

Plot-based aboveground biomass estimates - AfriSAR sites

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NB. All aboveground biomass (AGB) estimates are in Mg ha⁻¹. Calibration points with Area_code names including 'h', 'q' and 'c' represent 1ha, 0.25ha and 0.16ha, respectively.

Loading packages and datasets

```
# LOPE: nine 1 ha plots (100 x 100 m) from 10 cm DBH & three 0.5 ha plots (100 x 50 m) from 5 cm DBH
# MABOUNIE: twelve 1 ha plots (100 x 100 m) from 10 cm DBH
# MONDAH: fifteen 1 ha plots (100 x 100 m) from 5 cm DBH
# RABI: one 25 ha plot (500 x 500 m) from 1 cm DBH
# Plus GEM plots (2 in LOPE and 4 in MONDAH; data obtained from ForestPlots.net)

# Packages
library(BIOMASS)
library(oce) # to compute Earth magnetic declination
```

```
## Loading required package: gsw
```

```
library(lubridate) # convert ymd dates to decimal year
```

```
##
## Attaching package: 'lubridate'
```

```
## The following object is masked from 'package:base':
##
##      date
```

```
library(sp)

# Tree-level and botanical datasets
load("AfriSARstem.rdata")
load("AfriSARBota.rdata")
```

Getting wood density (WD) using names and accepted synonyms

```

dfsyn <- read.csv("taxo_synonymes_acceptes_230414_modNL.csv", sep = ";"); dfsyn$taxon_concept_
name_valid[which(dfsyn$taxon_concept_name_valid == "")] <- NA
old.list <- strsplit(as.character(dfsyn$taxon_name), " "); new.list <- strsplit(as.character(d
fsyn$taxon_concept_name_valid), " ")
dfsyn$oldname <- paste(lapply(old.list, function(x) x[1]), lapply(old.list, function(x) x[2]))
dfsyn$newname <- paste(lapply(new.list, function(x) x[1]), lapply(new.list, function(x) x[2]))
dfsyn$newgen <- unlist(lapply(new.list, function(x) x[1]))
dfsyn$newsp <- unlist(lapply(new.list, function(x) x[2]))

AfriSARstem$Genus <- dfbota$genusCorr[match(AfriSARstem$Name, dfbota$ID)]
AfriSARstem$Species <- dfbota$speciesCorr[match(AfriSARstem$Name, dfbota$ID)]
AfriSARstem$FamilyAPG <- dfbota$familyAPG[match(AfriSARstem$Name, dfbota$ID)]
AfriSARstem$NameCorr <- paste(AfriSARstem$Genus, AfriSARstem$Species)

# Some trees (n=48) were identified at family level in the field; we manually fill the family
column
AfriSARstem$FamilyAPG[which(is.na(AfriSARstem$FamilyAPG) & !(is.na(AfriSARstem$Info_fam)))] <-
  AfriSARstem$Info_fam[which(is.na(AfriSARstem$FamilyAPG) & !(is.na(AfriSARstem$Info_fam)))]

AfriSARstem$Stand <- ifelse(AfriSARstem$Site == "RABI", "RABI", as.character(AfriSARstem$Plot_
code))
dataWD <- getWoodDensity(genus=AfriSARstem$Genus, species=AfriSARstem$Species, family=AfriSARs
tem$FamilyAPG, stand=AfriSARstem$Stand)

```

```

## The reference dataset contains 16467 wood density values
## Your taxonomic table contains 549 taxa

```

```

AfriSARstem$WD <- dataWD$meanWD
AfriSARstem$sdWD <- dataWD$sdWD
AfriSARstem$levelWD <- dataWD$levelWD
#table(dataWD$levelWD)

AfriSARstem$SynGen <- dfsyn$newgen[match(AfriSARstem$NameCorr, dfsyn$oldname)] # , nomatch = N
A, incomparables = ???
AfriSARstem$SynSp <- dfsyn$newsp[match(AfriSARstem$NameCorr, dfsyn$oldname)]

dataWDsyn <- getWoodDensity(genus=AfriSARstem$SynGen, species=AfriSARstem$SynSp, stand=AfriSAR
stem$Stand, family=AfriSARstem$FamilyAPG)

```

```

## The reference dataset contains 16467 wood density values
## Your taxonomic table contains 88 taxa

```

```

AfriSARstem$WDsyn <- dataWDsyn$meanWD
AfriSARstem$sdWDsyn <- dataWDsyn$sdWD
AfriSARstem$levelWDsyn <- dataWDsyn$levelWD

# 316 WD changed using synonyms
AfriSARstem$WD[which(!(AfriSARstem$levelWD %in% c("genus", "species")) & AfriSARstem$levelWDsyn
%in% c("genus", "species"))] <- AfriSARstem$WDsyn[which(!(AfriSARstem$levelWD %in% c("genus", "
species")) & AfriSARstem$levelWDsyn %in% c("genus", "species"))]
AfriSARstem$sdWD[which(!(AfriSARstem$levelWD %in% c("genus", "species")) & AfriSARstem$levelWDs

```

```

yn %in% c("genus", "species"))] <- AfriSARstem$sdWDSyn[which(!(AfriSARstem$levelWD %in% c("genus", "species")) & AfriSARstem$levelWDSyn %in% c("genus", "species"))]
AfriSARstem$levelWD[which(!(AfriSARstem$levelWD %in% c("genus", "species")) & AfriSARstem$levelWDSyn %in% c("genus", "species"))] <- AfriSARstem$levelWDSyn[which(!(AfriSARstem$levelWD %in% c("genus", "species")) & AfriSARstem$levelWDSyn %in% c("genus", "species"))]
#table(AfriSARstem$levelWD)

```

Refining permanent plot georeferencing

```

# Preliminary work in order to georeference the data
load("AfriSARplotcoord.rdata")

pattern <- c("SAV1", "SAV2", "SAV3", "COL1", "COL2", "COL3", "OKO1", "OKO2", "OKO3", "MAR1", "MAR2", "MIX1")
replacement <- c("LOP01", "LOP02", "LOP03", "LOP04", "LOP05", "LOP06", "LOP07", "LOP08", "LOP09", "LOP10", "LOP11", "LOP12")
for (i in (1:length(pattern))) {
  coordplot.afri$Point <- gsub(pattern[i], replacement[i], coordplot.afri$Point) # ignore.case = FALSE, perl = FALSE, fixed = FALSE, useBytes = FALSE
  coordplot.afri$Plot_code <- gsub(pattern[i], replacement[i], coordplot.afri$Plot_code) # ignore.case = FALSE, perl = FALSE, fixed = FALSE, useBytes = FALSE
}

FP.coord <- read.csv("Plot_coord_FP.csv", sep = ";") # Corresponds to coordinates of ForestPlots.net plots that were added to the study
coordplot.afri <- rbind(coordplot.afri, FP.coord)
magdev <- magneticField(coordplot.afri$long, coordplot.afri$lat, decimal_date(ymd(coordplot.afri$Time_census)))
coordplot.afri$True_bearing <- round((coordplot.afri$Compass_bearing - magdev$declination) %% 360, 0) # modulus operator %%

coordplot.afri$X_utm <- NA; coordplot.afri$Y_utm <- NA

coord.geoS <- SpatialPoints(cbind(coordplot.afri$long[which(coordplot.afri$lat < 0)], coordplot.afri$lat[which(coordplot.afri$lat < 0)], proj4string = CRS("+proj=longlat"))
coord.utmS <- spTransform(coord.geoS, CRS("+proj=utm +zone=32 +south +datum=WGS84 +units=m +no_defs +ellps=WGS84 +towgs84=0,0,0")) #; str(coord.utmS)
coordplot.afri$X_utm[which(coordplot.afri$lat < 0)] <- coord.utmS@coords[,1]
coordplot.afri$Y_utm[which(coordplot.afri$lat < 0)] <- coord.utmS@coords[,2]

coord.geoN <- SpatialPoints(cbind(coordplot.afri$long[which(coordplot.afri$lat > 0)], coordplot.afri$lat[which(coordplot.afri$lat > 0)], proj4string = CRS("+proj=longlat"))
coord.utmN <- spTransform(coord.geoN, CRS("+proj=utm +zone=32 +north +datum=WGS84 +units=m +no_defs +ellps=WGS84 +towgs84=0,0,0")) #; str(coord.utmN)
coordplot.afri$X_utm[which(coordplot.afri$lat > 0)] <- coord.utmN@coords[,1]
coordplot.afri$Y_utm[which(coordplot.afri$lat > 0)] <- coord.utmN@coords[,2]

## Get "true" bearing from coordinates for RABI, MONDAH, MABOUNIE and LOPE GEM (bearing from compass only for other LOPE plots)
afrisubplot <- c(as.character(unique(coordplot.afri$Plot_code[-which(coordplot.afri$Site == "LOPE")])), "LPG01")
coordplot.afri$Loc <- substring(coordplot.afri$Point, 6)

```

```

coordplot.afri$Loc[which(coordplot.afri$Plot_code %in% c("MON21A","RABI"))] <- c("a","b","c","d","a","b","d","c")

for (i in (1:length(afrisubplot))) {
  swe.afri <- (atan2(coordplot.afri$X_utm[which(coordplot.afri$Plot_code == afrisubplot[i] & c
coordplot.afri$Loc == "b")]) - coordplot.afri$X_utm[which(coordplot.afri$Plot_code == afrisubplo
t[i] & coordplot.afri$Loc == "a")]),
              coordplot.afri$Y_utm[which(coordplot.afri$Plot_code == afrisubplot[i] & c
coordplot.afri$Loc == "b")]) - coordplot.afri$Y_utm[which(coordplot.afri$Plot_code == afrisubplo
t[i] & coordplot.afri$Loc == "a")]))*180/pi)

  nwe.afri <- (atan2(coordplot.afri$X_utm[which(coordplot.afri$Plot_code == afrisubplot[i] & c
coordplot.afri$Loc == "c")]) - coordplot.afri$X_utm[which(coordplot.afri$Plot_code == afrisubplo
t[i] & coordplot.afri$Loc == "d")]),
              coordplot.afri$Y_utm[which(coordplot.afri$Plot_code == afrisubplot[i] & c
coordplot.afri$Loc == "c")]) - coordplot.afri$Y_utm[which(coordplot.afri$Plot_code == afrisubplo
t[i] & coordplot.afri$Loc == "d")]))*180/pi)

  coordplot.afri$True_bearing[which(coordplot.afri$Plot_code == afrisubplot[i])] <- round(mean
(c(swe.afri, nwe.afri)),1) + 270
}
coordplot.afri$True_bearing <- round((coordplot.afri$True_bearing) %% 360,1) # modulus operato
r %%
coordplot.afri$True_bearing[which(coordplot.afri$Plot_code == "LPG02")] <- unique(coordplot.af
ri$True_bearing[which(coordplot.afri$Plot_code == "LPG01")])

# Converting "true" bearing in (1) radians, and then in (2) plot rotation
coordplot.afri$TB_rad <- (pi/2 - (coordplot.afri$True_bearing*pi/180)) %% pi # TB stands for t
rue bearing
coordplot.afri$RotAng_rad <- (coordplot.afri$TB_rad - pi/2)

AfriSARstem$TreeRad <- sqrt(AfriSARstem$X_rel^2 + AfriSARstem$Y_rel^2)
AfriSARstem$TreeAng_rel <- atan2(AfriSARstem$X_rel, AfriSARstem$Y_rel)

# Assigning plot rotation to each stem
AfriSARstem$PlotAng <- ifelse(AfriSARstem$Site == "RABI",
                             coordplot.afri$RotAng_rad[which(coordplot.afri$Site == "RABI")],
                             coordplot.afri$RotAng_rad[match(AfriSARstem$Plot_code, coordplot
.afri$Plot_code)])

# Computing new stem coordinates after plot rotation
AfriSARstem$Xrot_rel <- AfriSARstem$X_rel * cos(AfriSARstem$PlotAng) - AfriSARstem$Y_rel * sin
(AfriSARstem$PlotAng) # x' = x * cos(theta) - y * sin(theta)
AfriSARstem$Yrot_rel <- AfriSARstem$X_rel * sin(AfriSARstem$PlotAng) + AfriSARstem$Y_rel * cos
(AfriSARstem$PlotAng) # y' = x * sin(theta) + y * cos(theta)

# Works because it selects the first value in the data.frame and that value is the one we need
(x=0; y=0)
AfriSARstem$X_abs <- ifelse(AfriSARstem$Site == "RABI",
                             coordplot.afri$X_utm[match(AfriSARstem$Site, coordplot.afri$Plot_c
ode)] + AfriSARstem$Xrot_rel,
                             coordplot.afri$X_utm[match(AfriSARstem$Plot_code, coordplot.afri$P
lot_code)] + AfriSARstem$Xrot_rel)

