

The treatment of uncertainties in global water models

R code

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1 Preliminary functions

```
# PRELIMINARY FUNCTIONS -----

# 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(
  "bibliometrix", "tidyverse", "data.table", "scales", "pdfsearch", "pdfutils",
  "openxlsx", "cowplot", "wesanderson", "sjmisc", "ggpubr"))

# 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),
          strip.background = element_rect(fill = "white"),
          legend.margin = margin(0.5, 0.1, 0.1, 0.1),
          legend.box.margin=margin(0.2,-2,-7,-7))
}

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

checkpoint("2022-05-11",
          R.version = "4.2.0",
          checkpointLocation = getwd())
```

2 Bibliometric analysis

```
# VECTOR WITH NAME OF MODELS -----

models <- c("WaterGAP", "PCR-GLOBWB", "MATSIRO", "H08", "JULES-W1", "MPI-HM",
           "MHM", "LPJmL", "CWatM", "CLM", "DBHM", "ORCHIDEE")

models_vec <- paste(models, "_ref.bib", sep = "")
```

2.1 Analysis

```
# RUN FOR LOOP -----
output <- results <- years <- journals <- dt <- list()

for (i in 1:length(models_vec)) {

  output[[i]] <- convert2df(file = models_vec[i],
                           dbsource = "wos",
                           format = "bibtex")

  # Extract title -----

  title <- output[[i]]$TI

  # Extract Authors, Countries and Universities -----

  # Authors
  tmp.authors <- output[[i]]$AU
  first.author <- sub(" *\\;.+", "", tmp.authors)
  last.author <- sub(".*\\;", "", tmp.authors)

  # First author affiliation and country
  country.first <- sub(".*\\;", "", output[[i]]$RP)
  university.first <- sub(" *\\;.+", "", output[[i]]$affiliations)

  # Last author affiliation and country
  last.affiliation <- sub(".*\\;", "", output[[i]]$C1)
  country.last <- sub("\\.", "", sub(".*\\;", "", last.affiliation))
  university.last <- sub(".*\\;", "", output[[i]]$affiliations)

  # Extract keywords -----

  keywords <- gsub(";;", ";", output[[i]]$DE)
  keywords.plus <- gsub(";;", ";", output[[i]]$ID)

  # Create data.table -----
  dt[[i]] <- data.table("WOS" = output[[i]]$UT,
```

```

        "title" = title,
        "year" = output[[i]]$PY,
        "keywords" = keywords,
        "keywords.plus" = keywords.plus,
        "first.author" = first.author,
        "last.author" = last.author,
        "country.first" = country.first,
        "country.last" = country.last,
        "university.first" = university.first,
        "university.last" = university.last,
        "abstract" = output[[i]]$AB)

# Retrieve analysis bibliometrix -----
results[[i]] <- biblioAnalysis(output[[i]], sep = ";")
years[[i]] <- data.table(results[[i]]$Years)
journals[[i]] <- data.table(results[[i]]$Sources) %>%
  .[, SO:= str_to_title(SO)]
}

names(years) <- models
names(journals) <- models
names(dt) <- models

# ARRANGE DATA -----

# Correct for USA and China
colsName <- c("country.first", "country.last")
full.dt <- rbindlist(dt, idcol = "Model") %>%
  .[, (colsName):= lapply(.SD, function(x)
    ifelse(grepl("USA", x), "USA", x)), .SDcols = colsName] %>%
  .[, (colsName):= lapply(.SD, function(x)
    ifelse(grepl("CHINA", x), "CHINA", x)), .SDcols = colsName]

# Check which studies have "Uncertainty" or "Sensitivity" in keywords,
# keywords.plus or abstract
keywords_search <- c("UNCERTAINTY|SENSITIVITY")

tmp <- cbind(str_detect(full.dt$abstract, keywords_search),
  str_detect(full.dt$keywords.plus, keywords_search),
  str_detect(full.dt$keywords, keywords_search))

full.dt[, "uncertainty.sensitivity":= apply(tmp, 1, function(x) any(x, na.rm = TRUE))]

# Write dataset-
write.xlsx(full.dt, "full.dt.xlsx")

# Write dataset only with studies that have the keywords

```

```
full.dt[uncertainty.sensitivity == TRUE] %>%
  write.xlsx(., "full.dt.uncertain.xlsx")
```

