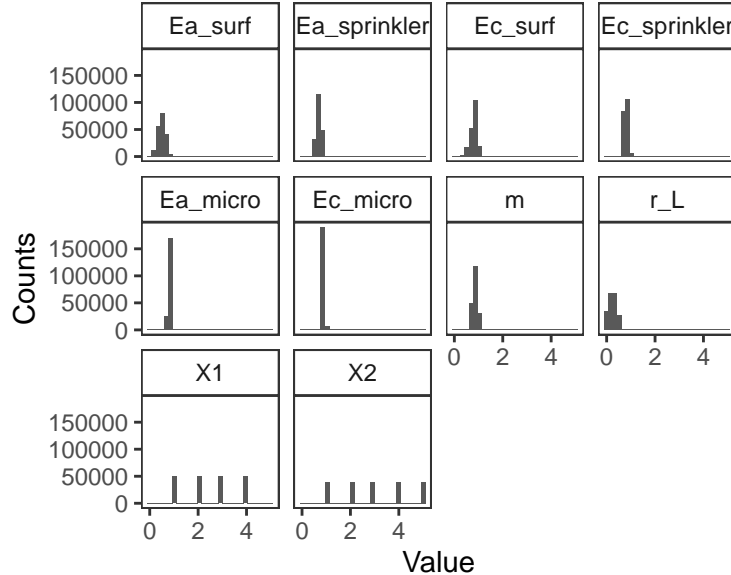
melt(mat, measure.vars = colnames(mat)) %>%
  ggplot(., aes(value)) +
  geom_histogram() +
  labs(x = "Value", y = "Counts") +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  facet_wrap(~variable) +
  theme_AP()

```

```

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

```



```

# EXTRACT SOBOLO' INDICES -----

```

```

ind <- lapply(y$`Rohwer et al. 2007`, function(x) x[["indices"]])$results)
names(ind) <- rohwer$Country
ind <- rbindlist(ind, idcol = "Country")

```

```

ind[, Continent:= countrycode(ind[, Country], origin = "country.name",

```

```

destination = "continent"]

## Warning in countrycode_convert(sourcevar = sourcevar, origin = origin, destination = dest,
tmp.ift <- split(rohwer, rohwer$IFT)

out <- list()
for(i in names(tmp.ift)) {
  out[[i]] <- ind[Country %in% tmp.ift[[i]][, Country]]
}

# PLOT SOBOL' INDICES -----

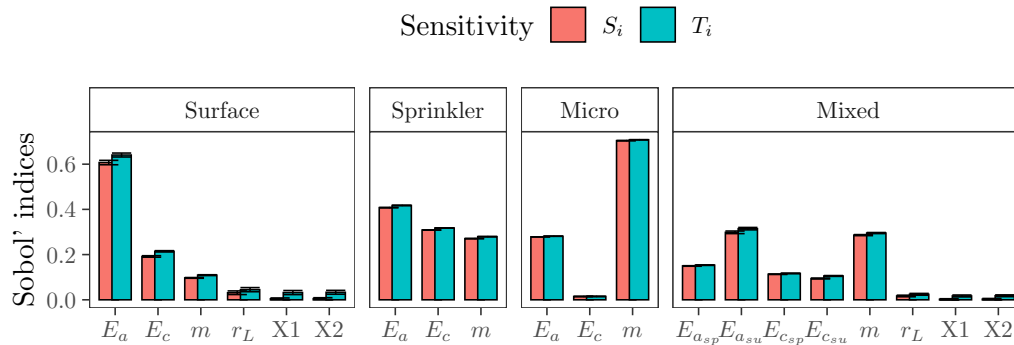
ind.dt <- rbindlist(out, idcol = "IFT") %>%
  .[, IFT:= factor(IFT, levels = c("Surface", "Sprinkler", "Micro", "Mixed"))]

tmp <- ind.dt[, .(mean = mean(original), sd = sd(original)),
               .(sensitivity, parameters, IFT)]

tmp2 <- tmp[!IFT == "Mixed"][, parameters:= ifelse(parameters == "Ea_surf", "$E_a$",
                                                    ifelse(parameters == "Ec_surf", "$E_c$",
                                                          ifelse(parameters == "Ea_sprinkler",
                                                                ifelse(parameters == "Ec_sprinkler",
                                                                      ifelse(parameters == "I
                                                                ifelse(parameters == "I

