

# Package ‘CropRotationViz’

February 11, 2025

**Title** Interactive Tool for Crop Rotation Sequence Analysis and Visualization

**Version** 0.0.1

**Description** An interactive Shiny application for analyzing and visualizing crop rotation sequences from agricultural field data. The package provides tools for processing field geometries, analyzing crop rotations across multiple years, and creating interactive visualizations. Key features include field intersection analysis, support for multiple spatial file formats (SHP, GeoJSON, FlatGeobuf, GeoPackage), and customizable visualization options. Designed for agricultural researchers and practitioners working with spatiotemporal crop rotation data.

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**URL** <https://github.com/franz-geoeco/CropRotationViz>

**BugReports** <https://github.com/franz-geoeco/CropRotationViz/issues>

**Depends** R (>= 3.5.0)

**Imports** biscale,  
cowplot,  
dplyr,  
data.table,  
forcats,  
htmltools,  
plyr,  
purrr,  
stringr,  
tidyr,  
shiny,  
sf (>= 1.0.12),  
leaflet (>= 2.1.2),  
ggplot2,  
plotly (>= 4.10.1),  
data.table,  
DT,  
ggalluvial,  
rmapshaper,  
shinyBS,  
shinyFiles,  
shinyWidgets,  
shinyalert,  
shinycssloaders,

shinythemes,  
sortable,  
units

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knitr (>= 1.42),  
rmarkdown (>= 2.21),  
biscake,  
DiagrammeR,  
furry,  
future,  
geodata,  
ggridges,  
ggmap,  
ggpubr,  
ggrepel (>= 0.9.3),  
ggridges,  
gridExtra (>= 2.3),  
grDevices,  
lattice (>= 0.20.45),  
leaflet.minicharts (>= 0.6.2),  
osmdata (>= 0.2.3),  
progress (>= 1.2.2),  
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rworldmap,  
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tidyverse,  
tools

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**Arguments**

intersected	sf object containing intersected fields
years	Numeric vector of years to process
crop_codes	crop codes for aggregaton
display_names	names corresponding to the codes

**Value**

sf object with added aggregated columns

---

clean_thin_polygons	<i>Clean Thin Polygons</i>
---------------------	----------------------------

---

**Description**

Clean Thin Polygons

**Usage**

```
clean_thin_polygons(sf_data, min_width = 1)
```

**Arguments**

sf_data	SF object containing polygons
min_width	Minimum width in meters

**Value**

SF object with cleaned polygons

---

create_crop_rotation_sankey	<i>Create Crop Rotation Sankey Diagram</i>
-----------------------------	--

---

**Description**

Creates a Sankey diagram visualizing crop rotation patterns

**Usage**

```
create_crop_rotation_sankey(
  data,
  min_area = 1,
  exclude_crops = c("grassland", "forest", "flowering area", "fallow", "fruits",
    "permanent/tree", "Meerettich"),
  output_path = NULL,
  width = 14,
  height = 10,
  resolution = 200,
  color = NULL
)
```

**Arguments**

data	sf object containing crop rotation data with Aggregated_* columns for each year
min_area	Minimum area threshold in hectares (default: 1)
exclude_crops	Character vector of crop types to exclude
output_path	Optional file path for saving the plot
width	Plot width in inches (default: 14)
height	Plot height in inches (default: 10)
resolution	Plot resolution in dpi (default: 200)

**Details**

The function creates a Sankey diagram showing crop rotation patterns over time. Area values are converted from square meters to hectares in the visualization.

**Value**

A ggplot object containing the Sankey diagram visualizing crop rotations over time

**Examples**

```
## Not run:
library(sf)
# Assuming 'crop_data' is an sf object with Aggregated_* columns
plot <- create_crop_rotation_sankey(
  data = crop_data,
  min_area = 1,
  exclude_crops = c("grassland", "forest")
)

## End(Not run)
```

---

create\_multi\_year\_donut

*Create a Multi-Year Donut Chart with Equal-Width Rings*

---

**Description**

This function generates a multi-year donut chart where each year is represented as a concentric ring. All rings have equal width and are spaced evenly. Additionally, one specific year can be highlighted with a different border color.