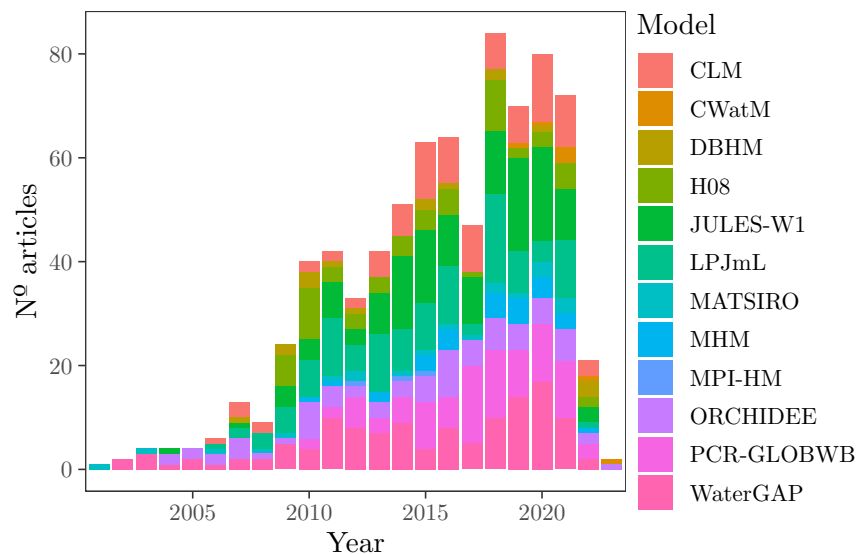
```
# PLOT -----
names(years) <- models
tmp <- rbindlist(years, idcol = "Model")[, .N, .(V1, Model)]

# Print total number of studies
tmp[, sum(N)]
```

```
## [1] 778
```

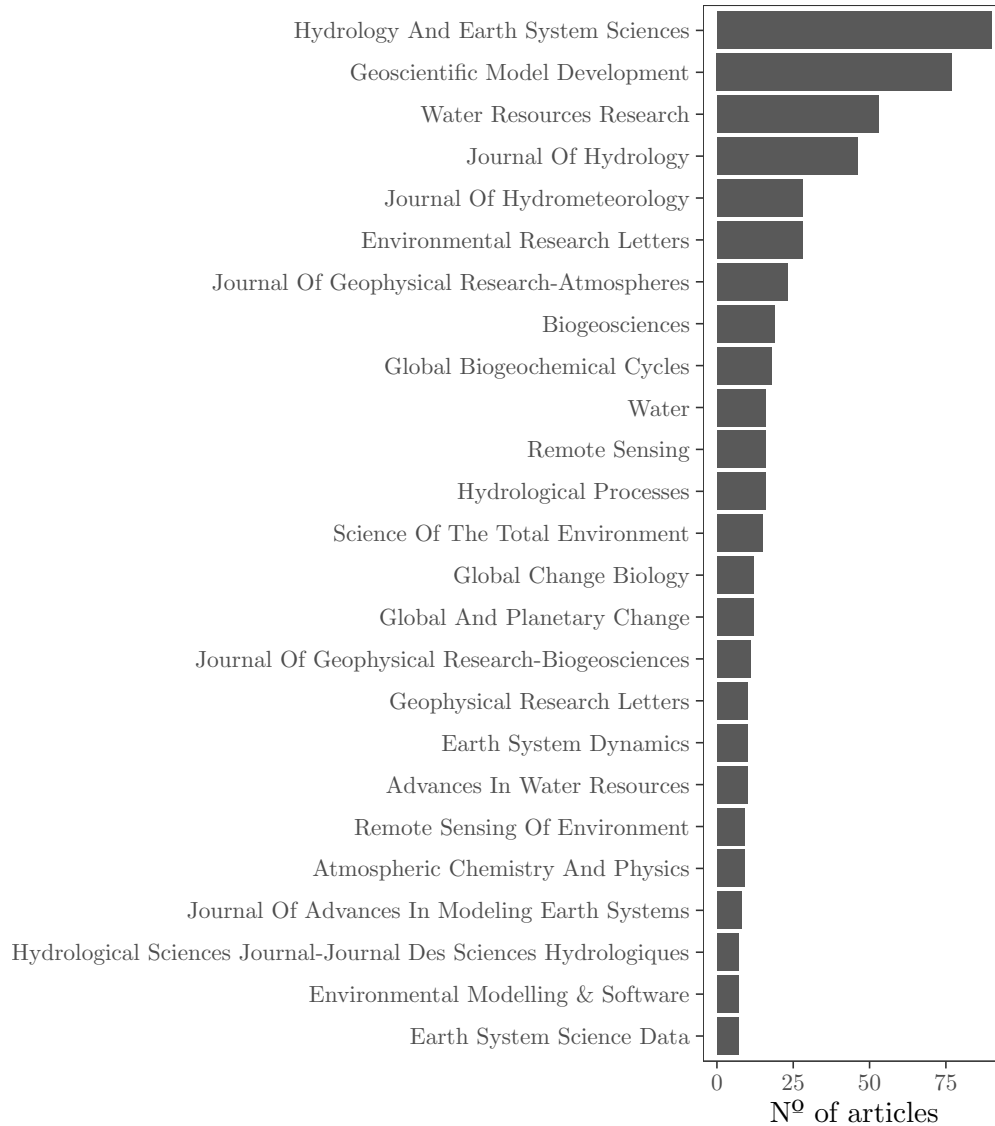
```
plot.time <- tmp %>%
  .[, V1:= as.factor(V1)] %>%
  ggplot(., aes(V1, N, fill = Model)) +
  geom_col() +
  scale_x_discrete(breaks = pretty_breaks(n = 3)) +
  labs(x = "Year", y = "N° articles") +
  theme_AP()
```

```
plot.time
```



```
# PLOT JOURNALS -----
rbindlist(journals, idcol = "Models") %>%
  .[, sum(N), S0] %>%
  .[order(-V1)] %>%
  .[, .SD[1:25]] %>%
  na.omit() %>%
  ggplot(., aes(x = reorder(S0, V1), y = V1)) +
  geom_bar(stat = "identity") +
  coord_flip() +
  labs(x = "", y = "N° of articles") +
```

```
theme_AP()
```