```

```
AfriSARstem$Y_abs <- ifelse(AfriSARstem$Site == "RABI",
                           coordplot.afri$Y_utm[match(AfriSARstem$Site, coordplot.afri$Plot_c
ode)] + AfriSARstem$Yrot_rel,
                           coordplot.afri$Y_utm[match(AfriSARstem$Plot_code, coordplot.afri$P
lot_code)] + AfriSARstem$Yrot_rel)

# CHANGING COORDINATES AFTER VISUAL INSPECTION OF BIG TREES LOCATION AND LIDAR-DERIVED CHM
# RABI, LOPE AND MONDAH OK; NOT POSSIBLE TO DO IT WITH MABOUNIE BECAUSE NO TREE-LEVEL COORDINA
TES
plotID.lope <- as.character(sort(unique(AfriSARstem$Plot_code[which(AfriSARstem$Site == "LOPE"
)])))
plotID.mondah <- as.character(sort(unique(AfriSARstem$Plot_code[which(AfriSARstem$Site == "MON
DAH")]])))

df.changcoord <- data.frame(plot = as.character(c(plotID.lope, plotID.mondah)),
                           modX = c(2,-4,-3,-3,-2,-3,-1,-6,5,-1,5,0,0,2,3,1,6,0,-1,6,-3,3,-1,
-2,0,-4,2,3,0,-2,0,-14),
                           modY = c(-3,1,2,3,8,3,1,-4,-1,-1,3,0,0,-2,1,3,7,0,2,-3,4,-3,1,2,0,
3,3,4,0,7,0,4),
                           stringsAsFactors=F)

AfriSARstem$X_absCORR <- AfriSARstem$X_abs + df.changcoord$modX[match(AfriSARstem$Plot_code, d
f.changcoord$plot)]
AfriSARstem$Y_absCORR <- AfriSARstem$Y_abs + df.changcoord$modY[match(AfriSARstem$Plot_code, d
f.changcoord$plot)]

AfriSARstem$X_absCORR[which(AfriSARstem$Site == "RABI")] <- AfriSARstem$X_abs[which(AfriSARste
m$Site == "RABI")] - 114
AfriSARstem$Y_absCORR[which(AfriSARstem$Site == "RABI")] <- AfriSARstem$Y_abs[which(AfriSARste
m$Site == "RABI")] + 42

coordplot.afri$X_utmCORR <- coordplot.afri$X_utm + df.changcoord$modX[match(coordplot.afri$Plo
t_code, df.changcoord$plot)]
coordplot.afri$Y_utmCORR <- coordplot.afri$Y_utm + df.changcoord$modY[match(coordplot.afri$Plo
t_code, df.changcoord$plot)]

coordplot.afri$X_utmCORR[which(coordplot.afri$Site == "RABI")] <- coordplot.afri$X_utm[which(c
oordplot.afri$Site == "RABI")] - 114
coordplot.afri$Y_utmCORR[which(coordplot.afri$Site == "RABI")] <- coordplot.afri$Y_utm[which(c
oordplot.afri$Site == "RABI")] + 42

coordplot.afri$X_utmCORR[which(coordplot.afri$Site == "MABOUNIE")] <- coordplot.afri$X_utm[whi
ch(coordplot.afri$Site == "MABOUNIE")]
coordplot.afri$Y_utmCORR[which(coordplot.afri$Site == "MABOUNIE")] <- coordplot.afri$Y_utm[whi
ch(coordplot.afri$Site == "MABOUNIE")]

coordplot.afri$X_utmCORR[which(coordplot.afri$Plot_code == "LOP01")] <- coordplot.afri$X_utm[w
hich(coordplot.afri$Plot_code == "LOP01")]
coordplot.afri$Y_utmCORR[which(coordplot.afri$Plot_code == "LOP01")] <- coordplot.afri$Y_utm[w
hich(coordplot.afri$Plot_code == "LOP01")]
```

Creating georeferenced sets of calibration points

(at 1ha and 0.16-0.25ha)

```

site = c("RABI", "LOPE", "MABOUNIE", "MONDAH")
fullscale = c(100, 50, 20)
smallplot = c("LOP04", "LOP05", "LOP06")
suffixe = c("h", "q")

# Creating dataframe to georeference quarter hectare features (n=620)
coord_orig_q <- coordplot.afri[which(coordplot.afri$X_rel == 0 & coordplot.afri$Y_rel == 0),]
coord_orig_q$full_lengthX <- NA; coord_orig_q$full_lengthY <- NA
coord_orig_q$full_lengthX[-which(coord_orig_q$Plot_code == "LPG02")] <- coordplot.afri$X_rel[w
high(coordplot.afri$Loc == "b")]
coord_orig_q$full_lengthY[-which(coord_orig_q$Plot_code == "LPG02")] <- coordplot.afri$Y_rel[w
high(coordplot.afri$Loc == "d")]
coord_orig_q$full_lengthX[which(is.na(coord_orig_q$full_lengthX))] <- 100
coord_orig_q$full_lengthY[which(is.na(coord_orig_q$full_lengthY))] <- 100
coord_orig_temp <- coord_orig_q

# Creating dataframe to georeference hectare features (n=119)
coord_orig_h <- coord_orig_temp
coord_orig_h <- coord_orig_h[-which(coord_orig_h$full_lengthX < 100 | coord_orig_h$full_length
Y < 100),] # Removing 50x50m plots

scale.list <- list(); site.list <- list() # plot.df <- data.frame(); # Yet, plot.df already de
fined later in the loops
for (i in (1:length(site))) {
  if (site[i] %in% c("RABI", "LOPE", "MONDAH")) scale = fullscale[1:2] else scale = fullscale[
c(1,3)]

  for (j in (1:length(scale))) {
    if (j == 1) coord_orig = coord_orig_h else coord_orig = coord_orig_q
    plot.df <- data.frame()
    tempoplot <- as.character(coord_orig$Plot_code[which(coord_orig$Site == site[i])]); tempop
lot # used to be sort(as.character(...)) but plot order dealt with cf. coord_orig
    #tempoplot <- if (scale[j] == 100) tempoplot[!tempoplot %in% smallplot] else tempoplot; te
mpoplot # NB. ifelse() can't return vectors !

    for (k in (1:length(tempoplot))) {
      lengthX <- coord_orig$full_lengthX[which(coord_orig$Plot_code == tempoplot[k])]; lengthX
      lengthY <- coord_orig$full_lengthY[which(coord_orig$Plot_code == tempoplot[k])]; lengthY

      incrX_h <- cos(coord_orig$RotAng_rad[which(coord_orig$Plot_code == tempoplot[k])]) * sca
le[j] # increment for X coordinates horizontally
      incrY_h <- sin(coord_orig$RotAng_rad[which(coord_orig$Plot_code == tempoplot[k])]) * sca
le[j] # increment for Y coordinates horizontally

      incrX_v <- cos(coord_orig$RotAng_rad[which(coord_orig$Plot_code == tempoplot[k])] + pi/2
) * scale[j] # increment for X coordinates vertically; also equals (-incrY_h)
      incrY_v <- sin(coord_orig$RotAng_rad[which(coord_orig$Plot_code == tempoplot[k])] + pi/2
) * scale[j] # increment for Y coordinates vertically; also equals incrX_h

      nbptX <- length(seq(0, lengthX, scale[j]))

```

```
nbptY <- length(seq(0, lengthY, scale[j]))
incrX.mat <- matrix(rep(0:(nbptX-1),nbptY), nrow=nbptY, ncol=nbptX, byrow = T); incrX.ma
t
incrY.mat <- matrix(rep(rev(0:(nbptY-1)),nbptX), nrow=nbptY, ncol=nbptX); incrY.mat

XX <- coord_orig$X_utmCORR[which(coord_orig$Plot_code == tempoplot[k])] + incrX_h * incr
X.mat + incrX_v * incrY.mat
YY <- coord_orig$Y_utmCORR[which(coord_orig$Plot_code == tempoplot[k])] + incrY_h * incr
X.mat + incrY_v * incrY.mat
#plot(as.vector(YY) ~ as.vector(XX))

XX_SW.mat <- XX[2:nbptY, 1:(nbptX-1)]; YY_SW.mat <- YY[2:nbptY, 1:(nbptX-1)]
XX_NW.mat <- XX[1:(nbptY-1), 1:(nbptX-1)]; YY_NW.mat <- YY[1:(nbptY-1), 1:(nbptX-1)]
XX_SE.mat <- XX[2:nbptY, 2:nbptX]; YY_SE.mat <- YY[2:nbptY, 2:nbptX]
XX_NE.mat <- XX[1:(nbptY-1), 2:nbptX]; YY_NE.mat <- YY[1:(nbptY-1), 2:nbptX]

XX_SW.vect <- as.vector(XX_SW.mat); YY_SW.vect <- as.vector(YY_SW.mat)
XX_NW.vect <- as.vector(XX_NW.mat); YY_NW.vect <- as.vector(YY_NW.mat)
XX_SE.vect <- as.vector(XX_SE.mat); YY_SE.vect <- as.vector(YY_SE.mat)
XX_NE.vect <- as.vector(XX_NE.mat); YY_NE.vect <- as.vector(YY_NE.mat)

for (l in (1:(nbptX-1))) {
  XX_SW.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(XX_SW.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
  XX_NW.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(XX_NW.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
  XX_SE.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(XX_SE.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
  XX_NE.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(XX_NE.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
}
for (l in (1:(nbptX-1))) {
  YY_SW.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(YY_SW.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
  YY_NW.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(YY_NW.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
  YY_SE.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(YY_SE.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
  YY_NE.vect[((l-1)*(nbptY-1)+1):(l*(nbptY-1))] <- rev(YY_NE.vect[((l-1)*(nbptY-1)+1):(l
*(nbptY-1))])
}

if(site[i] == "MABOUNIE" & j == 2) {
  XX_SW.vect <- XX_SW.vect[c(1,4,16,19)]; YY_SW.vect <- YY_SW.vect[c(1,4,16,19)]
  XX_NW.vect <- XX_NW.vect[c(2,5,17,20)]; YY_NW.vect <- YY_NW.vect[c(2,5,17,20)]
  XX_SE.vect <- XX_SE.vect[c(6,9,21,24)]; YY_SE.vect <- YY_SE.vect[c(6,9,21,24)]
  XX_NE.vect <- XX_NE.vect[c(7,10,22,25)]; YY_NE.vect <- YY_NE.vect[c(7,10,22,25)]
  nbptX <- length(XX_SW.vect) - 1; nbptY <- length(YY_SW.vect) - 1
}

templot.df <- data.frame(Site = rep(site[i], (nbptX-1) * (nbptY-1)),
  Area_code = if (site[i] == "RABI") paste(sprintf(fmt=paste("RAB
%0",nchar((nbptX-1)*(nbptY-1)),"d",sep=""),
c(1:(nbptX-1)
```



```

*(nbptY-1))), suffixe[j],sep="")
                                else if (scale[j] == 100) paste(tempoplot[k], suffixe[j], sep="
")
                                else if (site[i] == "MABOUNIE" & j == 2) paste(tempoplot[k], "c
", c(1:((nbptX-1)*(nbptY-1))), sep="")
                                else paste(tempoplot[k], suffixe[j], c(1:((nbptX-1)*(nbptY-1)))
, sep=""),
                                Plot_code = rep(tempoplot[k], (nbptX-1) * (nbptY-1)),
                                Scale = paste(ifelse(site[i] == "MABOUNIE" & j == 2, scale[j] *
2, scale[j])^2/10^4,"ha", sep=""), # scale[j]
                                sw_x = XX_SW.vect, sw_y = YY_SW.vect,
                                nw_x = XX_NW.vect, nw_y = YY_NW.vect,
                                se_x = XX_SE.vect, se_y = YY_SE.vect,
                                ne_x = XX_NE.vect, ne_y = YY_NE.vect)
                                plot.df <- rbind(plot.df, templot.df)
                                }
                                scale.list[[j]] <- plot.df
                                }
                                site.list[[i]] <- scale.list
                                }
                                #site.list

