

# ImagesToLARModel, a tool for creation of three-dimensional models from a stack of images

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November 6, 2015

## **Abstract**

Here we will present a software for creating a three-dimensional model from a stack of images. This can be useful because of the simplicity of these type of representations. In particular a scope of use can be offered by medicine, where there is an enormous number of images but with very complex two-dimensional representations.

This work will use the LAR representation ([[CL13](#)]) with the Julia language, because of its simplicity, showing how it can be used for quickly process image data.

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# 1 Introduction

This work has the aim to transform a two-dimensional representation of a model (based on a stack of images) into a three-dimensional representation based on the LAR schema. In particular, it will produce a single obj model which can be viewed with standard graphics softwares.

In the past were developed other softwares using same principles (see [PDFJ15]). However, they were optimized for speed and cannot be able to accept huge amounts of data. With the rise of the big data era, we now have more and more data available for research purposes, so softwares must be able to deal with them. A typical hardware environment is based on a cluster of computers where computation can be distributed among a lot of different processes. However, as stated by *Amdahl's law*, the speedup of a program using multiple processors is limited by the time needed for the sequential fraction of the program. So use of parallel techniques for dealing with big data is not important for time performance gain but for memory space gain. In fact, our biggest problem is lack of memory, due to model sizes. As a consequence, every parts of this software is written with the clear objective of minimizing memory usage at the cost of losing something in terms of time performance. So, for example, images will be converted in blocks determined by a grid size (see section 5) among different processes and different machines of the cluster



Figure 1: Amdahl's law

## 1.1 Why Julia

Ricordare che precedenti versioni erano in python

Semplicità

Efficienza

Capacità di realizzare programmi paralleli con poco sforzo

## 2 Software structure

### 2.1 Julia packages

This software will be distributed as a Julia Package. For the actual release (Julia 0.4) a package is a simple git project with the structure showed in figure 2

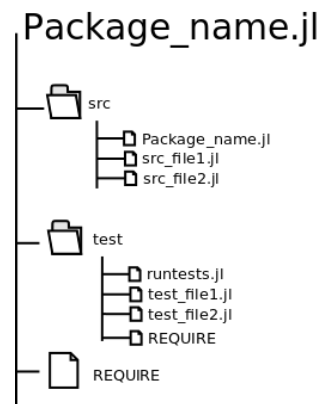


Figure 2: Julia module structure

Source code must be in folder `src`, while in `test` folder there are module tests with a `runtests.jl` for executing them and with a `REQUIRE` file for specifying tests dependencies. For listing dependencies for the entire project, there is another `REQUIRE` file in main folder. As an example in figure 3 there is the `REQUIRE` file for `ImagesToLARMModel.jl`.

After creating this structure for a project it can be pushed on a git repository and installed on Julia systems. The usual installation procedure use this syntax:

```
Pkg.add("Package-name")
```

This will check for that package in METADATA.jl repository on github where there are all official Julia package. However it is also possible to install an unofficial package (on a public git repository) using this syntax:

```
julia 0.3
JSON
Logging
PyCall
Images
Colors
```

Figure 3: REQUIRE contents for `ImagesToLARModel.jl`

```
Pkg.clone("git://repository-address.git")
```

This will install the package on your system with all the dependencies listed in REQUIRE file.

## 2.2 Architecture of ImagesToLARModel

In previous section we have seen how to create a Julia package for distribute our application. Now we focus on the structure of our application. In `src` folder we can find the following modules:

**ImagesToLARModel.jl**: main module for the software, it takes input parameters and start images conversion

**ImagesConversion.jl**: it is called by `ImagesToLARModel.jl` module and controls the entire conversion process calling all other modules

**GenerateBorderMatrix.jl**: it generates the boundary operator for grid specified in input, saving it in a JSON file

**PngStack2Array3dJulia.jl**: it is responsible of images loading and conversion into computable data

**Lar2Julia.jl**: it contains a small subset of LAR functions written in Julia language

**LARUtils.jl**: it contains utility functions for manipulation of LAR models

**Model2Obj.jl**: it contains function that manipulates obj files

**larcc.py**: python larcc module for boundary computation. In next releases of the software it will be rewritten in Julia language

In figure 4 there is a simple schema of dependencies between modules.

Next sections of this document will explain in details all these modules showing also the code involved in conversion

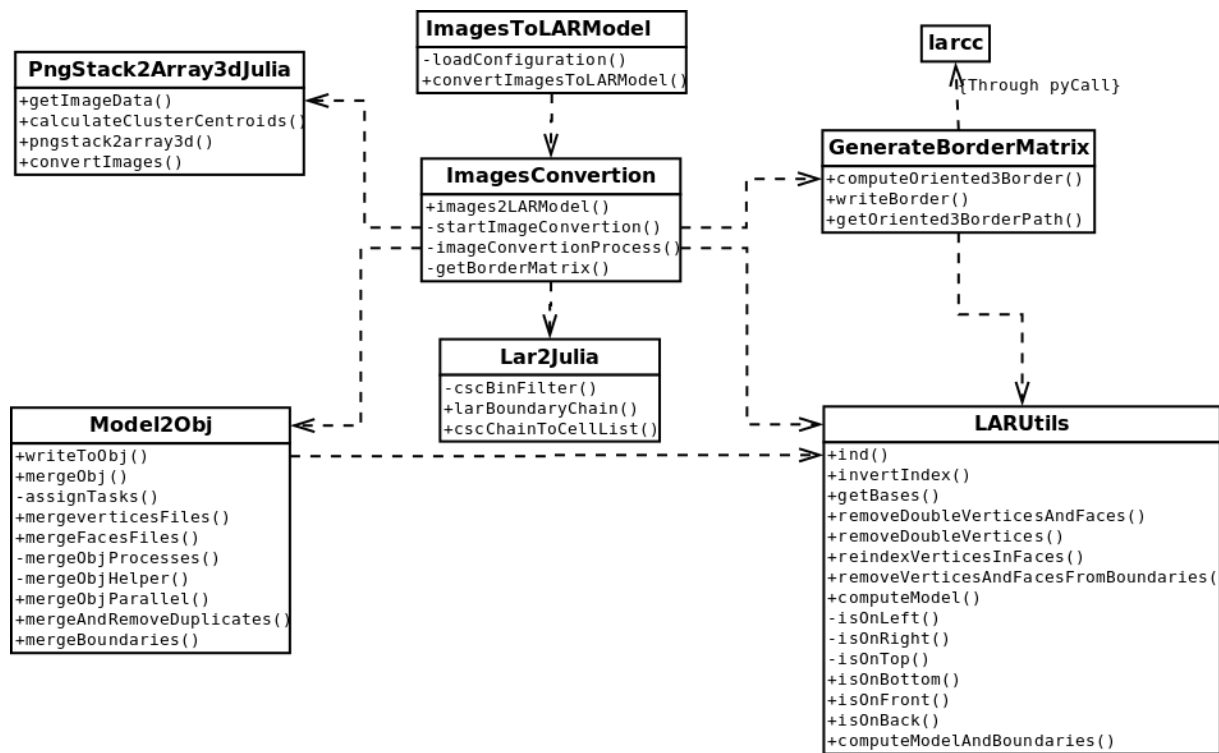


Figure 4: Schema of module dependencies of ImagesToLARModel

### 3 ImagesToLARModel

This is the main module for the application; it takes the input data and start conversion calling `ImagesConversion.jl`.

#### 3.1 Calling modules

As we have already said, this first module has the responsibility of starting the conversion calling all other modules in the package. In Julia calling modules requires that they are in a path specified by `LOAD_PATH` array. So at the beginning of this module we need to add this line:

```

< update load path 6 > ≡
    push!(LOAD_PATH, Pkg.dir("ImagesToLARModel/src"))
    ◇

```

Fragment referenced in 32a.