2.2 Institutional legacy in models

```
# ANALYSE DATASET -----

dt.use <- full.dt[, .N, .(Model, university.first)] %>%
  dcast(., university.first~ Model, value.var = "N")

for(j in seq_along(dt.use)){
  set(dt.use, i = which(is.na(dt.use[[j]]) & is.numeric(dt.use[[j]])), j = j, value = 0)
}

# Total number each institute uses a model
dt.use[, total:= rowSums(.SD), .SDcols = models]
```

```

# First 50
dt.50 <- copy(dt.use)[order(-total)][1:50]

# Turn lowercase of institutions except acronyms
exceptions <- c("USA", "UK", "CNRS", "IIASA", "DOE", "PCSHE", "IIT", "NCAR",
               "NOAA")
pattern <- sprintf("(?:%s)(*SKIP)(*FAIL)|\\b([A-Z])(\\w+)",
                  paste0(exceptions, collapse = "|"))
dt.50 <- dt.50[, university.first:= gsub(pattern, "\\1\\L\\2",
                                       university.first, perl = TRUE)]

tmp <- dt.50[, lapply(.SD, function(x) x / total), .SDcols = models] %>%
  .[, lapply(.SD, round, 2), .SDcols = models]

# RETRIEVE MAX VALUES PER INSTITUTE -----

matrix.50 <- as.matrix(tmp)
colIndex <- apply(matrix.50, 1, which.max)

out <- vector()
for(i in 1:length(colIndex)) {
  out[i] <- matrix.50[[i, colIndex[i]]]
}

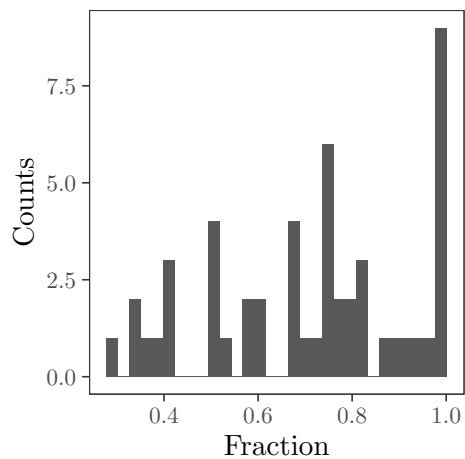
# Compute some statistics on the vector
f <- c(mean, median, min, max)
sapply(f, function(f) f(out, na.rm = TRUE))

## [1] 0.712 0.750 0.300 1.000

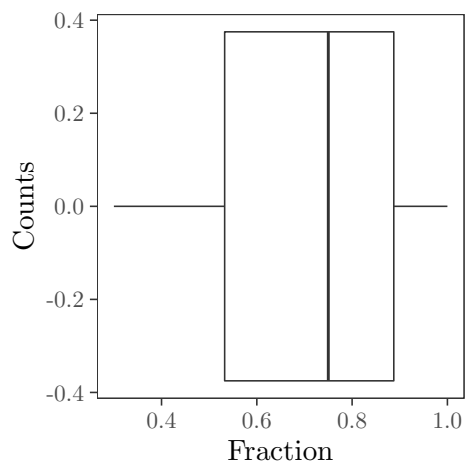
# Plot
data.table(out) %>%
  ggplot(., aes(out)) +
  geom_histogram() +
  labs(x = "Fraction", y = "Counts") +
  theme_AP()

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