```

Assigning trees to hectares (1ha), quarters (0.25ha) and corners (0.16ha) based on Plot_code and relative XY

```

AfriSARstem$Hect_code <- ifelse(AfriSARstem$Plot_code %in% c("MND01","MND02"), NA, p
aste(AfriSARstem$Plot_code,"h",sep=""))

AfriSARstem$quartX <- ifelse(AfriSARstem$X_rel == 0, 1, ceiling(AfriSARstem$X_rel/50))
AfriSARstem$quartY <- ifelse(AfriSARstem$Y_rel == 0, 1, ceiling(AfriSARstem$Y_rel/50))
AfriSARstem$plotnbQ <- ifelse(AfriSARstem$Site == "RABI", (AfriSARstem$quartX-1)*10 + AfriSARs
tem$quartY,
                                ifelse(AfriSARstem$Plot_code %in% smallplot, (AfriSARstem$quartX
-1)*1 + AfriSARstem$quartY, (AfriSARstem$quartX-1)*2 + AfriSARstem$quartY))

AfriSARstem$Quart_code <- ifelse(is.na(AfriSARstem$plotnbQ), NA,
                                ifelse(AfriSARstem$Site == "RABI", paste(sprintf(fmt="RAB%03d
",AfriSARstem$plotnbQ),"q",sep=""), paste(AfriSARstem$Plot_code,"q",AfriSARstem$plotnbQ,sep="
"))))

AfriSARstem$Corn_code <- NA
cornerblock <- list(c(1, 2, 6, 7), c(4, 5, 9, 10), c(16, 17, 21, 22), c(19, 20, 24, 25))
for (i in (1:length(cornerblock))) {
    AfriSARstem$Corn_code[which(AfriSARstem$Site == "MABOUNIE" & AfriSARstem$Info_loc %in% corne
rblock[[i]])] <- paste(AfriSARstem$Plot_code[which(AfriSARstem$Site == "MABOUNIE" & AfriSARste
m$Info_loc %in% cornerblock[[i]])], "c", i, sep="")
}

```


Estimating H from Feldpausch H:D relationship

```
range(AfriSARstem$Diameter)

## [1] 10.0 251.6

dataHfeld <- retrieveH(D=AfriSARstem$Diameter, region = "CAfrica"); range(dataHfeld) # H ranges
  from 6.2 - 49.7 m

## [1] 6.17700 49.69018

AfriSARstem$Hfeld <- dataHfeld$H
AfriSARstem$HfeldRSE <- dataHfeld$RSE
```

Developing local H:D relationships (5 in total: 1 per site + 1 for savanna specie SAV.SP)

```
# Subset data for HD model building (NB. sav sp. excluded)
AfriSARforHD <- AfriSARstem[-which(is.na(AfriSARstem$Height) | (AfriSARstem$NameCorr %in% c("C
rossopteryx febrifuga", "Sarcocephalus latifolius"))),] # | AfriSARstem$Plot == "SAV3"
AfriSARforHD <- AfriSARforHD[which(AfriSARforHD$H_info %in% c("a", "ae", "e", "efg", "el", "eq", "es
", "i", "l", "lq", "q", "s", NA)),]

# Compute site-specific H:D models
HDmodelPerSite <- by(AfriSARforHD, AfriSARforHD$Site,
  function(x) modelHD(D=x$Diameter, H=x$Height, method="michaelis", useWeight
    =T),
  simplify=FALSE)
RSEmodels <- sapply(HDmodelPerSite, function(x) x$RSE)
Coeffmodels <- lapply(HDmodelPerSite, function(x) x$coefficients)
ResHD <- data.frame(Site=names(unlist(RSEmodels)),
  a=round(unlist(sapply(Coeffmodels, "[", 1)), 3),
  b=round(unlist(sapply(Coeffmodels, "[", 2)), 3),
  RSE=round(unlist(RSEmodels), 3))

# Retrieve predicted height values in the database
AfriSARstem$Hlocal <- AfriSARstem$Height # keeping directly measured trees
AfriSARstem$HlocRSE <- 1 # to be refined?! Assume a 1-m error on directly measured trees
AfriSARstem$levelHloc <- "FIELD"

Site=as.character(ResHD$Site)
for(i in 1:length(ResHD$Site)){
  filt<-AfriSARstem$Site==Site[i] & is.na(AfriSARstem$Hlocal)
  AfriSARstem$Hlocal[filt]<-retrieveH(D=AfriSARstem$Diameter[filt], model=HDmodelPerSite[[Site[
i]]])$H
  AfriSARstem$HlocRSE[filt]<-HDmodelPerSite[[Site[i]]]$RSE
  AfriSARstem$levelHloc[filt]<-Site[i]
```

```

}

# Model for savanna species
# NB. from lope.csv keeping only sav sp. with flag1 %in% c("a","ae","e","i") and Diameter < 15
# (but DBH < 10, otherwise can't build allometry because n too small)
dfsavsp <- read.csv("savsp4hd.csv", sep=";", stringsAsFactors=T)

HDmodel.sav <- modelHD(D=dfsavsp$Diameter, H=dfsavsp$Height, drawGraph = FALSE, useWeight=TRUE
, method="michaelis")
coefHDmodel.sav <- HDmodel.sav$coefficients

dataHlocal.sav <- retrieveH(D=AfriSARstem$Diameter[which(!(AfriSARstem$levelHloc == "FIELD") &
  AfriSARstem$NameCorr %in% c("Crossopteryx febrifuga","Sarcocephalus latifolius"))], model = H
Dmodel.sav)
AfriSARstem$Hlocal[which(!(AfriSARstem$levelHloc == "FIELD") & AfriSARstem$NameCorr %in% c("Cr
ossopteryx febrifuga","Sarcocephalus latifolius"))] <- dataHlocal.sav$H
AfriSARstem$HlocRSE[which(!(AfriSARstem$levelHloc == "FIELD") & AfriSARstem$NameCorr %in% c("C
rossopteryx febrifuga","Sarcocephalus latifolius"))] <- dataHlocal.sav$RSE
AfriSARstem$HlocRSE[which((AfriSARstem$levelHloc == "FIELD") & AfriSARstem$NameCorr %in% c("Cr
ossopteryx febrifuga","Sarcocephalus latifolius"))] <- 0.1
AfriSARstem$levelHloc[which(!(AfriSARstem$levelHloc == "FIELD") & AfriSARstem$NameCorr %in% c(
"Crossopteryx febrifuga","Sarcocephalus latifolius"))] <- "SAVASP"

table(is.na(AfriSARstem$Hlocal)) # all stems have Hloc

```

Assigning mean plot coordinates to trees to get environmental factor E

```

longitude <- tapply(coordplot.afri$long, coordplot.afri$Plot_code, mean)
latitude <- tapply(coordplot.afri$lat, coordplot.afri$Plot_code, mean)
meancoord <- data.frame(Plot_code=names(longitude), long=as.numeric(longitude), lat=as.numeric
(latitude))

AfriSARstem$long <- ifelse(AfriSARstem$Site == "RABI",
                          meancoord[match(AfriSARstem$Site, meancoord$Plot_code),"long"],
                          meancoord[match(AfriSARstem$Plot_code, meancoord$Plot_code),"long"]
)
AfriSARstem$lat <- ifelse(AfriSARstem$Site == "RABI",
                          meancoord[match(AfriSARstem$Site, meancoord$Plot_code),"lat"],
                          meancoord[match(AfriSARstem$Plot_code, meancoord$Plot_code),"lat"])

```

Compute AGB at hectare/quarter/corner level using 3 different models

```

AfriSARstem <- AfriSARstem[with(AfriSARstem, order(Site, decreasing = c(F), method = "radix"))
,]
resolAGB <- c("Hect_code", "Quart_code", "Corn_code")
coefmult <- c(1, 4, 25/4)

```

AGB FELDPAUSCH (agb_fph)

```
AGB_fph.list <- list()

rm(resultMC_FeldGB); gc()
resultMC_FeldGB <- by(AfriSARstem, AfriSARstem["Site"],
                      function(x) AGBmonteCarlo(D=x$Diameter, WD=x$WD, errWD=x$sdWD, H=x$Hfeld
                      ,
                      errH=x$HfeldRSE, Dpropag="chave2004"), simplif
y=F)

tempLOP <- as.data.frame(resultMC_FeldGB$LOPE$AGB_simu)
tempMAB <- as.data.frame(resultMC_FeldGB$MABOUNIE$AGB_simu)
tempMON <- as.data.frame(resultMC_FeldGB$MONDAH$AGB_simu)
tempRAB <- as.data.frame(resultMC_FeldGB$RABI$AGB_simu)
tempAFRI <- rbind(tempLOP,tempMAB,tempMON,tempRAB)
Afriprop_FELD <- cbind(AfriSARstem, tempAFRI)

for (i in (1:length(resolAGB))) {
  tempocalc <- by(Afriprop_FELD, Afriprop_FELD[,resolAGB[i]],
                  function(x) list(meanAGB = mean(apply(x[,46:1045], 2, sum, na.rm = T)),
                                   #medAGB = median(apply(x[,46:1045], 2, sum, na.rm = T)),
                                   #sdAGB = sd(apply(x[,46:1045], 2, sum, na.rm = T)),
                                   credibilityAGB = quantile(apply(x[,46:1045], 2, sum, na.rm
= T), probs = c(0.025,0.975))))

  AGB_fph.list[[i]] <- data.frame(Area_code = names(tempocalc),
                                agb_fph = round(as.numeric(sapply(tempocalc,"[,1))*coefmult
[i],1),
                                cred_fph_2.5 = round(as.numeric(lapply(sapply(tempocalc,"[,
2), function(x) x[1]))*coefmult[i],1),
                                cred_fph_97.5 = round(as.numeric(lapply(sapply(tempocalc,"[
,2), function(x) x[2]))*coefmult[i],1), stringsAsFactors = F)
  rownames(AGB_fph.list[[i]]) <- NULL
}
AGB_fph.list
AGB_fph.df <- Reduce(rbind, AGB_fph.list)
AGB_fph.df
```

##	Area_code	agb_fph	cred_fph_2.5	cred_fph_97.5
## 1	LOP02h	1.2	0.8	2.1
## 2	LOP03h	29.2	21.5	40.6
## 3	LOP07h	342.8	316.0	373.4
## 4	LOP08h	306.0	287.0	326.6
## 5	LOP09h	366.0	330.5	405.3
## 6	LOP10h	417.3	376.7	464.0
## 7	LOP11h	405.1	361.9	458.9
## 8	LOP12h	382.7	349.8	416.9
## 9	LPG01h	474.6	397.5	580.8
## 10	LPG02h	597.1	518.4	687.2

