

# OTNuEC\_data\_and\_calibration

November 25, 2025

## 1 0) Environment

### 1.1 0.0) Packages

```
[49]: options(warn = -1)

knitr::opts_chunk$set(
  warning = FALSE,
  message = FALSE
)

quiet <- function(x) suppressWarnings(suppressMessages(x))

suppressPackageStartupMessages({
  library(sf); library(terra); library(dplyr); library(purrr); library(sidrar);
  ↪library(geobr); library(ggplot2); library(osrm)
})
```

### 1.2 0.1) Internal data paths

```
[50]: DATA <- list(
  viirs_blackmarble_2016 = "data_viirs/viirs_blackmarble_br_2016.tif",
  etopo1_elevation        = "ETOPO1_Ice_g_geotiff.tif",
  erosao_2020_ibge        = "Brasil_vulnerabilidade_2020/
  ↪BR_vulnerabilidade_solos_erosao_2020.tif",

  solos_wrb = c(
    "soils_brazil_wrb_wgs84.shp",
    "soils_brazil_wrb_wgs84.dbf",
    "soils_brazil_wrb_wgs84.shx",
    "soils_brazil_wrb_wgs84.prj",
    "soils_brazil_wrb_wgs84.cst"
  ),

  mapbiomas_10m_2023 = "data/mapbiomas_near_1000m.
  ↪tif"#"mapbiomas_10m_collection2_integration_v1-classification_2023.tif"
)
```

```

# BC250
bc250_layers <- c(
  "rod_trecho_rodoviario_1",
  "fer_trecho_ferroviario_1",
  "hdv_trecho_hidroviario_1",
  "hdv_complexo_portuario_p",
  "hdv_eclusa_p",
  "tra_ponte_1",
  "tra_tunel_1",
  "tra_travessia_1",
  "enc_trecho_energia_1",
  "enc_hidreletrica_p",
  "enc_termeletrica_p",
  "enc_subest_transm_distrib_energia_eletrica_p"
)
bc250_ext <- c(".shp", ".dbf", ".shx", ".prj")

DATA$bc250 <- as.vector(outer(
  file.path("bc250_completo", bc250_layers),
  bc250_ext,
  paste0
))

```

### 1.3 0.2) Check datasets

```

[51]: check_dataset <- function(label, paths) {
  ok <- file.exists(paths)
  if (all(ok)) message("[OK] ", label)
  else {
    message("[FALTA] ", label)
    walk(paths[!ok], ~ message(" - ", .x))
  }
}

walk(names(DATA), DATA, check_dataset)

```

### 1.4 0.3) Sidrar API data paths

```

[52]: # PIB municipal (mil R$ correntes) e POP (pessoas)
pib_sidra <- sidrar::get_sidra(5938, variable = 37, period = "last",
geo = "City", format = 3) |>
transmute(
cod_mun = as.integer(`Município (Código)`),
pib_mil_rs = suppressWarnings(as.numeric(Valor))
)

```

```

pop_est <- sidrar::get_sidra(6579, variable = 9324, period = "2025",
geo = "City", format = 3) |>
transmute(
cod_mun = as.integer(`Município (Código)`),
populacao = suppressWarnings(as.numeric(Valor)))
)

```

Considering all categories once 'classific' was set to 'all' (default)

Considering all categories once 'classific' was set to 'all' (default)

## 1.5 0.4) Geobr data paths

```

[53]: # Brasil continental (aprox.)
clip_mainland_ll <- function(x) {
  sf::st_crop(
    x,
    xmin = -75, # oeste
    xmax = -34, # leste (corta Fernando de Noronha, ~ -32.4)
    ymin = -35, # sul
    ymax = 6 # norte
  )
}
micro_br <- geobr::read_micro_region(
  year = 2013,
  simplified = TRUE
) |>
  clip_mainland_ll() # recorte Brasil continental

#micro_br <- geobr::read_micro_region(
#  year = 2013,
#  simplified = TRUE
#)

ggplot() +
  geom_sf(
    data = micro_br,
    fill = NA,
    color = "grey40",
    linewidth = 0.2
  ) +
  coord_sf(expand = FALSE) +
  theme_void() +
  labs(title = "Microrregiões do Brasil (IBGE / geobr)")

```

Using year/date 2013



```
[54]: str(micro_br)
summary(micro_br)
```

```
Classes 'sf' and 'data.frame': 557 obs. of 6 variables:
 $ code_state  : num  11 11 11 11 11 11 11 11 12 12 ...
 $ abbrev_state: chr  "RO" "RO" "RO" "RO" ...
 $ name_state  : chr  "Rondônia" "Rondônia" "Rondônia" "Rondônia" ...
 $ code_micro   : num  11005 11003 11006 11008 11002 ...
 $ name_micro   : chr  "Alvorada D'oeste" "Ariquemes" "Cacoal" "Colorado Do
Oeste" ...
 $ geom         : sfc_GEOMETRY of length 557; first list element: List of 1
  ..$ : num [1:1168, 1:2] -63.6 -63.6 -63.6 -63.6 -63.6 ...
  ..- attr(*, "class")= chr [1:3] "XY" "POLYGON" "sfg"
- attr(*, "sf_column")= chr "geom"
- attr(*, "agr")= Factor w/ 3 levels "constant","aggregate",...: NA NA NA NA NA
  ..- attr(*, "names")= chr [1:5] "code_state" "abbrev_state" "name_state"
"code_micro" ...
#> #> code_state      abbrev_state      name_state      code_micro
#> #> Min.    :11.00  Length:557      Length:557      Min.    :11001
#> #> 1st Qu.:24.00  Class  :character  Class  :character  1st Qu.:24007
#> #> Median  :31.00  Mode   :character  Mode   :character  Median  :31028
#> #> Mean    :31.31
#> #> 3rd Qu.:41.00
#> #> Max.    :53.00
#> #> 
#> #> name_micro      geom
#> #> Length:557      MULTIPOLYGON : 26
#> #> Class  :character  POLYGON      :531
#> #> Mode   :character  epsg:4674     :  0
```

```
+proj=long...: 0
```

## 2 1) Nodes Positions

```
[55]: # -----
# 1) Ler raster VIIRS
# -----
ntl <- terra::rast(DATA$viirs_blackmarble_2016)
names(ntl) <- "value"

# -----
# 2) Função minimalista: ponto mais iluminado por microrregião
# -----
build_nodes_simple <- function(regions_sf, ntl_rast) {

    # garantir mesmo CRS
    regions_ll <- st_transform(regions_sf, st_crs(ntl_rast))

    # extrair valores do raster dentro de cada polígono
    vals <- terra::extract(ntl_rast, terra::vect(regions_ll), cells = TRUE)

    best <- vals |>
        dplyr::group_by(ID) |>
        dplyr::filter(!is.na(value)) |>
        dplyr::slice_max(order_by = value, n = 1, with_ties = FALSE) |>
        dplyr::ungroup()

    xy <- terra::xyFromCell(ntl_rast, best$cell)

    sf::st_as_sf(
        data.frame(
            id = regions_ll$code_micro[best$ID],
            lon = xy[,1],
            lat = xy[,2]
        ),
        coords = c("lon", "lat"),
        crs = st_crs(ntl_rast)
    )
}

# -----
# 3) Construir os nós
# -----
nodes_micro <- build_nodes_simple(micro_br, ntl)
```

```

# -----
# 4) Plot leve
# -----
ggplot() +
  geom_sf(data = micro_br, fill = NA, color = "grey70", linewidth = 0.2) +
  geom_sf(data = nodes_micro, color = "#2E7D32", size = 1) +
  theme_void() +
  labs(title = "Nós = pixel mais iluminado por microrregião")

```

Nós = pixel mais iluminado por microrregião



### 3 2) Contiguity

```

[56]: build_edges_contiguity <- function(regions_sf, nodes_sf,
                                         crs_metric = 5880,
                                         snap_size = 5,
                                         buffer_dist = 5) {

  # detectar automaticamente a coluna de ID
  id_col_regions <- "code_micro"
  id_col_nodes   <- "id"

  # transformar para métrico
  reg <- regions_sf |>
    st_transform(crs_metric) |>
    st_set_precision(1e6) |>
    lwgeom::st_snap_to_grid(size = snap_size) |>
    st_buffer(0) |>
    st_make_valid()

  # ordenar nós pela ordem das regiões

```

```

nodes_m <- nodes_sf |>
  st_transform(crs_metric) |>
  arrange(match(.data[[id_col_nodes]], reg[[id_col_regions]]))

stopifnot(nrow(nodes_m) == nrow(reg))

coords <- st_coordinates(nodes_m)

buf <- st_buffer(reg, buffer_dist)

edge_pairs <- purrr::imap_dfr(
  st_intersects(buf, buf),
  ~ tibble::tibble(i = .y, j = .x)
) |>
  filter(i != j) |>
  mutate(
    a = pmin(i, j),
    b = pmax(i, j)
  ) |>
  distinct(a, b) |>
  transmute(i = a, j = b)

edge_lines <- purrr::map2(
  edge_pairs$i,
  edge_pairs$j,
  ~ st_linestring(rbind(coords[.x, ], coords[.y, ])))
)

st_sf(
  tibble::tibble(
    id_i = reg[[id_col_regions]][edge_pairs$i],
    id_j = reg[[id_col_regions]][edge_pairs$j],
    geometry = st_sfc(edge_lines, crs = crs_metric)
  )
)
}
}

```

```

[57]: edges_micro <- build_edges_contiguity(micro_br, nodes_micro)

ggplot() +
  geom_sf(data = micro_br, fill = NA, color = "grey80", linewidth = 0.2) +
  geom_sf(data = edges_micro, color = "#1E88E5", linewidth = 0.15) +
  geom_sf(data = nodes_micro, color = "#2E7D32", size = 1) +
  theme_void()

```



## 4 3) Contiguity with PA

### 4.1 3.1) Simple PA

```
[58]: read_protected_areas_simple <- function(crs_target = 5880) {

  uc <- geobr::read_conservation_units() |>
    st_make_valid() |>
    st_transform(crs_target)

  ti <- geobr::read_indigenous_land() |>
    st_make_valid() |>
    st_transform(crs_target)

  bind_rows(
    uc |> mutate(src = "UC"),
    ti |> mutate(src = "TI")
  ) |>
    st_make_valid()
}

filter_edges_by_pa <- function(edges_sf, protected_sf, thr = 0.95) {

  # criar um id explícito para cada aresta
  edges_sf <- edges_sf |>
    dplyr::mutate(edge_id = dplyr::row_number())

  inter <- suppressWarnings(sf::st_intersection(edges_sf, protected_sf))
}
```

```

total_len <- as.numeric(sf::st_length(edges_sf))

protected_len <- numeric(nrow(edges_sf))

if (nrow(inter) > 0) {
  # mesmo id das arestas originais
  id    <- inter$edge_id
  len   <- as.numeric(sf::st_length(inter))

  # soma do comprimento protegido por aresta
  agg <- tapply(len, id, sum)

  protected_len[as.integer(names(agg))] <- agg
}

ratio <- protected_len / total_len
keep  <- ratio < thr

list(
  edges_clean = dplyr::select(edges_sf[keep, ], -edge_id),
  removed_idx = !keep,
  n_removed   = sum(!keep),
  n_original   = nrow(edges_sf)
)
}

plot_edges_pa_filter <- function(regions_sf, nodes_sf,
                                    edges_before, edges_after,
                                    prot_sf = NULL,
                                    title = "Arestas com corte por áreas
← protegidas") {

  p <- ggplot() +
    geom_sf(data = regions_sf,
            fill = NA, color = "grey80", linewidth = 0.2)

  if (!is.null(prot_sf)) {
    p <- p +
      geom_sf(data = prot_sf,
              fill = "#FFCDD2", color = "#D32F2F",
              alpha = 0.6, linewidth = 0.2)
  }
}

```

```

p +
  geom_sf(data = edges_before, color = "#9E9E9E",
           linewidth = 0.25, alpha = .4) +
  geom_sf(data = edges_after,  color = "#1E88E5",
           linewidth = 0.35) +
  geom_sf(data = nodes_sf,     color = "#2E7D32",
           size = 1) +
  theme_void() +
  labs(
    title = title,
    subtitle = paste0(
      "Original: ", nrow(edges_before),
      " | Removidas: ", nrow(edges_before) - nrow(edges_after),
      " | Final: ", nrow(edges_after)
    )
  )
}

```

```

[59]: # 1) Ler PAs
prot <- read_protected_areas_simple(crs_target = st_crs(edges_micro))

mask_reg <- micro_br |>
  sf::st_union() |>
  sf::st_transform(sf::st_crs(prot))

prot <- suppressWarnings(
  sf::st_intersection(prot, mask_reg)
)

# 2) Filtrar arestas
pa_clean <- filter_edges_by_pa(edges_sf = edges_micro,
                                 protected_sf = prot,
                                 thr = 0.75)  # thr representa a porcentagem dau
  ↪fronteira que está em PA

edges_clean <- pa_clean$edges_clean

cat("Original:", pa_clean$n_original,
    "| Removidas:", pa_clean$n_removed,
    "| Final:", nrow(edges_clean), "\n")

# 3) Plot
plot_edges_pa_filter(
  regions_sf    = micro_br,
  nodes_sf      = nodes_micro,
  edges_before  = edges_micro,

```

```

    edges_after  = edges_clean,
    prot_sf      = prot
)

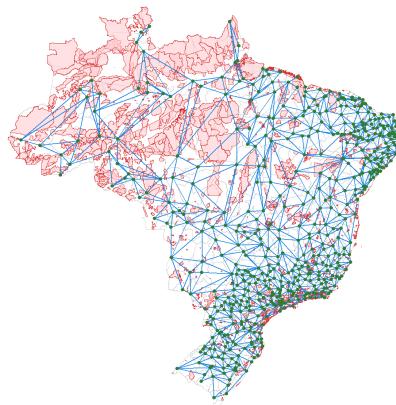
```

Using year/date 201909

Using year/date 201907

Original: 1547 | Removidas: 37 | Final: 1510

Arestas com corte por áreas protegidas  
Original: 1547 | Removidas: 37 | Final: 1510



## 4.2 3.2) PA including forests

```
[60]: library(terra); library(sf); library(dplyr)

crs_target     <- st_crs(edges_micro)
forest_codes   <- c(3L, 4L, 5L)
buf_m          <- 3200    # buffer em metros
min_patch_km2 <- 10      # tamanho mínimo da mancha de floresta (ajuste)

# 1) Raster binário
r <- rast("data/mapbiomas_near_1000m.tif")
r[r == 0] <- NA
r_bin <- classify(r, rbind(
  cbind(forest_codes, forest_codes, 1),
  c(-Inf, min(forest_codes) - 0.001, 0),
  c(max(forest_codes) + 0.001, Inf, 0)
))

# 2) Projeta para métrico
tmp <- tempfile(fileext = ".tif")
```

```

r_bin_m <- project(r_bin, crs_target$wkt, method = "near",
                     filename = tmp, overwrite = TRUE)

# 3) Remove manchas pequenas
cell_area_km2 <- prod(res(r_bin_m)) / 1e6
min_cells      <- ceiling(min_patch_km2 / cell_area_km2)

cl   <- patches(ifel(r_bin_m == 1, 1, NA), directions = 8, zeroAsNA = TRUE)
ftbl <- as.data.frame(freq(cl, digits = 0)) |> filter(!is.na(value))

big_ids <- ftbl$value[ftbl$count >= min_cells]

if (length(big_ids) == 0) {
  message("Nenhum patch >= min_patch_km2; mantendo floresta original.")
  r_forest_clean <- ifel(r_bin_m == 1, 1, NA)
} else {
  r_forest_clean <- ifel(cl %in% big_ids, 1, NA)
}

# 4) Buffer só da floresta (não considera 0 como "dentro")
dist_forest  <- distance(r_forest_clean)
r_forest_buf <- ifel(dist_forest <= buf_m, 1, NA)

# 5) PAs no mesmo buffer e na mesma grade
pa <- read_protected_areas_simple(crs_target = crs_target) |>
  st_make_valid() |>
  st_buffer(buf_m)

if (nrow(pa) == 0) {
  pa_r <- setValues(r_forest_buf, NA)
} else {
  pa_r <- rasterize(vect(pa), r_forest_buf,
                     field = 1, background = NA, touches = TRUE)
}

# garante numérico
pa_r <- as.numeric(pa_r)

# 6) União PA + floresta buffered (cover = pega o 1 do primeiro não-NA)
# máscara Brasil
mask_reg <- micro_br |>
  st_union() |>
  st_transform(crs_target) |>
  st_make_valid()
mask_v <- vect(mask_reg)

```

```

# recorta raster combinado
pa_forest_r <- cover(pa_r, r_forest_buf)
pa_forest_sf <- as.polygons(pa_forest_r, dissolve = TRUE) |>
  st_as_sf() |>
  st_transform(crs_target) |>
  mutate(src = "PA2")

saveRDS(pa_forest_sf, "data/pa2_pa_floresta.rds")

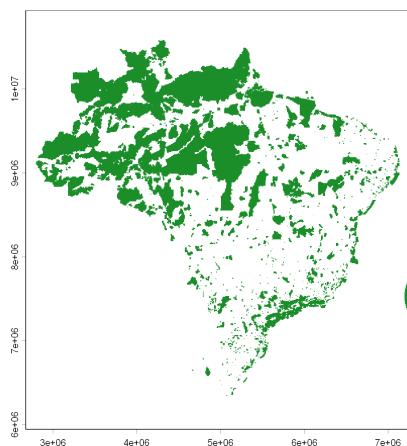
```

Nenhum patch  $\geq$  min\_patch\_km2; mantendo floresta original.