```



```
data.table(out) %>%
  ggplot(., aes(out)) +
  geom_boxplot() +
  labs(x = "Fraction", y = "Counts") +
  theme_AP()
```



PLOT USE OF MODELS PER INSITUTE -----

```
a <- melt(dt.50, measure.vars = models) %>%
  na.omit() %>%
  .[, variable:= factor(variable, levels = sort(models))] %>%
  ggplot(., aes(value, university.first, fill = variable)) +
  scale_y_discrete(limits = rev) +
  labs(x = "Nº of articles", y = "") +
  scale_fill_discrete(name = "Model") +
  geom_bar(position = "stack", stat = "identity") +
  theme_AP() +
  theme(legend.position = "none",
        axis.text.y = element_text(size = 9))

b <- melt(dt.50, measure.vars = models) %>%
  na.omit() %>%
```

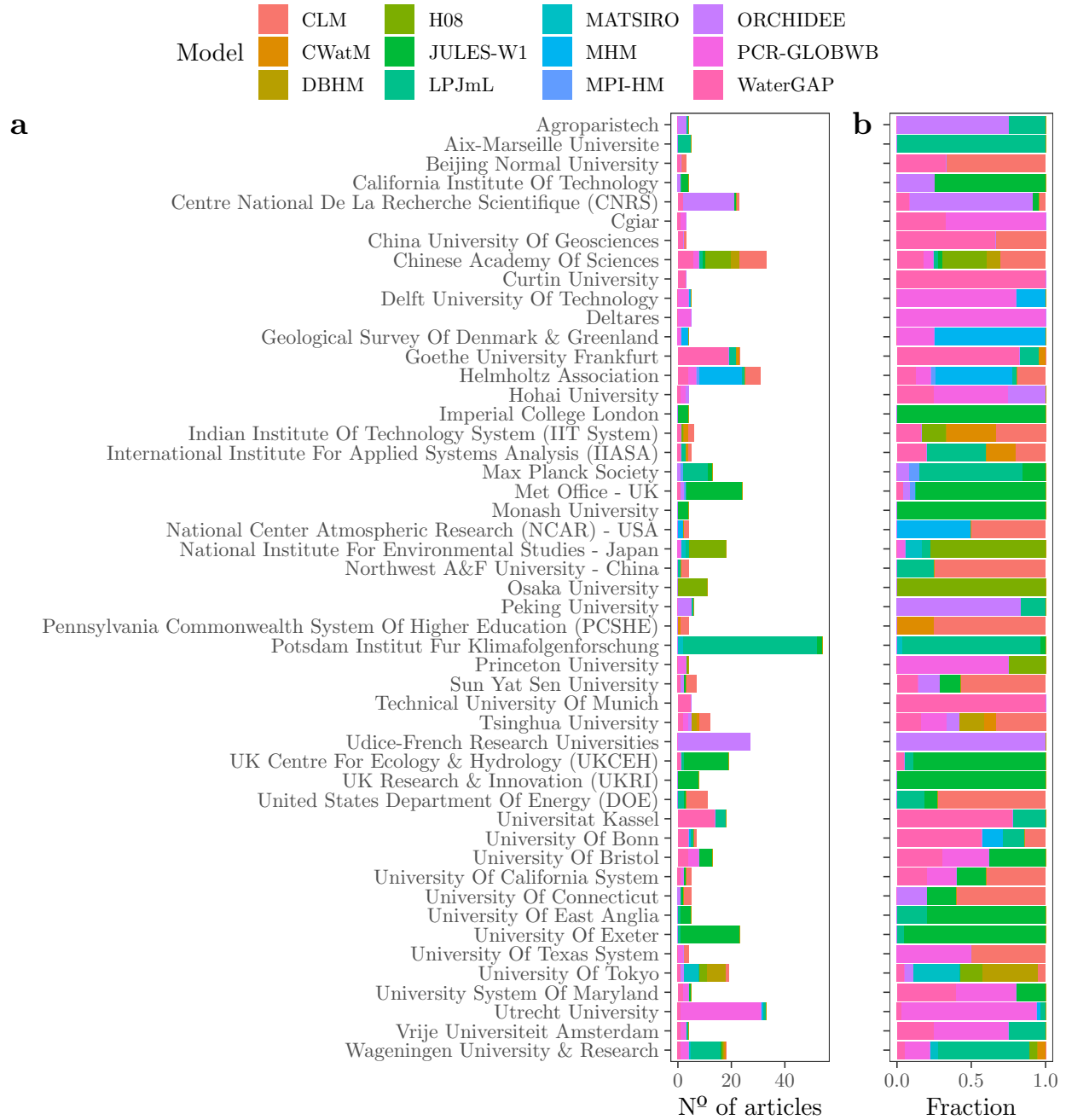


```

.[, variable:= factor(variable, levels = sort(models))] %>%
ggplot(., aes(value, university.first, fill = variable)) +
scale_y_discrete(limits = rev) +
labs(x = "Fraction", y = "") +
scale_fill_discrete(name = "Model") +
scale_x_continuous(breaks = pretty_breaks(n = 3)) +
geom_bar(position = "fill", stat = "identity") +
theme_AP() +
theme(axis.text.y = element_blank(),
      legend.position = "none")

legend <- get_legend(a + theme(legend.position = "top"))
bottom <- plot_grid(a, b, ncol = 2, labels = "auto", rel_widths = c(0.79 , 0.21))
ggarrange(legend, bottom, nrow = 2, heights = c(0.1, 0.9))