Plot-based aboveground biomass estimates - AfriSAR sites

##	11	MAB01h	376.3	343.1	412.4
##	12	MAB02h	349.3	312.0	389.5
##	13	MAB03h	378.6	337.4	432.5
##	14	MAB04h	501.9	440.7	572.3
##	15	MAB05h	481.5	420.5	572.2
##	16	MAB06h	356.9	319.0	402.9
##	17	MAB07h	286.0	258.5	321.5
##	18	MAB08h	350.5	318.2	388.8
##	19	MAB09h	466.4	421.0	516.1
##	20	MAB10h	393.9	348.1	446.9
##	21	MAB11h	495.1	450.8	544.3
##	22	MAB12h	216.0	196.7	238.9
##	23	MNG03h	528.3	484.4	574.4
##	24	MNG04h	460.4	421.9	508.7
##	25	MON01h	31.9	26.0	39.8
##	26	MON02h	338.5	278.6	422.6
##	27	MON03h	62.1	55.2	70.2
##	28	MON05h	111.7	99.5	125.8
##	29	MON09h	3.0	2.2	4.0
##	30	MON10h	125.8	107.3	150.4
##	31	MON11h	40.3	36.1	44.8
##	32	MON13h	282.2	251.5	323.2
##	33	MON14h	175.9	154.7	202.1
##	34	MON19h	4.2	3.4	5.2
##	35	MON20h	87.5	70.1	113.9
##	36	MON21Ah	171.7	158.5	186.0
##	37	MON21h	2.5	1.9	3.3
##	38	MON22h	342.7	305.0	389.5
##	39	MON23h	150.3	133.0	171.3
##	40	RAB01h	264.5	233.3	301.5
##	41	RAB02h	327.3	289.5	373.3
##	42	RAB03h	343.7	315.9	376.0
##	43	RAB04h	343.5	314.1	377.7
##	44	RAB05h	367.7	320.4	424.8
##	45	RAB06h	337.6	305.5	376.9
##	46	RAB07h	477.6	415.3	550.6
##	47	RAB08h	244.1	225.5	265.1
##	48	RAB09h	317.2	280.5	361.6
##	49	RAB10h	368.5	318.3	431.8
##	50	RAB11h	337.5	297.4	389.9
##	51	RAB12h	358.1	321.6	399.8
##	52	RAB13h	293.0	266.4	325.3
##	53	RAB14h	292.1	266.2	321.2
##	54	RAB15h	361.1	324.1	404.1
##	55	RAB16h	298.8	256.8	356.9
##	56	RAB17h	309.3	279.4	343.8
##	57	RAB18h	303.3	259.4	358.6
##	58	RAB19h	308.0	276.5	348.6
##	59	RAB20h	409.0	356.9	473.7
##	60	RAB21h	346.6	306.3	395.5
##	61	RAB22h	282.7	258.9	309.4
##	62	RAB23h	339.0	294.9	390.7
##	63	RAB24h	576.8	509.1	658.6
##	64	RAB25h	344.8	308.6	388.5

Plot-based aboveground biomass estimates - AfriSAR sites

##	65	LOP02q2	0.7	0.2	1.3
##	66	LOP02q3	2.5	0.8	5.6
##	67	LOP02q4	1.8	1.1	3.0
##	68	LOP03q1	15.6	11.9	21.0
##	69	LOP03q2	93.8	62.8	139.1
##	70	LOP03q3	3.1	2.2	4.2
##	71	LOP03q4	4.3	2.0	8.3
##	72	LOP04q1	312.3	269.5	362.2
##	73	LOP04q2	150.5	127.4	177.6
##	74	LOP05q1	53.7	44.0	64.9
##	75	LOP05q2	77.4	67.1	89.8
##	76	LOP06q1	166.3	149.2	184.5
##	77	LOP06q2	214.6	193.6	238.1
##	78	LOP07q1	382.2	321.6	452.5
##	79	LOP07q2	346.4	299.9	393.5
##	80	LOP07q3	312.5	267.2	363.3
##	81	LOP07q4	330.2	281.8	393.7
##	82	LOP08q1	310.5	272.7	351.2
##	83	LOP08q2	316.3	282.8	349.6
##	84	LOP08q3	293.2	257.0	334.4
##	85	LOP08q4	304.0	267.2	349.9
##	86	LOP09q1	464.8	373.7	582.5
##	87	LOP09q2	294.7	254.2	340.6
##	88	LOP09q3	266.9	225.0	319.0
##	89	LOP09q4	437.7	367.2	534.1
##	90	LOP10q1	371.5	318.9	434.0
##	91	LOP10q2	355.2	292.3	427.5
##	92	LOP10q3	540.9	444.8	661.6
##	93	LOP10q4	401.4	316.3	526.6
##	94	LOP11q1	483.6	395.6	614.7
##	95	LOP11q2	313.0	263.3	380.2
##	96	LOP11q3	446.5	341.5	600.7
##	97	LOP11q4	377.4	300.0	477.1
##	98	LOP12q1	427.7	369.4	489.8
##	99	LOP12q2	351.0	301.8	414.0
##	100	LOP12q3	494.2	411.6	609.2
##	101	LOP12q4	257.8	222.8	299.0
##	102	LPG01q1	404.1	332.0	495.7
##	103	LPG01q2	411.8	329.7	524.9
##	104	LPG01q3	273.0	226.4	328.7
##	105	LPG01q4	808.4	543.2	1202.0
##	106	LPG02q1	461.9	358.4	598.3
##	107	LPG02q2	948.3	735.6	1247.7
##	108	LPG02q3	296.0	234.4	373.7
##	109	LPG02q4	676.8	552.4	851.9
##	110	MND01q1	433.0	368.0	511.3
##	111	MND01q2	431.3	360.5	517.2
##	112	MND01q3	457.6	381.6	538.3
##	113	MND01q4	462.6	386.9	552.0
##	114	MND02q1	332.7	282.1	393.4
##	115	MND02q2	486.7	394.5	595.8
##	116	MND02q3	512.3	436.8	607.5
##	117	MND02q4	572.4	493.6	668.7
##	118	MNG03q1	574.6	495.5	670.7

Plot-based aboveground biomass estimates - AfriSAR sites

##	119	MNG03q2	546.2	461.2	653.9
##	120	MNG03q3	426.2	360.7	502.0
##	121	MNG03q4	566.2	480.7	666.0
##	122	MNG04q1	628.7	524.8	753.8
##	123	MNG04q2	380.0	332.1	438.4
##	124	MNG04q3	360.5	300.1	436.0
##	125	MNG04q4	472.3	399.9	568.9
##	126	MON01q1	15.5	11.5	20.7
##	127	MON01q2	7.2	5.4	9.8
##	128	MON01q3	40.4	27.8	59.0
##	129	MON01q4	64.6	46.7	92.8
##	130	MON02q1	212.0	163.1	271.2
##	131	MON02q2	139.6	103.7	190.0
##	132	MON02q3	823.2	604.9	1153.7
##	133	MON02q4	179.4	153.1	211.1
##	134	MON03q1	30.5	21.9	45.2
##	135	MON03q2	19.4	13.7	28.0
##	136	MON03q3	76.8	63.7	94.0
##	137	MON03q4	121.7	103.4	142.1
##	138	MON05q1	182.7	146.7	226.3
##	139	MON05q2	184.2	158.9	221.0
##	140	MON05q3	31.6	22.8	43.0
##	141	MON05q4	48.4	37.4	63.3
##	142	MON09q1	2.5	1.3	4.4
##	143	MON09q2	1.6	0.9	2.4
##	144	MON09q3	0.7	0.4	1.2
##	145	MON09q4	7.0	4.6	10.5
##	146	MON10q1	115.4	67.5	198.5
##	147	MON10q2	132.5	104.8	167.7
##	148	MON10q3	75.8	58.6	101.4
##	149	MON10q4	179.4	139.3	233.3
##	150	MON11q1	24.8	19.5	31.3
##	151	MON11q2	36.7	28.9	47.2
##	152	MON11q3	36.0	27.8	46.8
##	153	MON11q4	63.7	55.1	73.2
##	154	MON13q1	368.6	308.2	443.5
##	155	MON13q2	294.5	250.9	344.6
##	156	MON13q3	187.5	149.2	241.9
##	157	MON13q4	278.2	204.3	401.7
##	158	MON14q1	285.8	227.4	368.0
##	159	MON14q2	217.0	179.0	269.2
##	160	MON14q3	132.4	99.6	179.4
##	161	MON14q4	68.4	60.9	77.9
##	162	MON19q1	10.2	7.7	13.2
##	163	MON19q2	0.4	0.1	0.8
##	164	MON19q3	5.8	3.9	8.3
##	165	MON19q4	0.6	0.2	1.1
##	166	MON20q1	176.7	114.3	284.4
##	167	MON20q2	134.2	111.1	166.1
##	168	MON20q3	26.3	19.8	34.8
##	169	MON20q4	12.9	9.9	16.9
##	170	MON21Aq1	142.4	123.8	164.8
##	171	MON21Aq2	95.7	79.3	117.7
##	172	MON21Aq3	249.7	221.1	281.4

Plot-based aboveground biomass estimates - AfriSAR sites

##	173	MON21Aq4	198.9	171.7	230.2
##	174	MON21q1	0.3	0.1	0.8
##	175	MON21q3	5.5	3.6	8.3
##	176	MON21q4	4.3	3.0	6.0
##	177	MON22q1	150.8	131.2	173.6
##	178	MON22q2	521.7	416.6	647.6
##	179	MON22q3	150.4	130.3	174.3
##	180	MON22q4	547.9	453.1	677.6
##	181	MON23q1	360.7	306.2	425.4
##	182	MON23q2	186.7	146.9	236.7
##	183	MON23q3	21.2	11.7	36.3
##	184	MON23q4	32.6	19.9	52.4
##	185	RAB001q	294.5	242.3	365.4
##	186	RAB002q	192.8	165.2	230.0
##	187	RAB003q	429.7	356.3	525.3
##	188	RAB004q	246.6	179.9	362.7
##	189	RAB005q	330.3	284.5	379.7
##	190	RAB006q	268.3	229.0	316.5
##	191	RAB007q	414.6	350.8	490.8
##	192	RAB008q	328.5	280.0	386.0
##	193	RAB009q	485.0	385.2	628.7
##	194	RAB010q	495.8	360.4	683.5
##	195	RAB011q	233.4	197.9	274.9
##	196	RAB012q	337.3	246.5	475.2
##	197	RAB013q	358.7	277.8	465.3
##	198	RAB014q	270.9	225.4	330.3
##	199	RAB015q	372.0	316.3	443.0
##	200	RAB016q	404.0	339.5	494.8
##	201	RAB017q	267.5	221.8	323.0
##	202	RAB018q	363.1	306.5	432.8
##	203	RAB019q	235.8	193.3	288.9
##	204	RAB020q	254.3	214.6	307.9
##	205	RAB021q	277.7	240.4	322.4
##	206	RAB022q	414.6	341.4	510.1
##	207	RAB023q	424.6	328.6	555.1
##	208	RAB024q	734.1	565.1	955.1
##	209	RAB025q	284.2	251.5	319.5
##	210	RAB026q	182.2	155.4	214.3
##	211	RAB027q	216.6	182.4	261.8
##	212	RAB028q	375.8	304.5	474.8
##	213	RAB029q	517.2	417.4	661.2
##	214	RAB030q	209.0	168.1	272.1
##	215	RAB031q	353.0	285.6	453.6
##	216	RAB032q	305.0	259.3	359.5
##	217	RAB033q	530.5	415.7	667.0
##	218	RAB034q	221.3	185.1	268.2
##	219	RAB035q	256.1	219.7	303.7
##	220	RAB036q	253.8	214.5	305.9
##	221	RAB037q	390.2	294.5	532.0
##	222	RAB038q	286.4	246.3	334.8
##	223	RAB039q	287.0	238.6	357.8
##	224	RAB040q	461.0	314.7	678.9
##	225	RAB041q	383.4	283.6	542.0
##	226	RAB042q	309.3	248.7	392.5