**Usage**

```
create_multi_year_donut(
  data,
  year_columns,
  title = "Crop Distribution",
  highlight_year = NULL,
  colors
)
```

Arguments

data	A dataframe containing frequency values for each year and crop category.
year_columns	A vector of column names (as strings) from the data dataframe that represent years.
title	A string representing the title of the chart (default is "Crop Distribution").
highlight_year	A string specifying which year should be highlighted with a different border color. If NULL, no year is highlighted.

Value

A plotly donut chart with multiple concentric rings representing different years.

Examples

```
# Example usage of the function
create_multi_year_donut(data = my_data,
                        year_columns = c("2018", "2019", "2020", "2021"),
                        highlight_year = "2020")
```

---

diversity_mapper	Create a Bivariate Choropleth Map for Agricultural Diversity
------------------	--

---

Description

This function generates an interactive leaflet map visualizing agricultural diversity patterns using a bivariate color scheme. It displays the relationship between unique crops and crop transitions across different geographical units (Districts or River Basins).

Usage

```
diversity_mapper(data, type)
```

Arguments

data	A list containing the following elements: <ul style="list-style-type: none"><li>• BISCALE: Spatial data frame with diversity metrics</li><li>• color_pal: Color palette function for bivariate mapping</li><li>• labels1: List with bi_x and bi_y labels for the legend</li></ul>
type	Character string indicating the geographic unit type ("District" or "River Basin")

Value

- A leaflet map object with:
- Choropleth layer showing bivariate relationships
  - Custom legend displaying the bivariate color scheme
  - Popups with detailed information for each geographic unit

Examples

```
## Not run:
diversity_mapper(processed_data, type = "District")
diversity_mapper(processed_data, type = "River Basin")

## End(Not run)
```

---

diversity_mapping	Create Bivariate Maps for Agricultural Diversity Analysis
-------------------	---

---

Description

Generate bivariate maps showing the relationship between crop diversity (unique crop count) and crop rotation transitions at district and river basin levels.

Usage

```
diversity_mapping(input, agg_cols, districts, EZGs = NA, AOIs = NA)
```

Arguments

input	An sf object containing agricultural field data with crop information over multiple years
agg_cols	Character vector specifying the column names containing crop information for each year
districts	sf object containing district-level polygons with at least "District" and "geometry" columns
EZGs	Optional sf object containing river basin (EZG) polygons. Default NA

Details

This function creates bivariate choropleth maps showing the relationship between two agricultural diversity metrics:

1. Number of unique crops grown in each spatial unit
2. Number of crop transitions (crop changes between consecutive years)

The metrics are calculated first at field level, then aggregated to district and river basin level using both simple means and area-weighted means. The resulting maps use a 3x3 bivariate color scheme to show the relationship between these metrics.

Value

List containing two leaflet map objects:

- District\_Map: Bivariate choropleth map at district level
- EZG\_Map: Bivariate choropleth map at river basin level (if EZGs provided)

## Examples

```
## Not run:
# Create maps for districts only
maps <- diversity_mapping(
  input = crop_data,
  agg_cols = c("crop_2020", "crop_2021", "crop_2022"),
  districts = district_polygons
)

# Create maps for both districts and river basins
maps <- diversity_mapping(
  input = crop_data,
  agg_cols = c("Name_2020", "Name_2021", "Name_2022"),
  districts = district_polygons,
  EZGs = river_basin_polygons
)

## End(Not run)
```

---

dummy\_field\_creator      *Create Dummy Agricultural Fields*

---

## Description

Generate a set of dummy agricultural fields over multiple years in a region of your choice.

## Usage

```
dummy_field_creator(
  output_dir,
  base_location = c(2.3, 49.2),
  field_count = 100,
  years = 2020:2023,
  min_field_size = 0.003,
  max_field_size = 0.005
)
```

## Arguments

output_dir	Character string specifying the directory to save shapefiles
base_location	Numeric vector of length 2 with longitude and latitude coordinates. Default c(2.3, 49.2) for Northern France
field_count	Integer specifying number of fields to generate. Default 100
years	Numeric vector specifying years to generate data for. Default 2020:2023
min_field_size	Numeric value for minimum field size in degrees. Default 0.0003
max_field_size	Numeric value for maximum field size in degrees. Default 0.0005

## Details

Creates a set of irregular agricultural fields with crop rotations and NC codes. Fields are generated around a specified base location with realistic sizes for European agriculture.