```



2.3 Keywords and keywords plus

```
# ANALYSE FREQUENCY OF KEYWORDS AND KEYWORDS PLUS -----

# Keywords
strsplit(full.dt$keywords, ";", fixed = TRUE) %>%
  unlist(lapply(., str_trim)) %>%
  data.table("keywords" = .) %>%
  .[, .N, keywords] %>%
  .[order(-N)] %>%
```

```
head(50)
```

```
##               keywords      N
##  1:                <NA> 367
##  2:          CLIMATE CHANGE 39
##  3:                GRACE 28
##  4:                GRACE 24
##  5:                MODEL 22
##  6:          CLIMATE CHANGE 20
##  7:          SOIL MOISTURE 18
##  8:          GROUNDWATER 15
##  9:          LAND SURFACE MODEL 15
## 10:    COMMUNITY LAND MODEL 15
## 11:          EVAPOTRANSPIRATION 14
## 12:                MODIS 13
## 13:                WGHM 12
## 14:          WATERGAP 11
## 15:          HYDROLOGY 11
## 16:          IRRIGATION 11
## 17:          PCR-GLOBWB 11
## 18:                WATER 10
## 19:          DROUGHT 10
## 20:          WATER SCARCITY 9
## 21:          MODELLING 9
## 22:          UNCERTAINTY 9
## 23:          DATA ASSIMILATION 9
## 24:                RUNOFF 8
## 25:          WATER RESOURCES 8
## 26:          SOIL MOISTURE 8
## 27:          WATER STRESS 7
## 28:          WATER USE 7
## 29:          PRECIPITATION 7
## 30:          MODELING 7
## 31:          REMOTE SENSING 7
## 32:          VEGETATION 6
## 33:                GLDAS 6
## 34:    GLOBAL HYDROLOGICAL MODEL 6
## 35:          CALIBRATION 6
## 36:                DATA 6
## 37:    TERRESTRIAL WATER STORAGE 6
## 38:                SOIL 6
## 39:                LPJML 6
## 40:    COMMUNITY LAND MODEL 6
## 41:          HYDROLOGICAL MODEL 5
## 42:          SCENARIOS 5
## 43:                GLOBAL 5
## 44:          WATER STORAGE 5
```

```
## 45:      HYDROLOGICAL MODELS  5
## 46:      ASSIMILATION        5
## 47:      LAND SURFACE MODELS  5
## 48:      VARIABILITY         5
## 49:      CHINA               5
## 50:      DROUGHT             5
##                keywords      N
```

```
# Keywords plus
```

```
strsplit(full.dt$keywords.plus, ";", fixed = TRUE) %>%
  unlist(lapply(., str_trim)) %>%
  data.table("keywords.plus" = .) %>%
  .[, .N, keywords.plus] %>%
  .[order(-N)] %>%
  head(50)
```

```
##                keywords.plus  N
##  1:      VARIABILITY 135
##  2:      MODEL 122
##  3:      CLIMATE 112
##  4:      WATER 95
##  5:      CLIMATE-CHANGE 93
##  6:      PRECIPITATION 72
##  7:      VEGETATION 68
##  8: ENVIRONMENT SIMULATOR JULES 67
##  9:      SOIL-MOISTURE 60
## 10:      IMPACT 59
## 11:      IMPACTS 57
## 12:      DYNAMICS 51
## 13:      VALIDATION 50
## 14:      RUNOFF 48
## 15:      RESOURCES 44
## 16:      CLIMATE-CHANGE 41
## 17:      EVAPOTRANSPIRATION 40
## 18:      UNCERTAINTY 40
## 19:      MODEL DESCRIPTION 40
## 20:      DROUGHT 39
## 21:      CARBON 37
## 22:      SIMULATION 36
## 23:      SENSITIVITY 35
## 24:      FLUXES 35
## 25:      CO2 34
## 26:      ENERGY 34
## 27:      TEMPERATURE 33
## 28:      SYSTEM 31
## 29:      AVAILABILITY 30
## 30:      PARAMETERIZATION 30
## 31:      BALANCE 29
```

```
## 32:          TRENDS 28
## 33:      ASSIMILATION 28
## 34:      STREAMFLOW 27
## 35:      PHOTOSYNTHESIS 27
## 36:  STOMATAL CONDUCTANCE 26
## 37:          BASIN 25
## 38:          SOIL 25
## 39:      REPRESENTATION 25
## 40:      MANAGEMENT 24
## 41:      CALIBRATION 24
## 42:      GROUNDWATER 24
## 43:          LAND 23
## 44:      EVAPORATION 23
## 45:      IRRIGATION 21
## 46:  INTEGRATED MODEL 21
## 47:          SURFACE 21
## 48:      PERFORMANCE 21
## 49:          FOREST 21
## 50:      PRODUCTIVITY 21
##          keywords.plus  N
```