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##	227	RAB043q	353.7	287.8	442.4
##	228	RAB044q	360.1	293.5	451.7
##	229	RAB045q	278.2	235.5	331.5
##	230	RAB046q	255.3	215.2	302.4
##	231	RAB047q	380.9	317.4	466.3
##	232	RAB048q	342.7	286.7	408.6
##	233	RAB049q	264.7	220.4	320.5
##	234	RAB050q	408.2	332.4	513.0
##	235	RAB051q	352.5	295.1	426.0
##	236	RAB052q	304.9	236.5	406.1
##	237	RAB053q	267.0	224.4	322.4
##	238	RAB054q	451.6	372.8	553.9
##	239	RAB055q	393.1	320.4	491.6
##	240	RAB056q	245.4	207.1	294.3
##	241	RAB057q	202.9	163.7	252.7
##	242	RAB058q	242.0	206.0	287.1
##	243	RAB059q	404.1	323.6	520.7
##	244	RAB060q	367.5	300.4	443.6
##	245	RAB061q	409.2	290.9	618.9
##	246	RAB062q	175.1	144.7	212.5
##	247	RAB063q	321.4	267.9	400.4
##	248	RAB064q	369.3	297.6	470.4
##	249	RAB065q	272.3	206.0	374.8
##	250	RAB066q	344.9	249.2	505.6
##	251	RAB067q	310.9	248.0	394.1
##	252	RAB068q	196.1	168.2	232.1
##	253	RAB069q	563.5	417.1	773.1
##	254	RAB070q	416.3	335.9	525.7
##	255	RAB071q	282.9	219.6	374.1
##	256	RAB072q	327.4	266.2	406.7
##	257	RAB073q	274.4	228.9	335.6
##	258	RAB074q	271.9	230.8	325.4
##	259	RAB075q	207.9	177.3	241.4
##	260	RAB076q	388.1	292.2	544.3
##	261	RAB077q	317.1	264.0	391.4
##	262	RAB078q	407.7	330.3	524.0
##	263	RAB079q	400.3	325.4	490.2
##	264	RAB080q	252.9	198.4	330.2
##	265	RAB081q	419.9	344.5	515.6
##	266	RAB082q	364.5	287.6	476.7
##	267	RAB083q	306.7	261.5	365.1
##	268	RAB084q	320.5	264.6	401.9
##	269	RAB085q	220.7	173.7	282.0
##	270	RAB086q	360.3	287.3	460.6
##	271	RAB087q	795.2	607.0	1039.3
##	272	RAB088q	479.2	388.7	616.6
##	273	RAB089q	360.7	277.8	469.3
##	274	RAB090q	245.1	204.6	297.4
##	275	RAB091q	396.4	310.2	536.4
##	276	RAB092q	205.7	167.1	265.2
##	277	RAB093q	209.3	181.0	241.3
##	278	RAB094q	293.1	253.5	339.2
##	279	RAB095q	340.1	264.9	444.0
##	280	RAB096q	434.5	333.2	597.0

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##	281	RAB097q	492.9	410.6	593.9
##	282	RAB098q	539.8	432.6	679.3
##	283	RAB099q	415.5	343.1	507.4
##	284	RAB100q	357.9	292.8	455.3
##	285	MAB01c1	245.6	197.7	300.5
##	286	MAB01c2	384.1	315.7	464.7
##	287	MAB01c3	341.4	285.6	405.4
##	288	MAB01c4	291.1	247.2	346.4
##	289	MAB02c1	335.3	267.5	435.2
##	290	MAB02c2	202.6	162.1	246.3
##	291	MAB02c3	317.2	240.9	408.2
##	292	MAB02c4	592.3	457.2	760.4
##	293	MAB03c1	244.6	193.2	316.3
##	294	MAB03c2	324.6	261.2	414.0
##	295	MAB03c3	832.0	627.4	1117.8
##	296	MAB03c4	371.2	297.2	475.7
##	297	MAB04c1	504.9	391.5	670.2
##	298	MAB04c2	446.5	334.1	624.0
##	299	MAB04c3	626.0	485.7	825.6
##	300	MAB04c4	301.5	238.5	384.5
##	301	MAB05c1	501.6	398.3	635.7
##	302	MAB05c2	321.6	257.7	401.4
##	303	MAB05c3	331.2	272.6	410.8
##	304	MAB05c4	391.3	306.2	494.4
##	305	MAB06c1	320.3	247.3	427.2
##	306	MAB06c2	404.9	316.4	521.8
##	307	MAB06c3	210.0	167.9	272.9
##	308	MAB06c4	463.2	364.6	596.0
##	309	MAB07c1	242.8	201.2	293.4
##	310	MAB07c2	264.7	218.9	323.2
##	311	MAB07c3	212.3	176.7	255.8
##	312	MAB07c4	416.8	310.2	592.0
##	313	MAB08c1	318.0	244.7	433.6
##	314	MAB08c2	370.8	282.6	508.7
##	315	MAB08c3	278.7	227.9	346.2
##	316	MAB08c4	354.2	290.6	438.2
##	317	MAB09c1	461.4	374.9	565.8
##	318	MAB09c2	404.7	335.6	488.4
##	319	MAB09c3	451.7	347.6	588.9
##	320	MAB09c4	657.1	526.6	822.4
##	321	MAB10c1	383.9	247.1	598.0
##	322	MAB10c2	258.1	218.4	307.1
##	323	MAB10c3	565.3	426.3	759.9
##	324	MAB10c4	674.1	533.0	865.6
##	325	MAB11c1	399.6	327.7	492.6
##	326	MAB11c2	406.7	310.9	546.2
##	327	MAB11c3	882.6	735.8	1056.4
##	328	MAB11c4	314.4	253.5	381.8
##	329	MAB12c1	319.4	264.0	394.1
##	330	MAB12c2	175.9	139.9	227.3
##	331	MAB12c3	169.4	137.4	212.7
##	332	MAB12c4	175.3	144.3	213.9

AGB USING ENVIRONMENTAL FACTOR E (agb_chv)

```
AGB_chv.list <- list()

rm(resultMC_ChaveGB); gc()
resultMC_ChaveGB <- by(AfriSARstem, AfriSARstem[, "Site"],
                      function(x) AGBmonteCarlo(D=x$Diameter, WD=x$WD, errWD=x$sdWD, coord=cb
ind(x$long,x$lat),

                      Dpropag="chave2004"), simplify=F)

tempLOP <- as.data.frame(resultMC_ChaveGB$LOPE$AGB_simu)
tempMAB <- as.data.frame(resultMC_ChaveGB$MABOUNIE$AGB_simu)
tempMON <- as.data.frame(resultMC_ChaveGB$MONDAH$AGB_simu)
tempRAB <- as.data.frame(resultMC_ChaveGB$RABI$AGB_simu)
tempAFRI <- rbind(tempLOP,tempMAB,tempMON,tempRAB)
Afriprop_CHAV <- cbind(AfriSARstem, tempAFRI)

for (i in (1:length(resolAGB))) {
  tempocalc <- by(Afriprop_CHAV, Afriprop_CHAV[,resolAGB[i]],
                 function(x) list(meanAGB = mean(apply(x[,46:1045], 2, sum, na.rm = T)),
                                credibilityAGB = quantile(apply(x[,46:1045], 2, sum, na.rm
= T), probs = c(0.025,0.975))))

  AGB_chv.list[[i]] <- data.frame(Area_code = names(tempocalc),
                                agb_chv = round(as.numeric(sapply(tempocalc,"[,1))*coefmult
[i],1),
                                cred_chv_2.5 = round(as.numeric(lapply(sapply(tempocalc,"[,
2), function(x) x[1]))*coefmult[i],1),
                                cred_chv_97.5 = round(as.numeric(lapply(sapply(tempocalc,"[
,2), function(x) x[2]))*coefmult[i],1), stringsAsFactors = F)
  rownames(AGB_chv.list[[i]]) <- NULL
}
AGB_chv.list
AGB_chv.df <- Reduce(rbind, AGB_chv.list)
AGB_chv.df
```

##	Area_code	agb_chv	cred_chv_2.5	cred_chv_97.5
## 1	LOP02h	1.2	0.7	2.0
## 2	LOP03h	27.5	19.6	39.3
## 3	LOP07h	331.3	302.2	365.0
## 4	LOP08h	288.0	265.7	310.4
## 5	LOP09h	357.5	317.3	403.4
## 6	LOP10h	403.3	357.9	461.6
## 7	LOP11h	391.4	339.5	453.5
## 8	LOP12h	364.6	328.5	406.4
## 9	LPG01h	492.8	387.0	685.8
## 10	LPG02h	602.1	500.7	722.9
## 11	MAB01h	344.8	310.4	385.0
## 12	MAB02h	327.5	284.0	378.3
## 13	MAB03h	352.8	303.6	415.2

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##	14	MAB04h	468.4	403.9	548.9
##	15	MAB05h	458.0	388.1	566.5
##	16	MAB06h	328.2	286.2	375.0
##	17	MAB07h	260.1	231.0	292.5
##	18	MAB08h	320.2	285.7	364.3
##	19	MAB09h	425.5	380.6	481.7
##	20	MAB10h	366.5	314.1	433.1
##	21	MAB11h	456.1	408.7	508.0
##	22	MAB12h	196.5	177.1	218.6
##	23	MNG03h	497.9	448.8	553.4
##	24	MNG04h	423.8	380.0	472.8
##	25	MON01h	29.4	23.3	37.1
##	26	MON02h	330.1	259.8	432.0
##	27	MON03h	56.7	50.3	65.1
##	28	MON05h	102.0	88.9	117.4
##	29	MON09h	2.7	2.1	3.6
##	30	MON10h	116.2	97.6	141.5
##	31	MON11h	36.6	32.9	41.0
##	32	MON13h	258.5	225.1	299.3
##	33	MON14h	164.1	142.9	192.4
##	34	MON19h	3.9	3.1	4.7
##	35	MON20h	81.9	64.0	109.2
##	36	MON21Ah	155.4	142.3	169.4
##	37	MON21h	2.3	1.8	2.9
##	38	MON22h	314.6	275.2	363.1
##	39	MON23h	139.2	121.2	162.1
##	40	RAB01h	228.6	197.8	273.9
##	41	RAB02h	284.8	247.7	337.3
##	42	RAB03h	294.9	265.7	335.2
##	43	RAB04h	291.5	265.1	322.3
##	44	RAB05h	329.2	277.1	401.6
##	45	RAB06h	290.0	258.6	328.4
##	46	RAB07h	427.1	361.2	513.9
##	47	RAB08h	204.9	186.9	225.9
##	48	RAB09h	275.3	241.1	322.1
##	49	RAB10h	329.1	274.5	406.7
##	50	RAB11h	295.8	254.5	354.7
##	51	RAB12h	311.5	275.2	354.3
##	52	RAB13h	250.9	224.9	281.0
##	53	RAB14h	247.8	221.4	278.6
##	54	RAB15h	313.0	273.3	355.0
##	55	RAB16h	262.8	220.2	326.9
##	56	RAB17h	265.6	236.4	299.1
##	57	RAB18h	267.9	223.2	327.6
##	58	RAB19h	267.1	235.3	304.9
##	59	RAB20h	363.7	308.1	440.4
##	60	RAB21h	303.1	260.5	359.0
##	61	RAB22h	241.5	218.1	269.3
##	62	RAB23h	295.4	255.0	348.8
##	63	RAB24h	513.9	444.9	602.2
##	64	RAB25h	300.3	263.8	349.5
##	65	LOP02q2	0.6	0.3	1.1
##	66	LOP02q3	2.4	0.9	5.3
##	67	LOP02q4	1.7	1.1	2.6