`Pkg.dir()` function gives us the path of the Julia installation, so `Pkg.dir("ImagesToLARModel/src")` returns “ $\langle Julia - path \rangle / ImagesToLARModel / src$ ”

After this line we can now import all modules defined here and export public functions:

```
 $\langle modules import ImagesToLARModel 7a \rangle \equiv$   
    import JSON  
    import ImagesConversion  
  
    using Logging  
  
    export convertImagesToLARModel  
  
     $\diamond$ 
```

Fragment referenced in 32a.

## 3.2 Input loading

Images conversion takes several parameters:

- `inputDirectory`: The path of the directory containing the stack of images
- `outputDirectory`: The path of the directory containing the output
- `bestImage`: Image chosen for centroid computation (see section 4)
- `nx`, `ny`, `nz`: Sizes of the grid chosen for image segmentation (see section 4)
- `DEBUG.LEVEL`: Debug level for Julia logger
- `parallelMerge` (experimental): Choose between sequential or parallel merge of files (see section 9)

Because of their number it has been realized a function for simply loading them from a JSON configuration file; this is the code:

```
 $\langle load JSON configuration 7b \rangle \equiv$   
function loadConfiguration(configurationFile)  
    """  
        load parameters from JSON file  
  
        configurationFile: Path of the configuration file  
    """
```



```

configuration = JSON.parse(configurationFile)

DEBUG_LEVELS = [DEBUG, INFO, WARNING, ERROR, CRITICAL]

parallelMerge = false
try
    if configuration["parallelMerge"] == "true"
        parallelMerge = true
    else
        parallelMerge = false
    end
catch
end

return configuration["inputDirectory"], configuration["outputDirectory"],
        configuration["bestImage"],
        configuration["nx"], configuration["ny"], configuration["nz"],
        DEBUG_LEVELS[configuration["DEBUG_LEVEL"]],
        parallelMerge

end
◇

```

Fragment referenced in 32a.

A valid JSON file has the following structure:

```

{
    "inputDirectory": "Path of the input directory",
    "outputDirectory": "Path of the output directory",
    "bestImage": "Name of the best image (with extension) ",
    "nx": x grid size,
    "ny": y grid size,
    "nz": border z,
    "DEBUG_LEVEL": julia Logging level (can be a number from 1 to 5)
    "parallelMerge": "true" or "false"
}

```

For example, we can write:

```

{
    "inputDirectory": "/home/juser/IMAGES/"
}

```

```

    "outputDirectory": "/home/juser/OUTPUT/",
    "bestImage": "0009.tiff",
    "nx": 2,
    "ny": 2,
    "nz": 2,
    "DEBUG_LEVEL": 2
}

```

As we can see, in a valid JSON configuration file `DEBUG_LEVEL` can be a number from 1 to 5. Instead, when we explicitly define parameters, `DEBUG_LEVEL` can only be one of the following Julia constants:

- `DEBUG`
- `INFO`
- `WARNING`
- `ERROR`
- `CRITICAL`

### 3.3 Starting conversion

As we have already said, this module has the only responsibility to collect data input and starts other modules. These are the functions that start the process and the only exposed to the application users:

```

⟨ Start conversion from JSON file 9 ⟩ ≡
function convertImagesToLARModel(configurationFile)
    """
        Start conversion of a stack of images into a 3D model
        loading parameters from a JSON configuration file

        configurationFile: Path of the configuration file
    """
    inputDirectory, outputDirectory, bestImage, nx, ny, nz,
        DEBUG_LEVEL, parallelMerge = loadConfiguration(open(configurationFile))
    convertImagesToLARModel(inputDirectory, outputDirectory, bestImage,
        nx, ny, nz, DEBUG_LEVEL, parallelMerge)
end
◇

```

Fragment referenced in 32a.

$\langle \textit{Start manual conversion 10} \rangle \equiv$

```
function convertImagesToLARModel(inputDirectory, outputDirectory, bestImage,
                                nx, ny, nz, DEBUG_LEVEL = INFO, parallelMerge = false)
    """
    Start conversion of a stack of images into a 3D model

    inputDirectory: Directory containing the stack of images
    outputDirectory: Directory containing the output
    bestImage: Image chosen for centroids computation
    nx, ny, nz: Border dimensions (Possibly the biggest power of two of images dimensions)
    DEBUG_LEVEL: Debug level for Julia logger. It can be one of the following:
        - DEBUG
        - INFO
        - WARNING
        - ERROR
        - CRITICAL
    """
    # Create output directory
    try
        mkpath(outputDirectory)
    catch
    end

    Logging.configure(level=DEBUG_LEVEL)
    ImagesConversion.images2LARModel(nx, ny, nz, bestImage,
                                     inputDirectory, outputDirectory, parallelMerge)
end
◇
```

Fragment referenced in 32a.

## 4 PngStack2Array3dJulia

This module has the responsibility of convert a png image into an array of values that will be passed to other modules

### 4.1 Module imports

These are modules needed for this part of the package and the public functions exported

```

⟨ modules import PngStack2Array3dJulia 11a ⟩ ≡
    using Images # For loading png images
    using Colors # For grayscale images
    using PyCall # For including python clustering
    using Logging
    @pyimport scipy.ndimage as ndimage
    @pyimport scipy.cluster.vq as cluster

    NOISE_SHAPE_DETECT=10

    export calculateClusterCentroids, pngstack2array3d, getImageData, convertImages
    ◇

```

Fragment referenced in 57.

We need `Images` and `Colors` packages for manipulating png images and `PyCall` for using Python functions for clustering and filtering images. As a consequence, we need a python environment with `scipy` to be able to run the package

## 4.2 Convert input to png

First thing to do in our program is getting our input folder and convert the stack of images into png format. This process lets us to avoid managing an enormous variety of formats during computation, simplifying code used for transformation.

Conversion needs the following parameters:

- `inputPath`: path of the folder containing the original images
- `outputPath`: path where we will save png images
- `bestImage`: name of the image chosen for centroids computing (see section 4.4)

After conversion `outputPath` will contain our png images and the function will return the new name chosen for the best image.

Now we can examine single parts of conversion process. First of all we need to specify a new name for images, keeping the right order between them; so we need to define a prefix based on number of images:

```

⟨ Define string prefix 11b ⟩ ≡

```

```

imageFiles = readdir(inputPath)
numberOfImages = length(imageFiles)
outputPrefix = ""
for i in 1: length(string(numberOfImages)) - 1
    outputPrefix = string(outputPrefix,"0")
end ◇

```

Fragment referenced in 14.

Next we need to open the single image doing the following operations:

1. Open images using `Images` library (which relies on `ImageMagick`) and save them in greyscale png format
2. if one or both dimensions of the image are odd we need to remove one row (or column) of pixels to make it even. This will be more clear when we will introduce the grid for parallel computation (see section 5)

*⟨ Greyscale conversion 12a ⟩ ≡*

```

rgb_img = convert(Image{ColorTypes.RGB}, img)
gray_img = convert(Image{ColorTypes.Gray}, rgb_img) ◇

```

Fragment referenced in 14.

As we can see, we first need to convert image to RGB and then reconvert to greyscale. Without the RGB conversion these rows will return a stackoverflow error due to the presence of alpha channel

*⟨ Image resizing 12b ⟩ ≡*

```

# resizing images if they do not have even dimensions
dim = size(img)
if(dim[1] % 2 != 0)
    debug("Image has odd x; resizing")
    xrange = 1: dim[1] - 1
else
    xrange = 1: dim[1]
end

if(dim[2] % 2 != 0)
    debug("Image has odd y; resizing")
    yrange = 1: dim[2] - 1
end

```

```

else
    yrange = 1: dim[2]
end

img = subim(gray_img, xrange, yrange) ◇

```

Fragment referenced in 14.