**Value**

List of sf objects, one for each year

---

extract_aggregation	<i>Extract Aggregation Status from Processing Summary</i>
---------------------	---

---

**Description**

This function extracts the aggregation status from a processing summary text. It searches for a line beginning with "Aggregation:" and returns its value. The function assumes the text contains a properly formatted Options section with an Aggregation field.

**Usage**

```
extract_aggregation(text)
```

**Arguments**

text	Character string containing the processing summary text
------	---

**Value**

Character string indicating the aggregation status ("Yes" or "No")

**Examples**

```
text <- "Options\nAggregation: Yes\nIntersection: complete"
extract_aggregation(text) # Returns "Yes"
```

---

fast_ui	<i>Main Application UI with Data Loader</i>
---------	---

---

**Description**

Creates the main application UI with a dynamic loader interface that switches to the visualization UI once data is loaded. Provides a file upload interface for loading RData files containing crop rotation data.

**Usage**

```
fast_ui(app_data)
```

**Arguments**

app_data	List containing application data including: <ul style="list-style-type: none"> <li>Input_App_data - List with loaded environment and other configuration</li> </ul>
----------	---

## Details

The UI transitions through two stages:

- Initial loader interface:
  - RData file upload
  - Variable validation
  - Progress indication
- Main visualization interface:
  - Full visualization capabilities
  - Interactive components
  - Data analysis tools

## Value

A Shiny UI object with dynamic loading capabilities

## Examples

```
## Not run:  
shinyApp(ui = ui, server = server)  
  
## End(Not run)
```

---

fast\_viz\_server

*Interactive District/River Catchment Map Server*

---

## Description

Server logic for an interactive mapping application that enables users to load environmental data and compare visualizations between different districts or river catchments. The server handles data loading, map interactions, and dynamic image rendering.

## Usage

```
fast_viz_server(input, output, session, app_data, input_dir)
```

## Arguments

input	Shiny input object
output	Shiny output object
session	Shiny session object
app_data	Application data list
input_dir	Optional directory path for data loading. Default is NA.

## Details

The server functionality includes:

- Data Loading Interface:
  - RData file upload capability
  - Validation of required variables
  - Dynamic UI transitions based on data state
- Map Interaction Handling:
  - District/Catchment selection tracking
  - Interactive highlighting of selected regions
  - Toggle selection functionality
- Visualization Management:
  - Dynamic loading of region-specific images
  - Conditional panel display logic
  - Title updates based on selections

## Value

A Shiny server function that manages the interactive mapping interface

## Required Variables

The following variables must be present in the loaded RData file:

- district\_CropRotViz\_intersection
- cropping\_area
- Crop\_choices
- Districts

## Examples

```
## Not run:
# Run the app with this server
shinyApp(ui = fast_viz_ui, server = fast_viz_server)

## End(Not run)
```

---

fast\_viz\_ui

---

*Interactive District/River Catchment Map UI*


---

## Description

Creates an interactive mapping interface that allows users to compare districts or river catchments through a side-by-side visualization layout. The interface includes a central map with selectable regions and panels for displaying related images.

## Usage

```
fast_viz_ui(input_dir = NA)
```

## Arguments

`input_dir` Optional input directory path. Default is NA.