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##	68	LOP03q1	14.6	11.1	19.5
##	69	LOP03q2	88.5	57.7	134.3
##	70	LOP03q3	3.0	2.2	4.1
##	71	LOP03q4	3.8	1.9	7.0
##	72	LOP04q1	301.2	257.8	352.0
##	73	LOP04q2	138.8	117.2	164.1
##	74	LOP05q1	49.7	41.0	60.8
##	75	LOP05q2	71.8	62.1	82.9
##	76	LOP06q1	154.4	137.6	172.3
##	77	LOP06q2	197.9	176.3	220.6
##	78	LOP07q1	375.5	312.5	464.6
##	79	LOP07q2	331.8	284.9	383.1
##	80	LOP07q3	301.0	254.2	350.2
##	81	LOP07q4	316.7	269.0	382.3
##	82	LOP08q1	292.7	254.9	335.0
##	83	LOP08q2	296.7	264.1	335.2
##	84	LOP08q3	276.3	242.0	316.6
##	85	LOP08q4	286.3	247.1	329.3
##	86	LOP09q1	466.6	360.8	597.3
##	87	LOP09q2	287.3	245.3	338.8
##	88	LOP09q3	259.5	213.4	318.5
##	89	LOP09q4	416.7	338.7	513.1
##	90	LOP10q1	347.9	300.1	411.2
##	91	LOP10q2	333.7	271.5	408.9
##	92	LOP10q3	534.8	423.5	671.3
##	93	LOP10q4	396.9	306.4	534.1
##	94	LOP11q1	465.2	371.8	596.3
##	95	LOP11q2	297.0	244.5	363.0
##	96	LOP11q3	438.5	323.8	604.7
##	97	LOP11q4	365.1	282.5	467.0
##	98	LOP12q1	412.0	349.8	492.1
##	99	LOP12q2	330.5	280.0	393.7
##	100	LOP12q3	474.4	385.3	598.1
##	101	LOP12q4	241.6	207.2	285.2
##	102	LPG01q1	389.9	312.6	492.1
##	103	LPG01q2	408.6	315.0	549.0
##	104	LPG01q3	260.7	209.5	327.2
##	105	LPG01q4	911.0	547.2	1636.7
##	106	LPG02q1	460.7	340.9	616.9
##	107	LPG02q2	990.2	704.8	1383.9
##	108	LPG02q3	288.5	224.4	376.3
##	109	LPG02q4	663.9	523.1	857.7
##	110	MND01q1	410.1	337.4	494.6
##	111	MND01q2	411.7	332.7	506.8
##	112	MND01q3	438.4	362.0	537.3
##	113	MND01q4	443.6	364.3	543.0
##	114	MND02q1	306.2	252.2	375.7
##	115	MND02q2	469.5	370.0	588.5
##	116	MND02q3	483.4	407.3	575.0
##	117	MND02q4	553.5	461.9	672.3
##	118	MNG03q1	541.5	457.5	645.9
##	119	MNG03q2	512.6	426.0	628.1
##	120	MNG03q3	400.1	331.6	482.8
##	121	MNG03q4	537.6	441.9	647.9

##	122	MNG04q1	587.8	479.8	723.5
##	123	MNG04q2	343.1	299.6	391.5
##	124	MNG04q3	335.0	271.9	413.5
##	125	MNG04q4	429.3	354.6	526.0
##	126	MON01q1	13.9	10.1	18.8
##	127	MON01q2	6.6	5.0	8.7
##	128	MON01q3	37.2	24.9	54.9
##	129	MON01q4	59.8	42.7	88.2
##	130	MON02q1	197.6	149.4	258.1
##	131	MON02q2	129.9	94.8	190.4
##	132	MON02q3	831.7	564.2	1232.4
##	133	MON02q4	161.2	138.2	191.3
##	134	MON03q1	27.2	19.6	39.7
##	135	MON03q2	17.7	12.8	25.6
##	136	MON03q3	70.7	58.0	89.1
##	137	MON03q4	111.2	94.4	132.4
##	138	MON05q1	166.2	130.5	213.6
##	139	MON05q2	169.4	139.9	205.1
##	140	MON05q3	28.4	20.4	39.2
##	141	MON05q4	43.9	33.8	58.6
##	142	MON09q1	2.3	1.2	4.0
##	143	MON09q2	1.5	1.0	2.2
##	144	MON09q3	0.6	0.4	1.0
##	145	MON09q4	6.4	4.3	9.5
##	146	MON10q1	107.2	60.2	192.9
##	147	MON10q2	122.7	96.6	156.4
##	148	MON10q3	70.1	53.2	91.1
##	149	MON10q4	164.8	127.5	216.3
##	150	MON11q1	22.4	17.8	28.5
##	151	MON11q2	33.2	26.7	43.0
##	152	MON11q3	32.9	25.7	42.7
##	153	MON11q4	58.0	50.5	67.2
##	154	MON13q1	335.2	277.7	411.5
##	155	MON13q2	267.1	225.0	318.6
##	156	MON13q3	172.4	133.9	227.8
##	157	MON13q4	259.1	180.6	383.7
##	158	MON14q1	270.6	210.8	356.8
##	159	MON14q2	199.8	160.2	252.1
##	160	MON14q3	122.6	91.5	174.1
##	161	MON14q4	63.4	55.6	72.5
##	162	MON19q1	9.3	7.1	12.1
##	163	MON19q2	0.4	0.2	0.7
##	164	MON19q3	5.3	3.7	7.5
##	165	MON19q4	0.5	0.3	1.0
##	166	MON20q1	166.8	101.9	272.7
##	167	MON20q2	124.5	100.6	158.8
##	168	MON20q3	24.2	18.4	32.8
##	169	MON20q4	12.0	9.3	15.3
##	170	MON21Aq1	128.6	111.1	151.6
##	171	MON21Aq2	86.5	70.9	106.2
##	172	MON21Aq3	224.3	197.7	255.2
##	173	MON21Aq4	182.3	153.2	218.8
##	174	MON21q1	0.3	0.1	0.6
##	175	MON21q3	4.9	3.5	7.2

##	176	MON21q4	3.9	2.9	5.3
##	177	MON22q1	134.8	116.2	155.4
##	178	MON22q2	485.7	380.5	627.6
##	179	MON22q3	133.8	116.1	154.6
##	180	MON22q4	503.9	402.7	643.9
##	181	MON23q1	334.8	283.3	405.1
##	182	MON23q2	172.6	135.9	231.3
##	183	MON23q3	19.7	11.0	33.1
##	184	MON23q4	29.9	18.2	49.5
##	185	RAB001q	253.2	201.8	323.7
##	186	RAB002q	163.0	137.7	200.3
##	187	RAB003q	367.3	300.8	455.9
##	188	RAB004q	223.8	152.5	359.4
##	189	RAB005q	278.6	236.7	326.9
##	190	RAB006q	227.4	191.6	273.4
##	191	RAB007q	353.4	299.8	425.7
##	192	RAB008q	276.3	234.9	325.0
##	193	RAB009q	430.0	325.6	574.9
##	194	RAB010q	472.7	316.3	695.4
##	195	RAB011q	196.5	165.5	231.8
##	196	RAB012q	301.5	203.7	456.2
##	197	RAB013q	314.9	237.2	427.8
##	198	RAB014q	230.5	187.5	288.2
##	199	RAB015q	318.2	266.6	403.0
##	200	RAB016q	355.6	281.2	461.0
##	201	RAB017q	224.6	185.7	276.7
##	202	RAB018q	311.8	253.3	382.2
##	203	RAB019q	200.0	160.6	251.7
##	204	RAB020q	213.9	178.4	263.2
##	205	RAB021q	234.6	197.9	276.2
##	206	RAB022q	354.6	285.7	446.2
##	207	RAB023q	371.8	281.1	502.7
##	208	RAB024q	678.3	501.8	934.9
##	209	RAB025q	236.9	207.3	269.6
##	210	RAB026q	153.5	132.0	183.5
##	211	RAB027q	184.8	154.9	223.4
##	212	RAB028q	323.8	257.5	428.1
##	213	RAB029q	458.0	352.2	628.6
##	214	RAB030q	178.8	141.8	236.4
##	215	RAB031q	311.1	245.3	407.8
##	216	RAB032q	259.6	214.9	314.4
##	217	RAB033q	470.2	355.0	622.9
##	218	RAB034q	188.3	157.9	231.0
##	219	RAB035q	215.0	180.5	263.4
##	220	RAB036q	214.4	177.9	261.2
##	221	RAB037q	348.5	256.1	501.4
##	222	RAB038q	244.0	207.8	292.2
##	223	RAB039q	244.2	198.7	310.3
##	224	RAB040q	435.3	282.2	663.3
##	225	RAB041q	353.4	246.6	535.9
##	226	RAB042q	270.1	212.1	349.7
##	227	RAB043q	308.6	238.9	398.6
##	228	RAB044q	315.7	251.1	410.8
##	229	RAB045q	237.1	197.7	286.0

##	230	RAB046q	217.2	182.8	263.6
##	231	RAB047q	324.5	267.7	407.1
##	232	RAB048q	289.8	238.6	358.7
##	233	RAB049q	224.8	184.1	280.1
##	234	RAB050q	352.4	278.5	451.6
##	235	RAB051q	297.8	245.0	373.9
##	236	RAB052q	262.1	197.8	365.5
##	237	RAB053q	228.3	189.0	279.2
##	238	RAB054q	393.6	310.5	500.7
##	239	RAB055q	342.2	271.8	431.9
##	240	RAB056q	207.2	176.2	245.9
##	241	RAB057q	172.8	136.6	225.8
##	242	RAB058q	204.1	168.1	242.4
##	243	RAB059q	357.9	277.6	468.0
##	244	RAB060q	316.7	257.6	392.9
##	245	RAB061q	374.1	251.0	614.4
##	246	RAB062q	146.4	121.2	178.2
##	247	RAB063q	272.2	221.1	338.3
##	248	RAB064q	326.4	257.2	423.2
##	249	RAB065q	233.1	170.8	332.7
##	250	RAB066q	314.4	221.0	487.0
##	251	RAB067q	277.4	211.5	373.1
##	252	RAB068q	165.5	138.8	197.3
##	253	RAB069q	522.8	364.7	791.4
##	254	RAB070q	372.4	289.2	493.6
##	255	RAB071q	250.9	188.0	348.5
##	256	RAB072q	279.2	228.0	353.4
##	257	RAB073q	232.8	190.5	290.3
##	258	RAB074q	231.1	191.1	286.1
##	259	RAB075q	175.0	148.6	209.2
##	260	RAB076q	349.2	247.6	523.4
##	261	RAB077q	271.9	223.0	333.3
##	262	RAB078q	353.7	282.5	462.3
##	263	RAB079q	339.9	275.2	421.9
##	264	RAB080q	217.2	167.2	282.7
##	265	RAB081q	361.4	291.6	454.0
##	266	RAB082q	319.4	247.0	431.7
##	267	RAB083q	261.6	222.3	313.7
##	268	RAB084q	274.9	219.8	356.0
##	269	RAB085q	191.5	148.5	250.9
##	270	RAB086q	311.8	248.1	400.6
##	271	RAB087q	727.0	535.0	1003.5
##	272	RAB088q	425.4	330.4	560.5
##	273	RAB089q	321.6	243.4	449.2
##	274	RAB090q	207.6	170.5	255.4
##	275	RAB091q	352.6	258.0	505.5
##	276	RAB092q	178.9	137.4	246.6
##	277	RAB093q	177.6	151.9	211.9
##	278	RAB094q	251.0	213.0	303.4
##	279	RAB095q	296.3	223.0	401.7
##	280	RAB096q	381.5	285.1	533.3
##	281	RAB097q	421.1	343.1	525.5
##	282	RAB098q	482.0	377.3	631.4
##	283	RAB099q	361.9	296.5	454.7

##	284	RAB100q	309.9	245.9	410.1
##	285	MAB01c1	223.2	177.2	279.4
##	286	MAB01c2	349.6	283.5	442.3
##	287	MAB01c3	309.4	258.4	377.9
##	288	MAB01c4	263.8	218.9	322.3
##	289	MAB02c1	308.5	245.3	396.9
##	290	MAB02c2	183.7	147.7	232.0
##	291	MAB02c3	298.1	224.5	400.4
##	292	MAB02c4	557.9	414.2	746.7
##	293	MAB03c1	219.4	172.6	283.5
##	294	MAB03c2	295.7	236.1	375.0
##	295	MAB03c3	813.3	567.8	1163.6
##	296	MAB03c4	341.9	261.2	453.5
##	297	MAB04c1	464.1	350.8	635.1
##	298	MAB04c2	424.1	313.3	606.2
##	299	MAB04c3	580.2	451.1	783.5
##	300	MAB04c4	274.7	216.7	355.3
##	301	MAB05c1	462.3	364.6	597.0
##	302	MAB05c2	289.2	223.1	366.5
##	303	MAB05c3	301.9	245.4	384.7
##	304	MAB05c4	362.4	273.8	475.7
##	305	MAB06c1	294.5	222.3	406.3
##	306	MAB06c2	373.7	284.6	509.9
##	307	MAB06c3	192.2	149.1	247.4
##	308	MAB06c4	421.8	323.4	543.3
##	309	MAB07c1	219.6	182.4	268.9
##	310	MAB07c2	239.4	193.1	299.7
##	311	MAB07c3	190.3	158.7	231.7
##	312	MAB07c4	384.9	271.3	551.4
##	313	MAB08c1	295.6	217.3	411.8
##	314	MAB08c2	340.9	248.2	494.8
##	315	MAB08c3	252.8	204.5	318.9
##	316	MAB08c4	320.4	257.7	400.9
##	317	MAB09c1	421.3	334.9	537.7
##	318	MAB09c2	361.2	295.0	447.0
##	319	MAB09c3	411.7	307.4	567.3
##	320	MAB09c4	607.1	476.8	773.9
##	321	MAB10c1	374.2	232.1	624.3
##	322	MAB10c2	232.6	197.0	278.3
##	323	MAB10c3	529.5	383.1	742.0
##	324	MAB10c4	624.7	466.3	843.9
##	325	MAB11c1	362.9	287.6	452.1
##	326	MAB11c2	379.7	282.8	519.3
##	327	MAB11c3	813.3	664.4	986.2
##	328	MAB11c4	282.5	222.4	360.5
##	329	MAB12c1	291.3	238.4	361.6
##	330	MAB12c2	160.5	126.1	205.4
##	331	MAB12c3	154.0	124.4	195.7
##	332	MAB12c4	158.6	128.8	192.7

