Simple Translation

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In [ ]:
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# To follow along, run this cell first.

options(warn = -1)
suppressMessages(library(neotoma2))
suppressMessages(library(sf))
suppressMessages(library(geojsonsf))
suppressMessages(library(dplyr))
suppressMessages(library(ggplot2))
suppressMessages(library(leaflet))
options(dplyr.summarise.inform = FALSE)

cz_dl <- readRDS('data/czDownload.RDS')
allSamp <- samples(cz_dl)</pre>
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In [ ]:
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taxaSites <- allSamp %>%
  group_by(variablename, siteid) %>%
  group_by(variablename) %>%
  summarise(sites = length(unique(siteid)))
DT::datatable(data = head(taxaSites, n = 15), rownames = FALSE)
```

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In [ ]:

taxaSamples <- allSamp %>%
    group_by(variablename) %>%
    summarise(samples = n())
DT::datatable(data = head(taxaSamples, n = 8), rownames = FALSE)
```

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In [ ] :
    taxaplots <- taxaSites %>% inner_join(taxaSamples, by = "variablename")
    DT::datatable(data = head(taxaplots, n = 20), rownames = FALSE)
```

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In [ ] :
    translation <- readr::read_csv("data/taxontable.csv", show_col_types = FALSE)
    DT::datatable(data = head(translation, n = 8), rownames = FALSE)</pre>
```