## Details

The UI consists of three main components:

- Control Panel:
  - Map type selector (Districts/Catchment)
- Central Interactive Map:
  - Leaflet-based map for region selection
  - Interactive selection capabilities
- Comparison Panels:
  - Left panel for first selection visualization
  - Right panel for second selection visualization
  - Conditional display based on user selections

## Value

A Shiny UI object containing a fluid page layout with map and comparison panels

## Theme

Uses the "cyborg" theme with custom styling for:

- Title formatting
- Conditional panel headers (red/orange)
- Layout spacing and alignment

## Examples

```
## Not run:  
# Run the app with this UI  
shinyApp(ui = fast_viz_ui, server = server)  
  
## End(Not run)
```

---

group\_overlapping\_polygons  
*Group Overlapping Polygons*

---

## Description

Assigns group IDs to overlapping polygons in a spatial dataset

## Usage

```
group_overlapping_polygons(polygons_sf)
```

**Arguments**

`polygons_sf`      sf object containing polygons to be grouped

**Value**

sf object with added `group_id` column

---

handle\_diversity\_mapping

*Calculate and Map Diversity Metrics with Error Handling*

---

**Description**

different administrative and ecological boundaries. It includes comprehensive error handling and data validation.

**Usage**

```
handle_diversity_mapping(
  list_intersect_with_borders,
  CropRotViz_intersection,
  agg_cols,
  Districts,
  EZGs = NULL,
  AOIs = NULL
)
```

**Arguments**

<code>list_intersect_with_borders</code>	List containing intersection data with different boundary types Must contain either 'EZG_inter' or 'borders_inter' elements
<code>CropRotViz_intersection</code>	Data frame containing crop rotation visualization data
<code>agg_cols</code>	Character vector of column names to use for aggregation
<code>Districts</code>	sf object containing district boundaries
<code>EZGs</code>	sf object containing ecological zone boundaries (optional)
<code>AOIs</code>	sf object containing areas of interest boundaries (optional)
<code>min_rows</code>	Minimum number of rows required for sufficient data (default: 9)
<code>min_years</code>	Minimum number of years required for sufficient data (default: 3)

**Value**

Returns either: - A spatial object with diversity metrics if successful - NULL if warnings occurred during processing - NA if errors occurred or insufficient data was available

## Examples

```
## Not run:
# Basic usage with only Districts
result <- handle_diversity_mapping(
  list_intersect_with_borders = list(borders_inter = borders_data),
  CropRotViz_intersection = crop_data,
  agg_cols = c("year", "crop_type"),
  Districts = district_sf
)

# Usage with all boundary types
result_full <- handle_diversity_mapping(
  list_intersect_with_borders = list(
    EZG_inter = ezg_data,
    borders_inter = borders_data
  ),
  CropRotViz_intersection = crop_data,
  agg_cols = c("year", "crop_type"),
  Districts = district_sf,
  EZGs = ezg_sf,
  AOIs = aoi_sf
)

## End(Not run)
```

---

Input\_App\_data

*Crop Rotation Application Input Data*


---

## Description

A comprehensive dataset containing crop codes, color mappings, and aggregated statistics for the CropRotationViz package. This dataset includes detailed information about different crop types, their classifications, reference codes, and visualization parameters.

## Usage

```
data(Input_App_data)
```

## Format

A list containing several components for crop rotation visualization:

**codierung\_all** A data frame with crop codes:

- NCNumeric crop code for unique identification
- KlarschriftCrop name in clear text, original designation

**crop\_codes** List of crop code mappings:

- Standard crop codes and their mappings
- Hierarchical structure of crop classifications
- Cross-references between different coding systems

**display\_names** Named vector of display names:

- Human-readable crop names for visualization
- Standardized naming conventions
- Multilingual support where applicable

**crop\_color\_mapping** Named character vector mapping crop types to hex color codes:

- Names: Crop type identifiers (e.g., "winter oil-plant", "winter durum")
- Values: Hex color codes (e.g., "#2e8b57", "#556b2f")
- Consistent color scheme for visualization

**all\_wrap\_number\_count\_small** Data frame with crop aggregation information:

- Name\_2023Character: Original crop name
- Aggregated\_2023Character: Aggregated crop category
- countInteger: Category identifier
- numberInteger: Count of occurrences

**loaded\_env** Environment object containing loaded runtime data

**EZG** River Basins from Germany (Bundesanstalt für Gewässerkunde)

## Crop Categories and Classification

The dataset organizes crops into multiple hierarchical levels:

- Primary Categories:
  - Cereals (winter wheat, winter rye, winter barley, etc.)
  - Oilseeds (winter oil-plant, summer oil-plant)
  - Legumes (protein plants, clover/lutzerne)
  - Root crops (potatoes, sugar beet)
- Secondary Categories:
  - Grassland and forage
  - Special crops (vegetables, fruits, ornamental plants)
  - Fallow and conservation areas
- Special Classifications:
  - Seasonal variants (winter/summer crops)
  - Land use types (agricultural/conservation)
  - Management intensity levels

## Color Mapping System

The visualization color scheme follows these principles:

- Winter Cereals: Dark shades (#556b2f, #7f0000)
- Summer Variants: Lighter complementary colors (#bdb76b, #4682b4)
- Grassland: Natural green tones (#2f4f4f)
- Special Crops: Distinct, vibrant colors
- Conservation Areas: Earth tones
- Consistent color families for related crops

## Data Aggregation Levels

The data supports multiple aggregation levels:

- Individual Crop Level:
  - Specific varieties and cultivars
  - Seasonal variants
  - Management practices
- Group Level:
  - Crop families
  - Usage categories
  - Growing seasons
- Administrative Level:
  - Land use categories
  - Policy relevant groupings
  - Statistical reporting units

## Usage Notes

- Crop codes (NC) should be used as primary keys for database operations
- Display names are optimized for visualization and user interfaces
- Color mappings are designed for maximum differentiation in plots
- Aggregation levels support various analysis scenarios

## Source

Data compiled from:

- Agricultural land use surveys
- Standardized crop classification systems
- Agricultural monitoring and reporting systems
- Expert-developed visualization schemes

## References

- Agricultural land use classification system
- Standard color coding schemes for crop visualization
- International crop coding standards
- Agricultural reporting guidelines

## Examples

```
## Not run:  
# Access basic crop information  
head(Input_App_data$codierung_all)  
  
# Get display name for a crop  
Input_App_data$display_names["wheat"]  
  
# Access color mapping
```



```

Input_App_data$crop_color_mapping["winter wheat"]

# Get aggregated statistics
subset(Input_App_data$all_wrap_number_count_small,
        Name_2023 == "winter wheat")

## End(Not run)

```

---

```

intersecting_check_spatial
      Check Spatial Intersections

```

---

**Description**

Performs spatial intersection checking using a tiled approach for large datasets

**Usage**

```
intersecting_check_spatial(input_list, intersection)
```

**Arguments**

`input_list`      List of sf objects to check  
`intersection`    sf object representing the intersection

**Value**

sf object with intersection results

---

```

intersect_fields      Intersect Multiple Field Layers

```

---

**Description**

Intersects multiple spatial layers while maintaining CRS consistency

**Usage**

```
intersect_fields(fields_list, max_area = 20000 * 1e+06)
```

**Arguments**

`fields_list`      List of sf objects to intersect

**Value**

sf object with intersected fields

---

```
intersect_fields_simple
```

*Simple Field Intersection*

---

### Description

Performs a simplified intersection of multiple spatial layers

### Usage

```
intersect_fields_simple(fields_list, max_area = 20000 * 1e+06)
```

### Arguments

`fields_list`      List of sf objects to intersect

### Value

sf object with intersected fields

---

```
intersect_with_borders
```

*Intersect Geometries with Administrative Borders*

---

### Description

Performs spatial intersection between input geometries and administrative borders at a specified level. The function first creates a central point from the input geometry's bounding box, identifies the corresponding country, and then intersects the input with administrative boundaries of that country.

### Usage

```
intersect_with_borders(input, level, EZG, aoi)
```

### Arguments

`input`              An sf object containing the input geometries to be intersected  
`level`                Numeric value specifying the administrative level for intersection (e.g., 1 for states/provinces, 2 for counties/districts)

### Details

The function follows these steps:

1. Creates a bounding box and center point from input geometry
2. Identifies the country using the center point
3. Retrieves administrative boundaries for the identified country
4. Performs spatial intersection

Special handling is implemented for Germany (DE) where it uses a predefined watershed boundary layer (EZG).