Using year/date 201909

Using year/date 201907

[61]: `plot(pa_forest_r, col = c(NA, "#1B8E2A"), legend = FALSE)`

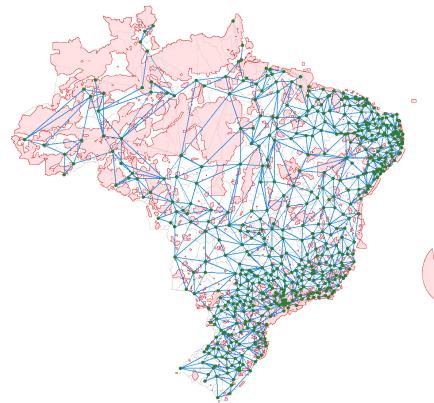


[62]: `pa_forest_clean <- filter_edges_by_pa(
 edges_sf = edges_micro,
 protected_sf = pa_forest_sf,
 thr = 0.75 # ajuste se quiser mais/menos estrito
)
edges_clean <- pa_forest_clean$edges_clean
cat("Original:", pa_forest_clean$n_original,
 "| Removidas:", pa_forest_clean$n_removed,
 "| Final:", nrow(edges_clean), "\n")`

Original: 1547 | Removidas: 106 | Final: 1441

```
[63]: plot_edges_pa_filter(
    regions_sf    = micro_br,
    nodes_sf      = nodes_micro,
    edges_before  = edges_micro,
    edges_after   = edges_clean,
    prot_sf       = pa_forest_sf
)
```

Arestas com corte por áreas protegidas  
Original: 1547 | Removidas: 106 | Final: 1441



## 5 4) Creating feasible edges

```
[64]: # file: R/edge_stat_from_roads_dijkstra_fix.R
suppressPackageStartupMessages({
  library(sf)
  library(dplyr)
  library(lwgeom)
  library(sfnetworks)
  library(tidygraph)
  library(tibble)
})

# --- 1) Sanitizador: deixa só LINESTRING válida e mensurável ---
clean_lines_for_network <- function(x) {
  x |>
    sf::st_zm(drop = TRUE, what = "ZM") |>                      # tirar Z/M (por quê: sfnetwork usa XY)
  ↵sfnetwork usa XY
    sf::st_make_valid() |>                                         # corrigir geometrias
  ↵inválidas
    sf::st_collection_extract("LINESTRING", warn = FALSE) |> # remover
  ↵não-linhas
```

```

sf::st_cast("LINESTRING", warn = FALSE) |> # explodir MULTI em
  ↪simples
  dplyr::filter(!sf::st_is_empty(geometry)) |>
  dplyr::mutate(.len = suppressWarnings(as.numeric(sf::st_length(geometry)))) |>
  ↪|
  dplyr::filter(is.finite(.len) & .len > 0) |>
  dplyr::select(-.len)
}

# --- 2) Endpoints seguros para LINESTRING ---
.edges_endpoints <- function(lines_sf) {
  lines_ok <- lines_sf |>
    sf::st_cast("LINESTRING", warn = FALSE) |>
    dplyr::filter(!sf::st_is_empty(geometry))
  starts <- lwgeom::st_startpoint(lines_ok$geometry)
  ends   <- lwgeom::st_endpoint(lines_ok$geometry)
  tibble(edge_id = seq_len(nrow(lines_ok)),
    from_geom = starts,
    to_geom   = ends) |>
    sf::st_as_sf(crs = sf::st_crs(lines_ok))
}

```

```

[65]: read_bc250_roads_from_DATA <- function(data_bc250,
                                              crs_target) {

  shp_paths <- data_bc250[grep1("\\.shp$", data_bc250)]

  rod_path <- shp_paths[basename(shp_paths) == "rod_trecho_rodoviario_l.shp"]
  if (!length(rod_path))
    stop("Shapefile 'rod_trecho_rodoviario_l.shp' não encontrado em DATA$bc250")

  terra::vect(rod_path[1]) |>
    sf::st_as_sf() |>
    sf::st_make_valid() |>
    sf::st_transform(crs_target)
}

roads_bc250 <- read_bc250_roads_from_DATA(
  DATA$bc250,
  crs_target = st_crs(edges_clean)
)

roads_bc250 <- roads_bc250 |>
  mutate(
    # 1) Pavimentação (você já tem algo assim)
    pav_pav = if_else(
      grep1("Sem revestimento", revestimen, ignore.case = TRUE),
      0, 1
  )

```

```

),
# 2) N° de faixas (já é numérico: nrfaias)

# 3) N° de pistas (simples x dupla)
pistas_duplas = if_else(nrpistas >= 2, 1, 0),

# 4) Concessão
conc_via = if_else(concession == "Sim", 1, 0),

# 5) Jurisdição "alta hierarquia"
jur_fed_est = case_when(
  jurisdicao %in% c("Federal", "Estadual/Distrital") ~ 1,
  jurisdicao == "Municipal" ~ 0,
  TRUE ~ NA_real_
),
# 6) Operacional / tráfego permanente
oper_ativa = if_else(operaciona == "Sim", 1, 0),
traj_perman = if_else(trajego == "Permanente", 1, 0)
)
edges_stats <- edges_clean

# --- 3) DROP-IN: igual à versão Dijkstra, mas com limpeza antes do grafo ---
edge_stat_from_roads <- function(edges_sf,
                                     roads_sf,
                                     var,
                                     buf_km = 25,           # mantido por LU
                                     ↵compatibilidade (ignorado)
                                     crs_metric = 5880) {
  if (!var %in% names(roads_sf)) {
    stop(sprintf("Coluna '%s' inexistente em roads_sf.", var))
  }

  # Projeção
  edges_m <- edges_sf |>
    sf::st_transform(crs_metric) |>
    dplyr::mutate(edge_id = dplyr::row_number())

  roads_m_raw <- roads_sf |>
    sf::st_transform(crs_metric)

  # **PASSO CRÍTICO**: limpar geometrias de rodovias (conserta o erro do print)
  roads_m <- clean_lines_for_network(roads_m_raw)
  if (nrow(roads_m) == 0) stop("roads_sf ficou vazio após limpeza; verifique a ↵camada.")
}

```

```

# Grafo
net <- sfnetworks::as_sfnetwork(roads_m, directed = FALSE)
net <- tidygraph::activate(net, "edges") |>
  dplyr::mutate(
    len_m = as.numeric(sf::st_length(geometry)),
    w      = len_m # custo = distância; troque por tempo se desejar
  )

# Endpoints das suas arestas e snap para nós
ep <- .edge_endpoints(edges_m)
g_nodes <- sf::st_as_sf(net, "nodes")
from_id <- sf::st_nearest_feature(ep$from_geom, g_nodes)
to_id   <- sf::st_nearest_feature(ep$to_geom,   g_nodes)

# Saída
col_name <- paste0("stat_", var)
edges_m[[col_name]] <- NA_real_

net_edges_sf <- sf::st_as_sf(net, "edges")
if (!var %in% names(net_edges_sf)) {
  stop(sprintf("Coluna '%s' não foi preservada na rede (cheque o pré-processamento).", var))
}

# Dijkstra por aresta  $i \rightarrow j$ 
for (i in seq_len(nrow(edges_m))) {
  fi <- from_id[i]; ti <- to_id[i]
  if (!is.finite(fi) || !is.finite(ti) || fi == ti) next

  paths <- tryCatch(
    sfnetworks::st_network_paths(net, from = fi, to = ti, weights = "w", type = "shortest"),
    error = function(e) NULL
  )
  if (is.null(paths) || length(paths$edge_paths) == 0 || length(paths$edge_paths[[1]]) == 0) next

  idx <- as.integer(paths$edge_paths[[1]])
  segs <- net_edges_sf[idx, , drop = FALSE] |>
    dplyr::mutate(
      len_km = as.numeric(sf::st_length(geometry)) / 1000,
      val     = .data[[var]]
    )

  if (!any(is.finite(segs$val))) next
  den <- sum(segs$len_km, na.rm = TRUE)
}

```

```

  if (!is.finite(den) || den <= 0) next

  edges_m[[col_name]][i] <- sum(segs$len_km * segs$val, na.rm = TRUE) / den
}

sf::st_transform(edges_m, sf::st_crs(edges_sf))
}

for (v in c("pav_pav", "nrfaixas", "pistas_duplas",
           "conc_via", "jur_fed_est", "oper_ativa", "traj_perman")) {

  edges_stats <- edge_stat_from_roads(
    edges_sf = edges_stats,    # reutiliza o que já tem
    roads_sf = roads_bc250,
    var       = v,
    buf_km   = 25
  )
}

edges_stats |>
  st_drop_geometry() |>
  summarise(
    across(starts_with("stat_"),
           list(
             n_non_na = ~sum(!is.na(.x)),
             mean     = ~mean(.x, na.rm = TRUE),
             sd       = ~sd(.x,   na.rm = TRUE)
           )),
    .names = "{.col}_{.fn}")
)

```

	stat_pav_pav_n_non_na	stat_pav_pav_mean	stat_pav_pav_sd	stat_nrfaixas_n_non
A tibble: 1 × 21	<int>	<dbl>	<dbl>	<int>
	1382	0.9254838	0.1776586	1379

```

[66]: plot_edge_stat <- function(edges_sf,
                                 regions_sf,
                                 var,
                                 title    = NULL,
                                 subtitle = NULL,
                                 limits   = NULL,
                                 palette  = "viridis",
                                 na.value = "grey80") {

  if (is.null(title)) {
    title <- paste("Rede -", var)
  }
}

```

```

    }

ggplot() +
  geom_sf(data = regions_sf,
           fill = NA, color = "grey90", linewidth = 0.2) +
  geom_sf(data = edges_sf,
           aes_string(color = var),
           linewidth = 0.6) +
  scale_color_viridis_c(
    option = palette,
    na.value = na.value,
    limits = limits
  ) +
  theme_void() +
  labs(
    title = title,
    subtitle = subtitle,
    color = var
  )
}
}

```

```

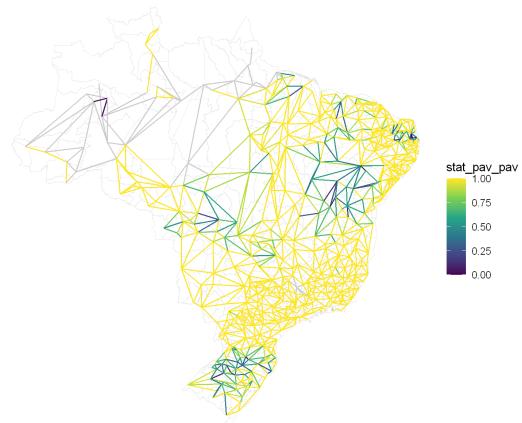
[67]: vars_to_plot <- c(
  "stat_pav_pav",
  "stat_nrfaixas",
  "stat_pistas_duplas",
  "stat_conc_via",
  "stat_jur_fed_est",
  "stat_oper_ativa",
  "stat_traf_perman"
)

plots <- lapply(vars_to_plot, function(v) {
  plot_edge_stat(edges_stats, micro_br, var = v)
})

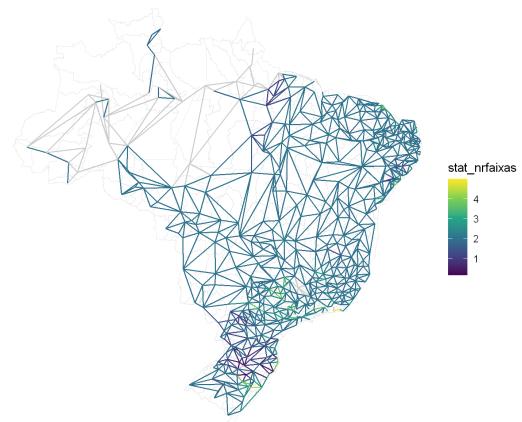
# Para ver um:
plots[]

```

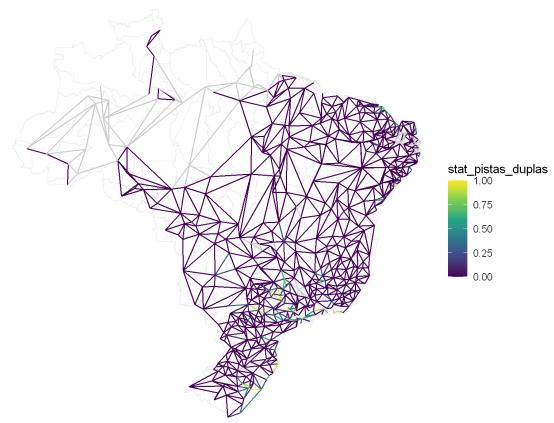
Rede - stat\_pav\_pav



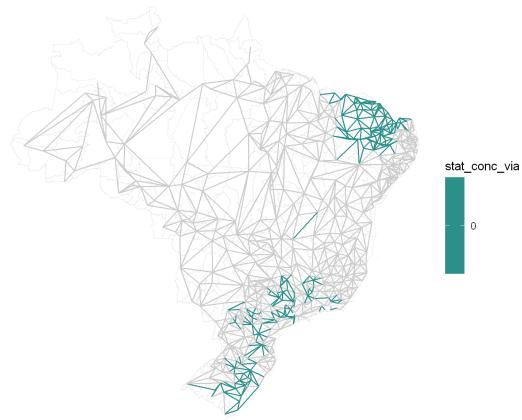
Rede - stat\_nrfaixas



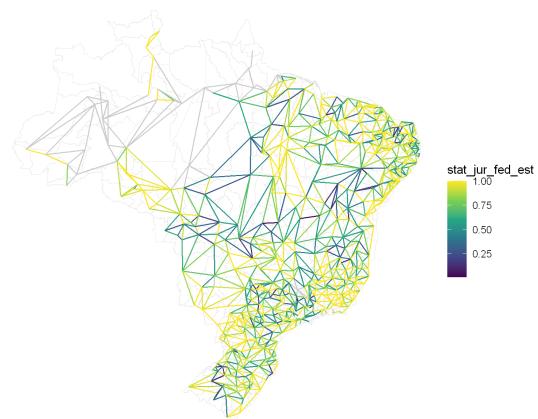
Rede - stat\_pistas\_duplas



Rede - stat\_conc\_via



Rede - stat\_jur\_fed\_est



[[1]]

