Estimate height

```
source(here::here("load_pkgs.R"))
```

Explore data

```
tree <- read_csv(here::here("data/tree_data.csv"), col_types = cols())

tree %>% group_by(sp) %>%
  count(is.na(h)) %>%
  spread('is.na(h)', n) %>%
  rename(n='FALSE', n_na = 'TRUE') %>%
  knitr::kable()

## group_by: one grouping variable (sp)

## count: now 9 rows and 3 columns, one group variable remaining (sp)

## spread: reorganized (is.na(h), n) into (FALSE, TRUE) [was 9x3, now 8x3]

## rename: renamed 2 variables (n, n_na)
```

sp	n	n_na
Adec	1	NA
Aopa	6	NA
Cmon	37	NA
Pdul	9	NA
Pter	9	NA
Qilex	NA	3
Qpyr	2738	664
Saria	18	NA

Trees with no values for height belongs to Q. ilex and Q. pyrenaica. Height data for computation of Q. ilex biomass is not needed according to allometric equation proposed by Ruiz-Peinado, Montero, and Del Rio (2012). For Q. pyrenaica and estimation of height values was done using non-linear squares model. We used the best equation according to Adame, del Río, and Cañellas (2008)

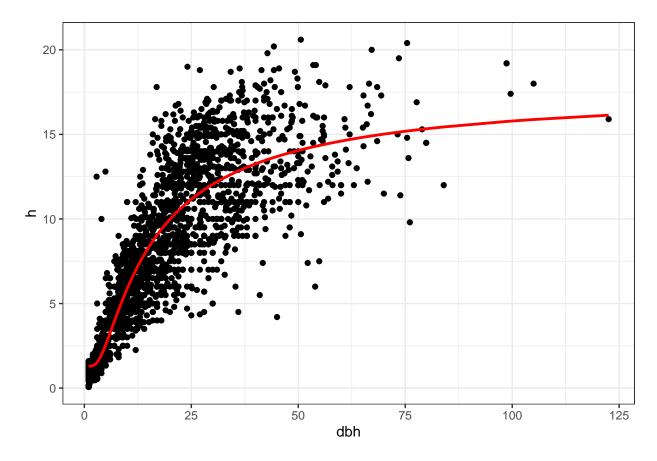
$$h = 1.3 + a * \exp(b/dbh)$$

Model height-diameter

```
\textit{\# see pag. 391 from @PommereningGrabarnik2019IndividualbasedMethods}
# See @Adameetal2008MixedNonlinear for equations. The best equation is
\# h \sim 1.3 + a*e^{(b/D)}
# start values from @Adameetal2008MixedNonlinear
# Remove data withouth dbh
df_model <- tree %>%
 filter(sp == "Qpyr") %>%
filter(!is.na(h))
## filter: removed 83 rows (2%), 3,402 rows remaining
## filter: removed 664 rows (20%), 2,738 rows remaining
# NonLinear Least Squares
m \leftarrow nls(h \sim 1.3 + a * exp(b / dbh),
         data = df_model,
         start = list(a = 15, b = -10), trace = TRUE)
## 12375.66 : 15 -10
## 11678.52 : 16.15356 -12.35728
## 11664.34 : 16.44726 -12.77883
## 11664.27 : 16.46904 -12.80886
## 11664.27 : 16.47026 -12.81070
```

term	estimate	std.error	statistic	p.value
a	16.47026	0.1853400	88.86515	0
b	-12.81070	0.2445795	-52.37845	0

sigma	isConv	finTol	logLik	AIC	BIC	deviance	df.residual
2.064765	TRUE	8.6e-06	-5869.149	11744.3	11762.04	11664.27	2736



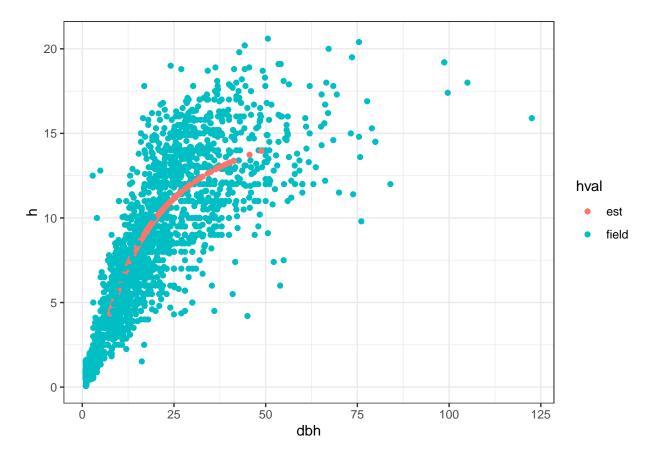
Compute the estimate values

```
tree %<>%
  mutate(h = ifelse(
    sp == "Qpyr" & hval == "est",
    predict(m, list(dbh = dbh)),
    h))
```

mutate: changed 662 values (19%) of 'h' (662 fewer NA)

```
# (1.3 + coef(m)[1]*exp(coef(m)[2]/dbh))
```

Warning: Removed 5 rows containing missing values (geom_point).



Export data

write_csv(tree, here::here("data/tree_data_full.csv"))

References

Adame, Patricia, Miren del Río, and Isabel Cañellas. 2008. "A Mixed Nonlinear Height Diameter Model for Pyrenean Oak (Quercus Pyrenaica Willd.)." Forest Ecology and Management 256 (1-2): 88–98. https://doi.org/10.1016/j.foreco.2008.04.006.

Ruiz-Peinado, R., G. Montero, and M. Del Rio. 2012. "Biomass Models to Estimate Carbon Stocks for Hardwood Tree Species." Forest Systems 21 (1): 42. https://doi.org/10.5424/fs/2112211-02193.