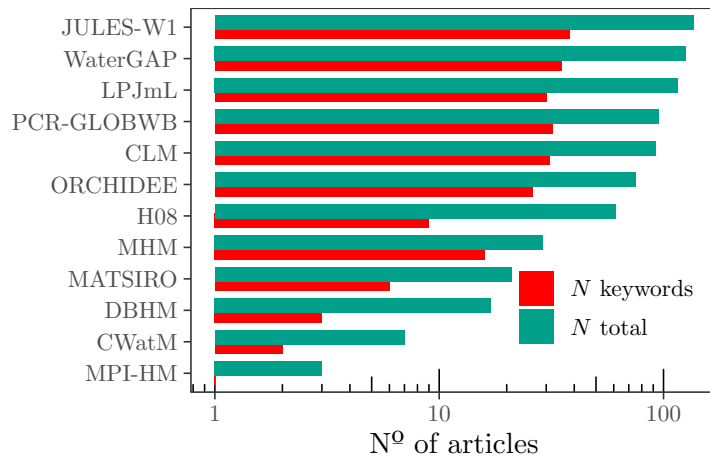
```
# PLOT TOTAL NUMBER OF STUDIES AND TOTAL NUMBER STUDIES WITH KEYWORDS -----
```

```
total.studies <- tmp[, sum(N), Model]
setnames(total.studies, "V1", "Total")

plot.bars <- full.dt[uncertainty.sensitivity == "TRUE"] %>%
  .[, sum(uncertainty.sensitivity), Model] %>%
  merge(., total.studies, by = "Model") %>%
  melt(., measure.vars = c("V1", "Total")) %>%
  ggplot(., aes(reorder(Model, value), value, fill = variable)) +
  coord_flip() +
  labs(y = "N° of articles", x = "") +
  scale_fill_manual(values = wes_palette(2, name = "Darjeeling1"),
                    name = "",
                    labels = c("$N$ keywords",
                              "$N$ total")) +

  scale_y_log10() +
  annotation_logticks(sides = "b") +
  geom_bar(stat = "identity", position = position_dodge(width = 0.6)) +
  theme_AP() +
  theme(legend.position = c(0.8, 0.3))

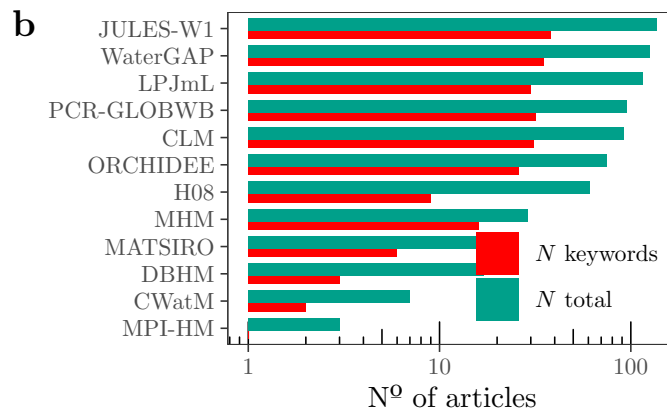
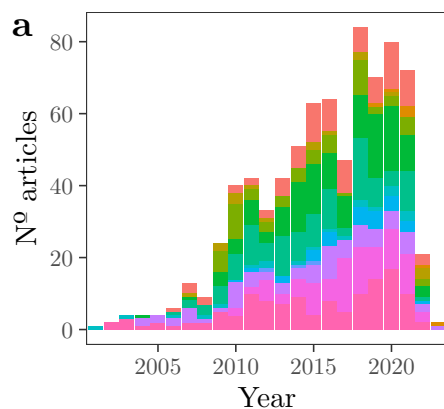
plot.bars
```



MERGE PLOTS -----

```
legend <- get_legend(plot.time + theme(legend.position = "top"))

bottom <- plot_grid(plot.time + theme(legend.position = "none"),
  plot.bars, ncol = 2, labels = "auto",
  rel_widths = c(0.4, 0.6))
plot_grid(legend, bottom, ncol = 1, rel_heights = c(0.3, 0.7))
```



KEYWORDS SEARCH -----

```
# Define vectors for search -----
directory <- "/Users/arnalduy/Documents/papers/ghms_bibliometric/"
directory_vec <- paste(directory, models, "_pdfs", sep = "")
keywords_vec <- c("sensitivity analysis", "uncertainty analysis", "uncertainty")
filename_keywords <- paste(models, "keywords", sep = "_")

# Loop -----
dt <- result <- list()
```

```

for (i in 1:length(directory_vec)) {

  result[[i]] <- keyword_directory(directory_vec[i],
                                   keyword = keywords_vec,
                                   split_pdf = TRUE)

  dt[[i]] <- data.table("name" = result[[i]]$pdf_name,
                       "keyword" = result[[i]]$keyword,
                       "text" = result[[i]]$line_text)

  fwrite(dt[[i]], file = paste(filename_keywords[i], ".csv", sep = ""))

}
names(result) <- models
names(dt) <- models

```

PLOT HISTOGRAMS WITH KEYWORDS -----

```

dt.keywords <- rbindlist(dt, idcol = "Model") %>%
  .[, .N, .(Model, name, keyword)] %>%
  .[, keyword:= str_to_title(keyword)]

plot.keywords.histogram <- dt.keywords %>%
  ggplot(., aes(N, fill = keyword, color = keyword)) +
  geom_histogram(position = "identity", alpha = 0.5) +
  facet_wrap(~Model, ncol = 6) +
  scale_y_continuous(breaks = pretty_breaks(n = 2)) +
  scale_x_continuous(breaks = pretty_breaks(n = 3)) +
  labs(x = "N° of mentions", y = "N° of papers") +
  theme_AP() +
  theme(legend.position = "top",
        strip.text.x = element_text(size = 8))

plot.keywords.histogram

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

```

