```
# ---- Install roadDB ----
# Github installation
install.packages("devtools")
devtools::install_github("sommergeo/roadDB")
# CRAN installation (will be made available later)
# install.packages(roadDB)
# ---- Load library ----
# Load the library
library(roadDB)
# ---- Install external packages ----
# Install packages (needs to be done only once)
install.packages("tidyverse")
install.packages("sf")
install.packages("tmap")
install.packages("rcarbon")
# ---- Load dependencies ----
# Load libraries (every time you run this tutorial)
library(roadDB)
library(tidyverse)
library(sf)
library(tmap)
library(rcarbon)
# ---- First query ----
# Now you are ready to run your first query and inspect the results directly in the R console:
my_query <- road_get_localities(cultural_period='Middle Paleolithic')</pre>
# ---- Argument examples ----
# Single argument
road_get_localities(cultural_period='MSA')
# Multiple arguments
road_get_localities(cultural_period='MSA', countries='South Africa')
# Lists of arguments
road_get_localities(cultural_period=c('MSA','LSA'), countries=c('South Africa','Namibia'))
# ---- Helper functions ----
# Examples
help(road get localities)
help(road_get_lithic_typology)
help(road_get_dates)
# Examples of listing valid argument values
road_list_argument_values("countries")
road_list_argument_values("cultural_periods")
road_list_argument_values("human_genus")
# Example summarizing term appearance
road_summarize_archaeology('egg shell')
head(road_summarize_archaeology('egg shell'), 4)
# ---- Example #1: Localities ----
# Query all localities
localities <- road_get_localities()</pre>
head(localities) # head() shows the first 6 rows of a dataset
# Query selected localities
african_msa_localities <- road_get_localities(continents = "Africa", cultural_periods = "MSA")
head(african msa localities)
```

```
# Load geospatial libraries
               # for spatial objects
library(sf)
library(tmap)
               # for visualization
# Use coordinate columns to create a spatial dataset
african_msa_sf <- st_as_sf(african_msa_localities,</pre>
                            coords = c("coord_x", "coord_y"),
                            crs = 4326) # WGS84 (lat/lon)
# Choose mapping engine
tmap_mode("plot") # static map mode
# Create map
tm_shape(african_msa_sf) +
  tm_basemap("OpenStreetMap") +
  tm_dots()
# ---- Technocomplex map ----
road_list_argument_values("technocomplexes")
# Query Mousterian sites
mousterian <- road get localities(technocomplexes = c("MP/ Mousterian - Eurasia", "MP/ Mousterian
- Levant"))
# Create spatial dataset
mousterian_sf <- st_as_sf(mousterian,</pre>
                           coords = c("coord_x", "coord_y"),
                           crs = 4326) # WGS84 (lat/lon)
# Display map
tmap_mode("plot") # static map mode
tm_shape(mousterian_sf) +
  tm_basemap("OpenStreetMap") +
  tm_dots(
    col = "technocomplexes",
    palette = "Set2") +
  tm_layout(legend.outside = TRUE)
# ---- Example #2: Assemblages ----
# We can query all assemblages in ROAD
assemblages <- road_get_assemblages()</pre>
# Or we can make more specific queries
lithic_typology <- road_get_lithic_typology()</pre>
lithic_raw_mat <- road_get_lithic_raw_material()</pre>
organic_tools <- road_get_organic_tools()</pre>
symbolic <- road_get_symbolic_artifacts()</pre>
feature <- road_get_feature()</pre>
mix <- road_get_miscellaneous_finds()</pre>
people <- road_get_human_remains()</pre>
animals <- road_get_paleofauna()</pre>
plants <- road_get_plantremains()</pre>
# ---- Lithic Tools ----
techno <- road_list_argument_values("technocomplexes")</pre>
# Filter for Mousterian technocomplexes
df_mousterian <- techno[grepl("mousterian", techno$technocomplex, ignore.case = TRUE), ]</pre>
# Gather lithic typology data for Mousterian contexts
mous_tools <- road_get_lithic_typology(technocomplexes = df_mousterian)</pre>
# List available tool types
```

```
road_list_argument_values("tool_list")
# Gather Levallois technology examples
mous_levallois <- road_get_lithic_typology(</pre>
  technocomplexes = df mousterian,
  tool_list = "levallois"
)
# Label data sources
mous levallois$source <- "Levallois"
                     <- "Mousterian"
mous_tools$source
# Combine data
combined <- rbind(mous_levallois, mous_tools)</pre>
# Plot comparison
ggplot(combined, aes(x = fct_infreq(country),
                     fill = source)) +
  geom_bar(position = "dodge") +
  coord_flip() +
 labs(
   title = "Levallois vs. Mousterian",
    x = NULL
    y = "Assemblages (n=)",
   fill = "Key"
  scale y continuous(breaks = scales::pretty breaks(n = 10)) +
  theme_minimal(base_size = 12)
# ---- Ostrich Egg Shells ----
road_summarize_archaeology('ostrich egg shell')
# Query from miscellaneous finds
eggshell_misc <- road_get_miscellaneous_finds()</pre>
eggshell_misc <- eggshell_misc[grepl("ostrich egg shell",
eggshell_misc$miscellaneous_finds_material), ]
# Query from symbolic artifacts
eggshell_symbolic <- road_get_symbolic_artifacts()</pre>
eggshell_symbolic <- eggshell_symbolic[grepl("ostrich egg shell",
eggshell_symbolic$symbolic_artifacts_material), ]
# Combine datasets
eggshell_combined <- merge(
 eggshell misc,
  eggshell symbolic,
  by = c("locality id", "assemblage id", "continent", "subcontinent", "country",
         "locality_types", "coord_x", "coord_y", "assemblage_name", "categories",
         "age_min", "age_max", "cultural_periods", "technocomplexes"),
  all = TRUE
# Plot ostrich egg shell distribution
ggplot(eggshell_combined, aes(y = cultural_periods)) +
  geom_bar() +
  labs(
   title = "Ostrich egg shells",
   x = "Number of assemblages",
    y = "Cultural period"
  )
# ---- Example #3: Dates ----
# Load all dates
all_dates <- road_get_dates()</pre>
```

```
# Filter for "Hohle Fels"
hohle_fels_dates <- all_dates[grepl("Hohle Fels", all_dates$locality_id), ]</pre>
# Filter for archaeological layer Vb and symbolic artifacts
hohle_fels_Vb <- hohle_fels_dates[grepl("AH Vb", hohle_fels_dates$archlayer), ]</pre>
hohle_fels_Vb <- hohle_fels_Vb[grepl("symbolic artifacts", hohle_fels_Vb$categories), ]</pre>
# Load radiocarbon library
library(rcarbon)
# Calibrate radiocarbon ages
hohle_fels_Vb_cal <- calibrate(</pre>
  x = hohle_fels_Vb$age,
  errors = hohle_fels_Vb$positive_standard_deviation,
  calCurves = 'intcal20'
# Display calibrated ages
summary(hohle_fels_Vb_cal)
# Visualize first date
plot(hohle_fels_Vb_cal, 1)
# Visualize all calibrated dates
multiplot(
  hohle_fels_Vb_cal,
  decreasing = TRUE,
  rescale = TRUE,
  HPD = TRUE,
  label.pos = 0.9,
  label.offset = -200
)
```