rbind(tmp[IFT == "Mixed"], tmp2) %>%
  ggplot(. , aes(parameters, mean, fill = sensitivity), color = black) +
  geom_bar(stat = "identity", position = position_dodge(0.6), color = "black") +
  geom_errorbar(aes(ymin = mean - sd, ymax = mean + sd), position = position_dodge(0.6)) +
  scale_x_discrete(labels = label_facets) +
  scale_fill_discrete(name = "Sensitivity", labels = c("$S_i$", "$T_i$")) +
  labs(x = "", y = "Sobol' indices") +
  facet_grid(~IFT, space = "free_x", scale = "free_x") +
  theme_AP()

```



```

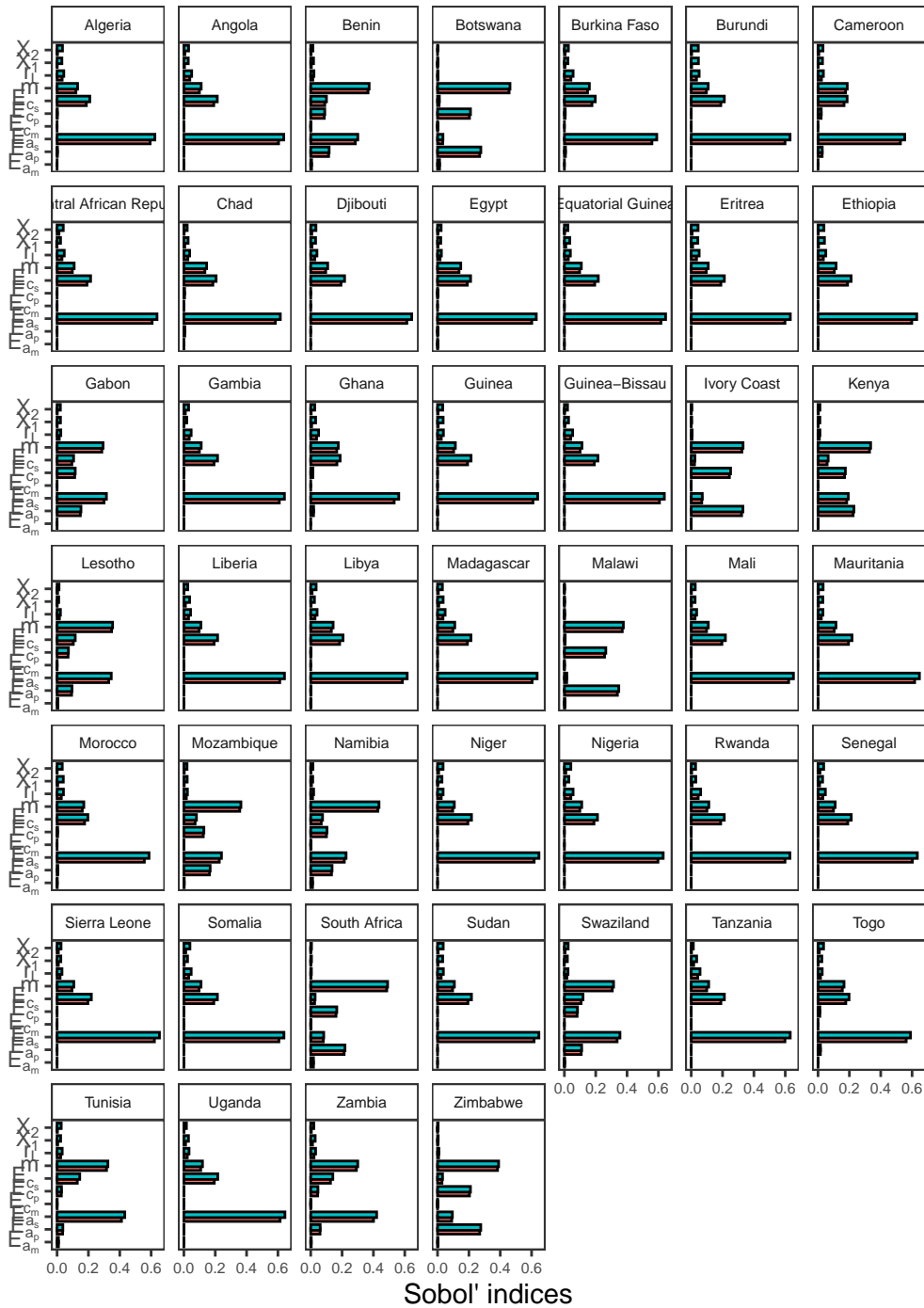
# EXTRACT SOBOL' INDICES FOR JAGER -----