[[2]]

[[3]]

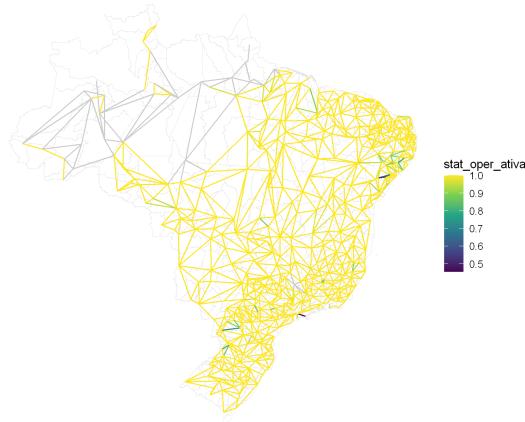
[[4]]

[[5]]

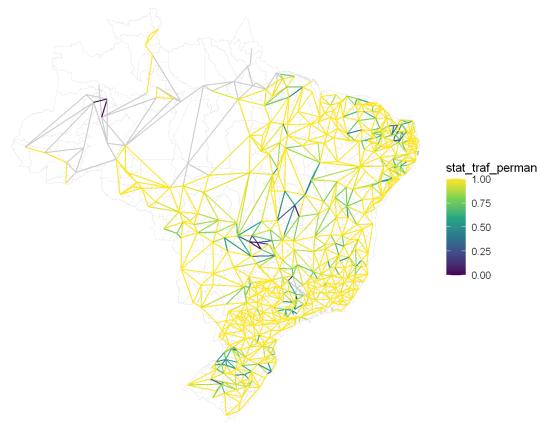
[[6]]

[[7]]

Rede - stat\_oper\_ativa



Rede - stat\_traf\_perman



## 6 5) The Speed Challenge

### 6.1 5.1) Google API

```
[68]: library(httr2)
library(jsonlite)
library(dplyr)

key <- "MY_KEY"

gmaps_speed_live <- function(origin, destination, api_key) {
```

```

url <- "https://maps.googleapis.com/maps/api/directions/json"

resp <- request(url) |>
  req_url_query(
    origin      = origin,
    destination = destination,
    mode        = "driving",
    key         = api_key
  ) |>
  req_perform() |>
  resp_body_json()

leg <- resp$routes[[1]]$legs[[1]]

dist_km <- leg$distance$value / 1000
dur_h   <- leg$duration$value / 3600

tibble(
  origin,
  destination,
  dist_km,
  dur_h,
  speed_kmh = dist_km / dur_h
)
}
}

```

[69]: `#gmaps_speed_live("Sao Paulo", "Campinas", key)`

## 6.2 5.2) Calibrating my infrastructure measure with roadspeed

I construct an infrastructure quality index  $I_{jk}$  as a function of simple road characteristics from BC250, such as pavement type and number of lanes. The functional form is parameterized, and its coefficients are calibrated using a representative subsample of Brazilian links, for which I estimate the mapping from these basic BC250 attributes to effective infrastructure quality. The estimated coefficients are then applied to all BC250 links in Brazil to obtain the nationwide infrastructure measure.

## 7 6) Infrastructure Measure following FS

[70]: `# 6) Infrastructure measure (FS-style)`

```

# Parâmetros tirados/derivados do FS OG
chi_lane   <- 0.355  # efeito de nº de faixas (aprox. da conta deles)
chi_use    <- 1.07   # custo maior em via não nacional
chi_paved  <- 1.35   # custo maior em estrada não pavimentada
chi_median <- 1.05   # custo maior sem canteiro/dupla pista

```

```

I_jk <- edges_stats |>
  mutate(
    # 1) Médias ao longo do caminho (já estão calculadas via Dijkstra)
    share_paved = stat_pav_pav,           # fração do caminho pavimentado
    lanes_mean = if_else(is.na(stat_nrfaixas) | stat_nrfaixas < 1,
                         1, stat_nrfaixas), # nº médio de faixas (mínimo 1)
    share_nat = stat_jur_fed_est,        # fração em via fed/estadual (proxy
    ↵ "national")
    share_median = stat_pistas_duplas,  # fração em pista dupla (proxy de
    ↵ median)

    # 2) Multiplicadores de custo em relação a uma referência "boa"
    mult_lane = lanes_mean^(-chi_lane),
    mult_use = chi_use^(1 - share_nat),
    mult_paved = chi_paved^(1 - share_paved),
    mult_median = chi_median^(1 - share_median),

    # 3) Custo relativo de transporte pela qualidade da infraestrutura
    infra_cost_mult = mult_lane * mult_use * mult_paved * mult_median,

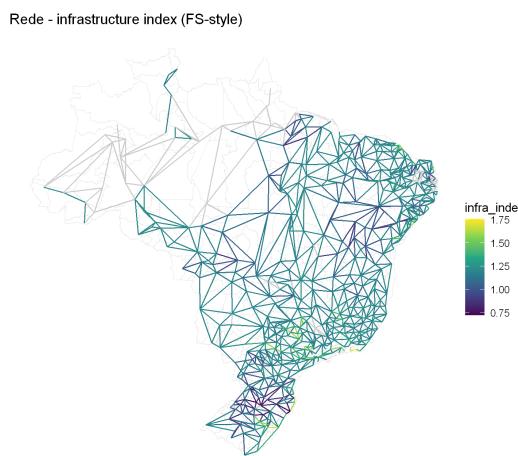
    # 4) Medida de infraestrutura: quanto maior, melhor a infraestrutura
    infra_index = 1 / infra_cost_mult
  )
)

```

```

[71]: plot_edge_stat(
  edges_sf = I_jk,
  regions_sf = micro_br,
  var = "infra_index",
  title = "Rede - infrastructure index (FS-style)"
)

```



## 8 7) Nodes Statistics

### 8.1 7.1) Population and GDP from SIDRA

```
[72]: # -----
# 7.1.1 Population (SIDRA 6579)
# -----
# POP_YEAR <- "2025"

pop_raw <- sidrar::get_sidra(
  6579,
  variable = 9324,
  period = POP_YEAR,
  geo = "City",
  format = 3
)

# Detect column names robustly (handles accents, name changes)
col_muni <- names(pop_raw)[
  grepl("Munic", names(pop_raw), ignore.case = TRUE) &
  grepl("C[oó]d", names(pop_raw), ignore.case = TRUE)
][1]

pop <- pop_raw |>
  transmute(
    cod_mun = suppressWarnings(as.integer(readr::parse_number(.[
      data[[col_muni]]])),
    populacao = suppressWarnings(as.numeric(Valor))
  )

lookup <- geobr::lookup_muni("all") |> select(code_muni, code_micro, abbrev_state)

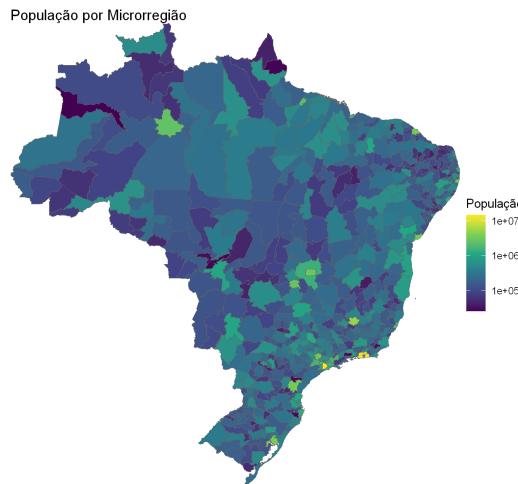
pop_micro <- pop |>
  left_join(lookup, by = c("cod_mun" = "code_muni")) |>
  group_by(code_micro, abbrev_state) |>
  summarise(populacao = sum(populacao, na.rm = TRUE), .groups = "drop")

micro_pop <- micro_br |>
  left_join(pop_micro, by = c("code_micro", "abbrev_state"))

ggplot(micro_pop) +
  geom_sf(aes(fill = populacao), color = "grey40", size = 0.05) +
  scale_fill_viridis_c(trans = "log10", na.value = "grey90") +
  coord_sf(expand = FALSE) +
  theme_void() +
```

```
  labs(fill = "População", title = "População por Microrregião")
```

Considering all categories once 'classific' was set to 'all' (default)



[73]:

```
# -----  
# 7.1.2 GDP (SIDRA 5938)  
# -----  
  
PIB_VAR <- 37  
  
pib_raw <- sidrar::get_sidra(  
  5938,  
  variable = PIB_VAR,  
  period = "last",  
  geo = "City",  
  format = 3  
)  
  
col_muni <- names(pib_raw)[  
  grepl("Munic", names(pib_raw), ignore.case = TRUE) &  
  grepl("C[oó]d", names(pib_raw), ignore.case = TRUE)  
][1]  
  
pib <- pib_raw |>  
  transmute(  
    cod_mun = suppressWarnings(as.integer(readr::parse_number(.  
      ↪ data[[col_muni]]))),  
    pib_mil_rs = suppressWarnings(as.numeric(Valor))  
)
```

```

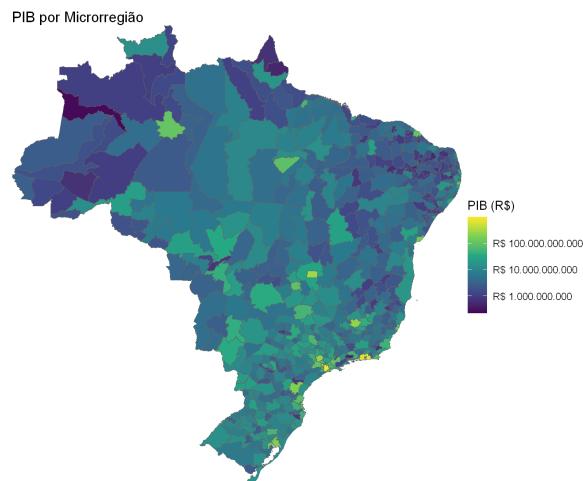
pib_micro <- pib |>
  left_join(lookup, by = c("cod_mun" = "code_muni")) |>
  group_by(code_micro, abbrev_state) |>
  summarise(pib_rs = sum(pib_mil_rs * 1000, na.rm = TRUE), .groups = "drop")

micro_pib <- micro_br |>
  left_join(pib_micro, by = c("code_micro", "abbrev_state"))

ggplot(micro_pib) +
  geom_sf(aes(fill = pib_rs), color = "grey40", size = 0.05) +
  scale_fill_viridis_c(
    trans = "log10",
    na.value = "grey90",
    labels = scales::label_number(big.mark = ".", decimal.mark = ",",
    prefix = "R$ ")
  ) +
  coord_sf(expand = FALSE) +
  theme_void() +
  labs(fill = "PIB (R$)", title = "PIB por Microrregião")

```

Considering all categories once 'classific' was set to 'all' (default)



```

[74]: library(tidyr)

# -----
# 7.1.3 Join population and GDP + percentile maps
# -----

micro_all <- micro_br |>
  left_join(pop_micro, by = c("code_micro", "abbrev_state")) |>

```

```

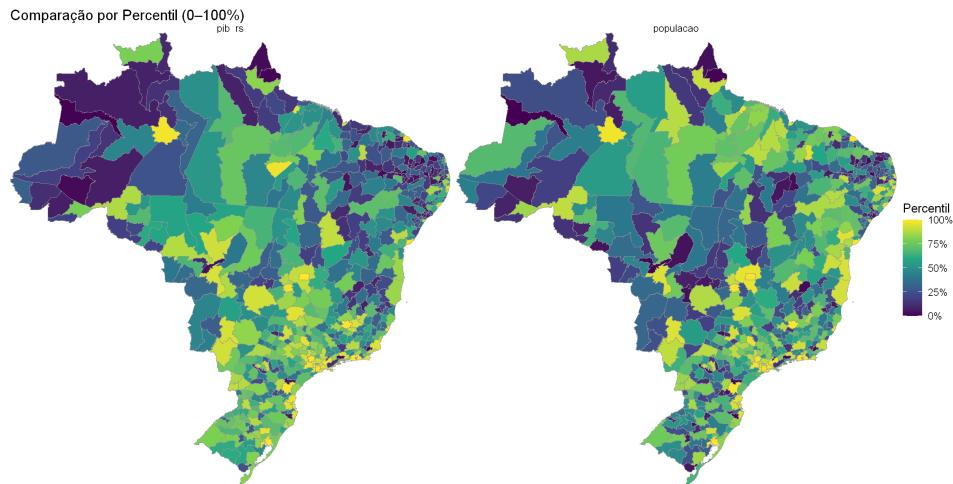
left_join(pib_micro, by = c("code_micro", "abbrev_state"))

mp_long <- micro_all |>
  select(code_micro, populacao, pib_rs) |>
  pivot_longer(
    c(populacao, pib_rs),
    names_to = "var",
    values_to = "val"
  )

mp_rel <- mp_long |>
  group_by(var) |>
  mutate(val_pct = percent_rank(val)) |>
  ungroup()

ggplot(mp_rel) +
  geom_sf(aes(fill = val_pct), color = "grey60", size = 0.06) +
  facet_wrap(~ var, nrow = 1) +
  scale_fill_viridis_c(
    limits = c(0,1),
    labels = scales::label_percent(accuracy = 1),
    name = "Percentil"
  ) +
  coord_sf(expand = FALSE) +
  theme_void() +
  labs(title = "Comparação por Percentil (0-100%)")

```



## 8.2 7.2) Population and GDP divided by sectors

### 8.2.1 7.2.1) SIDRA GDP per sector

```
[75]: baixar_pib_setor <- function(var_cod, nome_var) {  
  sidrar::get_sidra(  
    x = 5938,  
    variable = var_cod,  
    geo = "City",  
    period = "last",  
    format = 3  
  ) |>  
  transmute(  
    cod_mun = as.integer(`Município (Código)`),  
    !!nome_var := as.numeric(Valor)  
  )  
}
```

```
[76]: vars <- list(  
  pib_total = 37,  
  pib_agro = 513,  
  pib_imp = 514,  
  pib_serv = 6575,  
  pib_ind = 543  
)  
  
pib_mun <- purrr::map2_dfr(  
  vars,  
  names(vars),  
  baixar_pib_setor  
) |>  
  group_by(cod_mun) |>  
  summarise(across(everything()), ~ first(na.omit(.x))), .groups = "drop")
```

Considering all categories once 'classific' was set to 'all' (default)

Considering all categories once 'classific' was set to 'all' (default)

Considering all categories once 'classific' was set to 'all' (default)

Considering all categories once 'classific' was set to 'all' (default)