**Value**

An sf object containing the intersected geometries

**Examples**

```
## Not run:
# Create sample input geometry
input_geom <- st_read("input.shp")

# Intersect with level 1 administrative boundaries
result <- intersect_with_borders(input_geom, level = 1)

## End(Not run)
```

---

```
process_specific_transitions
```

*Process Crop Transitions for Specific Crop Type*

---

**Description**

Analyze and summarize temporal transitions into and out of a specific crop type across multiple years using aggregated agricultural data.

**Usage**

```
process_specific_transitions(data, crop)
```

**Arguments**

data	Data frame containing aggregated crop transition data with columns starting with "Aggregated_" followed by year numbers, and a freq column for transition frequencies
crop	Character string specifying the crop type to analyze transitions for

**Details**

This function processes temporal crop rotation patterns by identifying transitions where the specified crop appears as either the source or target. It handles multi-year transitions by pairing consecutive years and aggregates frequencies into percentages. The output distinguishes between transitions into the crop (marked as "before") and out of the crop (marked as "after").

**Value**

A data frame with columns:

- source: Origin crop type (with "(before)" suffix for transitions into target crop)
- target: Destination crop type (with "(after)" suffix for transitions from source crop)
- value: Percentage of transitions between the crop pairs

---

`run_fast_visualization_app`*Launch Interactive District/River Catchment Visualization App*

---

### Description

Initializes and launches the interactive mapping application for comparing district or river catchment visualizations. This function handles data loading, resource path configuration, and app deployment.

### Usage

```
run_fast_visualization_app(input_dir = NA)
```

### Arguments

<code>input_dir</code>	Optional directory path for data loading. If NA (default), the app will present a file upload interface.
------------------------	--

### Details

The function performs the following setup steps:

- Loads package data from CropRotationViz
- Creates an isolated environment for app data
- Configures resource paths for static assets
- Launches the Shiny application in the default browser

### Value

Launches a Shiny application instance

### Dependencies

Requires the following package components:

- CropRotationViz package with Input\_App\_data
- fast\_ui function for UI generation
- fast\_viz\_server function for server logic

### Note

The application will automatically open in the system's default web browser when launched.

## Examples

```
## Not run:
# Launch app with default file upload interface
run_fast_visualization_app()

# Launch app with specific data directory
run_fast_visualization_app(input_dir = "path/to/data")

## End(Not run)
```

---

run_processing_app	<i>Run the Crop Rotation Visualization Application</i>
--------------------	--

---

## Description

Launches the Shiny application for analyzing and visualizing agricultural crop rotation sequences. This application provides an interactive interface for loading spatial data, processing crop rotations, and generating visualizations and analysis outputs.

## Usage

```
run_processing_app(
  output_dir = NA,
  common_column = NA,
  start_year = NA,
  intersection_type = NA,
  preview = TRUE,
  vector_file = T
)
```

## Arguments

output_dir	Character string specifying the default output directory path. If NA (default), the user will be prompted to select an output directory through the application interface.
common_column	Character string specifying a default column name to be used for crop identification across all input files. If NA (default), column selection will be prompted for each file.
start_year	Numeric value indicating the initial year for the sequence analysis. If NA (default), years can be selected individually for each file. Otherwise, years will be auto-incremented from this starting value.
intersection_type	= "complete",
preview	logical. If True (default) you get a snapshot as png from your processed data as sankey chart in PNG format.
vector_file	logical. If True (default) you will get a vector file as an additional output with all intersected and aggregated field data.

## Details

The application provides a comprehensive interface for crop rotation analysis:

- Data Input
  - Support for multiple spatial file formats (.shp, .geojson, .fgb, .gpkg, .sqlite)
  - Capability to load up to 10 years of spatial data
  - Flexible column mapping for crop identification
- Processing Options
  - Choice between national coding (NC) or name-based crop identification
  - Optional crop class aggregation
  - Multiple intersection processing methods
- Output Generation
  - Multiple export format options
  - Comprehensive metadata documentation
  - Sankey diagram visualizations
  - Statistical summaries

## Value

A Shiny application object that can be run with `runApp()` or deployed to a Shiny server.

