
impro Documentation

Release a

Sebastian Reinhard

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Impro. is a package for data processing in super-resolution microscopy. It contains high performant GPU based visualization and algorithms for 2D and 3D data. Main features:

- Cuda accelerated 2D Alpha Shapes
- Automated image alignment via General Hough Transform
- Huge list of filters for localization datasets
- Customizable Opengl widget based on modelview perspective standard
- Pearson correlation

CHAPTER 1

References

A detailed explanation of the algorithm and it's functionality can be found at:

Reinhard S, Aufmkolk S, Sauer M, Doose S. Registration and Visualization of Correlative Super-Resolution Microscopy Data. *Biophys J.* 2019 Jun; 116(11) 2073-2078. doi:10.1016/j.bpj.2019.04.029. PMID: 31103233.

CHAPTER 2

Example

Basic usage example. Preprocessing is adapted to the dataset provided in Super-resolution correlator.

```
from impro.data.image_factory import ImageFactory
from impro.analysis.filter import Filter
from impro.analysis.facade import *

# Read and preprocess SIM image
image = ImageFactory.create_image_file(r"path_to_file.czi")

# Example for image preprocessing
image_array = image.data[:, 3] / 6
image_array = np.clip(image_array[0], 0, 255)
image_array = np.flipud(image_array)
image_array = (image_array).astype("uint8")[0:1400, 0:1400]
image_array = np.fliplr(image_array)

# Read dSTORM data
storm = ImageFactory.create_storm_file(r"path_to_file.txt")

# Preprocess dSTORM point data
indices = Filter.local_density_filter(storm.stormData, 100.0, 18)
storm_data = storm.stormData[indices]
# Render dSTORM data to image
im = create_alpha_shape(storm_data, 130)
col = int(im.shape[1]/200)
row = int(im.shape[0]/200)
source_points, target_points, overlay, results = find_mapping(sim, storm, n_row=row,
    ↪ n_col=col)
source_points, target_points = error_management(results, source_points, target_points,
    ↪ n_row=row)
M = transform.estimate_transform("affine", source_points, target_points)
correlation_index = pearson_correlation(sim, cv2.cvtColor(storm, cv2.COLOR_RGBA2GRAY),
    ↪ M)
```

2.1 Analysis module

Collection of useful analysis tools. For easy access use the *Analysis facade*

2.1.1 Analysis facade

Easy access to complex algorithmic concepts.

Author Sebastian Reinhard

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

```
impro.analysis.analysis_facade.create_alpha_shape(storm_data: numpy.ndarray,
                                                 pha_value: float, px_size=32.2,
                                                 line_width=1.0) → numpy.ndarray
```

Facade function to render alpha shape of dSTORM data

Image is rendered from (0,0) to the maximal dimension of the dSTORM data.

Parameters

- **storm_data** (*np.ndarray (nx5)*) – Point Cloud data to be rendered
- **px_size** (*float*) – Pixel size for the rendered image in nanometer per pixel
- **pha_value** (*float*) – Core value to compute the alpha shape in nanometer
- **line_width** (*float*) – Line width of the alpha shape lines in a.u.

Returns **image** – Rendered image

Return type *np.array*

Example

```
>>> points = np.random.randint(0,32000,(1000000, 2))
>>> image = create_alpha_shape(points, 130)
>>> image.shape
(994, 994)
```

```
impro.analysis.analysis_facade.create_storm(storm_data: numpy.ndarray, px_size=32.2,
                                             size=20, cluster=array(3.))
```

Facade function to render an image of dSTORM data.

Image is rendered from (0,0) to the maximal dimension of the dSTORM data.

Parameters

- **storm_data** (*np.array (nx5)*) – Point Cloud data to be rendered
- **px_size** (*float*) – Pixel size for the rendered image in nanometer per pixel
- **size** (*float*) – Size of the rendered points in nanometer
- **cluster** (*np.array (n)*) – Affiliation of the i th point to cluster[i] cluster

Returns **image** – Rendered image

Return type *np.array*

Example

```
>>> points = np.random.randint(0,32000,(1000000, 5))
>>> image = create_alpha_shape(points, 130)
>>> image.shape
(994, 994)
```

`impro.analysis.analysis_facade.error_management(result_list: list, source_points, target_points, n_row=5)`

Test the results of th weighted GHT for missmatches

Parameters

- **result_list** (*list*) – results of the Weighted General Hough Transform
- **source_points** (*np.array*) – Source points of affine transformation
- **target_points** (*np.array*) – Target points of affine transformation
- **n_row** (*int*) – number of rows used for the weighted GHT

Returns `source_points, target_points` – Filtered source and target points for affine transformation

Return type `np.array`

`impro.analysis.analysis_facade.find_mapping(target: numpy.ndarray, source_color: numpy.ndarray, n_col=5, n_row=5, offset=0) → numpy.ndarray`

Facade function to find a mapping from source_color image to gray scale target image.