jager.tmp <- lapply(y[["Jägermeyr et al. 2015"]], function(x) x$indices$results)

```


Africa

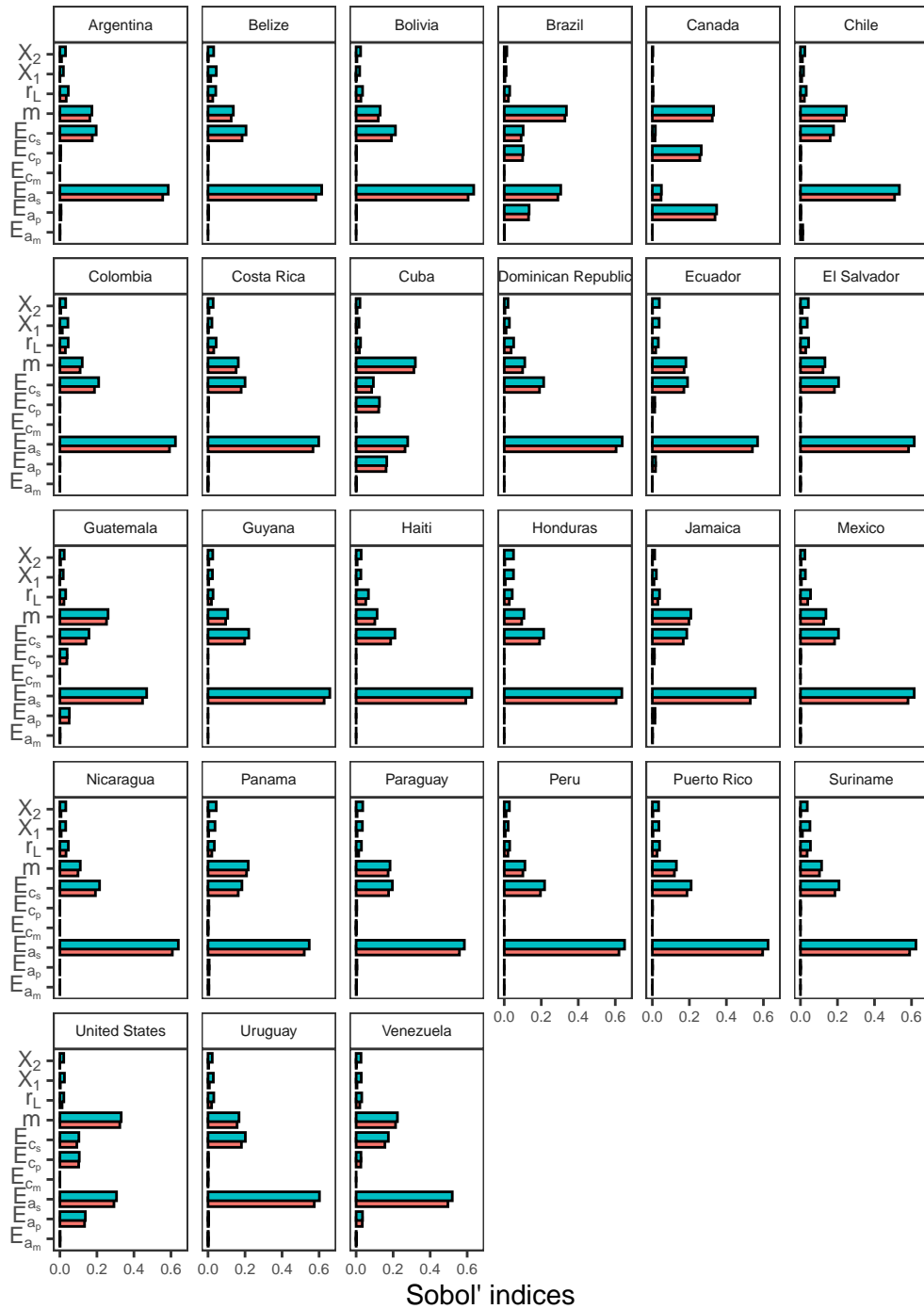