Next we just have to search for the best image and add one image if they are odd (for same reasons we need even image dimensions)

```

⟨ Search for best image 13a ⟩ ≡
    # Searching the best image
    if(imageFile == bestImage)
        newBestImage = string(outputPrefix[length(string(imageNumber)):end],
                                imageNumber, ".png")
    end
    imageNumber += 1 ◇

```

Fragment referenced in 14.

```

⟨ Add one image 13b ⟩ ≡
    # Adding another image if they are odd
    if(numberOfImages % 2 != 0)
        debug("Odd images, adding one")
        imageWidth, imageHeight = getImageData(string(outputPath, "/", newBestImage))

        if(imageWidth % 2 != 0)
            imageWidth -= 1
        end

        if(imageHeight % 2 != 0)
            imageHeight -= 1
        end

        imArray = zeros(UInt8, (imageWidth, imageHeight))
        img = grayim(imArray)
        outputFilename = string(outputPath, "/",
                                outputPrefix[length(string(imageNumber)):end], imageNumber, ".png")
        imwrite(img, outputFilename)
    end ◇

```

Fragment referenced in 14.

Finally this is the code for the entire function:

```
< Convert to png 14 > ≡
function convertImages(inputPath, outputPath, bestImage)
    """
    Get all images contained in inputPath directory
    saving them in outputPath directory in png format.
    If images have one of two odd dimensions, they will be resized
    and if folder contains an odd number of images another one will be
    added

    inputPath: Directory containing input images
    outputPath: Temporary directory containing png images
    bestImage: Image chosen for centroids computation

    Returns the new name for the best image
    """

< Define string prefix 11b >

    newBestImage = ""
    imageNumber = 0
    for imageFile in imageFiles
        img = imread(string(inputPath, imageFile))
        < Greyscale conversion 12a >
        < Image resizing 12b >
        outputFilename = string(outputPath, outputPrefix[length(string(imageNumber)):end],
                                imageNumber, ".png")
        imwrite(img, outputFilename)

        < Search for best image 13a >

    end

    < Add one image 13b >

    return newBestImage
end
◇
```

Fragment referenced in 57.

### 4.3 Getting data from a png

Now we need to load information data from png images. In particular we are interested in getting width and height of an image. As stated in [W3C] document, a standard PNG file contains a *signature* followed by a sequence of *chunks* (each one with a specific type).

The signature always contain the following values:

137 80 78 71 13 10 26 10

This signature indicates that the remainder of the datastream contains a single PNG image, consisting of a series of chunks beginning with an IHDR chunk and ending with an IEND chunk. Every chunk is preceded by four bytes indicating its length.

As we are interested in width and height we need to parse the IHDR chunk. It is the first chunk in PNG datastream and its type field contains the decimal values:

73 72 68 82

The header also contains:

|                    |         |
|--------------------|---------|
| Width              | 4 bytes |
| Height             | 4 bytes |
| Bit depth          | 1 bytes |
| Color type         | 1 byte  |
| Compression method | 1 byte  |
| Filter method      | 1 byte  |
| Interlace method   | 1 byte  |

So for reading width and height we need first 24 bytes; the first eight contain the signature, then we have four bytes for length, four bytes for the type field and eight bytes for information we are interested in. This is the code:

```
< Get image data 15 > ≡  
function getImageData(imageFile)  
    ""  
    Get width and height from a png image  
    ""  
  
    input = open(imageFile, "r")  
    data = readbytes(input, 24)  
  
    if (convert(Array{Int},data[1:8]) != reshape([137 80 78 71 13 10 26 10],8))  
        error("This is not a valid png image")  
    end
```



```

w = data[17:20]
h = data[21:24]

width = reinterpret{Int32, reverse(w)}[1]
height = reinterpret{Int32, reverse(h)}[1]

close(input)

return width, height
end
◇

```

Fragment referenced in 57.

## 4.4 Centroids computation

As we have seen above, this package uses greyscale images for conversion into three-dimensional models and for next steps we need binary images so we can distinguish between the background and the model we want to represent. We can use clustering techniques for obtaining this result. First step is centroids calculation from a chosen image (this choice must be made from the user, because we cannot know in advance what is the best image for finding clusters). Moreover we compute these centroids only for an image and then reuse them when we want to cluster all other images, saving processing time. Actually we need only two centroids, because next steps should only recognize between background and foreground pixels. This is the code used for centroid computation:

```

⟨ Centroid computation 16 ⟩ ≡
function calculateClusterCentroids(path, image, numberOfClusters = 2)
    """
    Loads an image and calculate cluster centroids for segmentation

    path: Path of the image folder
    image: name of the image
    numberOfClusters: number of desired clusters
    """
    imageFilename = string(path, image)

    img = imread(imageFilename) # Open png image with Julia Package

    imArray = raw(img)

    imageWidth = size(imArray)[1]

```

```

imageHeight = size(imArray)[2]

# Getting pixel values and saving them with another shape
image3d = Array{Array{UInt8,2}, 0}

# Inserting page on another list and reshaping
push!(image3d, imArray)
pixel = reshape(image3d[1], (imageWidth * imageHeight), 1)

centroids,_ = cluster.kmeans(pixel, numberOfClusters)

return centroids

end
◇

```

Fragment referenced in 57.

## 4.5 Transform pixels to three-dimensional array

Now we can study the most important part of this module, where images are converted into data usable by other modules for the creation of the three-dimensional model. The basic concept consists in transforming every single pixel in an integer value representing color, and then clustering them all using centroids computed earlier. So, we can obtain a matrix containing only two values (the two centroids) representing background and foreground of the image.

Now we will follow the code. This function uses four parameters

- path: Path of the images directory
- minSlice: First image to read
- maxSlice: Last image to read
- centroids: Array containing centroids for clustering

For every image we want to transform in the interval [minSlice, maxSlice) we have to read it from disk and save pixel informations into a multidimensional Array:

```

⟨ Read raw data 17 ⟩ ≡
    img = imread(imageFilename) # Open png image with Julia Package
    imArray = raw(img) # Putting pixel values into RAW 3d array ◇

```

Fragment referenced in 20.

The `Images.jl` `raw` function, get all pixel values saving them in an Array. In Figure 7 we can see how the array will be like for a sample greyscale image.

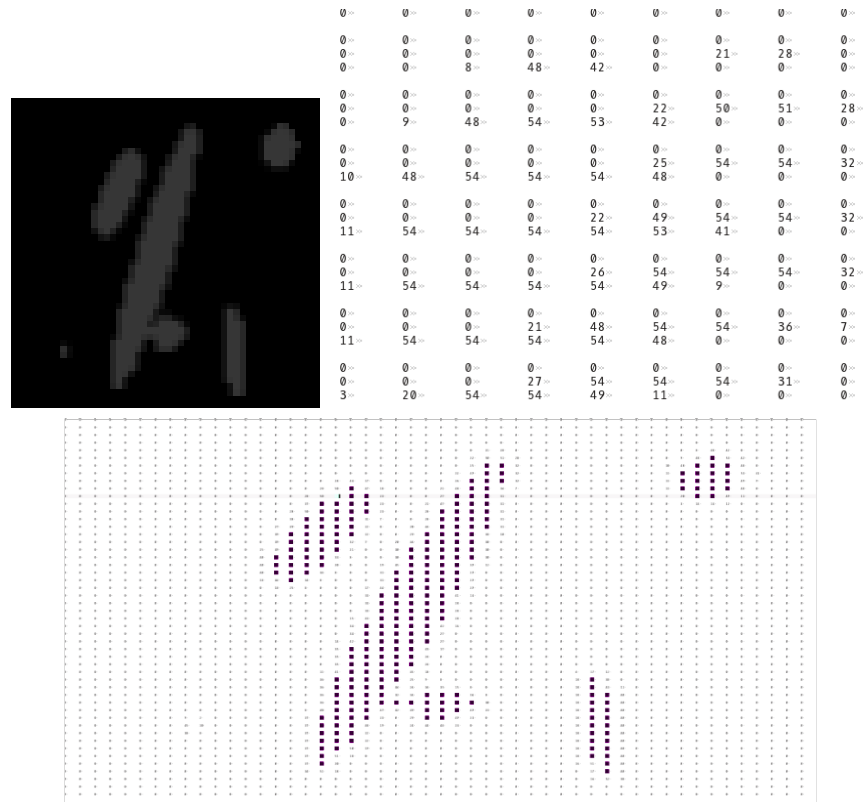


Figure 5: Reading raw data from image. (a) Original greyscale image (b) A view of raw data array (c) The entire raw data array with main color highlighted

Next we have to reduce noise on the image. The better choice is using a *median filter* from package `scipy.ndimage` because it preserves better the edges of the image:

```
⟨ Reduce noise 18 ⟩ ≡
    # Denoising
    image3d[page] = ndimage.median_filter(image3d[page], NOISE_SHAPE_DETECT) ◇
```

Fragment referenced in 20.