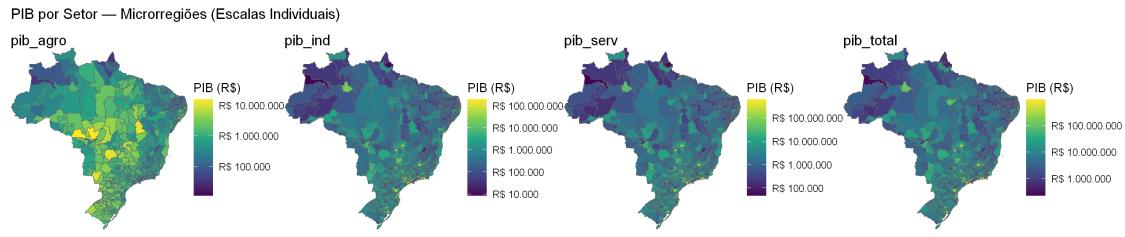
Considering all categories once 'classific' was set to 'all' (default)

```
[77]: pib_micro <- pib_mun |>  
  left_join(lookup, by = c("cod_mun" = "code_muni")) |>  
  group_by(code_micro, abbrev_state) |>
```

```
  summarise(across(starts_with("pib_"), ~ sum(.x, na.rm = TRUE)), .groups =  
  ↵"drop")
```

```
[78]: micro_pib_setores <- micro_br |>  
      left_join(pib_micro, by = c("code_micro", "abbrev_state"))
```

```
[79]: library(ggplot2)  
library(patchwork)  
  
plot_pib_setor <- function(df, var) {  
  ggplot(df) +  
    geom_sf(aes(fill = .data[[var]]), color = "grey40", size = 0.05) +  
    scale_fill_viridis_c(  
      trans = "log10",  
      na.value = "grey90",  
      #breaks = c(1e3, 1e4, 1e5, 1e6, 1e7, 1e8),  
      labels = scales::label_number(big.mark = ".", decimal.mark = ", ", prefix =  
      ↵= "R$ ")#, scale_cut = scales::cut_short_scale())  
    ) +  
    coord_sf(expand = FALSE) +  
    theme_void() +  
    labs(  
      title = var,  
      fill = "PIB (R$)"  
    )  
}  
  
p1 <- plot_pib_setor(micro_pib_setores, "pib_agro")  
#p2 <- plot_pib_setor(micro_pib_setores, "pib_imp")  
p3 <- plot_pib_setor(micro_pib_setores, "pib_ind")  
p4 <- plot_pib_setor(micro_pib_setores, "pib_serv")  
p5 <- plot_pib_setor(micro_pib_setores, "pib_total")  
  
(p1 | p3 | p4 | p5) +  
  plot_annotation(  
    title = "PIB por Setor - Microrregiões (Escalas Individuais)"  
  )
```



### 8.2.2 7.2.2) RAIS population per sector

```
[80]: library(readr)
library(dplyr)
library(tidyr)
library(geobr)
library(ggplot2)
library(scales)
library(stringr)
library(stringi)
library(patchwork)

# 1) Ler e preparar RAIS
colunas <- c(
  "municipio",
  "setor1_extrativa",
  "setor2_transformacao",
  "setor3_utilidade_publica",
  "setor4_construcao",
  "setor5_comercio",
  "setor6_servicos",
  "setor7_administracao",
  "setor8_agro_pesca",
  "setor9_ignorado",
  "total"
)
emprego_raw <- read_csv2(
  "consulta9293258.csv",
  skip = 1,
```

```

col_names = colunas,
locale = locale(encoding = "latin1"),
na = c("", "*", "-")
) |>
  mutate(across(-municipio, parse_number))

norm_nome <- function(x) {
  x |>
    stringi::stri_trans_general("Latin-ASCII") |>
    tolower() |>
    str_remove("^[a-z]{2}-") |>
    str_replace_all("[^a-z ]", " ") |>
    str_squish()
}

lookup <- geobr::lookup_muni("all") |>
  transmute(
    code_muni, code_micro, abbrev_state,
    nome_chave = norm_nome(name_muni)
  )

emprego_geo <- emprego_raw |>
  mutate(
    agro = setor8_agro_pesca,
    industria = setor1_extrativa + setor2_transformacao +
      setor3_utilidade_publica + setor4_construcao,
    servicos = setor5_comercio + setor6_servicos + setor7_administracao,
    nome_chave = norm_nome(municipio)
  ) |>
  left_join(lookup, by = "nome_chave")

emprego_micro <- emprego_geo |>
  group_by(code_micro, abbrev_state) |>
  summarise(across(c(agro, industria, servicos), sum, na.rm = TRUE), .groups = "drop")

micro_emprego <- geobr::read_micro_region(year = 2013) |>
  left_join(emprego_micro, by = c("code_micro", "abbrev_state"))

# 2) Plots com escalas independentes
plot_emprego_setor <- function(df, var, titulo) {
  ggplot(df, aes(fill = .data[[var]])) +
    geom_sf(color = "grey65", size = 0.05) +
    scale_fill_viridis_c(
      trans = "log10",
      labels = label_number(big.mark = ".", decimal.mark = ","),
      name = "Empregos"
}

```

```

) +
  labs(title = titulo) +
  theme_void() +
  theme(
    plot.title = element_text(hjust = 0.5, face = "bold"),
    legend.position = "right"
  )
}

p_agro <- plot_emprego_setor(micro_emprego, "agro", "Agropecuária")
p_ind  <- plot_emprego_setor(micro_emprego, "industria", "Indústria")
p_serv <- plot_emprego_setor(micro_emprego, "servicos", "Serviços")

(p_agro | p_ind | p_serv) +
  plot_layout(guides = "keep") +
  plot_annotation(title = "Emprego por Setor")

```

Using `','` as decimal and  
`'.'` as grouping mark. Use ``read_delim()`` for more control.

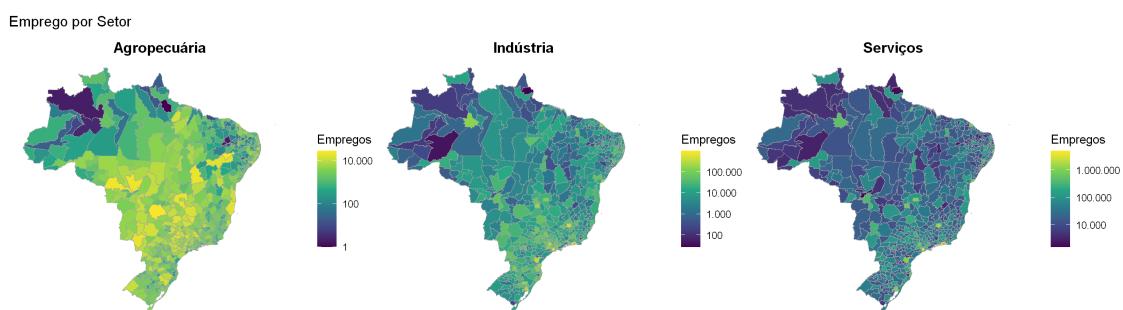
Rows: 5577 Columns: 11  
 Column specification

Delimiter: ";"  
`chr` (11): municipio, setor1\_extrativa, setor2\_transformacao,  
 setor3\_utilidad...

Use ``spec()`` to retrieve the full column specification for this data.

Specify the column types or set ``show_col_types = FALSE`` to quiet this message.

Using year/date 2013



### 8.2.3 7.2.3) Censo population and gdp per sector

[81]: #TBD

## 9 8) Cost of building infrastructure

### 9.1 8.1) delta FS

[82]: # 6) Delta<sup>~</sup>I\_jk - distance + ruggedness (FS/Collier)

```
library(terra)
library(purrr)

# 6.1 Ruggedness ao longo de cada aresta (ETOPO1)
ETOPO_PATH <- "ETOPO1_Ice_g_geotiff.tif" # ajuste se estiver em outro lugar
etopo <- terra::rast(ETOPO_PATH)
if (is.na(terra::crs(etopo))) terra::crs(etopo) <- "EPSG:4326"

compute_ruggedness <- function(edges_sf, etopo_rast) {
  edges_ll <- sf::st_transform(edges_sf, crs = terra::crs(etopo_rast))
  rugged_fun <- function(g) {
    pts <- sf::st_line_sample(g, density = 50) |> sf::st_coordinates()
    elev <- terra::extract(etopo_rast, pts[, c("X", "Y")])[,1]
    if (sum(!is.na(elev)) < 3) return(NA_real_)
    sd(diff(elev), na.rm = TRUE)
  }
  dplyr::mutate(edges_sf,
                rugged = purrr::map_dbl(edges_ll$geometry, rugged_fun))
}

rugged <- compute_ruggedness(I_jk, etopo)

# 6.2 Delta~I_jk usando exatamente os coeficientes do FS
compute_delta_geo <- function(edges_sf,
                                 rugged_col = "rugged",
                                 crs_metric = 5880) {

  edges_m <- sf::st_transform(edges_sf, crs_metric)
  dist_km <- as.numeric(sf::st_length(edges_m)) / 1000
  rugged <- edges_sf[[rugged_col]]

  delta0 <- 1
  delta1 <- -0.11 # dummy dist>50 km
  delta2 <- 0.12 # ln(ruggedness)
```

```

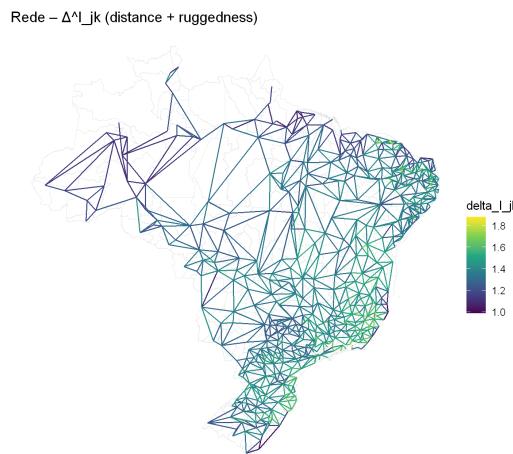
delta_geo <- exp(
  log(delta0) +
  delta1 * (dist_km > 50) +
  delta2 * log(pmax(rugged, 1e-4))
)

dplyr::mutate(edges_sf,
  dist_km      = dist_km,
  delta_I_jk = delta_geo
}

delta_I_jk <- compute_delta_geo(rugged)

```

```
[83]: plot_edge_stat(
  edges_sf  = delta_I_jk,
  regions_sf = micro_br,
  var        = "delta_I_jk",
  title      = "Rede - Δ^I_jk (distance + ruggedness)"
)
```



## 9.2 8.2) new delta (TBD)

```
[84]: #library(sf)
#library(dplyr)
#library(terra)

#rescale01 <- function(x) {
#  x <- as.numeric(x)
#  x_ok <- x[is.finite(x)]
```

```

#  if (!length(x_ok)) return(rep(0, length(x)))
#  rng <- range(x_ok, na.rm = TRUE)
#  if (diff(rng) == 0) return(rep(0, length(x)))
#  (x - rng[1]) / diff(rng)
#}

#build_env_nodes <- function(nodes_sf, DATA,
#                               crs_ll = 4326,
#                               soil_col = "soil_class") {
#
#  nodes_ll <- st_transform(nodes_sf, crs_ll)

## 1) Erosão IBGE (2.5 GB, mas só 1 extract em pontos)
#  eros_r <- rast(DATA$erosao_2020_ibge)
#  if (is.na(crs(eros_r))) crs(eros_r) <- sprintf("EPSG:%d", crs_ll)
#  eros_vals <- terra::extract(eros_r, vect(nodes_ll))[,1]
#  eros_vals[is.na(eros_vals)] <- median(eros_vals, na.rm = TRUE)
#  eros_idx <- rescale01(eros_vals)

## 2) Solo Embrapa WRB -> índice de fragilidade
#  soils <- st_read(DATA$solos_wrb[1], quiet = TRUE) />
#  st_make_valid() />
#  st_transform(crs_ll)

#  soil_join <- st_join(nodes_ll, soils)
#  soil_raw <- as.factor(soil_join[[soil_col]])
#  soil_num <- as.numeric(soil_raw)
#  soil_num[is.na(soil_num)] <- median(soil_num, na.rm = TRUE)
#  soil_frag <- rescale01(soil_num)

## 3) MapBiomas 10m -> raster agregado mais leve + share de floresta
#  mb <- rast(DATA$mapbiomas_10m_2023)
#  if (is.na(crs(mb))) crs(mb) <- sprintf("EPSG:%d", crs_ll)

#  forest_codes <- c(3L, 4L, 5L)
#  mb_forest <- mb %in% forest_codes

#  10m -> ~1km (ajuste o factor se quiser mais fino/grosso)
#  fact <- 100L
#  mb_forest_agg <- terra::aggregate(mb_forest, fact = fact, fun = mean, na.rm = TRUE)

#  forest_share <- terra::extract(mb_forest_agg, vect(nodes_ll))[,1]
#  forest_share[is.na(forest_share)] <- 0

#  env_nodes <- nodes_sf />
#  st_drop_geometry() />

```

```

#     transmute(
#       id,
#       eros_idx      = eros_idx,
#       soil_frag     = soil_frag,
#       forest_share  = pmin(pmax(forest_share, 0), 1)
#     )

# env_nodes
#}

# --- só roda UMA vez; depois é só ler o RDS ---
#env_nodes <- build_env_nodes(nodes_micro, DATA)
#dir.create("data", showWarnings = FALSE)
#saveRDS(env_nodes, "data/env_nodes.rds")

```

```

[85]: #env_nodes <- readRDS("data/env_nodes.rds")

# média i-j
#env_i <- env_nodes |> rename(id_i = id,
#                               eros_i = eros_idx,
#                               soil_i = soil_frag,
#                               forest_i = forest_share)
#env_j <- env_nodes |> rename(id_j = id,
#                               eros_j = eros_idx,
#                               soil_j = soil_frag,
#                               forest_j = forest_share)

#edges_env <- I_jk |>
#  left_join(env_i, by = "id_i") |>
#  left_join(env_j, by = "id_j") |>
#  mutate(
#    eros_mean    = rowMeans(cbind(eros_i, eros_j), na.rm = TRUE),
#    soil_mean    = rowMeans(cbind(soil_i, soil_j), na.rm = TRUE),
#    forest_mean  = rowMeans(cbind(forest_i, forest_j), na.rm = TRUE)
#  )

# distância + rugged (que já é rápido no seu código atual)
#edges_m <- st_transform(edges_env, 5880)
#dist_km <- as.numeric(st_length(edges_m)) / 1000
#rugged_m <- edges_env$rugged_mean  # se você já tem, ou use sua versão atual

#delta0      <- 1
#delta1_dist <- -0.11
#delta2_rug  <- 0.12
#theta_eros   <- 0.20
#theta_soil   <- 0.15
#theta_forest <- 0.25

```

```

#delta_I_jk_pedro <- exp(
#  log(delta0) +
#  delta1_dist * (dist_km > 50) +
#  delta2_rug * log(pmax(rugged_m, 1e-4)) +
#  theta_eros * rescale01(eros_mean) +
#  theta_soil * rescale01(soil_mean) +
#  theta_forest * forest_mean
#)

#delta_I_jk_pedro <- edges_env |>
#  mutate(
#    dist_km           = dist_km,
#    delta_I_jk_pedro = delta_I_jk_pedro
#  )

```

```

[86]: #plot_edge_stat
#  edges_sf  = delta_I_jk_pedro,
#  regions_sf = micro_br,
#  var        = "delta_I_jk",
#  title      = "Rede - Δ^I_jk (distance + ruggedness)"
#)

```

## 10 9) Creating Matrices and exporting to matlab

```

[87]: # -----
# 9) Matrizes + export + plots (hidro + aeroportos, sem mults)
# -----
# Fontes fixas:
#  - Solo: soils_brazil_wrb_wgs84.shp
#  - Hidrovias: bc250_completo/hdv_trecho_hidroviario_l.shp
#  - Eclusas: bc250_completo/hdv_eclusa_p.shp
#  - Portos: bc250_completo/hdv_complexo_portuario_p.shp
#  - Barragens: bc250_completo/hid_barragem_p.shp
#  - Aeroportos: bc250_completo/aer_complexo_aeroportuario_p.shp (sem pistas)

suppressPackageStartupMessages({
  library(dplyr); library(sf); library(terra); library(R.matlab)
  library(lwgeom)
  library(ggplot2); library(patchwork); library(scales); library(grid)
})
OUT_DIR <- "output"; crs_ll <- 4326; crs_m <- 5880

# -----
# Helpers
# -----