Sensitivity ■ Si ■ Ti



[[2]]

Americas

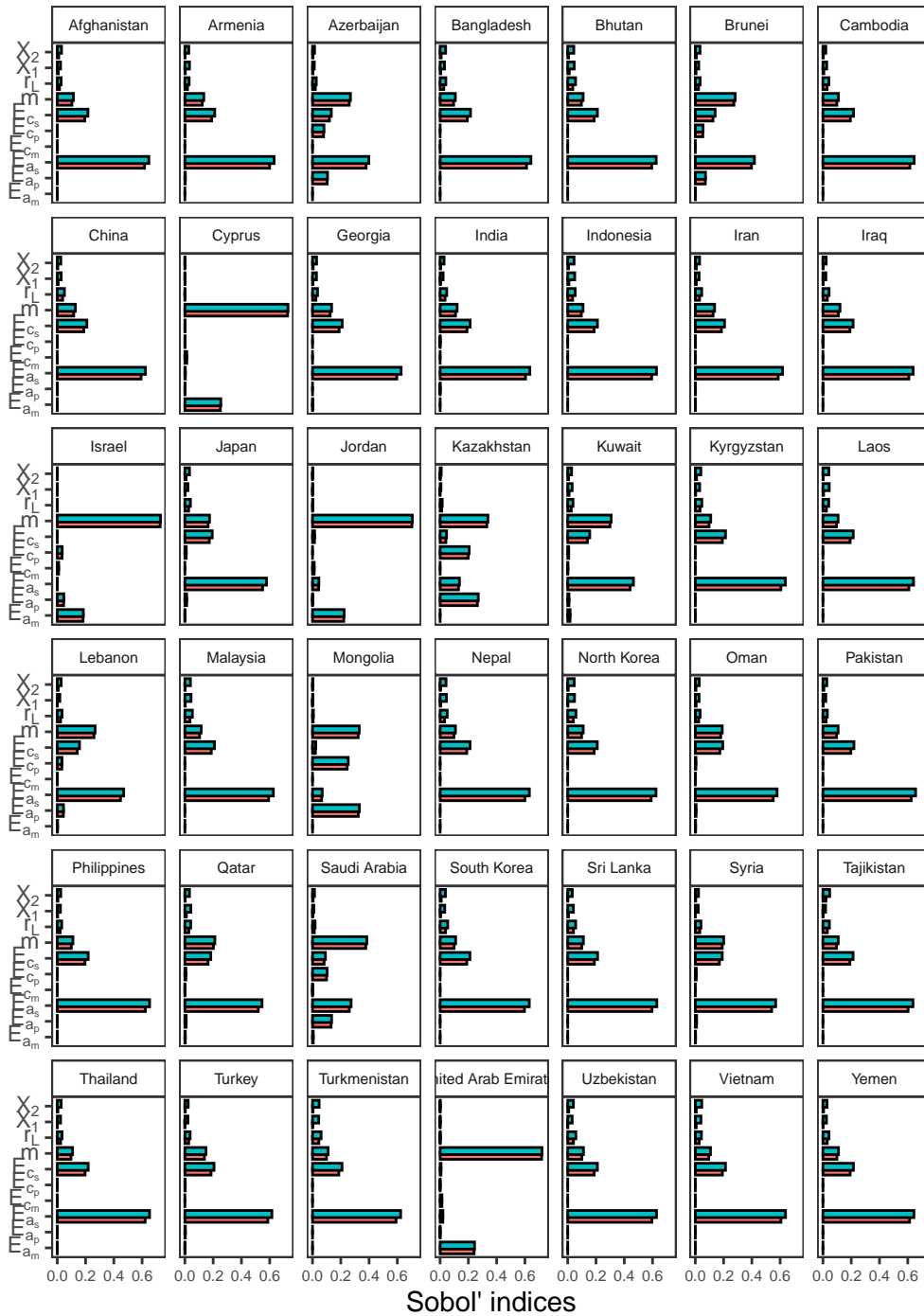
Sensitivity ■ Si ■ Ti



[[3]]

Asia

