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Required Downloads

Installing Anaconda

Mac Installation

1. Navigate to the Anaconda homepage, anaconda.com
2. Click the “Get Started”
3. Click the “Download Anaconda installers”
4. Locate the installers for Mac and click on the “64-Bit Graphical Installer” to download
5. Double click the downloaded file
6. Click the “Continue” on the pop up with the message “This package will run a program to determine if the software can be installed”
7. Click “Continue” on the first page of the installer menu
8. Click "Continue" on the Read Me page
9. Click "I Agree" to the terms of the software license agreement
10. Click "Continue" on the license page
11. On the Destination Select page choose one of the following options
 - a. “Install for me only”
 - i. Downloads anaconda for the current desktop profile
 - b. “Install on a specific disk” (administrator privileges required)
 - i. Downloads anaconda for all desktop profiles
 - ii. Choose the disk that contains the target desktop profiles
12. Click “Continue” on the Destination page
13. Click “Continue” on the PyCharm IDE page
14. Click “Close” on the Summary page

Windows Installation

1. Navigate to the Anaconda homepage, anaconda.com
2. Click the “Get Started” button
3. Click the “Download Anaconda installers” button
4. Locate the installers for your Windows and click on the “64-Bit Graphical Installer” to download
5. Double click the downloaded file to launch the installer
 1. Do not launch installer from Favorites folder
6. Click “Next” on the first page of the installer menu
7. Click “I Agree” to the terms of the software license agreement

8. On the Destination Select page choose one of the following options
 1. “Just Me” *Less prone to errors*
 - i. Downloads anaconda for the current desktop profile
 2. “Install on a specific disk” (administrator privileges required)
 - i. Downloads anaconda for all desktop profiles
 - ii. Choose the disk that contains the target desktop profiles
9. On the “Choose install Location page”, use the default destination folder to install anaconda in
 1. **Make sure the directory path doesn’t contain spaces or unicode characters**
 - i. Regular alphabetical characters are allowed
 - ii. See resources for unicode character table
10. Click “Next
11. On the Advanced Installation Options Page
 1. **Do not check** the box to add Anaconda3 your PATH environment variable
 2. **Check** the box to register Anaconda3 as your default python
12. Click “install”

Windows Troubleshooting

1. If antivirus software is active, temporarily disable it. Re-enable after installation concludes
2. If you installed for all users, OR under Administrative Privileges: uninstall anaconda (Uninstall directions in Resources) and reinstall for “Just me” without Administrative Privileges
3. **Double check that the Installation directory path doesn’t contain spaces or unicode characters. If it does then uninstall and reinstall in a new directory**

Anaconda Resources

- Installation Documentation: <https://docs.anaconda.com/anaconda/install/>
- Uninstall Documentation: <https://docs.anaconda.com/anaconda/install/uninstall/>
- User Guide: <https://docs.anaconda.com/anaconda/user-guide/>
- Unicode Character Chart: https://en.wikipedia.org/wiki/List_of_Unicode_characters

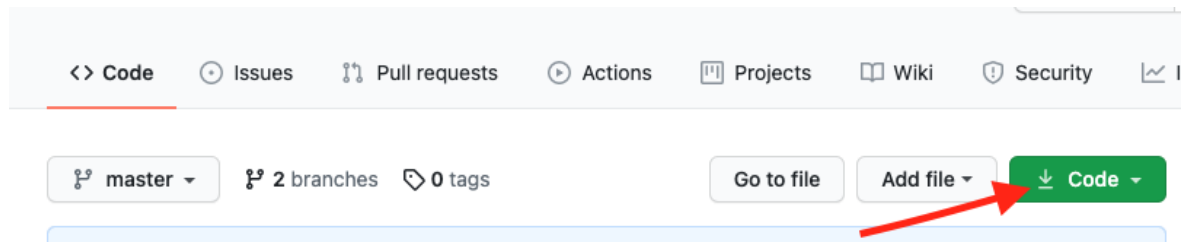
Downloading Pycharm

1. Choose the **community** download for your OS and follow download instructions found on <https://www.jetbrains.com/pycharm/download/>