Where `image3d` is an array containing all raw data from images

Finally we have to compute clusters (using `scipy`) obtaining images with only two values:

```

< Clustering images 19 > ≡
# Image Quantization
debug("page = ", page)
debug("image3d[page] dimensions: ", size(image3d[page])[1], "\t", size(image3d[page])[2])
pixel = reshape(image3d[page], size(image3d[page])[1] * size(image3d[page])[2] , 1)
qnt,_ = cluster.vq(pixel,centroids)

# Reshaping quantization result
centers_idx = reshape(qnt, size(image3d[page],1), size(image3d[page],2))
#centers_idx = reshape(qnt, size(image3d[page]))

# Inserting quantized values into 3d image array
tmp = Array{UInt8, size(image3d[page],1), size(image3d[page],2))

for j in 1:size(image3d[1],2)
    for i in 1:size(image3d[1],1)
        tmp[i,j] = centroids[centers_idx[i,j] + 1]
    end
end

image3d[page] = tmp ◇

```

Fragment referenced in 20.



Figure 6: Image transformation. (a) Original greyscale image (b) Denoised image (c) Two-colors image

This is the complete code:

*⟨ Pixel transformation 20 ⟩* ≡

```
function pngstack2array3d(path, minSlice, maxSlice, centroids)
    """
    Import a stack of PNG images into a 3d array

    path: path of images directory
    minSlice and maxSlice: number of first and last slice
    centroids: centroids for image segmentation
    """

    # image3d contains all images values
    image3d = Array{Array{UInt8,2}, 0}

    debug("maxSlice = ", maxSlice, " minSlice = ", minSlice)
    files = readdir(path)

    for slice in minSlice : (maxSlice - 1)
        debug("slice = ", slice)
        imageFilename = string(path, files[slice + 1])
        debug("image name: ", imageFilename)
        ⟨ Read raw data 17 ⟩
        debug("imArray size: ", size(imArray))

        # Inserting page on another list and reshaping
        push!(image3d, imArray)

    end

    # Removing noise using a median filter and quantization
    for page in 1:length(image3d)

        ⟨ Reduce noise 18 ⟩

        ⟨ Clustering images 19 ⟩

    end

    return image3d
end
◇
```

Fragment referenced in 57.

## 5 ImagesConversion

Now we will study the most important module for this package: `ImagesConversion`. It has the responsibility of doing the entire conversion process delegating tasks to the other modules.

### 5.1 General algorithm

Now we will examine, in a general way, the algorithm used for conversion from a two-dimensional to a three-dimensional representation of our biomedical models.

We have already seen in section 4 how to get information from a png image, obtaining arrays with only two values; one for the **background** color and one for **foreground** color. This is only the first step of the complete conversion process.

Now we focus only on a single image of the stack. Our two-dimensional representation, consists of pixels of two different colors (where only the one associated with foreground is significant); so we can obtain a three-dimensional representation simply replacing every foreground pixel with a small cube. Focusing on the entire stack of images, the full three-dimensional representation can be obtained simply overlapping all the image representations

This algorithm is very simple, however we does not considered problems concerning lack of memory. In fact, we could have images so big that we cannot build these models entirely in memory; moreover they would require a lot of CPU time for computation. A good solution to these problems consists in taking our representation based on images and divide according to a **grid**.

So, instead of converting the entire model with a unique process, we can subdivide the input among a lot of processes, where every process will execute the conversion process on a small number of **blocks** according to the grid subdivision.

Summing up we can define the following terms, which will be used in next parts of this documentation:

- **Grid:** It is the subdivision of the entire stack of images, with sizes defined by the user. They should be powers of two (for increasing performance during border matrix computation which we will see in section 6)
- **Block:** It is a single cell of the grid
- **xBlock:** It is the x-coordinate of a block
- **yBlock:** It is the y-coordinate of a block



Figure 7: The grid used for parallel computation (a) 2D grid on a single image (b) 3D grid for the stack of images

- **zBlock:** It is the z-coordinate of a block

$xBlock$  and  $yBlock$  are defined on a single image, while  $zBlock$  is defined on different images; in the code it will often be replaced by terms **StartImage** and **EndImage**, which indicate the first image and the last image of that block respectively.

In next subsections we will examine the conversion algorithm in detail, showing what happens for every block of the grid.

## 5.2 Module imports

These are modules needed for this part of the package and the public functions exported.

```
< modules import ImagesConversion 22 > ≡
import GenerateBorderMatrix
import PngStack2Array3dJulia
import Lar2Julia
import Model20bj
import LARUtils

import JSON

using Logging

export images2LARModel
```

◇

Fragment referenced in 32b.

### 5.3 Data preparation

As a first thing, we will see how to prepare our data for conversion process. Firstly we need to convert input images to greyscale png; so we need to create a temporary directory for saving them.

Later, we need to compute the LAR boundary operator for finding boundaries of our cells (for the generation see section 6) getting width and height from our images.

Finally we can start conversion with all these parameters calling `startImageConversion` function, which will be explained in next subsection.

```
< main function for ImagesConversion 23 > ≡
function images2LARModel(nx, ny, nz, bestImage,
                        inputDirectory, outputDirectory, parallelMerge)
    """
    Convert a stack of images into a 3d model
    """

    info("Starting model creation")

    numberOfClusters = 2 # Number of clusters for
                        # images segmentation

    info("Moving images into temp directory")
    try
        mkdir(string(outputDirectory, "TEMP"))
    catch
    end

    tempDirectory = string(outputDirectory, "TEMP/")

    newBestImage = PngStack2Array3dJulia.convertImages(inputDirectory, tempDirectory,
                                                         bestImage)

    imageWidth, imageHeight = PngStack2Array3dJulia.getImageData(
                                string(tempDirectory, newBestImage))
    imageDepth = length(readdir(tempDirectory))

    # Computing border matrix
    info("Computing border matrix")
```



```

try
    mkdir(string(outputDirectory, "BORDERS"))
catch
end
borderFilename = GenerateBorderMatrix.getOriented3BorderPath(
    string(outputDirectory, "BORDERS"), nx, ny, nz)

# Starting images conversion and border computation
info("Starting images conversion")
startImageConversion(tempDirectory, newBestImage, outputDirectory, borderFilename,
    imageHeight, imageWidth, imageDepth,
    nx, ny, nz,
    numberOfClusters, parallelMerge)

end
◇

```

Fragment referenced in 32b.