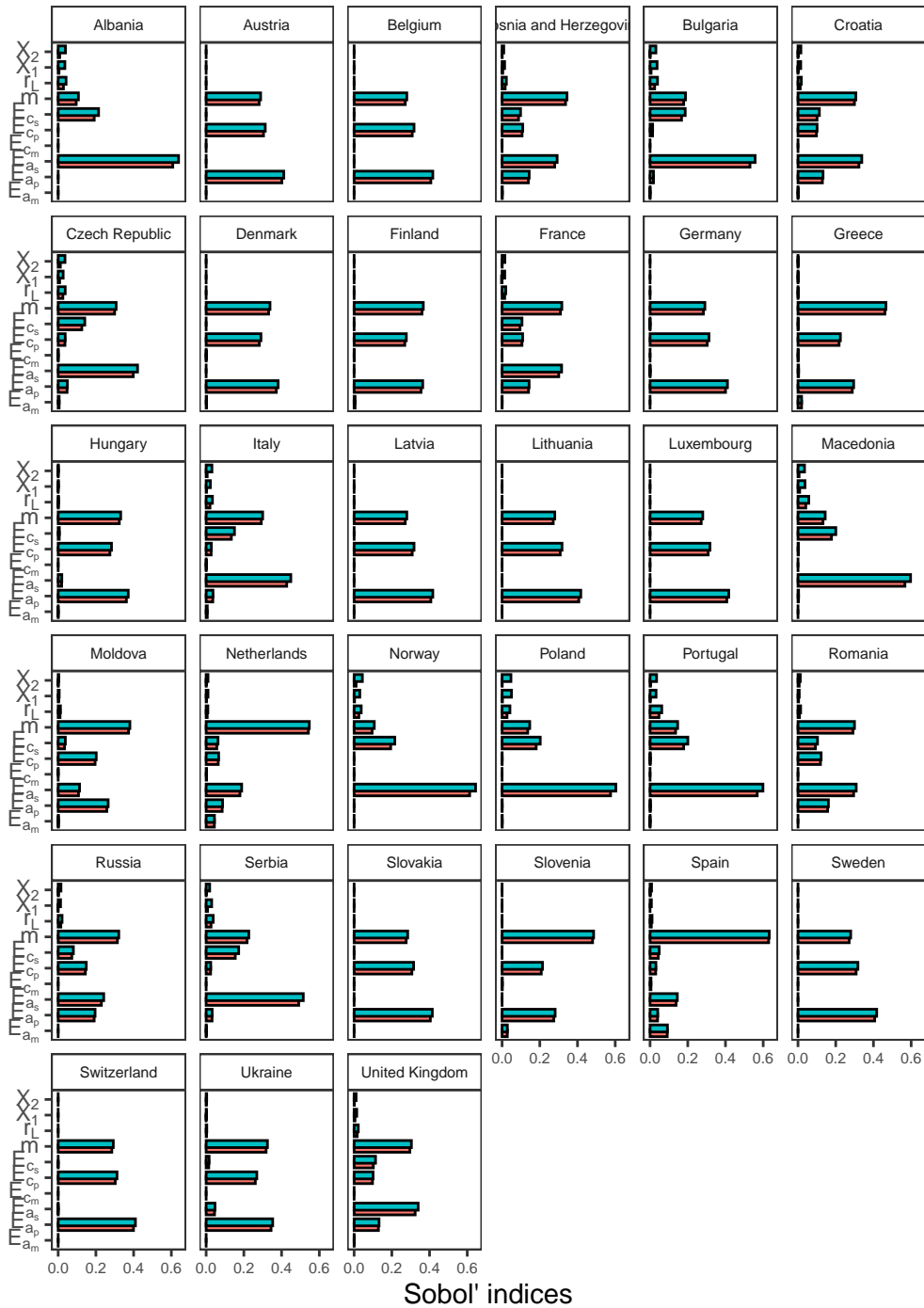
Sensitivity ■ Si ■ Ti



[[4]]

Europe

Sensitivity ■ Si ■ Ti



SESSION INFORMATION -----