Downloading the Pipeline

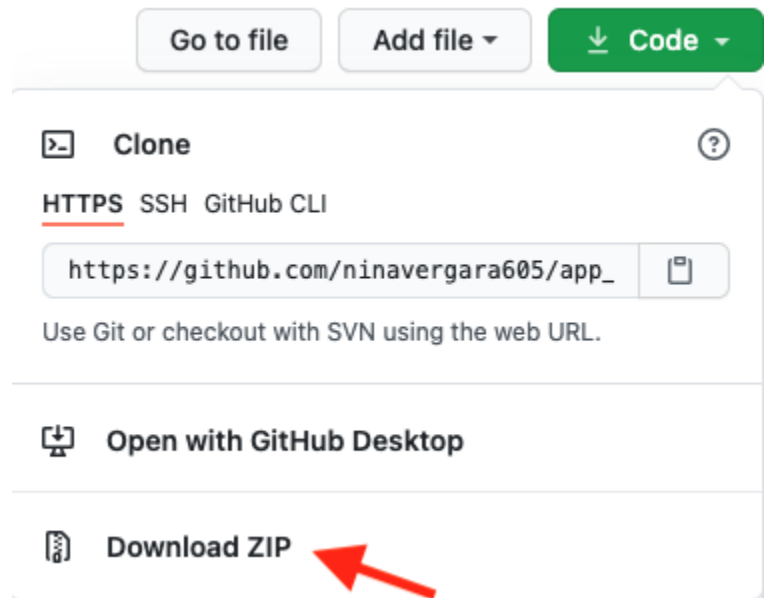
1. Navigate to https://github.com/ninavergara605/app_name-Null

2. Click
the
green
“Code” button



3. Click the “Download Zip” button

4. Unzip the downloaded file



GitHub Resources

- Downloading zip file:

https://www.itprotoday.com/development-techniques-and-management/how-do-i-download-files-github?reg_form=adv

- Cloning a Repository (advanced): <https://docs.github.com/en/github/creating-cloning-and-archiving-repositories/cloning-a-repository-from-github/cloning-a-repository>

Preparing the Data

Creating the ROI Template

Introduction

Fixations can be categorized and binned by **Regions of Interest** (ROI's). **ROI events** can be defined by their presentation and metadata attributes. Presentation attributes contain spatial and temporal information while metadata attributes contain identifying information.

ROI Event Attributes

Possible Attributes

The ROI templates allow for the following attributes to be tracked:

- roi_label (name)
- roi_id
- top_left_xy
- bottom_right_xy
- start
- stop

All coordinates must be comma separated, i.e. "x,y"

Required Attributes

The metadata attributes, "roi_label" and "roi_id", are required for all static and dynamic ROI. **These values must be unique.**

Optional Attributes

The coordinate (ending in "_xy") and temporal ("start" and "stop") attributes are optional. However, at least one pair needs to be filled in for every ROI in order to be correctly paired.

Note: Partially filling out spatial and temporal attributes isn't currently supported. This has not been tested and may lead to incorrect data pairing.

Static vs Dynamic ROI

A static ROI event appears in the same location and within the same time window for all trials and participants. A dynamic ROI event appears in a different location and/or time window across trials, blocks, or subjects.

Defining Static ROI

For static Roi, metadata and event attributes are listed under the header “static”.

static					
roi_label	roi_id	top_left_xy	bottom_right_xy	start	stop
static_1	1	x0,y0	x1,y1	t0	t1
static_2	2	x2,y2	x3,y3	t2	t3

Defining Dynamic ROI

Dynamic Event Keys

Dynamic event keys are associated with unique positions and timing attributes of dynamic ROI's. These key-presentation attribute relationships are listed under the 'dynamic_event_options' section of the ROI template.

dynamic_event_options				
key	top_left_xy	bottom_right_xy	start	stop
x	x4,y4	x5,y5	t0	t3
z	x1,y1	x3,y3	t2	t3
y	x2,y2	x4,y4	t1	t2

Dynamic Event Key Map

In a separate file, a column is created for each dynamic ROI.

The values of each column are populated with event keys that indicate the spatial temporal attributes of their respective ROI's for every subject trial.

subject	trial	dynamic_1_key	dynamic_2_key
1201	1	y	x
1201	2	z	y

Dynamic ROI Metadata

In addition the 'roi_label' and 'roi_id' metadata attributes, the dynamic ROI also contains an “event_key_map_column” section. These values indicate which column of the Dynamic Event Key Map should be associated with their respective dynamic ROI attributes.