## 5.4 Images conversion

Now we can see how conversion of images works. First of all we compute the centroids from the best image using module `PngStack2Array3dJulia` and then get the previously computed border matrix in csc sparse array format

```

⟨ compute centroids and get border matrix 24 ⟩ ≡
    # Create clusters for image segmentation
    info("Computing image centroids")
    debug("Best image = ", bestImage)
    centroidsCalc = PngStack2Array3dJulia.calculateClusterCentroids(sliceDirectory,
        bestImage, numberOfClusters)
    debug(string("centroids = ", centroidsCalc))

    try
        mkdir(string(outputDirectory, "BORDERS"))
    catch
    end
    debug("Opening border file: border_", imageDx, "-", imageDy, "-", imageDz, ".json")
    boundaryMat = GenerateBorderMatrix.getBorderMatrix(
        string(outputDirectory, "BORDERS/", "border_",
            imageDx, "-", imageDy, "-", imageDz, ".json"))
◇

```

Fragment referenced in 26.

Now we can start the conversion process.

First of all we need to iterate on the grid finding the zBlock coordinate; we saw earlier that the *imageDz* parameter must be a divisor of the image depth, so we will have exactly *imageDepth/imageDz* blocks on the z coordinate. Moreover, at every zBlock correspond a startImage and an endImage where *endImage - startImage = imageDz*.

Now we can simply parallelize the conversion process spawning a new process for every zBlock, so we open at most *imageDz* images for process.

This is the code for the first part of conversion.

```
< conversion parallelization 25a > ≡
    beginImageStack = 0
    endImage = beginImageStack

    info("Converting images into a 3d model")
    tasks = Array(RemoteRef, 0)
    for zBlock in 0:(imageDepth / imageDz - 1)
        startImage = endImage
        endImage = startImage + imageDz
        info("StartImage = ", startImage)
        info("endImage = ", endImage)

        task = @spawn imageConversionProcess(sliceDirectory, outputDirectory,
                                             beginImageStack, startImage, endImage,
                                             imageDx, imageDy, imageDz,
                                             imageHeight, imageWidth,
                                             centroidsCalc, boundaryMat)

        push!(tasks, task)
    end
    ◇
```

Fragment referenced in 26.

All processes produce a lot of files containing three-dimensional models of a block, so after their termination, we should merge the boundaries between blocks (see later for a better explanation) and merge all model files as we can see in this piece of code:

```
< final file merge 25b > ≡
```

```

info("Merging boundaries")
# Merge Boundaries files
Model2Obj.mergeBoundaries(string(outputDirectory, "MODELS"),
                           imageHeight, imageWidth, imageDepth,
                           imageDx, imageDy, imageDz)

info("Merging obj models")
if parallelMerge
  Model2Obj.mergeObjParallel(string(outputDirectory, "MODELS"))
else
  Model2Obj.mergeObj(string(outputDirectory, "MODELS"))
end
◇

```

Fragment referenced in 26.

This is the final code for this function:

*⟨ start conversion of images 26 ⟩*  $\equiv$

```

function startImageConversion(sliceDirectory, bestImage, outputDirectory, borderFilename,
                              imageHeight, imageWidth, imageDepth,
                              imageDx, imageDy, imageDz,
                              numberOfClusters, parallelMerge)
  """
  Support function for converting a stack of images into a model

  sliceDirectory: directory containing the image stack
  imageForCentroids: image chosen for centroid computation
  """

  ⟨ compute centroids and get border matrix 24 ⟩
  ⟨ conversion parallelization 25a ⟩
  # Waiting for tasks completion
  for task in tasks
    wait(task)
  end
  ⟨ final file merge 25b ⟩
  end
◇

```

Fragment referenced in 32b.

### 5.4.1 Images conversion (for single process)

After we have sketchily studied images conversion, now we will see in details what happens for every process. First thing to do is read an image calling the `PngStack2Array3dJulia`, after that is necessary to sort the centroid array for choosing correct background and foreground pixels.

```
< image read and centroids sort 27a > ≡
    info("Transforming png data into 3d array")
    theImage = PngStack2Array3dJulia.pngstack2array3d(sliceDirectory,
                                                    startImage, endImage, centroids)

    centroidsSorted = sort(vec(reshape(centroids, 1, 2)))
    background = centroidsSorted[1]
    foreground = centroidsSorted[2]
    debug(string("background = ", background, " foreground = ", foreground))
    ◇
```

Fragment referenced in 30b.

Now we can start iterating on other blocks of the grid getting the corresponding slice of the image:

```
< get image slice 27b > ≡
    # Getting a slice of theImage array

    image = Array{UInt8, (convert{Int, length(theImage)},
                          convert{Int, xEnd - xStart}, convert{Int, yEnd - yStart}))
    debug("image size: ", size(image))
    for z in 1:length(theImage)
        for x in 1 : (xEnd - xStart)
            for y in 1 : (yEnd - yStart)
                image[z, x, y] = theImage[z][x + xStart, y + yStart]
            end
        end
    end
    ◇
```

Fragment referenced in 30b.

Here  $xStart$  and  $yStart$  are the absolute coordinates of the model and are calculated from the block coordinates. At the end of this process, we have an array called `image` with size  $(imageDz, imageDx, imageDy)$ .

Now we can get the value of the single pixel into this array and, if it represents a foreground point, put it into an array called `chain3D`. This structure contains indexes of the linearized array created from the matrix. In Figure 8 there is a sample conversion from the matrix to the array

$$\begin{pmatrix} 0^0 & 0^2 \\ 0^1 & 0^3 \end{pmatrix} \begin{pmatrix} 46^4 & 0^6 \\ 46^5 & 46^7 \end{pmatrix} \rightarrow 0^0 \ 0^1 \ 0^2 \ 0^3 \ 46^4 \ 46^5 \ 0^6 \ 46^7$$

$$\begin{pmatrix} 0^0 & 0^2 \\ 0^1 & 0^3 \end{pmatrix} \begin{pmatrix} 0^4 & 46^6 \\ 46^5 & 46^7 \end{pmatrix} \rightarrow 0^0 \ 0^1 \ 0^2 \ 0^3 \ 0^4 \ 46^5 \ 46^6 \ 46^7$$

Figure 8: Transformation of a matrix resulting from a 2x2x2 grid into a linearized array (with cells indexes) (a) First example (b) Second example

As we can see from that figure, from a 2x2x2 grid we can obtain eight values for the single block (or **cell**), where the indexes for the foreground pixels represent indexes of non-empty cells in a 2x2x2 cuboidal geometry

This is the code for getting foreground pixels:

```

⟨ get foreground pixels 28 ⟩ ≡
  nz, nx, ny = size(image)
  chains3D = Array(UInt8, 0)
  zStart = startImage - beginImageStack
  for y in 0:(ny - 1)
    for x in 0:(nx - 1)
      for z in 0:(nz - 1)
        if(image[z + 1, x + 1, y + 1] == foreground)
          push!(chains3D, y + ny * (x + nx * z))
        end
      end
    end
  end
  end
  ◇

```

Fragment referenced in 30b.

Now we have full cells for the geometry, we can convert it into a full *LAR model*. In particular, we are interested in cell boundaries for every block (as we want to obtain only the boundaries for the final model) so we can call function `larBoundaryChain` from

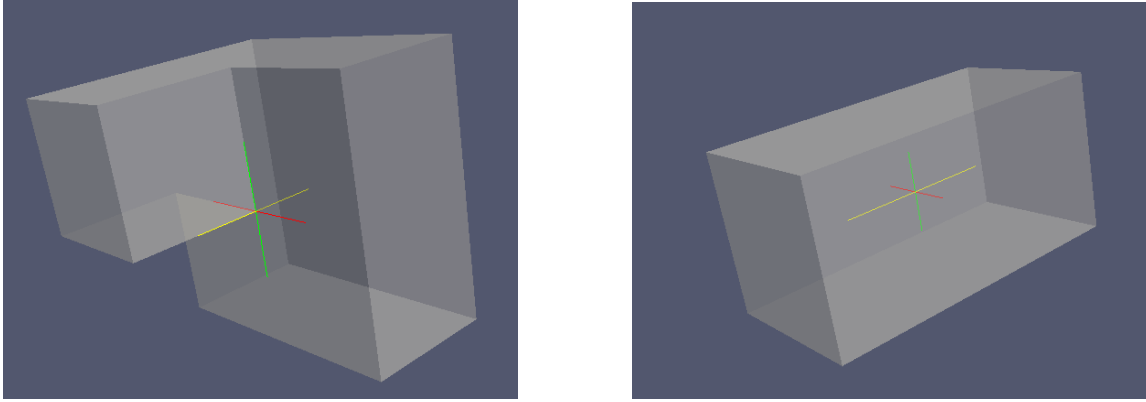


Figure 9: Sample models of 2x2x2 blocks

`Lar2Julia` module (which will be explained in section 7). In Figure 9 there are some examples of models extracted from a single 2x2x2 block.