```
sessionInfo()
```

```
## R version 4.0.3 (2020-10-10)
```

```

## Platform: x86_64-apple-darwin17.0 (64-bit)
## Running under: macOS Big Sur 10.16
##
## Matrix products: default
## BLAS:   /Library/Frameworks/R.framework/Versions/4.0/Resources/lib/libRblas.dylib
## LAPACK: /Library/Frameworks/R.framework/Versions/4.0/Resources/lib/libRlapack.dylib
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
##
## attached base packages:
## [1] parallel stats      graphics grDevices utils      datasets methods
## [8] base
##
## other attached packages:
## [1] checkpoint_1.0.0 benchmarkme_1.0.7 ncd4_1.17      rworldmap_1.3-6
## [5] sp_1.4-5          overlapping_1.6   testthat_3.0.4   scales_1.1.1
## [9] ggridges_0.5.3    countrycode_1.3.0 doParallel_1.0.16 iterators_1.0.13
## [13] foreach_1.5.1     cowplot_1.1.1    wesanderson_0.3.6 sensobol_1.0.3
## [17] forcats_0.5.1     stringr_1.4.0    dplyr_1.0.7      purrr_0.3.4
## [21] readr_2.0.1       tidyr_1.1.3      tibble_3.1.3     ggplot2_3.3.5
## [25] tidyverse_1.3.1   data.table_1.14.0
##
## loaded via a namespace (and not attached):
## [1] fs_1.5.0           lubridate_1.7.10    httr_1.4.2
## [4] tools_4.0.3        backports_1.2.1     utf8_1.2.2
## [7] R6_2.5.0           DBI_1.1.1           colorspace_2.0-2
## [10] withr_2.4.2        tidyrselect_1.1.1   gridExtra_2.3
## [13] compiler_4.0.3     cli_3.0.1           rvest_1.0.1
## [16] xml2_1.3.2         digest_0.6.27       foreign_0.8-81
## [19] rmarkdown_2.10     benchmarkmeData_1.0.4 pkgconfig_2.0.3
## [22] htmltools_0.5.1.1  highr_0.9           dbplyr_2.1.1
## [25] maps_3.3.0         rlang_0.4.11       readxl_1.3.1
## [28] rstudioapi_0.13    farver_2.1.0        generics_0.1.0
## [31] tikzDevice_0.12.3.1 jsonlite_1.7.2      magrittr_2.0.1
## [34] dotCall64_1.0-1    Matrix_1.3-4        Rcpp_1.0.7
## [37] munsell_0.5.0      fansi_0.5.0         viridis_0.6.1
## [40] lifecycle_1.0.0    stringi_1.7.3       yaml_2.2.1
## [43] plyr_1.8.6         grid_4.0.3          maptools_1.1-1
## [46] crayon_1.4.1       lattice_0.20-44     haven_2.4.3
## [49] hms_1.1.0          knitr_1.33          pillar_1.6.2
## [52] codetools_0.2-18   reprex_2.0.1        glue_1.4.2
## [55] evaluate_0.14      modelr_0.1.8        vctrs_0.3.8
## [58] spam_2.7-0         tzdb_0.1.2          Rdpack_2.1.2
## [61] cellranger_1.1.0   gtable_0.3.0        assertthat_0.2.1
## [64] xfun_0.25          rbibutils_2.2.3     broom_0.7.9
## [67] filehash_2.4-2     viridisLite_0.4.0   fields_12.5
## [70] ellipsis_0.3.2

```

```

## Return the machine CPU
cat("Machine:    "); print(get_cpu())$model_name)

## Machine:

## [1] "Intel(R) Core(TM) i9-9900K CPU @ 3.60GHz"

## Return number of true cores
cat("Num cores:  "); print(detectCores(logical = FALSE))

## Num cores:

## [1] 8

## Return number of threads
cat("Num threads: "); print(detectCores(logical = FALSE))

## Num threads:

## [1] 8

```