Parameters

- **target** (*np.array*) – Target image of affine transformation
- **source_color** (*np.array*) – Source image of affine transformation
- **n_col** (*int*) – number of collums to seperate the source image into
- **n_row** (*int*) – number of rows to seperate the source image into
- **offset** (*int*) – Start seperating the source image at (0+offset,0+offset). Offset is given in pixel

Returns

- **source_points, target_points** (*np.array*) – Source and target points for affine transformation
- **overlay** (*np.array*) – Target image overlayed with source image segments at the matching position
- **results** (*list*) – Results of the Weighted General Hough Transform

Example

```
>>> import cv2
>>> points = np.random.randint(0,32000,(1000000, 5))
>>> source = create_alpha_shape(points, 130)
>>> points = np.random.randint(0,120000,(1000000, 5))
>>> target = create_alpha_shape(points, 130)
>>> find_mapping(cv2.cvtColor(target, cv2.COLOR_RGB2GRAY), source)
```

```
impro.analysis.analysis_facade.pearson_correlation(target: numpy.ndarray,
                                                    source: numpy.ndarray, map:
                                                    numpy.ndarray) → float
```

Compute pearson correlation index after alignment

Parameters

- **target** (*np.array*) – target image in gray scale
- **source** (*np.array*) – source image in gray scale
- **map** (*sklearn.transformation*) – computed transformation

Returns **source_points, target_points** – Filtered source and target points for affine transformation

Return type *np.array*

Example

```
>>> import skimage.transform
>>> source = np.ones((1000,1000))
>>> target = np.ones((1000,1000))
>>> source_points = np.array([[1.0,1.0],[500,500],[700,500]])
>>> target_points = source_points
>>> M = transform.estimate_transform("affine",source_points,target_points)
>>> pearson_correlation(target, source, M)
(1.0)
```

2.1.2 Alpha Shape

The objective of the alpha shape algorithm is to deliver a formal meaning for the geometric notation of ‘shape’, in the context of finite point sets. The straciatella ice cream example gives a visual explanation of the concept: Consider a ball of ice cream with chocolate pieces. The chocolate pieces represent the distribution of points in space (ice cream). The alpha shape algorithm now tries to eat as much ice cream as possible without touching any chocolate pieces, using an arbitrary predefined spoon size, the α -value. Choosing a very small spoon size results in all ice cream being eaten, a very big spoon size in no ice cream being eaten at all (Convex Hull). But a size in between creates a concave hull representing the shape of the distributed chocolate pieces, the desired alpha shape.

References

- (1) Edelsbrunner, Herbert ; Mücke, Ernst P.: Three-dimensional Alpha Shapes. In: ACM Trans. Graph. 13 (1994), Januar, Nr. 1, 43–72. <http://dx.doi.org/10.1145/174462.156635>. – DOI 10.1145/174462.156635. – ISSN 0730-0301
- (2) Fischer, Kaspar: Introduction to Alpha Shapes. https://graphics.stanford.edu/courses/cs268-11-spring/handouts/AlphaShapes/as_fisher.pdf.