```

```

assert_cols <- function(x, cols, nm) {
  miss <- setdiff(cols, names(x))
  if (length(miss)) stop(nm, ": faltam colunas: ", paste(miss, collapse = ", "))
}

rescale01 <- function(x) {
  x <- as.numeric(x); ok <- is.finite(x)
  if (!any(ok)) return(rep(0, length(x)))
  r <- range(x(ok], na.rm = TRUE); if (diff(r) == 0) return(rep(0, length(x)))
  (x - r[1]) / diff(r)
}

fill_mat <- function(df, value_col, ids) {
  m <- matrix(0, length(ids), length(ids), dimnames = list(ids, ids))
  ii <- match(df$id_i, ids); jj <- match(df$id_j, ids); vals <- df[[value_col]]
  ok <- is.finite(ii) & is.finite(jj) & !is.na(vals)
  for (k in which(ok)) {
    i <- ii[k]; j <- jj[k]; v <- vals[k]
    m[i,j] <- v; m[j,i] <- v
  }
  diag(m) <- 0
  m
}

avg_edge <- function(varname, nodes_sf, edges_sf) {
  df_nodes <- st_drop_geometry(nodes_sf)
  assert_cols(df_nodes, c("code_micro", varname), "avg_edge")
  vals <- df_nodes[[varname]]
  idmap <- df_nodes$code_micro
  vi <- vals[match(edges_sf$id_i, idmap)]
  vj <- vals[match(edges_sf$id_j, idmap)]
  rowMeans(cbind(vi, vj), na.rm = TRUE)
}

norm_txt <- function(x) {
  x <- tolower(iconv(x, to = "ASCII//TRANSLIT"))
  trimws(x)
}

# -----
# Bases obrigatórias
# -----
assert_cols(nodes_micro, c("geometry"), "nodes_micro")
assert_cols(I_jk, c("id_i", "id_j", "geometry"), "I_jk")
if (!"code_micro" %in% names(nodes_micro)) {
  if ("id" %in% names(nodes_micro)) nodes_micro$code_micro <- nodes_micro$id
  else stop("nodes_micro precisa de code_micro ou id.")
}

# -----
# Coordenadas / nós

```

```

# -----
nodes_ll <- nodes_micro |> st_transform(crs_ll)
coords   <- st_coordinates(nodes_ll)
nodes_ll$lon <- coords[,1]
nodes_ll$lat <- coords[,2]

# -----
# Ruggedness (ETOPO TRI por n\ufe0f3)
# -----
tryCatch({
  etopo_path <- if (exists("DATA") && "etopo1_elevation" %in% names(DATA))
    DATA$etopo1_elevation else "ETOPO1_Ice_g_geotiff.tif"
  etopo_r <- rast(etopo_path)
  if (is.na(crs(etopo_r))) crs(etopo_r) <- sprintf("EPSG:%d", crs_ll)
  tri_rugged <- terra::terrain(etopo_r, v = "TRI", neighbors = 8)
  rugged_node <- terra::extract(tri_rugged, terra::vect(nodes_ll))[,1]
  rugged_node[is.na(rugged_node)] <- median(rugged_node, na.rm = TRUE)
  nodes_ll$rugged_tri <- rugged_node
  message("\u2714 Ruggedness (ETOPO TRI) por n\ufe0f3")
}, error = function(e) message("[ruggedness] ", e$message))

# -----
# Solo (WRB)
# -----
tryCatch({
  soils <- st_read("soils_brazil_wrb_wgs84.shp", quiet = TRUE) |>
    st_make_valid() |>
    st_transform(crs_ll)
  soil_join <- st_join(nodes_ll, soils)
  nodes_ll$soil_cat_idx <- rescale01(as.numeric(factor(soil_join$symbol_wrb)))
  message(" Solo: soils_brazil_wrb_wgs84.shp (indice categ. 0-1)")
}, error = function(e) message("[solo] ", e$message))

# -----
# Eclusas, Portos, Barragens (nós = contagem por microrregião)
# -----
tryCatch({
  locks_sf <- st_read("bc250_completo/hdv_eclusa_p.shp", quiet = TRUE) |>
    st_transform(crs_ll)
  locks_by_micro <- st_join(st_transform(micro_br, crs_ll), locks_sf, join = st_intersects) |>
    st_drop_geometry() |>
    count(code_micro, name = "n_locks")
  nodes_ll$n_locks <- locks_by_micro$n_locks[match(nodes_ll$code_micro, locks_by_micro$code_micro)]
  nodes_ll$n_locks[is.na(nodes_ll$n_locks)] <- 0
  message(" Eclusas: contagem por microrregião")
}

```

```

}, error = function(e) message("[eclusas] ", e$message))

tryCatch({
  ports_sf <- st_read("bc250_completo/hdv_complexo_portuario_p.shp", quiet = TRUE) |>
    st_transform(crs_ll)
  ports_by_micro <- st_join(st_transform(micro_br, crs_ll), ports_sf, join = st_intersects) |>
    st_drop_geometry() |>
    count(code_micro, name = "n_ports")
  nodes_ll$n_ports <- ports_by_micro$n_ports[match(nodes_ll$code_micro, ports_by_micro$code_micro)]
  nodes_ll$n_ports[is.na(nodes_ll$n_ports)] <- 0
  message(" Portos: contagem por microrregião")
}, error = function(e) message("[portos] ", e$message))

tryCatch({
  dams_sf <- st_read("bc250_completo/hid_barragem_p.shp", quiet = TRUE) |>
    st_transform(crs_ll)
  dams_by_micro <- st_join(st_transform(micro_br, crs_ll), dams_sf, join = st_intersects) |>
    st_drop_geometry() |>
    count(code_micro, name = "n_dams")
  nodes_ll$n_dams <- dams_by_micro$n_dams[match(nodes_ll$code_micro, dams_by_micro$code_micro)]
  nodes_ll$n_dams[is.na(nodes_ll$n_dams)] <- 0
  message(" Barragens: contagem por microrregião")
}, error = function(e) message("[barragens] ", e$message))

# -----
# Aeroportos (nós: dummy > 0)
# -----
tryCatch({
  aero_sf <- st_read("bc250_completo/aer_complexo_aeroportuario_p.shp", quiet = TRUE) |>
    st_make_valid() |>
    st_transform(crs_ll)

  coords <- st_coordinates(aero_sf)
  coords_rounded <- round(coords, 5)
  aero_sf$xy_round <- paste(coords_rounded[,1], coords_rounded[,2])
  aero_sf <- aero_sf[!duplicated(aero_sf$xy_round),]

  reg_aer <- st_transform(micro_br, crs_ll)
  aer_micro <- st_join(aero_sf, reg_aer["code_micro"], left = FALSE)
}

```

```

aer_counts <- aer_micro |>
  st_drop_geometry() |>
  count(code_micro, name = "n_airports")

nodes_ll$n_airports <- aer_counts$n_airports[match(nodes_ll$code_micro, □
  ↪aer_counts$code_micro)]
nodes_ll$n_airports[is.na(nodes_ll$n_airports)] <- 0
nodes_ll$n_airport_dummy <- as.numeric(nodes_ll$n_airports > 0)

  message(" Aeroportos: complexos dedup + intersect")
}, error = function(e) message("[aeroportos] ", e$message))

# -----
# Tabela de nós
# -----
assert_cols(micro_all,   c("code_micro", "abbrev_state", "populacao", "pib_rs"), □
  ↪"micro_all")
assert_cols(pib_micro,   c("code_micro", "abbrev_state"), □
  ↪"pib_micro")
assert_cols(micro_emprego, □
  ↪c("code_micro", "abbrev_state", "agro", "industria", "servicos"), □
  ↪"micro_emprego")

ids_micro <- sort(unique(c(I_jk$id_i, I_jk$id_j)))
J <- length(ids_micro)

nodes_tbl <- nodes_ll |>
  st_drop_geometry() |>
  left_join(micro_all |> st_drop_geometry() |>
    select(code_micro, abbrev_state, populacao, pib_rs),
    by = "code_micro") |>
  left_join(pib_micro, by = c("code_micro", "abbrev_state")) |>
  left_join(micro_emprego |> st_drop_geometry() |>
    select(code_micro, abbrev_state, agro, industria, servicos),
    by = c("code_micro", "abbrev_state")) |>
  mutate(
    pop_total = populacao,
    pib_total = coalesce(pib_total, pib_rs),
    pop_agro  = agro,
    pop_ind   = industria,
    pop_serv  = servicos
  ) |>
  filter(code_micro %in% ids_micro) |>
  arrange(match(code_micro, ids_micro))

df_nodes <- nodes_tbl |>
  transmute(

```

```

  id  = as.numeric(code_micro),
  pop = as.numeric(pop_total),
  pib = as.numeric(pib_total),
  lon = as.numeric(lon),
  lat = as.numeric(lat)
) |>
as.matrix()

pib_sector_cols <- setdiff(grep("pib_", names(nodes_tbl), value = TRUE), 
  ↪"pib_total")
pib_sector_mat  <- if (length(pib_sector_cols))
  as.matrix(nodes_tbl[, pib_sector_cols, drop = FALSE]) else NULL

pop_sector_cols <- intersect(c("pop_agro", "pop_ind", "pop_serv"), 
  ↪names(nodes_tbl))
pop_sector_mat  <- if (length(pop_sector_cols))
  as.matrix(nodes_tbl[, pop_sector_cols, drop = FALSE]) else NULL

node_airports <- if ("n_airport_dummy" %in% names(nodes_tbl))
  as.numeric(nodes_tbl$n_airport_dummy) else NULL

# -----
# Arestas / matrizes
# -----
edges_sf <- I_jk |>
  st_transform(crs_m) |>
  mutate(dist_km = as.numeric(st_length(geometry)) / 1000)

if ("share_nat" %in% names(edges_sf))
  edges_sf$share_fed_state <- edges_sf$share_nat

# Hidrovias como ARESTA: km de hidrovias do BC250 "próximas" à aresta
tryCatch({
  # carregar hidrovias operacionais
  hydro_m <- st_read("bc250_completo/hdv_trecho_hidroviario_l.shp", quiet = 
    ↪TRUE) |>
    st_make_valid() |>
    st_transform(crs_m) |>
    mutate(
      oper_norm = norm_txt(operaciona),
      sit_norm  = norm_txt(situacaofi)
    ) |>
    filter(
      oper_norm %in% c("sim", "operacional"),
      sit_norm  %in% c("construida", "em operacao")
    ) |>
  
```

```

dplyr::select(geometry)

# buffer em torno de cada aresta (corredor de navegação)
buf_dist <- 20000 # 20 km

edges_buf <- st_buffer(
  edges_sf |> dplyr::select(id_i, id_j, geometry),
  dist = buf_dist
)

# interseção: hidrovias que passam dentro do corredor da aresta
inter <- suppressWarnings(
  st_intersection(edges_buf, hydro_m)
)

if (nrow(inter) == 0) {
  edges_sf$waterway_km <- 0
} else {
  inter$len_km <- as.numeric(st_length(inter)) / 1000 # km

  water_by_edge <- inter |>
    st_drop_geometry() |>
    group_by(id_i, id_j) |>
    summarise(waterway_km = sum(len_km, na.rm = TRUE), .groups = "drop")

  key_edges <- paste(edges_sf$id_i, edges_sf$id_j)
  key_water <- paste(water_by_edge$id_i, water_by_edge$id_j)
  m <- match(key_edges, key_water)

  edges_sf$waterway_km <- 0
  ok <- !is.na(m)
  edges_sf$waterway_km[ok] <- water_by_edge$waterway_km[m[ok]]
}

# dummy só pra facilitar plot / matrizes binárias
edges_sf$waterway_dummy <- as.integer(edges_sf$waterway_km > 0)

message(
  " Hidrovias (aresta): km de hidrovias em buffer de ",
  buf_dist/1000, " km das arestas"
)

}, error = function(e) {
  message("[hidrovias-aresta] ", e$message)
  edges_sf$waterway_km <- 0
  edges_sf$waterway_dummy <- 0
})

```

```

# Atributos de aresta vindos dos nós
if ("rugged_tri" %in% names(nodes_ll))
  edges_sf$rugged_mean <- avg_edge("rugged_tri", nodes_ll, I_jk)
if ("soil_cat_idx" %in% names(nodes_ll))
  edges_sf$soil_cat_idx <- avg_edge("soil_cat_idx", nodes_ll, I_jk)
if ("n_ports" %in% names(nodes_ll))
  edges_sf$n_ports <- avg_edge("n_ports", nodes_ll, I_jk)
if ("n_locks" %in% names(nodes_ll))
  edges_sf$n_locks <- avg_edge("n_locks", nodes_ll, I_jk)
if ("n_dams" %in% names(nodes_ll))
  edges_sf$n_dams <- avg_edge("n_dams", nodes_ll, I_jk)

edge_comp_cols <- c(
  "share_paved", "lanes_mean", "share_fed_state", "share_median",
  "infra_index", "dist_km", "rugged_mean",
  "soil_cat_idx", "waterway_km",
  "n_ports", "n_locks", "n_dams"
)
edge_comp_cols <- edge_comp_cols[edge_comp_cols %in% names(edges_sf)]

edges_list <- edges_sf |>
  st_drop_geometry() |>
  select(id_i, id_j, all_of(edge_comp_cols)) |>
  distinct()

adj_mat <- edges_list |> mutate(val = 1) |> fill_mat("val", ids_micro)
if (!"infra_index" %in% names(edges_list))
  stop("Faltou infra_index em I_jk.")
infra_mat <- fill_mat(edges_list, "infra_index", ids_micro)
dist_mat <- fill_mat(edges_list, "dist_km", ids_micro)

# Aglomeracao por aresta: populacao de i, j e vizinhos
pop_vec <- setNames(as.numeric(nodes_tbl$pop_total), ids_micro)
pop_vec[is.na(pop_vec)] <- 0

neighbors_of <- function(node_id) {
  idx <- match(node_id, ids_micro)
  if (is.na(idx)) return(integer(0))
  ids_micro[which(adj_mat[idx, ] > 0)]
}

edge_agglom_pop <- vapply(
  seq_len(nrow(edges_list)),
  function(k) {
    i_id <- edges_list$id_i[k]; j_id <- edges_list$id_j[k]
    neigh_i <- neighbors_of(i_id)
  }
)

```

```

    neigh_j <- neighbors_of(j_id)
    nodes_use <- unique(c(i_id, j_id, neigh_i, neigh_j))
    sum(pop_vec[match(nodes_use, ids_micro)], na.rm = TRUE)
  },
  numeric(1)
)

edges_list$edge_agglom_pop <- edge_agglom_pop
edge_comp_cols <- unique(c(edge_comp_cols, "edge_agglom_pop"))

edge_keys_list <- paste(edges_list$id_i, edges_list$id_j)
edge_keys_sf <- paste(edges_sf$id_i, edges_sf$id_j)
edges_sf$edge_agglom_pop <- edge_agglom_pop[match(edge_keys_sf, edge_keys_list)]

extra_cols <- setdiff(edge_comp_cols, c("infra_index", "dist_km"))
extra_mats <- lapply(extra_cols, function(v) fill_mat(edges_list, v, ids_micro))
names(extra_mats) <- extra_cols

rugged_mat <- if ("rugged_mean" %in% names(extra_mats))
  extra_mats[["rugged_mean"]] else NULL
edge_agglom_mat <- if ("edge_agglom_pop" %in% names(extra_mats))
  extra_mats[["edge_agglom_pop"]] else NULL

find_first_existing <- function(cands, pool) cands[cands %in% pool][1]
A_solo <- if (!is.na(find_first_existing(c("soil_cat_idx"), names(extra_mats))))
  extra_mats[[find_first_existing(c("soil_cat_idx"), names(extra_mats))]] else NULL
A_hidro <- if (!is.na(find_first_existing(c("waterway_km"), names(extra_mats))))
  extra_mats[[find_first_existing(c("waterway_km"), names(extra_mats))]] else NULL
A_portos <- if (!is.na(find_first_existing(c("n_ports"), names(extra_mats))))
  extra_mats[[find_first_existing(c("n_ports"), names(extra_mats))]] else NULL
A_eclusas <- if (!is.na(find_first_existing(c("n_locks"), names(extra_mats))))
  extra_mats[[find_first_existing(c("n_locks"), names(extra_mats))]] else NULL
A_barragens <- if (!is.na(find_first_existing(c("n_dams"), names(extra_mats))))
  extra_mats[[find_first_existing(c("n_dams"), names(extra_mats))]] else NULL

emis <- rep(0, J)

# -----
# Export .mat
# -----
if (!dir.exists(OUT_DIR)) dir.create(OUT_DIR, recursive = TRUE)
mat_path <- file.path(OUT_DIR, "graph_BR.mat")