After model computation, next step is getting vertices and faces from model cells writing results to file. However, as we have already said, we are only interested in boundaries of the final model while now we have only boundaries of a single block. Consequently, we have to separate boundaries from the inner faces of the block on different files (boundaries separation will be explained in section 8). As we can see later, we will merge boundaries together deleting common faces on both block borders, obtaining a model without internal faces. These are pieces of code for getting the inner block model and the boundaries and for file writing:

*⟨get inner model and boundaries 29⟩* ≡

```
# IMPORTANT: inverting xStart and yStart for obtaining correct rotation of the model
models = LARUtils.computeModelAndBoundaries(imageDx, imageDy, imageDz,
                                              yStart, xStart, zStart, objectBoundaryChain)

V, FV = models[1][1] # inside model
V_left, FV_left = models[2][1]
V_right, FV_right = models[3][1] # right boundary
V_top, FV_top = models[4][1] # top boundary
V_bottom, FV_bottom = models[5][1] # bottom boundary
V_front, FV_front = models[6][1] # front boundary
V_back, FV_back = models[7][1] # back boundary
◇
```

Fragment referenced in 30b.

*< write block models to file 30a > ≡*

```
# Writing all models on disk
model_outputFilename = string(outputDirectory, "MODELS/model_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V, FV, model_outputFilename)

left_outputFilename = string(outputDirectory, "MODELS/left_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V_left, FV_left, left_outputFilename)

right_outputFilename = string(outputDirectory, "MODELS/right_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V_right, FV_right, right_outputFilename)

top_outputFilename = string(outputDirectory, "MODELS/top_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V_top, FV_top, top_outputFilename)

bottom_outputFilename = string(outputDirectory, "MODELS/bottom_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V_bottom, FV_bottom, bottom_outputFilename)

front_outputFilename = string(outputDirectory, "MODELS/front_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V_front, FV_front, front_outputFilename)

back_outputFilename = string(outputDirectory, "MODELS/back_output_", xBlock,
                              "-", yBlock, "_", startImage, "_", endImage)
Model2Obj.writeToObj(V_back, FV_back, back_outputFilename)
```

◇

Fragment referenced in 30b.

*< image conversion process 30b > ≡*

```
function imageConversionProcess(sliceDirectory, outputDirectory,
                               beginImageStack, startImage, endImage,
                               imageDx, imageDy, imageDz,
                               imageHeight, imageWidth,
                               centroids, boundaryMat)

    ""
    Support function for converting a stack of image on a single
    independent process
    ""
```

*⟨ image read and centroids sort 27a ⟩*

```
for xBlock in 0:(imageHeight / imageDx - 1)
  for yBlock in 0:(imageWidth / imageDy - 1)
    yStart = xBlock * imageDx
    xStart = yBlock * imageDy
    #xEnd = xStart + imageDx
    #yEnd = yStart + imageDy
    xEnd = xStart + imageDy
    yEnd = yStart + imageDx
    debug("*****")
    debug(string("xStart = ", xStart, " xEnd = ", xEnd))
    debug(string("yStart = ", yStart, " yEnd = ", yEnd))
    debug("theImage dimensions: ", size(theImage)[1], " ",
          size(theImage[1])[1], " ", size(theImage[1])[2])
```

*⟨ get image slice 27b ⟩*

*⟨ get foreground pixels 28 ⟩*

```
if(length(chains3D) != 0)
  # Computing boundary chain
  debug("chains3d = ", chains3D)
  debug("Computing boundary chain")
  objectBoundaryChain = Lar2Julia.larBoundaryChain(boundaryMat, chains3D)
  debug("Converting models into obj")
  try
    mkdir(string(outputDirectory, "MODELS"))
  catch
  end
  ⟨ get inner model and boundaries 29 ⟩
```

*⟨ write block models to file 30a ⟩*

```
else
  debug("Model is empty")
end
end
end
end ◇
```

Fragment referenced in 32b.



## 6 GenerateBorderMatrix

## 7 Lar2Julia

## 8 LARUtils

## 9 Model2Obj

## 10 Exporting the library

### ImagesToLARModel

```
"src/ImagesToLARModel.jl" 32a≡  
    module ImagesToLARModel  
  
        < update load path 6 >  
  
        < modules import ImagesToLARModel 7a >  
        < load JSON configuration 7b >  
        < Start conversion from JSON file 9 >  
        < Start manual conversion 10 >  
    end  
    ◇
```

### ImagesConversion

```
"src/ImagesConversion.jl" 32b≡  
    module ImagesConversion  
  
        < modules import ImagesConversion 22 >  
  
        < main function for ImagesConversion 23 >  
  
        < start conversion of images 26 >  
  
        < image conversion process 30b >  
    end  
    ◇
```

## GenerateBorderMatrix

```
"src/GenerateBorderMatrix.jl" 33≡
module GenerateBorderMatrix

    type MatrixObject
        ROWCOUNT
        COLCOUNT
        ROW
        COL
        DATA
    end

    export computeOriented3Border, writeBorder, getOriented3BorderPath

    import LARUtils
    using PyCall

    import JSON

    @pyimport sys
    unshift!(PyVector(pyimport("sys")["path"]), "") # Search for python modules in folder
    # Search for python modules in package folder
    unshift!(PyVector(pyimport("sys")["path"]), Pkg.dir("ImagesToLARModel/src"))
    @pyimport larcc # Importing larcc from local folder
    @pyimport scipy.sparse as Pysparse

    # Compute the 3-border operator
    function computeOriented3Border(nx, ny, nz)
        """
            Compute the 3-border matrix using a modified
            version of larcc
        """
        V, bases = LARUtils.getBases(nx, ny, nz)
        boundaryMat = larcc.signedCellularBoundary(V, bases)
        return boundaryMat
    end

    function writeBorder(boundaryMatrix, outputFile)
        """
            Write 3-border matrix on json file

            boundaryMatrix: matrix to write on file
            outputFile: path of the outputFile
        """
    end
```

```

rowcount = boundaryMatrix[:shape][1]
colcount = boundaryMatrix[:shape][2]

row = boundaryMatrix[:indptr]
col = boundaryMatrix[:indices]
data = boundaryMatrix[:data]

# Writing informations on file
outfile = open(outputFile, "w")

matrixObj = MatrixObject(rowcount, colcount, row, col, data)
JSON.print(outfile, matrixObj)
close(outfile)

end

function getOriented3BorderPath(borderPath, nx, ny, nz)
    """
    Try reading 3-border matrix from file. If it fails matrix
    is computed and saved on disk in JSON format

    borderPath: path of border directory
    nx, ny, nz: image dimensions
    """

    filename = string(borderPath, "/border_", nx, "-", ny, "-", nz, ".json")
    if !isfile(filename)
        border = computeOriented3Border(nx, ny, nz)
        writeBorder(border, filename)
    end
    return filename
end

function getBorderMatrix(borderFilename)
    """
    TO REMOVE WHEN PORTING OF LARCC IN JULIA IS COMPLETED

    Get the border matrix from json file and convert it in
    CSC format
    """

    # Loading borderMatrix from json file
    borderData = JSON.parsefile(borderFilename)
    row = Array{Int64, length(borderData["ROW"])}
    col = Array{Int64, length(borderData["COL"])}