Author Sebastian Reinhard

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2018.03.09

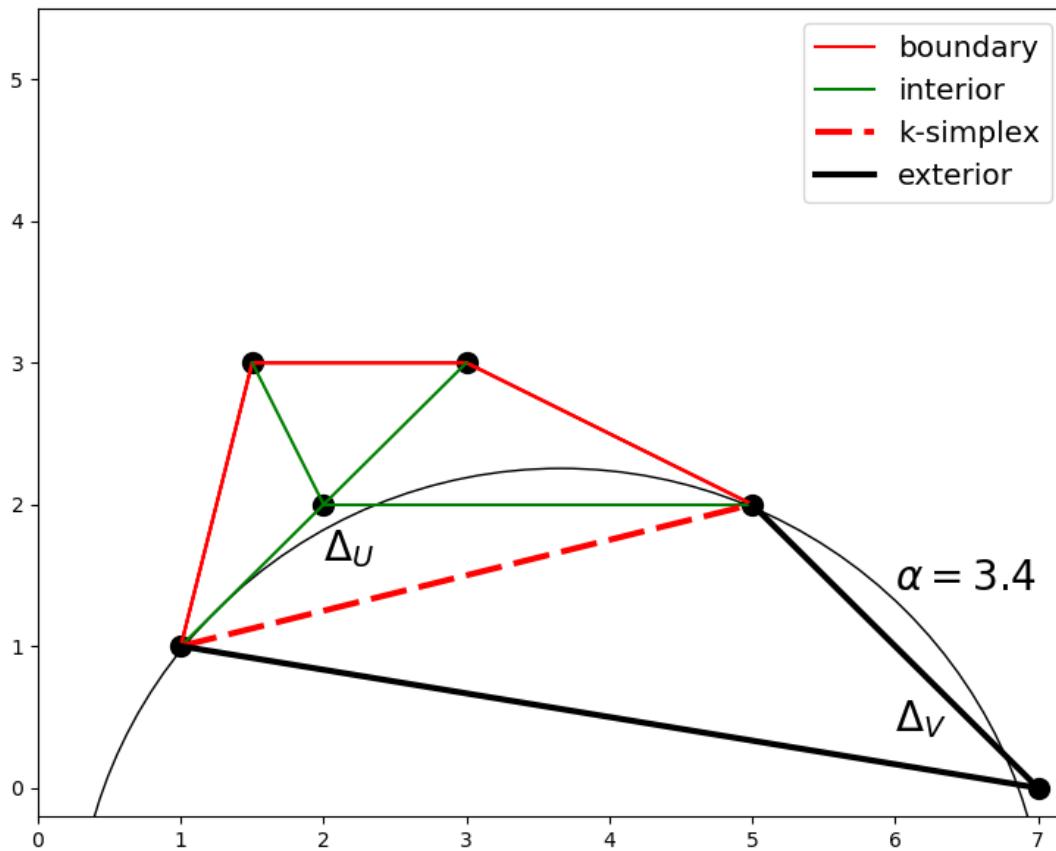


Fig. 1: Considering the red dotted line l , the limit a is defined as the minimum of the diameter of the circumcircles around Δ_U and Δ_V . The limit b is the maximum of circumcircles around Δ_U and Δ_V . Since the α -ball is smaller than b , but bigger than a , l is classified as boundary.

Example

```
>>> data = np.random.randint(0,32000,(1000000, 2))
>>> k_simplices = get_k_simplices(data[...,:2])[0]
```

```
class impro.analysis.alpha_shape_gpu.AlphaComplex(struct_ptr: int, indices: numpy.ndarray, points: numpy.ndarray, neighbors: numpy.ndarray)
```

Bases: object

Derive 2D alpha complex from scipy.spatial Delaunay

Class to create a alpha complex structure on GPU.

Parameters

- **struct_ptr** (*int*) – pointer to allocated memory for structure
- **indices** (*np.array*) – nx3 array containing point indices of the delaunay triangulation simplices
- **neighbors** (*np.array*) – nx3 array containing the indices of neighboring simplices
- **points** (*np.array*) – nx2 array of the points used for delaunay triangulation

get()

Returns nx5 of d=1 simplices containing: [index1, index2, dist, sigma1, sigma2] with sigma 1 < sigma 2

Return type numpy.array

memsize = 32

merge()

```
impro.analysis.alpha_shape_gpu.get_k_simplices(points: numpy.ndarray)
```

Parameters **points** (*np.array*) – nx2 array of points to use for alpha complex

Returns

- **alpha complex** (*mx5 array of d=1 simplices containing the upper and lower limits for a simplex to be interior/*)
- *on boundary of the alpha shape.*

2.1.3 Filter

Author Sebastian Reinhard

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

```
class impro.analysis.filter.Filter
```

Bases: object

```
static local_density_filter(points, r, n)
```

Get the indices of all points having a minimum of n neighbors in a radius of r

Parameters

- **points** (*np.array*) – Points to filter (X,Y,Z)