```

```

export_list <- list(
  con      = mat_path,
  A        = adj_mat,
  I        = infra_mat,
  df_nodes = df_nodes,
  ids      = as.numeric(ids_micro),
  J        = as.double(J),
  dist_mat = dist_mat,
  emis     = emis
)
if (!is.null(rugged_mat)) {
  export_list$rugged_mat <- rugged_mat
}
if (!is.null(edge_agglom_mat)) {
  export_list$edge_agglom_mat <- edge_agglom_mat
}
if (length(extra_mats)) {
  export_list$edge_mats      <- extra_mats
  export_list$edge_mats_names <- names(extra_mats)
}
if (!is.null(A_solo))      export_list$A_solo      <- A_solo
if (!is.null(A_hidro))     export_list$A_hidro     <- A_hidro
if (!is.null(A_portos))    export_list$A_portos    <- A_portos
if (!is.null(A_eclusas))   export_list$A_eclusas   <- A_eclusas
if (!is.null(A_barragens)) export_list$A_barragens <- A_barragens

export_list$node_pop_total <- as.numeric(nodes_tbl$pop_total)
export_list$node_pib_total <- as.numeric(nodes_tbl$pib_total)

if (!is.null(pib_sector_mat)) {
  export_list$node_pib_sector      <- pib_sector_mat
  export_list$node_pib_sector_names <- pib_sector_cols
}
if (!is.null(pop_sector_mat)) {
  export_list$node_pop_sector      <- pop_sector_mat
  export_list$node_pop_sector_names <- pop_sector_cols
}
if (!is.null(node_airports))
  export_list$node_airports <- node_airports

do.call(writeMat, export_list)
cat("OK: exportado", mat_path, " | J =", J,
  "| #arestas =", sum(adj_mat[upper.tri(adj_mat)]), "\n")

# -----
# Plots
# -----

```

```

edges_plot_ll <- edges_sf |> st_transform(crs_ll)
nodes_plot     <- nodes_tbl |> as.data.frame()

bbox_br <- list(xlim = c(-75, -34), ylim = c(-35, 6))
leg_small <- guides(color = guide_colorbar(title.position = "top",
                                             barheight = unit(18, "mm"),
                                             barwidth  = unit(5, "mm")))
map_theme <- theme_void(base_size = 10) +
  theme(legend.position = "right",
        legend.title  = element_text(size = 9),
        legend.text   = element_text(size = 8),
        plot.title    = element_text(face = "bold", size = 11, hjust = 0),
        plot.margin   = margin(4,4,4,4, "mm"))

edge_plot <- function(var, title, trans = NULL, name = var) {

  # CASO ESPECIAL: hidrovias
  if (var == "waterway_km") {
    e <- edges_plot_ll

    if (!"waterway_km" %in% names(e)) {
      stop("edges_plot_ll não tem coluna 'waterway_km'.")
    }

    # arestas que cruzam algum trecho hidroviário
    e$has_hidro <- e$waterway_km > 0

    return(
      ggplot() +
        geom_sf(data = micro_br, fill = NA, color = "grey85", linewidth = 0.2) +
        # rede completa em cinza claro
        geom_sf(data = e, color = "grey90", linewidth = 0.20, alpha = 0.4) +
        # apenas arestas com hidrovia em azul
        geom_sf(data = e[e$has_hidro, ], color = "#1565C0",
                linewidth = 0.55, alpha = 0.9) +
        geom_point(data = nodes_plot, aes(x = lon, y = lat),
                   color = "black", size = 0.8) +
        coord_sf(xlim = bbox_br$xlim, ylim = bbox_br$ylim, expand = FALSE) +
        map_theme + labs(title = title)
    )
  }

  # CASO GERAL
  sc <- if (is.null(trans)) scale_color_viridis_c(name = name)
         else scale_color_viridis_c(trans = trans, name = name)

  ggplot() +

```

```

  geom_sf(data = micro_br, fill = NA, color = "grey85", linewidth = 0.2) +
  geom_sf(data = edges_plot_ll, aes(color = .data[[var]]),
           linewidth = 0.35, alpha = 0.8) +
  geom_point(data = nodes_plot, aes(x = lon, y = lat),
             color = "black", size = 0.8) +
  sc + leg_small +
  coord_sf(xlim = bbox_br$xlim, ylim = bbox_br$ylim, expand = FALSE) +
  map_theme + labs(title = title)
}

node_plot <- function(var, title, trans = "log10", name = var) {
  if (var == "n_airport_dummy") {
    cols <- c("0" = "#cccccc", "1" = "#1f77b4")
    nodes_plot$._tmp_air_ <- factor(ifelse(nodes_plot[[var]] > 0, 1, 0))
    return(
      ggplot() +
      geom_sf(data = micro_br, fill = NA, color = "grey85", linewidth = 0.2) +
      geom_point(data = nodes_plot,
                 aes(x = lon, y = lat, color = ._tmp_air_),
                 size = 1.4) +
      scale_color_manual(values = cols, name = name,
                         labels = c("0" = "sem aeroporto", "1" = "tem_
aeroporto")) +
      leg_small +
      coord_sf(xlim = bbox_br$xlim, ylim = bbox_br$ylim, expand = FALSE) +
      map_theme + labs(title = title)
    )
  }
  sc <- if (is.null(trans))
    scale_color_viridis_c(labels = label_number(big.mark=".",
                                                decimal.
                                                mark=","),
                           name = name)
  else
    scale_color_viridis_c(trans = trans,
                           labels = label_number(big.mark=".",
                                                 decimal.mark=","),
                           name = name)

  ggplot() +
  geom_sf(data = micro_br, fill = NA, color = "grey85", linewidth = 0.2) +
  geom_point(data = nodes_plot,
             aes(x = lon, y = lat, color = .data[[var]]),
             size = 1.2) +
  sc + leg_small +
  coord_sf(xlim = bbox_br$xlim, ylim = bbox_br$ylim, expand = FALSE) +
  map_theme + labs(title = title)
}

```

```

view_plots <- function(plots, per_page = 4, ncol = 2) {
  n <- length(plots); if (!n) return(invisible(NULL))
  idx <- split(seq_len(n), ceiling(seq_len(n) / per_page))
  for (g in idx) print(wrap_plots(plots[g], ncol = ncol))
  invisible(NULL)
}

plots_edges_main <- lapply(
  intersect(c("infra_index", "dist_km", "rugged_mean"), names(edges_plot_ll)),
  function(v) edge_plot(
    v,
    paste("Arestas ", v),
    if (v == "dist_km") "log10" else NULL,
    if (v == "dist_km") "dist (km)" else v
  )
)

plots_edges_bc <- lapply(
  intersect(c("share_paved", "lanes_mean", "share_fed_state", "share_median"), names(edges_plot_ll)),
  function(v) edge_plot(v, paste("Arestas ", v))
)

plots_edges_env <- lapply(
  intersect(c("soil_cat_idx", "waterway_km", "n_ports", "n_locks", "n_dams"),
            names(edges_plot_ll)),
  function(v) edge_plot(v, paste("Arestas ", v))
)

plots_nodes_tot <- lapply(
  intersect(c("pop_total", "pib_total"), names(nodes_plot)),
  function(v) node_plot(v, paste("Nós ", v))
)

plots_nodes_pib <- lapply(
  intersect(c("pib_agro", "pib_ind", "pib_serv"), names(nodes_plot)),
  function(v) node_plot(v, paste("Nós ", v))
)

plots_nodes_pop <- lapply(
  intersect(c("pop_agro", "pop_ind", "pop_serv"), names(nodes_plot)),
  function(v) node_plot(v, paste("Nós ", v))
)

plots_nodes_air <- if ("n_airport_dummy" %in% names(nodes_plot)) {
  list(node_plot("n_airport_dummy", "Nós aeroportos (dummy)",
                trans = NULL, name = "dummy aeroportos"))
}

```

```

} else list()

message(">>> Plots principais")
view_plots(
  c(plots_edges_main, plots_edges_bc, plots_edges_env,
    plots_nodes_tot, plots_nodes_pib, plots_nodes_pop, plots_nodes_air),
  per_page = 4, ncol = 2
)

```

Ruggedness (ETOPO TRI) por nó

Solo: soils\_brazil\_wrb\_wgs84.shp (indice categ. 0-1)

Eclusas: contagem por microrregião

Portos: contagem por microrregião

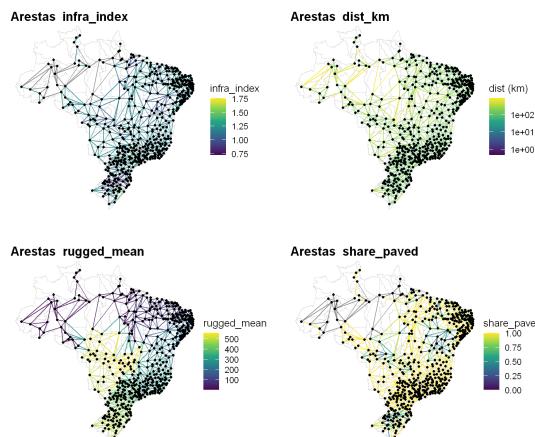
Barragens: contagem por microrregião

Aeroportos: complexos dedup + intersect

Hidrovias (aresta): km de hidrovias em buffer de 20 km das arestas

OK: exportado output/graph\_BR.mat | J = 555 | #arestas = 1441

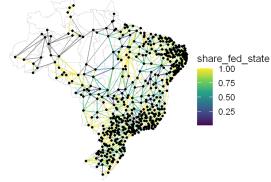
>>> Plots principais



Arestas lanes\_mean



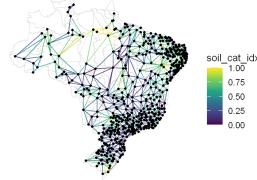
Arestas share\_fed\_state



Arestas share\_median



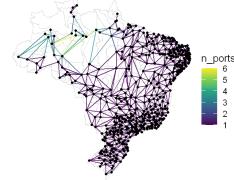
Arestas soil\_cat\_idx



Arestas waterway\_km



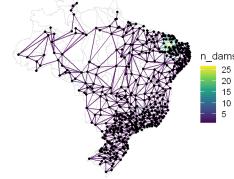
Arestas n\_ports



Arestas n\_locks



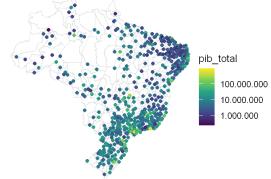
Arestas n\_dams



Nós pop\_total



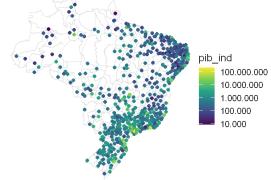
Nós pib\_total

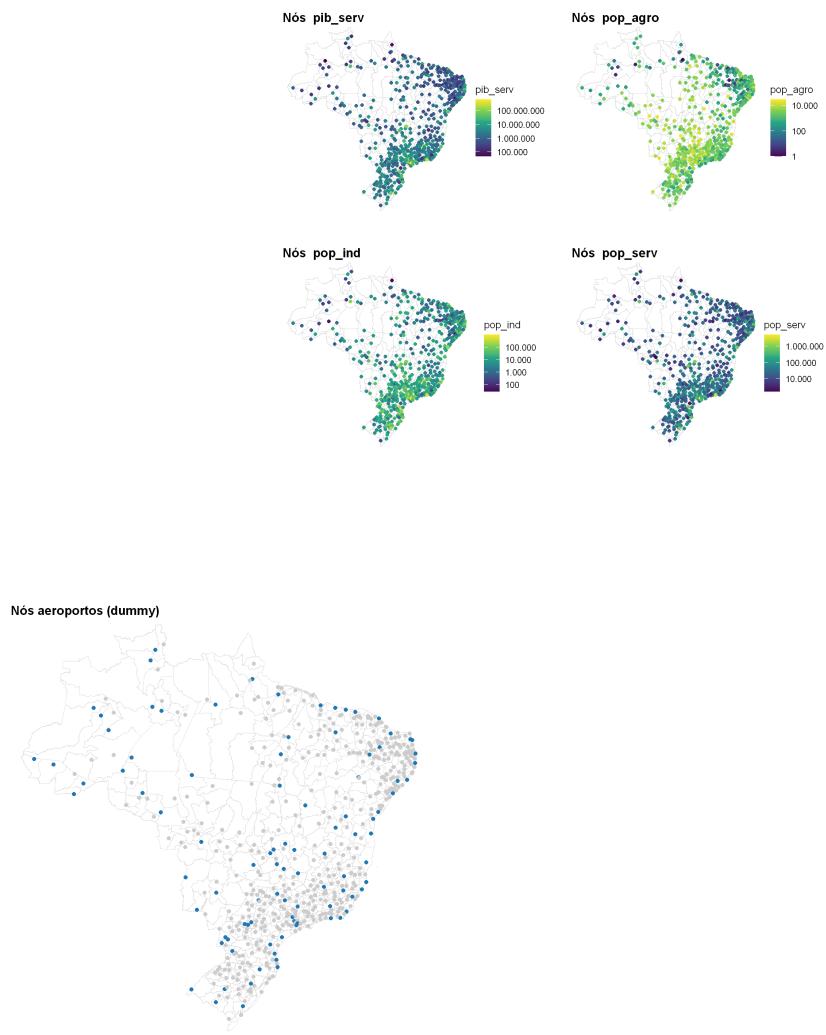


Nós pib\_agro



Nós pib\_ind





## 11 10) Missing Data

### 11.1 10.1) Mapbiomas Coverage

```
[88]: plot_mapbiomas_10m <- function(tif,
                                      aoi_path = NULL,
                                      title = "MapBiomas 10m - Cobertura do Solo",
                                      ↵(Brasil)) {
  r <- rast(tif)

  # Recorte opcional
```

```

if (!is.null(aoi_path) && file.exists(aoi_path)) {
  r <- crop(r, vect(aoi_path), mask = TRUE)
}

# === CORRIGIR FUNDO: 0 → NA ===
vals_raw <- unique(r[])
if (0 %in% vals_raw) {
  r[r == 0] <- NA
}

NAflag(r) <- NA

# Tabela oficial
mapbiomas_classes <- tibble::tribble(
  ~value, ~label, ~color,
  3, "Floresta", "#1A9850",
  4, "Formação Savânica", "#66BD63",
  5, "Savana Parque", "#A6D96A",
  6, "Campinarana / Vegetação Natural", "#D9EF8B",
  9, "Silvicultura", "#7F3C8D",
  11, "Pastagem", "#FDE725",
  12, "Agricultura", "#F46D43",
  13, "Agricultura Irrigada", "#F6C141",
  14, "Mosaico Agricultura e Pasto", "#BF812D",
  15, "Cana-de-açúcar", "#C51B7D",
  19, "Infraestrutura Urbana", "#999999",
  21, "Mineração", "#662506",
  22, "Outras Áreas Não Vegetadas", "#E6E6E6",
  23, "Afloramento Rochoso", "#BEBADA",
  24, "Praia e Duna", "#FFFFBF",
  25, "Outras Áreas Urbanas", "#F1B6DA",
  26, "Área Degradada", "#FDE0EF",
  27, "Solo Exposto", "#FDDBA2",
  29, "Cultivo Perene", "#FDAE61",
  30, "Pastagem Natural", "#A6DDBA",
  31, "Formações Campestres", "#1C9099",
  32, "Apicum", "#D1E5F0",
  33, "Mangue", "#2166AC",
  34, "Água", "#08306B",
  36, "Outras Áreas Aquáticas", "#41B6C4",
  49, "Glaciar", "#B3B3B3",
  50, "Neve", "#FFFFFF"
)

# Quais valores aparecem
vals <- sort(unique(na.omit(r[])))
df <- mapbiomas_classes[mapbiomas_classes$value %in% vals, ]