AGB USING LOCAL H:D RELATIONSHIP

(agb_loc)

```
AGB_loc.list <- list()

rm(resultMC_LocalGB); gc()
resultMC_LocalGB <- by(AfriSARstem, AfriSARstem["Site"],
  function(x) AGBmonteCarlo(D=x$Diameter, WD=x$WD, H=x$Hlocal, errWD=x$sd
WD,
  errH=x$HlocRSE, Dpropag ="chave2004"), simpli
fy=F)

tempLOP <- as.data.frame(resultMC_LocalGB$LOPE$AGB_simu)
tempMAB <- as.data.frame(resultMC_LocalGB$MABOUNIE$AGB_simu)
tempMON <- as.data.frame(resultMC_LocalGB$MONDAH$AGB_simu)
tempRAB <- as.data.frame(resultMC_LocalGB$RABI$AGB_simu)
tempAFRI <- rbind(tempLOP,tempMAB,tempMON,tempRAB)
Afriprop_LOCAL <- cbind(AfriSARstem, tempAFRI)

for (i in (1:length(resolAGB))) {
  tempocalc <- by(Afriprop_LOCAL, Afriprop_LOCAL[,resolAGB[i]],
    function(x) list(meanAGB = mean(apply(x[,46:1045], 2, sum, na.rm = T)),
      credibilityAGB = quantile(apply(x[,46:1045], 2, sum, na.rm
= T), probs = c(0.025,0.975))))

  AGB_loc.list[[i]] <- data.frame(Area_code = names(tempocalc),
    agb_loc = round(as.numeric(sapply(tempocalc,"[,1)*)coefmult
[i],1),
    cred_loc_2.5 = round(as.numeric(lapply(sapply(tempocalc,"[,
2), function(x) x[1]))*coefmult[i],1),
    cred_loc_97.5 = round(as.numeric(lapply(sapply(tempocalc,"[
,2), function(x) x[2]))*coefmult[i],1), stringsAsFactors = F)
  rownames(AGB_loc.list[[i]]) <- NULL
}
AGB_loc.list
AGB_loc.df <- Reduce(rbind, AGB_loc.list)
AGB_loc.df
```

##	Area_code	agb_loc	cred_loc_2.5	cred_loc_97.5
## 1	LOP02h	0.3	0.2	0.6
## 2	LOP03h	15.5	11.5	22.1
## 3	LOP07h	317.9	294.3	342.8
## 4	LOP08h	290.4	272.3	308.9
## 5	LOP09h	348.6	316.7	383.0
## 6	LOP10h	375.1	339.2	414.8
## 7	LOP11h	349.7	314.1	393.7
## 8	LOP12h	321.4	296.3	352.0
## 9	LPG01h	439.1	371.8	534.2
## 10	LPG02h	547.3	482.3	628.6
## 11	MAB01h	327.7	299.0	361.8
## 12	MAB02h	302.4	269.0	338.6
## 13	MAB03h	333.6	296.5	386.9
## 14	MAB04h	459.0	407.4	526.0

Plot-based aboveground biomass estimates - AfriSAR sites

##	15	MAB05h	438.8	377.9	518.5
##	16	MAB06h	309.4	279.4	344.6
##	17	MAB07h	247.0	225.7	272.7
##	18	MAB08h	290.9	265.3	318.7
##	19	MAB09h	411.0	375.0	454.5
##	20	MAB10h	344.8	304.7	393.9
##	21	MAB11h	523.2	477.4	576.0
##	22	MAB12h	171.8	156.8	186.9
##	23	MNG03h	510.6	468.5	561.3
##	24	MNG04h	444.0	403.2	487.7
##	25	MON01h	25.6	21.3	30.6
##	26	MON02h	294.7	248.7	357.8
##	27	MON03h	56.8	51.1	63.3
##	28	MON05h	93.2	83.8	104.5
##	29	MON09h	2.3	1.7	3.0
##	30	MON10h	103.4	88.1	121.4
##	31	MON11h	35.0	31.6	39.2
##	32	MON13h	247.8	222.9	275.9
##	33	MON14h	149.9	132.7	168.2
##	34	MON19h	3.4	2.8	4.2
##	35	MON20h	70.3	58.7	85.9
##	36	MON21Ah	160.3	148.3	174.1
##	37	MON21h	2.4	1.8	3.1
##	38	MON22h	289.1	258.2	322.5
##	39	MON23h	128.8	115.1	144.9
##	40	RAB01h	210.7	191.3	233.2
##	41	RAB02h	280.3	248.8	316.2
##	42	RAB03h	303.2	276.6	333.4
##	43	RAB04h	300.1	274.8	324.9
##	44	RAB05h	320.6	278.1	368.5
##	45	RAB06h	301.0	275.1	331.5
##	46	RAB07h	413.2	361.4	475.5
##	47	RAB08h	209.4	194.2	225.3
##	48	RAB09h	280.1	250.1	322.5
##	49	RAB10h	323.3	278.9	378.5
##	50	RAB11h	287.8	253.4	332.7
##	51	RAB12h	310.1	280.8	344.8
##	52	RAB13h	246.4	224.4	271.0
##	53	RAB14h	245.6	224.3	270.6
##	54	RAB15h	313.1	281.9	347.3
##	55	RAB16h	253.7	219.3	300.7
##	56	RAB17h	265.9	241.4	293.1
##	57	RAB18h	262.5	224.0	312.2
##	58	RAB19h	255.1	229.6	284.6
##	59	RAB20h	348.5	306.0	397.3
##	60	RAB21h	301.2	267.9	337.6
##	61	RAB22h	246.7	226.0	269.0
##	62	RAB23h	296.1	262.6	339.5
##	63	RAB24h	511.8	453.5	577.0
##	64	RAB25h	293.4	263.1	331.2
##	65	LOP02q2	0.2	0.1	0.3
##	66	LOP02q3	0.8	0.3	1.7
##	67	LOP02q4	0.4	0.3	0.6
##	68	LOP03q1	8.7	6.5	11.7

##	69	LOP03q2	50.1	34.3	76.3
##	70	LOP03q3	0.8	0.6	1.1
##	71	LOP03q4	2.4	1.2	4.4
##	72	LOP04q1	300.5	260.9	346.8
##	73	LOP04q2	138.0	120.9	159.3
##	74	LOP05q1	39.0	32.7	46.0
##	75	LOP05q2	63.3	55.1	72.0
##	76	LOP06q1	156.8	141.9	173.6
##	77	LOP06q2	206.1	187.9	226.9
##	78	LOP07q1	362.3	304.5	427.1
##	79	LOP07q2	324.6	286.9	364.8
##	80	LOP07q3	289.5	252.8	334.6
##	81	LOP07q4	295.1	256.7	339.5
##	82	LOP08q1	296.9	263.8	334.0
##	83	LOP08q2	302.9	273.9	338.0
##	84	LOP08q3	276.8	247.9	309.8
##	85	LOP08q4	285.0	249.7	324.9
##	86	LOP09q1	433.2	352.4	536.5
##	87	LOP09q2	299.6	256.6	345.7
##	88	LOP09q3	262.0	223.4	310.2
##	89	LOP09q4	399.5	336.7	470.4
##	90	LOP10q1	337.0	291.9	388.0
##	91	LOP10q2	300.4	250.3	357.3
##	92	LOP10q3	502.4	415.3	603.6
##	93	LOP10q4	360.4	287.5	478.6
##	94	LOP11q1	431.8	357.2	537.9
##	95	LOP11q2	273.7	231.9	328.6
##	96	LOP11q3	371.2	297.7	490.8
##	97	LOP11q4	322.2	264.9	399.3
##	98	LOP12q1	359.5	308.3	414.5
##	99	LOP12q2	303.2	263.4	352.7
##	100	LOP12q3	398.8	335.7	489.9
##	101	LOP12q4	224.1	197.3	254.8
##	102	LPG01q1	370.2	305.8	450.1
##	103	LPG01q2	384.7	310.6	503.1
##	104	LPG01q3	254.2	209.9	310.9
##	105	LPG01q4	746.2	499.7	1132.0
##	106	LPG02q1	423.7	324.9	548.8
##	107	LPG02q2	860.4	659.7	1109.2
##	108	LPG02q3	275.2	217.9	348.7
##	109	LPG02q4	624.8	506.4	764.7
##	110	MND01q1	416.6	351.9	506.8
##	111	MND01q2	402.5	330.6	486.3
##	112	MND01q3	429.1	358.4	507.7
##	113	MND01q4	454.5	378.2	543.5
##	114	MND02q1	319.1	266.4	381.2
##	115	MND02q2	484.2	396.1	592.7
##	116	MND02q3	496.8	416.7	585.9
##	117	MND02q4	572.5	481.8	673.6
##	118	MNG03q1	564.2	480.7	660.9
##	119	MNG03q2	516.1	431.2	624.6
##	120	MNG03q3	416.8	352.1	486.7
##	121	MNG03q4	545.3	454.5	644.2
##	122	MNG04q1	610.5	504.8	751.4

##	123	MNG04q2	362.2	315.0	415.7
##	124	MNG04q3	344.2	283.3	420.5
##	125	MNG04q4	459.2	390.6	544.1
##	126	MON01q1	13.8	10.1	18.4
##	127	MON01q2	6.8	4.8	9.3
##	128	MON01q3	31.1	22.0	43.3
##	129	MON01q4	50.6	38.2	67.3
##	130	MON02q1	183.5	145.9	230.2
##	131	MON02q2	151.5	108.5	222.1
##	132	MON02q3	674.8	514.3	927.7
##	133	MON02q4	169.0	143.7	200.9
##	134	MON03q1	29.9	21.4	43.0
##	135	MON03q2	16.5	12.1	22.2
##	136	MON03q3	69.0	57.6	81.8
##	137	MON03q4	111.9	96.3	129.0
##	138	MON05q1	143.0	115.3	179.1
##	139	MON05q2	153.7	133.3	177.9
##	140	MON05q3	29.9	21.3	41.7
##	141	MON05q4	46.2	34.8	61.8
##	142	MON09q1	1.8	1.0	3.1
##	143	MON09q2	1.5	0.9	2.4
##	144	MON09q3	0.7	0.3	1.3
##	145	MON09q4	5.1	3.3	7.6
##	146	MON10q1	91.0	58.3	144.6
##	147	MON10q2	111.4	89.8	139.1
##	148	MON10q3	61.4	49.4	77.8
##	149	MON10q4	149.7	116.0	189.8
##	150	MON11q1	18.4	14.4	23.5
##	151	MON11q2	29.6	24.4	36.5
##	152	MON11q3	34.3	26.0	44.8
##	153	MON11q4	57.7	50.3	67.0
##	154	MON13q1	329.2	277.1	395.8
##	155	MON13q2	256.9	217.5	305.1
##	156	MON13q3	165.7	131.5	208.2
##	157	MON13q4	239.4	182.9	317.6
##	158	MON14q1	226.6	182.8	280.1
##	159	MON14q2	184.9	154.9	223.5
##	160	MON14q3	122.9	88.9	170.1
##	161	MON14q4	65.1	56.6	75.2
##	162	MON19q1	8.7	6.5	11.3
##	163	MON19q2	0.4	0.1	0.9
##	164	MON19q3	4.1	2.9	5.9
##	165	MON19q4	0.6	0.2	1.2
##	166	MON20q1	128.5	88.8	189.2
##	167	MON20q2	114.9	95.2	140.4
##	168	MON20q3	25.2	18.5	33.5
##	169	MON20q4	12.4	9.4	16.0
##	170	MON21Aq1	133.9	114.8	157.0
##	171	MON21Aq2	87.8	72.5	105.0
##	172	MON21Aq3	233.0	203.8	267.4
##	173	MON21Aq4	186.4	160.2	220.6
##	174	MON21q1	0.3	0.1	0.6
##	175	MON21q3	5.1	3.4	7.4
##	176	MON21q4	4.1	2.9	5.7