- **r** (*float*) – Search for neighbors in a radius of r
- **n** (*int*) – Required neighbors in radius r to pass the filter

Returns `filt` – indices of entries passing the filter

Return type np.array

static values_filter (*value, minimum, maximum*)

Get the indices of an array values with value > minimum and value < maximum

Parameters

- **value** (*np.array*) – Array with values to filter
- **minimum** (*float*) – minimum value to pass the filter
- **maximum** (*float*) – maximum value to pass the filter

Returns `filt` – indices of entries passing the filter

Return type np.array

2.1.4 Hough Transform

The General Hough Transform (GHT) maps the orientation of edge points in a template image to a predefined origin (typically the middle pixel of the image). Comparing this map with an image gives each point a rating of likelihood for being a source point of that map.

Author Sebastian Reinhard

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

Example

```
>>> target = cv2.imread("path_to_file")
>>> template = cv2.imread("path_to_file")
>>> ght_target = GHTImage(target, blur=5, canny_lim=(130,180))
>>> H = HoughTransform()
>>> H.target = ght_target
>>> ght_template = TemplateImage(template)
>>> H.template = ght_template
>>> res = H.transform()
```

```
class impro.analysis.hough_transform_gpu.GHTImage(image, blur=4, canny_lim=(130, 200))
```

Bases: object

GHT specific image preprocessing

o_image

Input image

Type np.array

image

Resized and blurred input image

Type np.array

```
canny
    Canny Edge processed image
        Type np.array

gradient
    Gradient image
        Type np.array

class impro.analysis.hough_transform_gpu.HoughTransform(rotation_min=-10, rotation_max=10)
    Bases: object
    Perform Gradient Weighted General Hough Transform on the Graphics Processing Unit

rotation_min
    Start matching template at rotation_min
        Type float

rotation_max
    End matching template at rotation_max
        Type float

template
    Matching template image on target image
        Type np.array

target
    Matching template image on target image
        Type np.array

weighted
    Weight the GHT by Gradient density
        Type bool

transform()
    Perform GHT algorithm with given data

rotation_max
rotation_min
target
template
transform()

class impro.analysis.hough_transform_gpu.Matrix(array)
    Bases: object
    Wrapper class for matrix and matrixf struct on GPU:

    get()

class impro.analysis.hough_transform_gpu.TemplateImage(image, **kwargs)
    Bases: impro.analysis.hough\_transform\_gpu.GHTImage
    Extend GHTImage by tempte properties

    o_image
        Input image
```

Type np.array

image
Resized and blurred input image

Type np.array

canny
Canny Edge processed image

Type np.array

gradient
Gradient image

Type np.array

r_matrix_zero
Vector mapping of edge points to a predefined origin

Type np.array

r_matrix
Rotated r_matrix_zero

Type np.array

rotate_r_matrix(angle)
Rotate R-Matrix by angle rad

angle: float Angle to rotate matrix in rad

2.1.5 Module contents

2.2 Data module

2.2.1 Image Factory

Author Sebastian Reinhard

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

class impro.data.image_factory.**ImageFactory**
Bases: object

Interface to File readers

classmethod **create_image_file(path: str)**

classmethod **create_storm_file(path: str)**

2.2.2 Image

Author Sebastian Reinhard

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

```
class impro.data.image.ImageReader(path)
Bases: object

    Read Image file

    Possible formats are .czi, .lsm, .tiff, .png

    path
        Path to file
        Type str

    data
        Image data as array
        Type np.array

    metaData
        Meta Data of image file containing computational dimensions: {ShapeSizeX, ShapeSizeY, ShapeSizeZ, ShapeSizeC} and physical dimensions: SizeX(Pixel Size in X direction)...
        Type np.array
```

Example

```
>>> file = ImageReader("path_to_file.czi")
>>> file.parse()
>>> file.data.shape
(4, 103, 2430, 2430)
```

```
parse(calibration_px=1.0)
    Read data/Metadata from file

reset_data()
    resets data

set_calibration(px: float)
    Set pixel size manually

class impro.data.image.LocalisationReader(path: str)
Bases: object

    Read RapidSTORM files

    Rewritten class to read STORM/dSTORM data from a text file in a numpy array. Improved performance and logic. Can transform the localizations if a matrix is given. Receives metadata from file header.

    path
        Path to Rapidstorm .txt file
        Type str

    stormData
        Restructured data: X, Y, Precision, Z, Emission, Frame #todo switch Precision and Z
        Type np.array

    size
        X,Y size of data in nanometer
        Type np.array
```