Putting It All Together

An ROI template with static and dynamic ROI that includes all possible event attributes:

dynamic_roi_metadata		
event_key_map_column	roi_label	roi_id
dynamic_1_key	dynamic_1	3
dynamic_2_key	dynamic_2	4

static						dynamic_roi_metadata			dynamic_event_options					
roi_label	roi_id	top_left_xy	bottom_right_xy	start	stop	event_key_map_column	roi_label	roi_id	key	top_left_xy	bottom_right_xy	start	stop	
static_1	1	x0,y0	x1,y1	t0	t1	dynamic_1_key	dynamic_1	3	x	x4,y4	x5,y5	t0	t3	
static_2	2	x2,y2	x3,y3	t2	t3	dynamic_2_key	dynamic_2	4	z	x1,y1	x3,y3	t2	t3	
									Y	x2,y2	x4,y4	t1	t2	

File Naming

When File Naming is Important

File names are used to label imported If more than one matching file is found in the input path directory. A file is considered matching if the extension matches the default type specified for the input variable. See User Input for default values.

Splitting File Names

filenames are split on datatype transition and '-', '_' characters.

Ex:

- 1201study_3 -----> 1201, study, 3
- 598-2033carrot ----> 598, 2033, carrot

Metadata

Defining Metadata Keys

Metadata Keys Indicate relevant subject and experiment information that will be pulled from file names and/or behavioral data. These keys are used to label output and aid in pairing Eye-tracking data with Behavioral and ROI data.

Metadata Keys must be separately defined for behavioral, ASC filenames, and ROI Event data if they are to be used for analysis

Extracting Metadata from File names

Metadata is pulled from file names that have been split according to the in The File Naming section.

Dropping Unnecessary Information

If filenames have irrelevant information, the 'drop' keyword can be used to discard groups after splitting.

Ex:

if the metadata keys were defined as: ['Subject', 'drop', 'block']

...and a split file name is: ['1201', 'unnecessary', 'Test']

...the pairing would be: 'Subject': 1201, 'block': 'Test'

Standardizing split file name data

File names are standardized by the rules found in the Automatic Data Cleaning: Strings Section.

Extracting Metadata From DataFrames

If subject Behavioral or ROI Event data is stored in a combined file, the metadata keys must be present as column names.

Automatic Data Cleaning

Paths

Standardizing Paths

String paths are turned into Pathlib objects. These objects ensure that the path is in the correct format for the operating system. In addition, any directories that do not yet exist are created. For more Pathlib attributes, see the resources below.

File Names

If a filename does not split into the required number of groups to be paired with the metadata keys, then the full path will be printed out with the tag: "invalid path" and discarded.

Strings

Raw strings or strings found in dictionaries or lists are converted into a numeric datatype, if possible. If the string is found to contain uppercase alphabetical characters, then they will be lowered.

Tables

Tables are created from CSV and Excel file formats. The first row should contain column names.

String column names and column values are cleaned by the rules above. Any rows containing null values in a metadata columns will be dropped to prevent pairing errors in later analysis

ASC Files

ASC files will be filtered for 'EFIX', 'EBLINK', 'SSAC', and 'ESACC' values. A column labeled 'trial_id' will be generated from a cumulative count of each 'START' values found in the files

Resources

- Pathlib Library: <https://docs.python.org/3/library/pathlib.html>

User Input

ROI Options

Calc_roi_raster_coords: *boolean*

Boolean: default is False. Set to 'True' if roi template coordinates are in the cartesian coordinate system. The program is designed for a raster coordinate system.

Aspect_ratio: *tuple*

Should be a tuple of (screen_width, screen_height). Used if calc_raster_coords = True.

If no aspect ratio is provided and ASC files are present, aspect ratios will be extracted from the ASC files. This is especially helpful if participants were run with different aspect ratios

Resources:

Cartesian coordinate system description : https://mathinsight.org/cartesian_coordinates

Raster coordinate system description: https://en.wikipedia.org/wiki/Raster_graphics

Behavioral Options

behavior_test_path: *string*

String path leading to the behavioral data. This data can include test scores, roi event key maps, and other information that needs to be attached to output. This needs to be a '.xlsx' file.