```

```

data = Array{Int64, length(borderData["DATA"])}

for i in 1: length(borderData["ROW"])
    row[i] = borderData["ROW"][i]
end

for i in 1: length(borderData["COL"])
    col[i] = borderData["COL"][i]
end

for i in 1: length(borderData["DATA"])
    data[i] = borderData["DATA"][i]
end

# Converting csr matrix to csc
csrBorderMatrix = Pysparse.csr_matrix((data,col,row), shape=(borderData["ROWCOUNT"],borderData["COLCOUNT"]), dtype=Int64)
denseMatrix = pycall(csrBorderMatrix["toarray"],PyAny)

cscBoundaryMat = sparse(denseMatrix)

return cscBoundaryMat

end
end
◇

```

## Lar2Julia

```

"src/Lar2Julia.jl" 35≡
module Lar2Julia

export larBoundaryChain, cscChainToCellList

import JSON

using Logging

function larBoundaryChain(cscBoundaryMat, brcCellList)
    """
    Compute boundary chains
    """

    # Computing boundary chains

```

```

n = size(cscBoundaryMat)[1]
m = size(cscBoundaryMat)[2]

debug("Boundary matrix size: ", n, "\t", m)

data = ones(Int64, length(brcCellList))

i = Array{Int64, length(brcCellList)}
for k in 1:length(brcCellList)
    i[k] = brcCellList[k] + 1
end

j = ones{Int64, length(brcCellList)}

debug("cscChain rows length: ", length(i))
debug("cscChain columns length: ", length(j))
debug("cscChain data length: ", length(brcCellList))

debug("rows ", i)
debug("columns ", j)
debug("data ", data)

cscChain = sparse(i, j, data, m, 1)
cscmat = cscBoundaryMat * cscChain
out = cscBinFilter(cscmat)
return out
end

function cscBinFilter(CSCm)
    k = 1
    data = nonzeros(CSCm)
    sgArray = copysign(1, data)

    while k <= nnz(CSCm)
        if data[k] % 2 == 1 || data[k] % 2 == -1
            data[k] = 1 * sgArray[k]
        else
            data[k] = 0
        end
        k += 1
    end

    return CSCm
end

function cscChainToCellList(CSCm)

```

```

"""
Get a csc containing a chain and returns
the cell list of the "+1" oriented faces
"""

data = nonzeros(CSCm)
# Now I need to remove zero element (problem with Julia nonzeros)
nonzeroData = Array{Int64, 0}
for n in data
    if n != 0
        push!(nonzeroData, n)
    end
end

cellList = Array{Int64, 0}
for (k, theRow) in enumerate(findn(CSCm)[1])
    if nonzeroData[k] == 1
        push!(cellList, theRow)
    end
end
return cellList
end
end
◇

```

## LARUtils

```

"src/LARUtils.jl" 37≡
module LARUtils

using Logging

export ind, invertIndex, getBases, removeDoubleVerticesAndFaces, computeModel, computeModelAnd

function ind(x, y, z, nx, ny)
    """
    Transform coordinates into linearized matrix indexes
    """
    return x + (nx + 1) * (y + (ny + 1) * (z))
end

function invertIndex(nx, ny, nz)
    """

```

```

Invert indexes
"""
nx, ny, nz = nx + 1, ny + 1, nz + 1
function invertIndex0(offset)
    a0, b0 = trunc(offset / nx), offset % nx
    a1, b1 = trunc(a0 / ny), a0 % ny
    a2, b2 = trunc(a1 / nz), a1 % nz
    return b0, b1, b2
end
return invertIndex0
end

function getBases(nx, ny, nz)
    """
    Compute all LAR relations
    """

    function the3Dcell(coords)
        x,y,z = coords
        return [ind(x,y,z,nx,ny), ind(x+1,y,z,nx,ny), ind(x,y+1,z,nx,ny), ind(x,y,z+1,nx,ny), ind(x+1,y,z+1,nx,ny),
                ind(x+1,y,z+1,nx,ny), ind(x,y+1,z+1,nx,ny), ind(x+1,y+1,z+1,nx,ny)]
    end

    # Calculating vertex coordinates (nx * ny * nz)
    V = Array{Int64}[]
    for z in 0:nz
        for y in 0:ny
            for x in 0:nx
                push!(V, [x,y,z])
            end
        end
    end

    # Building CV relationship
    CV = Array{Int64}[]
    for z in 0:nz-1
        for y in 0:ny-1
            for x in 0:nx-1
                push!(CV, the3Dcell([x,y,z]))
            end
        end
    end

    # Building FV relationship
    FV = Array{Int64}[]

```

```

v2coords = invertIndex(nx,ny,nz)

for h in 0:(length(V)-1)
    x,y,z = v2coords(h)

    if (x < nx) && (y < ny)
        push!(FV, [h,ind(x+1,y,z,nx,ny),ind(x,y+1,z,nx,ny),ind(x+1,y+1,z,nx,ny)])
    end

    if (x < nx) && (z < nz)
        push!(FV, [h,ind(x+1,y,z,nx,ny),ind(x,y,z+1,nx,ny),ind(x+1,y,z+1,nx,ny)])
    end

    if (y < ny) && (z < nz)
        push!(FV, [h,ind(x,y+1,z,nx,ny),ind(x,y,z+1,nx,ny),ind(x,y+1,z+1,nx,ny)])
    end

end

# Building VV relationship
VV = map((x)->[x], 0:length(V)-1)

# Building EV relationship
EV = Array{Int64}[]
for h in 0:length(V)-1
    x,y,z = v2coords(h)
    if (x < nx)
        push!(EV, [h,ind(x+1,y,z,nx,ny)])
    end
    if (y < ny)
        push!(EV, [h,ind(x,y+1,z,nx,ny)])
    end
    if (z < nz)
        push!(EV, [h,ind(x,y,z+1,nx,ny)])
    end
end

# return all basis
return V, (VV, EV, FV, CV)
end

function lessThanVertices(v1, v2)
    """
    Utility function for comparing vertices coordinates
    """

```



```

    if v1[1] == v2[1]
        if v1[2] == v2[2]
            return v1[3] < v2[3]
        end
        return v1[2] < v2[2]
    end
    return v1[1] < v2[1]
end

function removeDoubleVerticesAndFaces(V, FV, facesOffset)
    """
    Removes double vertices and faces from a LAR model

    V: Array containing all vertices
    FV: Array containing all faces
    facesOffset: offset for faces indices
    """

    newV, indices = removeDoubleVertices(V)
    reindexedFaces = reindexVerticesInFaces(FV, indices, facesOffset)
    newFV = unique(FV)

    return newV, newFV
end

function removeDoubleVertices(V)
    """
    Remove double vertices from a LAR model

    V: Array containing all vertices of the model
    """

    # Sort the vertices list and returns the ordered indices
    orderedIndices = sortperm(V, lt = lessThanVertices, alg=MergeSort)

    orderedVerticesAndIndices = collect(zip(sort(V, lt = lessThanVertices),
                                             orderedIndices))

    newVertices = Array{Array{Int}, 0}
    indices = zeros{Int, length(V)}
    prevv = Nothing
    i = 1
    for (v, ind) in orderedVerticesAndIndices
        if v == prevv
            indices[ind] = i - 1
        else