Example

```
>>> file = LocalisationReader("path_to_Rapidstormfile.txt")
>>> file.parse()
>>> data = file.stormData
>>> data.shape
(100000, 6)
```

`parse()`

Read dSTORM data/Metadata from file

`reset_data()`

Resets dSTORM data

`transformAffine(path=None, src=None, dst=None)`

Transform dSTORM data affine with Landmarks file

Parameters

- `path (str, optional)` – path to file containing source points in column 1,2 (starting by 0) and target points in column 3,4
- `src ((, 2) ndarray)` – X,Y coordinates of source points
- `dst ((, 2) ndarray)` – X,Y coordinates of destination points

`class impro.data.image.StormReader(input_loc_path)`

Bases: object

Internal Class. Find dSTORM columns with regular expressions

`clear()`

`get_header_info()`

`readfile()`

`save_loc_file(path, file)`

2.2.3 Module contents

2.3 Render module

2.3.1 Common module

Common module

Buffer

A couple of buffer objects for GPU rendering.

`class impro.render.common.Buffer.Framebuffer`

Bases: object

`build(renderbuf, texbuf)`

`delete()`

```
class impro.render.common.Buffer.Renderbuffer
Bases: object

build(width, height)
delete()

class impro.render.common.Buffer.Texturebuffer
Bases: object

build(width, height)
delete()
```

Objects

A couple of mesh objects for GPU rendering.

```
class impro.render.common.Objects.Cube
Bases: object

class impro.render.common.Objects.Quad
Bases: object

class impro.render.common.Objects.Surface
Bases: object

class impro.render.common.Objects.Texture
Bases: object

class impro.render.common.Objects.UnitCube
Bases: object
```

Surface

A couple of texture mappings for GPU rendering.

```
class impro.render.common.Surface.CSurface
Bases: object

build_surface(width, height, numComponents, numTargets)

class impro.render.common.Surface.Surface
Bases: object

class impro.render.common.Surface.Volume(width, height, depth)
Bases: object
```

Textures

```
class impro.render.common.Textures.Create2DTexture
Bases: object

set_texture(data, interpolation)

class impro.render.common.Textures.Create3DTexture
Bases: object

set_texture(data)
```

```
class impro.render.common.Textures.CreateNoise
Bases: object
    get_texture_handle()
    set_texture(width, height)

class impro.render.common.Textures.CreateVolumeTexture
Bases: object
    set_texture(data)
```

Module contents

2.3.2 Render

Render multiple objects on GPU(points, image2D, image3D)

Author *Sebastian Reinhard*

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

2.3.3 Shader

Create and compile a program from vertex, fragment and geometry shader

Author *Sebastian Reinhard*

Organization Biophysics and Biotechnology, Julius-Maximilians-University of Würzburg

Version 2019.06.26

```
class impro.render.shader.shader(filename)
Bases: object

    Parameters filepath – path to vertex fragment and (optional) geometry shaders. should all have
        the same name

    Variables m_program – reference number to program

    attribute_loc(name)

    create_shader(shader, type)
        Compile OpenGL shaders string and check for shaders errors :param shader: String of shaders
        :param type: Type of shaders can be: GL_VERTEX_SHADER, GL_FRAGMENT_SHADER,
        GL_GEOMETRY_SHADER :return: Shader ID

    enums

    link_program()
        Link program :raises RuntimeError:

    load_shader(filename)
        Load shaders string from file :param filename: Full path to shaders :return: File as string

    set_uniform(name, data)

    uniform_loc(name)
```

```
validate_program()  
    Validate program :raises RuntimeError:
```

2.3.4 Module contents

2.4 Impro

2.4.1 impro package

Subpackages

impro.libs package

Submodules

impro.libs.czifile module

Read image and metadata from Carl Zeiss(r) ZISRAW (CZI) files.

CZI is the native image file format of the ZEN(r) software by Carl Zeiss Microscopy GmbH. It stores multidimensional images and metadata from microscopy experiments.