```

```

col_vec <- df$color
names(col_vec) <- df$value

oldpar <- par(no.readonly = TRUE)
on.exit(par(oldpar))

par(bg = "white", mar = c(3, 3, 2, 12), xaxs = "i", yaxs = "i")

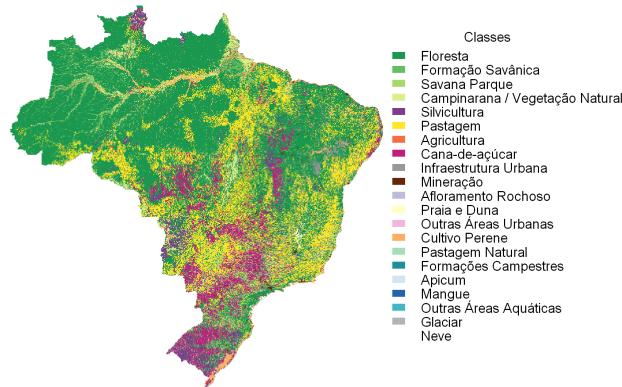
plot(
  r,
  col = col_vec[match(vals, df$value)],
  legend = FALSE,
  axes = FALSE,
  box = FALSE,
  colNA = "white",
  main = title
)

legend(
  "right",
  legend = df$label,
  fill = df$color,
  border = NA,
  cex = 1.0,
  bty = "n",
  xpd = TRUE,
  inset = c(-0.45, 0),
  title = "Classes"
)
}

plot_mapbiomas_10m(
  "C:/Users/Pedro/Desktop/OTNuEC/R/data/mapbiomas_near_1000m.tif"
)

```

MapBiomas 10m — Cobertura do Solo (Brasil)



```
[89]: plot_floresta_10m <- function(tif,
                                    aoi_path = NULL,
                                    title = "Cobertura Florestal - Brasil") {

  r <- rast(tif)

  # Recorte opcional
  if (!is.null(aoi_path) && file.exists(aoi_path)) {
    r <- crop(r, vect(aoi_path), mask = TRUE)
  }

  # Corrigir fundo: 0 → NA (fora do Brasil)
  if (0 %in% unique(r[])) {
    r[r == 0] <- NA
  }
  NAflag(r) <- NA

  # === Classes de floresta MapBiomas 10m ===
  # Seu dataset tem floresta apenas no código 3
  floresta_code <- 3

  # Criar raster com 2 classes:
  # NA → fundo branco (fora BR)
  # 0 → não floresta
  # 1 → floresta
  r2 <- classify(r, rbind(
    c(floresta_code, floresta_code, 1),      # floresta
    c(-Inf, floresta_code - 0.001, 0),        # não floresta
    c(floresta_code + 0.001, Inf, 0)
  ))
}
```

```

# Paleta:
cols <- c(
  "0" = "#E5E5E5",    # cinza claro
  "1" = "#008000"     # verde forte
)

oldpar <- par(no.readonly = TRUE)
on.exit(par(oldpar))

par(bg = "white", mar = c(3,3,2,8), xaxs = "i", yaxs = "i")

plot(
  r2,
  col = cols,
  legend = FALSE,
  axes = FALSE,
  box = FALSE,
  colNA = "white",      # FUNDO BRANCO REAL
  main = title
)

legend(
  "right",
  legend = c("Não Floresta", "Floresta"),
  fill = cols,
  border = NA,
  cex = 1.2,
  bty = "n",
  xpd = TRUE,
  inset = c(-0.15, 0),
  title = "Classes"
)

invisible(r2)
}
plot_floresta_10m(
  "C:/Users/Pedro/Desktop/OTNuEC/R/data/mapbiomas_near_1000m.tif"
)

```

Cobertura Florestal — Brasil



```
[90]: library(terra)
library(dplyr)

resumo_mapbiomas <- function(tif_path) {
  cat("\n===== Lendo raster =====\n")
  r <- rast(tif_path)

  cat("\n===== Metadados =====\n")
  print(r)

  # ---- Valores únicos ----
  vals <- unique(r[])
  vals <- sort(na.omit(as.numeric(vals)))

  cat("\n===== Valores únicos encontrados =====\n")
  print(vals)

  # ---- Pixel resolution ----
  res_m <- res(r)[1]           # tamanho do pixel em metros
  pixel_area_km2 <- (res_m^2) / 1e6

  # ---- Contagem de pixels ----
  cat("\n===== Contagem de pixels por classe =====\n")
  tab <- freq(r, digits = 0)
  tab <- tab[!is.na(tab$value), ]  # remove NA
  print(tab)

  # ---- Área por classe ----
  tab <- tab %>%
    mutate(
```

```

  area_km2 = count * pixel_area_km2,
  perc = 100 * area_km2 / sum(area_km2)
) %>%
arrange(desc(area_km2))

cat("\n==== Área por classe (km²) ====\n")
print(tab)

return(tab)
}
resumo <- resumo_mapbiomas(
  "C:/Users/Pedro/Desktop/OTNuEC/R/data/mapbiomas_near_1000m.tif"
)

```

===== Lendo raster =====

===== Metadados =====

```

class      : SpatRaster
size       : 4757, 4530, 1 (nrow, ncol, nlyr)
resolution : 0.009235355, 0.008996598 (x, y)
extent     : -74.8974, -33.06124, -34.85203, 7.94479 (xmin, xmax, ymin, ymax)
coord. ref. : lon/lat WGS 84 (EPSG:4326)
source     : mapbiomas_near_1000m.tif
name       : mapbiomas_10m_collection2_integration_v1-classification_2023
min value  :                               0
max value  :                               50

```

===== Valores únicos encontrados =====

```
[1] 0 3 4 5 6 9 11 12 15 19 21 23 24 25 29 30 31 32 33 36 49 50
```

===== Contagem de pixels por classe =====

layer	value	count
1	1	0 12950531
2	1	3 3504377
3	1	4 1073736
4	1	5 10059
5	1	6 347095
6	1	9 114506
7	1	11 254644
8	1	12 257428
9	1	15 1650241
10	1	19 600767
11	1	21 429846
12	1	23 4845
13	1	24 42215
14	1	25 49936
15	1	29 18655

```

16      1    30      3618
17      1    31      665
18      1    32      533
19      1    33    210844
20      1    36     15270
21      1    49     6720
22      1    50     2679

===== Área por classe (km2) =====
  layer value     count      area_km2        perc
1       1      0 12950531 1.104574e-03 60.097474571
2       1      3 3504377 2.988946e-04 16.262206364
3       1     15 1650241 1.407520e-04 7.658011593
4       1      4 1073736 9.158086e-05 4.982716304
5       1     19 600767 5.124049e-05 2.787884103
6       1     21 429846 3.666233e-05 1.994718136
7       1      6 347095 2.960435e-05 1.610708699
8       1     12 257428 2.195649e-05 1.194605278
9       1     11 254644 2.171904e-05 1.181686011
10      1     33 210844 1.798326e-05 0.978430300
11      1      9 114506 9.766422e-06 0.531369827
12      1     25 49936 4.259131e-06 0.231730073
13      1     24 42215 3.600593e-06 0.195900453
14      1     29 18655 1.591118e-06 0.086569299
15      1     36 15270 1.302406e-06 0.070861066
16      1      5 10059 8.579501e-07 0.046679205
17      1     49  6720 5.731608e-07 0.031184438
18      1     23  4845 4.132387e-07 0.022483423
19      1     30  3618 3.085857e-07 0.016789479
20      1     50  2679 2.284967e-07 0.012432010
21      1     31   665 5.671904e-08 0.003085960
22      1     32   533 4.546052e-08 0.002473409

```

## 11.2 10.2) IBGE vulnerability

```

[91]: library(terra)
plot_erosao <- function(path) {
  r <- rast(path)

  pal <- hcl.colors(30, "YlOrRd", rev = TRUE)

  plot(
    r,
    col = pal,
    main = "Vulnerabilidade à Erosão - IBGE (média ~1km)",
    axes = FALSE,
    box = FALSE
  )
}

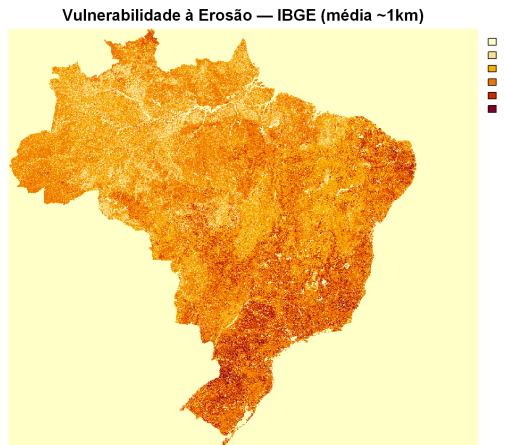
```

```

    )
}

plot_erosao("data/erosao_mean_1km.tif")

```



### 11.3 10.3) Embrapa soil

```

[92]: library(sf)
library(ggplot2)
library(dplyr)

plot_solos_wrb <- function(shp_path) {

  # 1. Ler shapefile
  soils <- sf::st_read(shp_path, quiet = TRUE)

  # 2. Remover coluna de geometria para inspecionar atributos
  attrs <- soils |> st_drop_geometry()

  # 3. Identificar colunas categóricas (factor ou character)
  cat_cols <- names(attrs)[sapply(attrs, \x) is.factor(x) || is.character(x))]

  if (length(cat_cols) == 0)
    stop("Nenhuma coluna categórica encontrada no shapefile!")

  # Usar a primeira coluna categórica
  col_to_plot <- cat_cols[1]

  # 4. Converter para fator
  soils[[col_to_plot]] <- factor(soils[[col_to_plot]])
}

```

```

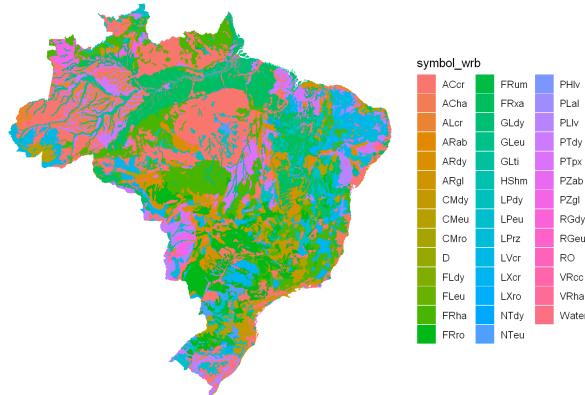
# 5. Criar paleta
n <- nlevels(soils[[col_to_plot]])
pal <- scales::hue_pal()(n)

# 6. Plot bonito estilo MapBiomass
ggplot(soils) +
  geom_sf(aes(fill = .data[[col_to_plot]]), color = NA) +
  scale_fill_manual(values = pal, name = col_to_plot) +
  labs(
    title = "Solos - WRB (Brasil)",
    subtitle = "Cada cor representa um tipo distinto de solo segundo a classificação WRB"
  ) +
  theme_void() +
  theme(
    legend.position = "right",
    plot.title = element_text(size = 20, face = "bold"),
    plot.subtitle = element_text(size = 14)
  )
}

shp_path <- "soils_brazil_wrb_wgs84.shp"
plot_solos_wrb(shp_path)

```

**Solos - WRB (Brasil)**  
Cada cor representa um tipo distinto de solo segundo a classificação WRB



## 12 11) Appendix

### 12.1 11.1) Mapbiomas compact

Sanity on full data

```
[93]: # file: R/sanity_mapbiomas_fix.R
suppressPackageStartupMessages(library(terra))

# --- helpers (por que: compatibilidade e precisão métrica) ---
is_ll <- function(x) isTRUE(tryCatch(is.lonlat(x), error = function(e) NA))

dtype_terra <- function(x) {
  # terra >= 1.5 usa `datatype()`. Fallback tenta `dataType()`.
  out <- tryCatch(datatype(x), error = function(e) NA_character_)
  if (is.na(out)) out <- tryCatch(dataType(x), error = function(e) NA_character_)
  out
}

epsg_code <- function(r) {
  d <- tryCatch(crs(r, describe = TRUE), error = function(e) NULL)
  if (is.null(d)) return(NA_character_)
  if (!is.null(d$code)) return(as.character(d$code))
  NA_character_
}

res_meters <- function(r) {
  rs <- res(r)
  if (!is_ll(r)) return(setNames(rs, c("x_m", "y_m")))
  yc <- (ymin(r) + ymax(r)) / 2
  x_m <- rs[1] * 111320 * cos(yc * pi/180)
  y_m <- rs[2] * 111132
  setNames(c(x_m, y_m), c("x_m", "y_m"))
}

pretty_bytes <- function(b) {
  if (is.na(b)) return(NA_character_)
  u <- c("B", "KB", "MB", "GB", "TB"); i <- 1
  while (b >= 1024 && i < length(u)) { b <- b/1024; i <- i+1 }
  sprintf("%.3f %s", b, u[i])
}

categorical_guess <- function(dtype, sample_vals) {
  sample_vals <- sample_vals[!is.na(sample_vals)]
  is_int_dtype <- grepl("(INT|BYTE)", dtype, ignore.case = TRUE)
  all_integer <- length(sample_vals) > 0 &&
    all(abs(sample_vals - round(sample_vals)) < .Machine$double.eps)
  nuniq <- length(unique(sample_vals))
  list(is_categorical = (is_int_dtype || (all_integer && nuniq <= 512)),
       unique_in_sample = nuniq)
}
```

```

suggest_output_dtype <- function(vals, categorical) {
  vals <- vals[!is.na(vals)]
  if (!length(vals)) return(if (categorical) "INT2U" else "FLT4S")
  vmin <- suppressWarnings(min(vals))
  vmax <- suppressWarnings(max(vals))
  if (categorical) {
    if (vmin >= 0 && vmax <= 255) return("INT1U")
    if (vmin >= 0 && vmax <= 65535) return("INT2U")
    return("INT4U")
  }
  "FLT4S"
}

agg_table_for_targets <- function(r, targets_m) {
  rm <- res_meters(r)
  fx <- pmax(1L, as.integer(round(targets_m / rm["x_m"])))
  fy <- pmax(1L, as.integer(round(targets_m / rm["y_m"])))
  data.frame(
    alvo_m    = as.integer(targets_m),
    fator_x  = as.integer(fx),
    fator_y  = as.integer(fy),
    fator     = as.integer(pmax(fx, fy)),
    ncol_out = as.integer(ceiling(ncol(r) / fx)),
    nrow_out = as.integer(ceiling(nrow(r) / fy)),
    outsize_x_percent = round(100/fx, 3),
    outsize_y_percent = round(100/fy, 3),
    stringsAsFactors = FALSE
  )
}

# --- SANITY principal ---
#' Diagnóstico rápido de um raster grande (sem varrer tudo)
#' @param path caminho do GeoTIFF
#' @param sample_n número de pixels para amostrar (rápido)
#' @param targets_m resoluções alvo em metros (vetor)
#' @param seed semente para reproduzibilidade
#' @return lista invisível com todos os resultados
raster_sanity <- function(path,
                           sample_n = 200000L,
                           targets_m = c(300L, 500L, 1000L),
                           seed = 1L) {
  stopifnot(file.exists(path))
  r <- rast(path)

  finfo <- file.info(path)
  rm <- res_meters(r)
  dtype <- dtype_terra(r)
}