##	177	MON22q1	137.8	119.3	159.8
##	178	MON22q2	417.8	339.9	517.3
##	179	MON22q3	142.2	122.8	164.9
##	180	MON22q4	458.7	380.9	550.2
##	181	MON23q1	309.2	268.3	358.1
##	182	MON23q2	154.7	126.0	190.1
##	183	MON23q3	20.5	11.2	33.8
##	184	MON23q4	30.9	18.1	51.5
##	185	RAB001q	252.6	206.8	311.2
##	186	RAB002q	161.2	139.3	187.8
##	187	RAB003q	368.6	315.0	435.5
##	188	RAB004q	215.3	161.2	296.1
##	189	RAB005q	287.5	252.6	333.7
##	190	RAB006q	231.2	198.3	271.5
##	191	RAB007q	369.2	315.6	431.1
##	192	RAB008q	282.7	244.0	326.8
##	193	RAB009q	410.3	330.5	524.9
##	194	RAB010q	441.0	325.5	603.3
##	195	RAB011q	200.3	170.0	235.6
##	196	RAB012q	228.7	184.7	296.7
##	197	RAB013q	304.7	232.8	405.7
##	198	RAB014q	229.6	195.7	270.8
##	199	RAB015q	326.4	275.7	399.6
##	200	RAB016q	367.6	305.3	444.0
##	201	RAB017q	232.3	196.7	278.3
##	202	RAB018q	315.9	264.2	373.8
##	203	RAB019q	205.3	168.1	254.0
##	204	RAB020q	225.9	190.9	269.7
##	205	RAB021q	243.4	211.8	279.6
##	206	RAB022q	376.9	314.2	453.9
##	207	RAB023q	348.9	270.1	452.3
##	208	RAB024q	650.6	503.4	854.1
##	209	RAB025q	242.4	217.2	271.8
##	210	RAB026q	156.9	137.1	183.1
##	211	RAB027q	184.8	161.8	215.4
##	212	RAB028q	339.0	273.8	418.9
##	213	RAB029q	478.8	386.7	611.8
##	214	RAB030q	183.8	149.6	232.6
##	215	RAB031q	313.1	255.0	398.9
##	216	RAB032q	270.6	228.5	319.2
##	217	RAB033q	472.4	377.5	598.7
##	218	RAB034q	180.8	154.3	213.4
##	219	RAB035q	217.7	188.7	258.5
##	220	RAB036q	220.8	185.1	264.1
##	221	RAB037q	343.3	261.2	478.1
##	222	RAB038q	253.2	222.0	295.9
##	223	RAB039q	248.7	206.0	305.6
##	224	RAB040q	381.9	265.3	556.4
##	225	RAB041q	311.4	233.5	432.3
##	226	RAB042q	257.0	208.0	316.8
##	227	RAB043q	300.0	244.9	374.7
##	228	RAB044q	318.3	257.6	390.7
##	229	RAB045q	230.2	197.6	271.3
##	230	RAB046q	215.0	181.4	258.8

Plot-based aboveground biomass estimates - AfriSAR sites

##	231	RAB047q	319.4	273.0	381.1
##	232	RAB048q	292.4	247.4	352.8
##	233	RAB049q	230.9	191.6	278.8
##	234	RAB050q	343.1	284.7	426.8
##	235	RAB051q	306.3	257.7	374.1
##	236	RAB052q	276.5	212.2	373.0
##	237	RAB053q	236.1	198.6	280.6
##	238	RAB054q	386.1	319.3	476.2
##	239	RAB055q	329.8	267.0	408.3
##	240	RAB056q	210.5	181.2	249.9
##	241	RAB057q	173.4	142.5	215.0
##	242	RAB058q	197.3	170.6	231.9
##	243	RAB059q	372.5	299.2	479.8
##	244	RAB060q	305.9	249.4	368.2
##	245	RAB061q	355.4	250.1	521.3
##	246	RAB062q	140.2	118.4	168.0
##	247	RAB063q	281.6	231.5	343.0
##	248	RAB064q	311.7	258.0	392.2
##	249	RAB065q	234.9	177.3	327.8
##	250	RAB066q	299.8	216.9	434.1
##	251	RAB067q	254.2	208.4	319.3
##	252	RAB068q	165.2	141.7	193.3
##	253	RAB069q	474.3	365.1	647.1
##	254	RAB070q	352.8	283.7	434.8
##	255	RAB071q	240.8	190.6	322.1
##	256	RAB072q	278.0	232.9	339.9
##	257	RAB073q	231.7	196.9	281.0
##	258	RAB074q	238.4	202.9	283.1
##	259	RAB075q	175.1	151.4	203.9
##	260	RAB076q	340.0	255.3	475.2
##	261	RAB077q	256.2	216.9	300.1
##	262	RAB078q	344.8	280.1	431.9
##	263	RAB079q	349.8	289.4	422.9
##	264	RAB080q	214.9	170.8	274.9
##	265	RAB081q	359.1	299.3	434.9
##	266	RAB082q	330.5	259.8	432.7
##	267	RAB083q	264.8	222.0	312.6
##	268	RAB084q	282.6	235.9	352.2
##	269	RAB085q	191.5	153.5	251.2
##	270	RAB086q	305.7	249.1	381.5
##	271	RAB087q	732.6	566.5	958.6
##	272	RAB088q	421.5	339.2	535.5
##	273	RAB089q	318.8	245.8	420.9
##	274	RAB090q	197.2	165.4	237.8
##	275	RAB091q	334.4	267.8	431.1
##	276	RAB092q	180.9	143.3	235.2
##	277	RAB093q	177.8	154.5	205.2
##	278	RAB094q	260.6	226.3	303.0
##	279	RAB095q	302.3	242.0	385.8
##	280	RAB096q	384.4	299.6	512.2
##	281	RAB097q	436.3	359.7	526.2
##	282	RAB098q	456.7	366.6	565.1
##	283	RAB099q	350.6	292.3	431.0
##	284	RAB100q	307.1	250.2	384.1

##	285	MAB01c1	208.5	168.6	253.1
##	286	MAB01c2	343.7	275.8	424.1
##	287	MAB01c3	284.2	240.3	336.6
##	288	MAB01c4	243.7	205.6	288.7
##	289	MAB02c1	295.8	238.1	370.0
##	290	MAB02c2	167.2	135.5	206.8
##	291	MAB02c3	272.0	207.0	350.8
##	292	MAB02c4	492.2	386.3	627.9
##	293	MAB03c1	219.7	175.3	279.3
##	294	MAB03c2	296.4	233.9	382.4
##	295	MAB03c3	737.9	550.7	1016.6
##	296	MAB03c4	310.0	247.7	382.8
##	297	MAB04c1	442.3	345.2	593.2
##	298	MAB04c2	416.5	308.3	594.5
##	299	MAB04c3	579.0	452.3	750.9
##	300	MAB04c4	265.4	210.0	336.1
##	301	MAB05c1	453.1	356.0	565.7
##	302	MAB05c2	283.7	225.2	353.0
##	303	MAB05c3	298.3	240.7	379.1
##	304	MAB05c4	360.5	281.0	465.7
##	305	MAB06c1	263.8	209.5	338.2
##	306	MAB06c2	351.6	277.7	448.0
##	307	MAB06c3	182.2	145.9	230.3
##	308	MAB06c4	406.1	316.5	515.2
##	309	MAB07c1	214.9	178.8	255.9
##	310	MAB07c2	218.8	180.8	268.7
##	311	MAB07c3	189.1	158.0	232.0
##	312	MAB07c4	341.8	258.8	468.6
##	313	MAB08c1	251.6	198.0	321.8
##	314	MAB08c2	298.8	238.4	392.7
##	315	MAB08c3	241.5	197.4	296.7
##	316	MAB08c4	304.4	243.1	374.1
##	317	MAB09c1	403.8	335.5	492.2
##	318	MAB09c2	379.9	317.2	460.8
##	319	MAB09c3	412.4	315.6	548.5
##	320	MAB09c4	567.5	450.2	706.6
##	321	MAB10c1	360.7	236.7	588.5
##	322	MAB10c2	231.7	194.1	274.9
##	323	MAB10c3	506.3	388.8	689.5
##	324	MAB10c4	559.3	453.2	700.8
##	325	MAB11c1	402.7	325.1	497.9
##	326	MAB11c2	424.3	324.3	582.8
##	327	MAB11c3	918.2	763.2	1103.4
##	328	MAB11c4	329.9	262.8	421.3
##	329	MAB12c1	263.0	219.8	320.7
##	330	MAB12c2	140.3	114.7	173.1
##	331	MAB12c3	145.0	116.8	189.6
##	332	MAB12c4	135.8	113.2	166.3

Reshaping the different information (estimates, coordinates) in a single object

```
# Convert list of georeferenced hectares/quarters/corners into a single data.frame
site.df <- Reduce(rbind, c(site.list[[1]], site.list[[2]], site.list[[3]], site.list[[4]]))
site.df <- site.df[order(site.df$Site),]
rownames(site.df) <- NULL

# Merge dataframes
AGB_FIN1 <- merge(site.df, AGB_fph.df, by="Area_code", sort = F, all=T)
AGB_FIN1$agb_fph[which(is.na(AGB_FIN1$agb_fph))] <- 0

AGB_FIN2 <- merge(AGB_FIN1, AGB_chv.df, by="Area_code", sort = F, all=T)
AGB_FIN2$agb_chv[which(is.na(AGB_FIN2$agb_chv))] <- 0

AGB_AfriSAR <- merge(AGB_FIN2, AGB_loc.df, by="Area_code", sort = F, all=T)
AGB_AfriSAR$agb_loc[which(is.na(AGB_AfriSAR$agb_loc) & AGB_AfriSAR$Site != "RABI")] <- 0

# Reorder columns
AGB_AfriSAR <- AGB_AfriSAR[c("Site", "Area_code", "Plot_code", "Scale", "sw_x", "sw_y", "nw_x", "nw_y",
"se_x", "se_y", "ne_x", "ne_y", "agb_fph", "cred_fph_2.5", "cred_fph_97.5", "agb_chv", "cred_chv_2.5",
"cred_chv_97.5", "agb_loc", "cred_loc_2.5", "cred_loc_97.5")]

# Reorder lines
AGB_AfriSAR$Site <- as.character(AGB_AfriSAR$Site)
AGB_AfriSAR$Scale <- as.character(AGB_AfriSAR$Scale)
AGB_AfriSAR$Plot_code <- as.character(AGB_AfriSAR$Plot_code)
AGB_AfriSAR$Area_code <- as.character(AGB_AfriSAR$Area_code)

AGB_AfriSAR <- AGB_AfriSAR[with(AGB_AfriSAR, order(Site, Scale, Plot_code, Area_code, decreasing = c(T,T,F,F), method = "radix")),]

AGB_AfriSAR$Site <- as.factor(AGB_AfriSAR$Site)
AGB_AfriSAR$Scale <- as.factor(AGB_AfriSAR$Scale)
AGB_AfriSAR$Plot_code <- as.factor(AGB_AfriSAR$Plot_code)
AGB_AfriSAR$Area_code <- as.factor(AGB_AfriSAR$Area_code)
rownames(AGB_AfriSAR) <- NULL

#AGB_AfriSAR
```