Author Christoph Gohlke

Organization Laboratory for Fluorescence Dynamics, University of California, Irvine

Version 2017.09.12

Requirements

- CPython 3.6 64-bit
- Numpy 1.13
- Scipy 0.19
- Tifffile.py 2017.09.12
- Czifile.pyx 2017.07.20 (for decoding JpegXrFile and JpgFile images)

Revisions

2017.09.12 Require tifffile.py 2017.09.12

2017.07.21 Use multi-threading in CziFile.asarray to decode and copy segment data. Always convert BGR to RGB.
Remove bgr2rgb options. Decode JpegXR directly from byte arrays.

2017.07.13 Add function to convert CZI file to memory-mappable TIFF file.

2017.07.11 Add ‘out’ parameter to CziFile.asarray. Remove memmap option from CziFile.asarray (backwards incompatible). Change spline interpolation order to 0 (backwards incompatible). Make axes return a string.
Require tifffile 2017.07.11.

2015.08.17 Require tifffile 2015.08.17.

2014.10.10 Read data into a memory mapped array (optional).

2013.12.04 Decode JpegXrFile and JpgFile via _czifile extension module. Attempt to reconstruct tiled mosaic images.

2013.11.20 Initial release.

Notes

The API is not stable yet and might change between revisions.

The file format design specification [1] is confidential and the licence agreement does not permit to write data into CZI files.

Only a subset of the 2012 specification is implemented in the initial release. Specifically, multifile images are not yet supported.

Tested on Windows with a few example files only.

References

- (1) ZISRAW (CZI) File Format Design specification Release Version 1.2.2. CZI 07-2016/CZI-DOC ZEN 2.3/DS_ZISRAW-FileFormat.pdf (confidential). Documentation can be requested at <http://microscopy.zeiss.com/microscopy/en_us/downloads/zen.html>
- (2) CZI The File Format for the Microscope | ZEISS International <http://microscopy.zeiss.com/microscopy/en_us/products/microscope-software/ zen-2012/czi.html>

Examples

```
>>> with CziFile('test.czi') as czi:  
...     image = czi.asarray()  
>>> image.shape  
(3, 3, 3, 250, 200, 3)  
>>> image[0, 0, 0, 0, 0]  
array([10, 10, 10], dtype=uint8)
```

impro.libs.czifile.**imread**(filename, *args, **kwargs)

Return image data from CZI file as numpy array.

‘args’ and ‘kwargs’ are arguments to the CziFile.asarray function.

Examples

```
>>> image = imread('test.czi')  
>>> image.shape  
(3, 3, 3, 250, 200, 3)  
>>> image.dtype  
dtype('uint8')
```

class impro.libs.czifile.CziFile(arg, multifile=True, filesize=None, detectmosaic=True)

Bases: object

Carl Zeiss Image (CZI) file.

header

Global file metadata such as file version and GUID.

Type FileHeaderSegment

metadata

Global image metadata in UTF-8 encoded XML format.

Type etree.ElementTree.Element

All attributes are read-only.

asarray (*resize=True, order=0, out=None, max_workers=None*)

Return image data from file(s) as numpy array.

Parameters

- **resize** (*bool*) – If True (default), resize sub/supersampled subblock data.
- **order** (*int*) – The order of spline interpolation used to resize sub/supersampled sub-block data. Default is 0 (nearest neighbor).
- **out** (*numpy.ndarray, str, or file-like object; optional*) – Buffer where image data will be saved. If numpy.ndarray, a writable array of compatible dtype and shape. If str or open file, the file name or file object used to create a memory-map to an array stored in a binary file on disk.
- **max_workers** (*int*) – Maximum number of threads to read and decode subblock data. By default up to half the CPU cores are used.

attachment_directory

Attribute whose value is computed on first access.

attachments()

Return iterator over all Attachment segments in file.

axes

Attribute whose value is computed on first access.

close()

dtype

Attribute whose value is computed on first access.

filtered_subblock_directory

Attribute whose value is computed on first access.

metadata

Attribute whose value is computed on first access.

save_attachments (*directory=None*)

Save all attachments to files.

segments (*kind=None*)

Return iterator over Segment data of specified kind.

Parameters **kind** (*bytestring or sequence thereof*) – Segment id(s) as listed in SEGMENT_ID. If None (default), all segments are returned.

shape

Attribute whose value is computed on first access.

start

Attribute whose value is computed on first access.

subblock_directory

Attribute whose value is computed on first access.

subblocks ()

Return iterator over all SubBlock segments in file.

Module contents

Module contents

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