## Data Requirements

The application expects:

- Spatial data files containing crop information
- Consistent coordinate reference systems across files
- Valid crop codes or names in specified columns

## Output Files

The application generates several output files:

- Spatial data file (in chosen format) containing intersection results
- RData file with processed data and statistics
- Processing information text file with metadata
- Sankey diagram visualization of crop sequences

## Examples

```
## Not run:
# Basic usage with default settings
run_sequencer_app()

# Specify output directory
run_processing_app(output_dir = "path/to/output")

# Specify output directory and common column name
run_processing_app(
  output_dir = "path/to/output",
  common_column = "crop_code"
```

```
)

# Full configuration with start year
run_processing_app(
  output_dir = "path/to/output",
  common_column = "crop_code",
  start_year = 2020,
  preview = T,
  vector_file = T
)

## End(Not run)
```

---

run\_visualization\_app *Run the Crop Rotation Visualization Application*

---

## Description

Launches the Shiny application for visualizing crop rotation plans and their impact. The application provides an interactive interface for exploring rotation patterns, analyzing sustainability metrics, and understanding environmental effects of different cropping sequences.

## Usage

```
run_visualization_app(input_dir = NA)
```

## Arguments

input_dir	Optional filepath to custom input data directory. If NA (default), you have to choose it manually in the app.
-----------	---

## Details

The function initializes the visualization environment by:

1. Loading built-in or custom input data
2. Setting up the resource paths for static assets
3. Launching the Shiny server with configured UI and server logic

## Value

A Shiny application object

## Examples

```
## Not run:
run_visualization_app()
run_visualization_app("path/to/custom/data")

## End(Not run)
```

---

transform\_rotation\_data

*Transform Rotation Data*


---

## Description

Transforms crop rotation data for visualization and analysis purposes

## Usage

```
transform_rotation_data(
  All_rot_big,
  distribution_df,
  input_area_range,
  choices,
  selected_crops,
  type,
  specific_crop = NULL,
  district = NULL,
  EZG = NULL
)
```

## Arguments

All_rot_big	Data frame containing crop rotation data with columns for different years and an 'area' column
input_area_range	Numeric vector of length 2 specifying the minimum and maximum area range in square kilometers
choices	Character vector containing all possible crop types
selected_crops	Character vector containing the subset of crops to include in the analysis
type	Character string specifying the type of transformation: either "basic" or "specific"
specific_crop	Character string specifying a particular crop to focus on when type = "specific"

## Value

A data frame with columns:

Area	Numeric, area in square kilometers
id	Integer, unique identifier for each rotation
value	Character, year identifier
key	Factor, crop type
year	Numeric, year of rotation



**Examples**

```
## Not run:
rotation_data <- transform_rotation_data(
  All_rot_big = my_rotation_data,
  input_area_range = c(0, 1000),
  choices = c("Wheat", "Corn", "Soybeans"),
  selected_crops = c("Wheat", "Corn"),
  type = "basic"
)

## End(Not run)
```

---

transform\_rotation\_summary

*Transform Rotation Summary*


---

**Description**

Creates a detailed summary of crop rotation patterns including area calculations and crop sequences across specified years

**Usage**

```
transform_rotation_summary(
  All_rot_big,
  area_range,
  type = c("basic", "specific"),
  choices = NULL,
  selected_crops = NULL,
  specific_crop = NULL,
  max_rows = 3000,
  years
)
```

**Arguments**

All_rot_big	Data frame containing crop rotation data with columns for different years and an 'area' column
area_range	Numeric vector of length 2 specifying the minimum and maximum area range in square kilometers
type	Character string specifying the type of transformation: either "basic" or "specific"
choices	Character vector containing all possible crop types (required when type = "basic")
selected_crops	Character vector containing the subset of crops to include (required when type = "basic")
specific_crop	Character string specifying a particular crop to focus on (required when type = "specific")
max_rows	Integer specifying the maximum number of rows to return in the summary (default: 3000)
years	Numeric vector specifying the years to include in the analysis

**Value**

A data frame with columns:

area_km2	Numeric, area in square kilometers
Crop_YYYY	Character, crop type for each year YYYY in the specified range

**Examples**

```
## Not run:
summary_data <- transform_rotation_summary(
  All_rot_big = my_rotation_data,
  area_range = c(0, 1000),
  type = "basic",
  choices = c("Wheat", "Corn", "Soybeans"),
  selected_crops = c("Wheat", "Corn"),
  max_rows = 1000,
  years = 2015:2020
)

## End(Not run)
```

---

ui

---

Main Application UI with Data Loader

---

**Description**

Creates the main application UI with a dynamic loader interface that switches to the visualization UI once data is loaded. Provides a file upload interface for loading RData files containing crop rotation data.