```

```

epsg  <- epsg_code(r)

# amostra rápida (1ª banda)
set.seed(seed)
v <- tryCatch({
  vv <- spatSample(r, size = sample_n, method = "random",
                    values = TRUE, na.rm = FALSE, warn = FALSE)
  # pega a primeira coluna de valores, qualquer que seja o nome
  as.numeric(vv[[1]])
}, error = function(e) numeric())

na_pct   <- if (length(v)) round(mean(is.na(v))*100, 3) else NA_real_
v_no_na <- v[!is.na(v)]
top_tbl  <- if (length(v_no_na)) sort(table(v_no_na), decreasing = TRUE) else
  ↪integer()
guess    <- categorical_guess(dtype, v)
outdtype <- suggest_output_dtype(v, guess$is_categorical)
aggs     <- agg_table_for_targets(r, unique(as.integer(targets_m)))

cat("\n===== SANITY =====\n")
cat("Arquivo: ", normalizePath(path, winslash="/"), "\n", sep = "")
cat("Tamanho: ", pretty_bytes(finfo$size), "\n", sep = "")
cat(sprintf("Camadas: %d | Dims: %d x %d (%.1f M células)\n",
            nlyr(r), ncol(r), nrow(r), ncell(r)/1e6))
cat(sprintf("CRS definido: %s | EPSG: %s | lon/lat: %s\n",
            !is.na(crs(r)), ifelse(is.na(epsg), "NA", epsg), is_ll(r)))
cat(sprintf("Res (CRS): %s\n", paste(signif(res(r), 8), collapse=" x ")))
cat(sprintf("Res aprox (m): x=%,.2f y=%,.2f | lat_centro=%,.5f\n",
            rm["x_m"], rm["y_m"], (ymin(r)+ymax(r))/2))
cat(sprintf("Datatype: %s | Tem levels?: %s\n",
            ifelse(is.na(dtype), "NA", dtype), any(lengths(levels(r)) > 0)))
if (length(v)) {
  cat(sprintf("Amostra: %d valores | NA: %.3f% | min/max(amostra)= %s /"
  ↪%s\n",
              length(v), na_pct,
              ifelse(length(v_no_na), as.character(min(v_no_na)), "NA"),
              ifelse(length(v_no_na), as.character(max(v_no_na)), "NA")))
  cat(sprintf("Categórico (guess): %s | únicos na amostra: %d\n",
              guess$is_categorical, guess$unique_in_sample))
  cat("\nTop 15 códigos (amostra):\n"); print(head(top_tbl, 15))
  cat(sprintf("\nSugestão datatype saída: %s (use LZW + TILED)\n", outdtype))
} else {
  cat("Amostra indisponível (segundo só com metadados).\n")
}
cat("\nAlvos e fatores sugeridos:\n"); print(aggs)
cat("Obs: use -outsize X% Y% do GDAL com outsize_x_percent /"
  ↪outsize_y_percent.\n")

```

```

gt <- Sys.which("gdal_translate"); gw <- Sys.which("gdalwarp")
cat(sprintf("\nGDAL: gdal_translate=%s | gdalwarp=%s\n",
           ifelse(nzchar(gt), "SIM", "NÃO"), ifelse(nzchar(gw), "SIM", "NÃO")))
cat("=====\\n\\n")

invisible(list(
  path = normalizePath(path, winslash="/"),
  size_bytes = finfo$size,
  dims = c(ncol=ncol(r), nrow=nrow(r), nlyr=nlyr(r), ncells=ncell(r)),
  crs = as.character(crs(r)),
  epsg = epsg,
  lonlat = is_ll(r),
  res_crs = res(r),
  res_m = unname(rm),
  dtype = dtype,
  sample = list(n=length(v), na_pct=na_pct,
                min=if(length(v_no_na)) min(v_no_na) else NA_real_,
                max=if(length(v_no_na)) max(v_no_na) else NA_real_,
                top=top_tbl),
  categorical_guess = guess,
  out_dtype_suggest = outdtype,
  targets = aggs,
  gdal = list(gdal_translate = nzchar(gt), gdalwarp = nzchar(gw)))
))
}

# ---- exemplo de uso ----
sanity <- raster_sanity(
  "mapbiomas_10m_collection2_integration_v1-classification_2023.tif",
  sample_n = 200000L,
  targets_m = c(300L, 500L, 1000L)
)

```

===== SANITY =====

Arquivo: C:/Users/Pedro/Desktop/OTNuEC/R/mapbiomas\_10m\_collection2\_integration\_v1-classification\_2023.tif

Tamanho: 5.122 GB

Camadas: 1 | Dims: 465718 x 476412 (221873.6 M células)

CRS definido: TRUE | EPSG: 4326 | lon/lat: TRUE

Res (CRS): 8.9831528e-05 x 8.9831528e-05

Res aprox (m): x=9.73 y=9.98 | lat\_centro=-13.45362

Datatype: INT1U | Tem levels?: TRUE

Amostra: 200000 valores | NA: 0.000% | min/max(amostra)= 0 / 50

Categórico (guess): TRUE | únicos na amostra: 22

Top 15 códigos (amostra):

v\_no\_na

0	3	15	4	19	21	6	12	11	33	9
118144	33925	15501	10382	5513	3984	3302	2418	2406	1931	1037
25	24	29	36							
513	355	173	152							

Sugestão datatype saída: INT1U (use LZW + TILED)

Alvos e fatores sugeridos:

	alvo_m	fator_x	fator_y	fator	ncol_out	nrow_out	outsize_x_percent
1	300	31	30	31	15024	15881	3.226
2	500	51	50	51	9132	9529	1.961
3	1000	103	100	103	4522	4765	0.971

	outsize_y_percent
1	3.333
2	2.000
3	1.000

Obs: use -outsize X% Y% do GDAL com outsize\_x\_percent / outsize\_y\_percent.

GDAL: gdal\_translate=NÃO | gdalwarp=NÃO

=====

Compressing the data:

```
[94]: # file: R/compactar_mapbiomas_hyperfast_fix2.R
#suppressPackageStartupMessages(library(terra))

# ---- utils ----
#is_ll <- function(x) isTRUE(tryCatch(is.lonlat(x), error = function(e) NA))
#deg_from_m <- function(r, target_m) {
#  yc <- (ymin(r) + ymax(r)) / 2
#  c(deg_x = target_m / (111320 * cos(yc * pi/180)),
#    deg_y = target_m / 111132)
#}
#make_template <- function(r, deg_x, deg_y) {
#  ex <- ext(r)
#  ncol_out <- as.integer(ceiling((xmax(ex) - xmin(ex)) / deg_x))
#  nrow_out <- as.integer(ceiling((ymax(ex) - ymin(ex)) / deg_y))
#  rast(xmin = xmin(ex), xmax = xmax(ex), ymin = ymin(ex), ymax = ymax(ex),
#        ncol = ncol_out, nrow = nrow_out, crs = crs(r))
#}
#locate_gdal <- function() {
#  first_ok <- function(v) { v <- v[file.exists(v) & nzchar(v)]; if (length(v) <
#    normalizePath(v[1], winslash="/") else "" }
#  t_candidates <- c(Sys.which("gdal_translate"),
#                    Sys.glob("C:/OSGeo4W*/bin/gdal_translate.exe"),
#                    Sys.glob("C:/Program Files/QGIS*/bin/gdal_translate.exe"))
#  a_candidates <- c(Sys.which("gdaladdo"),
```

```

#           Sys.glob("C:/OSGeo4W*/bin/gdaladdo.exe"),
#           Sys.glob("C:/Program Files/QGIS*/bin/gdaladdo.exe"))
# list(translate = first_ok(t_candidates), addo = first_ok(a_candidates))
#}
#wopt_int2 <- list(
#  datatype = "INT2U", # por quê: seus códigos >255
#  gdal = c("COMPRESS=LZW", "PREDICTOR=2", "TILED=YES",
#          "BLOCKXSIZE=512", "BLOCKYSIZE=512", "BIGTIFF=IF_SAVER")
#)

# ---- principal ----
#compactar_fast <- function(in_path, out_path, target_m = 1000L, aoi_path = "") {
#  stopifnot(file.exists(in_path))
#  dir.create(dirname(out_path), recursive = TRUE, showWarnings = FALSE)
#  terraOptions(memfrac = 0.9, todisk = TRUE)

#  r0 <- rast(in_path)
#  if (!is_ll(r0)) stop("Este runner assume WGS84 (graus).")
#  if (nzchar(aoi_path) & file.exists(aoi_path)) r0 <- crop(r0, vect(aoi_path), mask = TRUE)

#  td <- deg_from_m(r0, target_m)
#  ex <- ext(r0)
#  ncol_out <- ceiling((xmax(ex) - xmin(ex)) / td["deg_x"])
#  nrow_out <- ceiling((ymax(ex) - ymin(ex)) / td["deg_y"])
#  outsize_x <- (ncol_out / ncol(r0)) * 100
#  outsize_y <- (nrow_out / nrow(r0)) * 100

#  g <- locate_gdal()
#  if (nzchar(g$translate)) {
#    message(sprintf("GDAL: gdal_translate (nearest) → outsize %.6f%% x %.6f%%",
#                  outsize_x, outsize_y))
#    if (nzchar(g$addo)) {
#      ovs <- c(2,4,8,16,32,64,128,256)
#      try(system2(g$addo, c("-r", "nearest", shQuote(in_path), paste(ovs)), stdout = "", stderr = ""),
#           silent = TRUE)
#    }
#    args <- c("-r", "nearest",
#              sprintf("%.6f%%", outsize_x), sprintf("%.6f%%", outsize_y),
#              "-co", "COMPRESS=LZW", "-co", "TILED=YES", "-co", "BIGTIFF=IF_SAVER",
#              shQuote(in_path), shQuote(out_path))
#    st <- system2(g$translate, args, stdout = "", stderr = "")
#    if (!identical(st, 0L)) stop("gdal_translate falhou.")
#    return(normalizePath(out_path, winslash="/"))
#  }
#}

```

```

# message("GDAL não encontrado: usando terra::resample('near')")
# tmpl <- make_template(r0, td["deg_x"], td["deg_y"])
# # IMPORTANTE: escrever apenas uma vez (evita 'source and target filename
# ↪cannot be the same')
# resample(r0, tmpl, method = "near",
#           filename = out_path, overwrite = TRUE, wopt = wopt_int2)
# normalizePath(out_path, winslash="/")
#}

# ---- exemplo ----
# out <- compactar_fast(
#   in_path = "C:/Users/Pedro/Desktop/OTNuEC/R/
# ↪mapbiomas_10m_collection2_integration_v1-classification_2023.tif",
#   out_path = "data/mapbiomas_near_1000m.tif",
#   target_m = 1000L
# )
# cat("Saída:", out, "\n")

```

## 12.2 11.2) Erosao IBGE compact

```

[95]: #library(terra)
#compactar_erosao <- function(in_path, out_path, alvo_m = 1000) {
#
#   r <- rast(in_path)
#
#   # Supondo que está em graus (WGS84) como no IBGE
#   if (is.lonlat(r)) {
#     # converter graus para metros
#     yc <- (ymin(r) + ymax(r))/2
#     px_m <- 111320 * cos(yc*pi/180)
#   } else {
#     px_m <- res(r)[1]
#   }
#
#   fact <- max(1, round(alvo_m / px_m))

#   cat("Res original (m):", px_m, "| Fator:", fact, "\n")

#   r_agg <- aggregate(
#     r,
#     fact = fact,
#     fun = mean,
#     na.rm = TRUE,
#     filename = out_path,
#     overwrite = TRUE,
#     wopt = list(

```

```

#           datatype = "FLT4S",
#           gdal = c("COMPRESS=LZW", "PREDICTOR=2", "TILED=YES")
#       )
#   )
#
#   return(r_agg)
#}

#eros_simplif <- compactar_erosao(
#  "Brasil_vulnerabilidade_2020/BR_vulnerabilidade_solos_erosao_2020.tif",
#  "data/erosao_mean_1km.tif",
#  alvo_m = 1000
#)

```

```

[96]: sanity_raster <- function(path, sample_n = 100000, seed = 123) {
  r <- rast(path)

  cat("\n===== SANITY -", basename(path), "=====\\n")
  cat("Tamanho:", round(file.info(path)$size/1024^2, 2), "MB\\n")
  cat("Dim:", ncol(r), "x", nrow(r), "pixels\\n")
  cat("Res:", paste(res(r), collapse=" x "), "(graus)\\n")

  # resolução aproximada em metros
  if (is.lonlat(r)) {
    yc <- (ymin(r) + ymax(r))/2
    res_m <- c(
      res(r)[1] * 111320 * cos(yc*pi/180),
      res(r)[2] * 111132
    )
  } else {
    res_m <- res(r)
  }

  cat("Res (m):", round(res_m, 2), "\\n")
  cat("CRS:", crs(r), "\\n")
  cat("Extent:", as.vector(ext(r)), "\\n\\n")

  # Amostra -----
  set.seed(seed)
  v <- spatSample(r, size = sample_n, method = "random", values = TRUE)[,1]

  cat("Amostra:", length(v), "pixels\\n")
  cat(sprintf("NA: %.3f%%\\n", mean(is.na(v)) * 100))

  v2 <- v[!is.na(v)]
  cat("Min-Max:", min(v2), "->", max(v2), "\\n")
  cat("Média:", mean(v2), "\\n")
}

```

```

cat("Desvio-padrão:", sd(v2), "\n")

# Histograma textual
h <- hist(v2, plot = FALSE, breaks = 20)
cat("\nHistograma (20 classes):\n")
print(data.frame(mid = h$mid, freq = h$counts))

cat("\n=====\\n\\n")
}

sanity_raster("data/erosao_mean_1km.tif")

```

```

===== SANITY - erosao_mean_1km.tif =====
Tamanho: 3159.8 MB
Dim: 167557 x 144805 pixels
Res: 0.000269494585235856 x 0.000269494585235856 (graus)
Res (m): 29.08 29.95
CRS: GEOGCRS["WGS 84",
  ENSEMBLE["World Geodetic System 1984 ensemble",
    MEMBER["World Geodetic System 1984 (Transit)"],
    MEMBER["World Geodetic System 1984 (G730)"],
    MEMBER["World Geodetic System 1984 (G873)"],
    MEMBER["World Geodetic System 1984 (G1150)"],
    MEMBER["World Geodetic System 1984 (G1674)"],
    MEMBER["World Geodetic System 1984 (G1762)"],
    MEMBER["World Geodetic System 1984 (G2139)"],
    MEMBER["World Geodetic System 1984 (G2296)"],
    ELLIPSOID["WGS 84",6378137,298.257223563,
      LENGTHUNIT["metre",1]],
    ENSEMBLEACCURACY[2.0]],
  PRIMEM["Greenwich",0,
    ANGLEUNIT["degree",0.0174532925199433]],
  CS[ellipsoidal,2,
    AXIS["geodetic latitude (Lat)",north,
      ORDER[1],
      ANGLEUNIT["degree",0.0174532925199433]],
    AXIS["geodetic longitude (Lon)",east,
      ORDER[2],
      ANGLEUNIT["degree",0.0174532925199433]],
  USAGE[
    SCOPE["Horizontal component of 3D system."],
    AREA["World."],
    BBOX[-90,-180,90,180]],
  ID["EPSG",4326]]
Extent: -73.99109 -28.83538 -33.75177 5.272392

Amostra: 100000 pixels
NA: 0.000%

```

Min-Max: 0 → 5

Média: 0.97177

Desvio-padrão: 1.345151

Histograma (20 classes):

	mid	freq
1	0.1	61294
2	0.3	0
3	0.5	0
4	0.7	0
5	0.9	4535
6	1.1	0
7	1.3	0
8	1.5	0
9	1.7	0
10	1.9	15311
11	2.1	0
12	2.3	0
13	2.5	0
14	2.7	0
15	2.9	14308
16	3.1	0
17	3.3	0
18	3.5	0
19	3.7	0
20	3.9	3664
21	4.1	0
22	4.3	0
23	4.5	0
24	4.7	0
25	4.9	888

=====