Behavior_test_trial_col: *string*

The name of the trial id column located in the behavior data. This is needed to pair the ASC data with the behavior test data, since a trial id column is automatically created when ASC data is imported.

Behavior_metadata_keys: *list of strings*

The column names of the metadata columns found in the behavior data. Should be in list format.

Attach_behavior_cols: *list of strings*

Column names that should be attached to all output.

Eye-Tracking Options

Asc_directory_path: *string*

String path leading to the asc file directory. Asc files can be nested within the directory and grouped with other files. Any file containing an '.asc' extension flagged for import.

Attach_movement_cols: *list of strings*

Column names that will be attached to all output. Columns are generated from eprime output.

Possible column names are:

- 'file_position'
- 'type'
- 'eye'
- 'block_relative_start'
- 'block_relative_stop'
- 'start'
- 'stop'
- 'duration'
- 'type_count'

Asc_metadata_keys: *list of strings*

The labels of the metadata extracted from the ASC filenames. See Metadata section for more information.

Analysis Options

Time bin size: *integer*

The size of the time bins for Response-locked and Stimulus-locked binning analysis.

Summary filter out: *dictionary*

A dictionary containing column names and values to be excluded when calculating proportion of view binning summary.

Ex: Wanting to exclude rows based on the 'roi_label' column. We don't want to include rows associated with 'scene' and 'whole_screen' roi

Summary_filter_out = {'roi_label': ['scene', 'whole_screen']}

Summary filter for: *dictionary*

A dictionary containing column names and values to be included when calculating proportion of view binning summary.

ex: Wanting to include rows based on the 'roi_label' column. We want to calculate the proportion of view for the two people roi.

summary_filter_out={'roi_label': ['person_1', 'person_2']}

Roi Event Map Options

Roi_event_map_path: *string*

Takes a directory containing ROI event Key map files. Each file should be in CSV format and labeled with subject block metadata.

At this time, all key maps must contain a 'trial_id' column for successful pairing with the Eye-tracking data

Roi_event_map_filename_contains: *string*

Used to identify roi event map files. Files will be chosen based on partial string matches

Ex: filenames have 'event_map' in their names like '1201_1_event_map.csv'

Roi_event_map_filename_contains = 'event_map'

Roi_event_map_metadata_keys: *list of strings*

Labels for the metadata values that will be extracted from the event map filenames.

ex: ['subject_id', 'block_id']

Roi_event_map_import_skip_rows: *integer or list of integers*

Specifies row(s) to skip while importing the event maps. Integers should be zero indexed, so the first row of a file is 0, the second is 1...

Ex: skip first row because it's blank:

`Roi_event_map_import_skip_rows = 0`

Attach_event_cols: *list of strings*

Columns present in the event maps that need to be attached to all export files.

ex: `['col_1', 'col_2']`

Export Options

Output_directory_path: *string*

A path leading to the desired storage directory for results. Any folders that are specified in the path and don't exist will automatically be created.

Output_folder_name: *string*

The desired name for the output data folder. Default is 'processed_data'.

Running the Application

Opening the Application

1. Open the pycharm application
2. Click the 'open' folder icon
3. Click on the PeyeTracker folder and select 'OK'
4. Navigate to 'main.py' located in './PeyeTracker/PeyeTracker/main.py'

Activating the Virtual Environment

1. To sync anaconda environment, navigate to the preferences panel
2. Select 'Project: PeyeTracker' in the side panel
3. Under project settings click 'Python Interpreter'
4. Click on the gear icon next to the 'python interpreter' dropdown menu
5. In the left-hand panel, select 'Conda Environment'
6. Select the 'Existing environment' bullet point and navigate to anaconda's python 3.8. interpreter
 - 6.1. Mine was located in 'opt/anaconda3/bin/python3.8'
7. Select 'OK'

8. In the preferences panel, ensure that 'pandas' is present in the 'package' column

Specifying Runtime Configuration

1. In the top right corner select 'add configuration'
2. Double click on 'add new run configuration'
3. Select 'python'
4. Specify the configuration name
5. For script path navigate to 'main.py'
6. Under the 'execution' dropdown check 'run with python console'
7. On the bottom of the window check 'activate tool window'

Run

1. press the green play button in the upper right-hand corner

Test Data

Roi templates, event key maps, and asc files are provided. The required configurations for each test dataset is provided in the 'test_data' folder.