**Usage**

```
ui(app_data)
```

**Arguments**

app_data	List containing application data including: <ul style="list-style-type: none"> <li>• Input_App_data - List with loaded environment and other configuration</li> </ul>
----------	---

**Details**

The UI transitions through two stages:

- Initial loader interface:
  - RData file upload
  - Variable validation
  - Progress indication
- Main visualization interface:
  - Full visualization capabilities
  - Interactive components
  - Data analysis tools

**Value**

A Shiny UI object with dynamic loading capabilities

**Examples**

```
## Not run:
shinyApp(ui = ui, server = server)

## End(Not run)
```

---

viz_server	<i>Crop Rotation Visualization Server Logic</i>
------------	---

---

**Description**

Implements comprehensive server-side logic for the crop rotation visualization application. Handles data loading, transformations, and creates reactive visualizations including Sankey diagrams, tables, and summary plots.

**Usage**

```
viz_server(input, output, session, app_data, input_dir)
```

**Arguments**

input	Shiny input object
output	Shiny output object
session	Shiny session object
app_data	List containing application data including crop mappings and settings

**Value**

A Shiny server function that manages the complete visualization workflow

**Data Loading**

Handles the loading and validation of RData files containing:

- CropRotViz\_intersection - Main rotation data
- cropping\_area - Area calculations
- Crop\_choices - Available crop choices
- District\_choices - Available district choices
- EZG\_choices - Available river basin choices

## Visualization Components

Creates and manages multiple visualization types:

- Percentage Plots:
  - Area distribution visualization
  - Dynamic updates based on selection
- Sankey Diagrams:
  - Crop rotation flow visualization
  - Crop-specific pattern analysis
- Data Tables:
  - Color-coded rotation summaries
  - Interactive filtering and sorting
- Plotly Integration:
  - Interactive Sankey diagrams
  - Custom styling and layout

## Reactive Flow

- Data Loading Stage:
  - File validation
  - Environment setup
  - Variable checking
- Data Processing Stage:
  - Rotation data transformation
  - Area calculations
  - Color mapping
- Visualization Stage:
  - Plot generation
  - Table formatting
  - Interactive updates

## Examples

```
## Not run:  
# Run the complete Shiny application  
shinyApp(ui = ui, server = viz_server)  
  
## End(Not run)
```

---

`viz_ui`*Crop Rotation Visualization UI*

---

### Description

Creates an interactive visualization interface for crop rotation patterns, featuring multiple visualization types including Sankey diagrams, tables, and plots. The UI is split into two main tabs: general sequence plotting and crop-specific analysis.

### Usage

```
viz_ui(input_dir = NA)
```

### Details

The UI includes several components:

- Plot Sequence Tab:
  - Area range selector (0-4000 km<sup>2</sup>)
  - Crop selection interface with multi-select capability
  - Percentage area visualization
  - Sankey diagram of crop rotations
  - Interactive data table with color-coded rotations
  - Crop class merging visualization
- Plot Crop Specific Sequence Tab:
  - Specific crop selector
  - Area range configuration
  - Crop-specific Sankey diagram
  - Detailed rotation patterns table

### Value

A Shiny UI object containing the complete visualization interface

### Theme

Uses the "cyborg" theme with custom CSS modifications for:

- DataTables styling
- Tab navigation appearance
- Popup content formatting

### Note

This UI requires the following global variables to be present:

- choices - Vector of crop choices
- app\_data - List containing Input\_App\_data

**Examples**

```
## Not run:  
# Run the app with this UI  
shinyApp(ui = viz_ui, server = server)  
  
## End(Not run)
```

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