```

```

        push!(newVertices, v)
        indices[ind] = i
        i += 1
        prevv = v
    end
end
return newVertices, indices
end

function reindexVerticesInFaces(FV, indices, offset)
    """
    Reindex vertices indices in faces array

    FV: Faces array of the LAR model
    indices: new Indices for faces
    offset: offset for faces indices
    """

    for f in FV
        for i in 1: length(f)
            f[i] = indices[f[i] - offset] + offset
        end
    end
    return FV
end

function removeVerticesAndFacesFromBoundaries(V, FV)
    """
    Remove vertices and faces duplicates on
    boundaries models

    V,FV: lar model of two merged boundaries
    """

    newV, indices = removeDoubleVertices(V)
    uniqueIndices = unique(indices)

    # Removing double faces on both boundaries
    FV_reindexed = reindexVerticesInFaces(FV, indices, 0)
    FV_unique = unique(FV_reindexed)

    FV_cleaned = Array{Array{Int}, 0}
    for f in FV_unique
        if(count((x) -> x == f, FV_reindexed) == 1)
            push!(FV_cleaned, f)
        end
    end
end

```

```

end

# Creating an array of faces with explicit vertices
FV_vertices = Array(Array{Array{Int}}, 0)

for i in 1 : length(FV_cleaned)
    push!(FV_vertices, Array(Array{Int}, 0))
    for vtx in FV_cleaned[i]
        push!(FV_vertices[i], newV[vtx])
    end
end

V_final = Array(Array{Int}, 0)
FV_final = Array(Array{Int}, 0)

# Saving only used vertices
for face in FV_vertices
    for vtx in face
        push!(V_final, vtx)
    end
end

V_final = unique(V_final)

# Renumbering FV
for face in FV_vertices
    tmp = Array{Int, 0}
    for vtx in face
        ind = findfirst(V_final, vtx)
        push!(tmp, ind)
    end
    push!(FV_final, tmp)
end

return V_final, FV_final
end

function computeModel(imageDx, imageDy, imageDz,
                      xStart, yStart, zStart,
                      facesOffset, objectBoundaryChain)
    """
    Takes the boundary chain of a part of the entire model
    and returns a LAR model

    imageDx, imageDy, imageDz: Boundary dimensions
    xStart, yStart, zStart: Offset of this part of the model
    """

```

```

facesOffset: Offset for the faces
objectBoundaryChain: Sparse csc matrix containing the cells
"""

V, bases = getBases(imageDx, imageDy, imageDz)
FV = bases[3]

V_model = Array{Array{Int}, 0}
FV_model = Array{Array{Int}, 0}

vertex_count = 1

#b2cells = Lar2Julia.cscChainToCellList(objectBoundaryChain)
# Get all cells (independently from orientation)
b2cells = findn(objectBoundaryChain)[1]

debug("b2cells = ", b2cells)

for f in b2cells
    old_vertex_count = vertex_count
    for vtx in FV[f]
        push!(V_model, [convert{Int}(V[vtx + 1][1] + xStart),
                        convert{Int}(V[vtx + 1][2] + yStart),
                        convert{Int}(V[vtx + 1][3] + zStart)])
        vertex_count += 1
    end

    push!(FV_model, [old_vertex_count + facesOffset, old_vertex_count + 1 + facesOffset, old_v
    push!(FV_model, [old_vertex_count + facesOffset, old_vertex_count + 3 + facesOffset, old_v
end

# Removing double vertices
return removeDoubleVerticesAndFaces(V_model, FV_model, facesOffset)

end

function isOnLeft(face, V, nx, ny, nz)
    """
    Check if face is on left boundary
    """

    for(vtx in face)
        if(V[vtx + 1][2] != 0)
            return false
        end
    end
end

```

```

    return true

end

function isOnRight(face, V, nx, ny, nz)
    ""
    Check if face is on right boundary
    ""

    for(vtx in face)
        if(V[vtx + 1][2] != ny)
            return false
        end
    end
    return true
end

function isOnTop(face, V, nx, ny, nz)
    ""
    Check if face is on top boundary
    ""

    for(vtx in face)
        if(V[vtx + 1][3] != nz)
            return false
        end
    end
    return true
end

function isOnBottom(face, V, nx, ny, nz)
    ""
    Check if face is on bottom boundary
    ""

    for(vtx in face)
        if(V[vtx + 1][3] != 0)
            return false
        end
    end
    return true
end

function isOnFront(face, V, nx, ny, nz)
    ""

```

```

Check if face is on front boundary
"""

for(vtx in face)
    if(V[vtx + 1][1] != nx)
        return false
    end
end
return true
end

function isOnBack(face, V, nx, ny, nz)
    """
    Check if face is on back boundary
    """

    for(vtx in face)
        if(V[vtx + 1][1] != 0)
            return false
        end
    end
    return true
end

function computeModelAndBoundaries(imageDx, imageDy, imageDz,
                                    xStart, yStart, zStart,
                                    objectBoundaryChain)
    """
    Takes the boundary chain of a part of the entire model
    and returns a LAR model splitting the boundaries

    imageDx, imageDy, imageDz: Boundary dimensions
    xStart, yStart, zStart: Offset of this part of the model
    objectBoundaryChain: Sparse csc matrix containing the cells
    """

    function addFaceToModel(V_base, FV_base, V, FV, face, vertex_count)
        """
        Insert a face into a LAR model

        V_base, FV_base: LAR model of the base
        V, FV: LAR model
        face: Face that will be added to the model
        vertex_count: Indices for faces vertices
        """
        new_vertex_count = vertex_count
    end

```

```

    for vtx in FV_base[face]
        push!(V, [convert(Int, V_base[vtx + 1][1] + xStart),
                  convert(Int, V_base[vtx + 1][2] + yStart),
                  convert(Int, V_base[vtx + 1][3] + zStart)])
        new_vertex_count += 1
    end
    push!(FV, [vertex_count, vertex_count + 1, vertex_count + 3])
    push!(FV, [vertex_count, vertex_count + 3, vertex_count + 2])

    return new_vertex_count
end

V, bases = getBases(imageDx, imageDy, imageDz)
FV = bases[3]

V_model = Array{Array{Int}, 0}
FV_model = Array{Array{Int}, 0}

V_left = Array{Array{Int}, 0}
FV_left = Array{Array{Int}, 0}

V_right = Array{Array{Int}, 0}
FV_right = Array{Array{Int}, 0}

V_top = Array{Array{Int}, 0}
FV_top = Array{Array{Int}, 0}

V_bottom = Array{Array{Int}, 0}
FV_bottom = Array{Array{Int}, 0}

V_front = Array{Array{Int}, 0}
FV_front = Array{Array{Int}, 0}

V_back = Array{Array{Int}, 0}
FV_back = Array{Array{Int}, 0}

vertex_count_model = 1
vertex_count_left = 1
vertex_count_right = 1
vertex_count_top = 1
vertex_count_bottom = 1
vertex_count_front = 1
vertex_count_back = 1

#b2cells = Lar2Julia.cscChainToCellList(objectBoundaryChain)
# Get all cells (independently from orientation)

```

```

b2cells = findn(objectBoundaryChain)[1]

debug("b2cells = ", b2cells)

for f in b2cells
    old_vertex_count_model = vertex_count_model
    old_vertex_count_left = vertex_count_left
    old_vertex_count_right = vertex_count_right
    old_vertex_count_top = vertex_count_top
    old_vertex_count_bottom = vertex_count_bottom
    old_vertex_count_front = vertex_count_front
    old_vertex_count_back = vertex_count_back

    # Choosing the right model for vertex
    if(isOnLeft(FV[f], V, imageDx, imageDy, imageDz))
        vertex_count_left = addFaceToModel(V, FV, V_left, FV_left, f, old_vertex_count_left)
    elseif(isOnRight(FV[f], V, imageDx, imageDy, imageDz))
        vertex_count_right = addFaceToModel(V, FV, V_right, FV_right, f, old_vertex_count_right)
    elseif(isOnTop(FV[f], V, imageDx, imageDy, imageDz))
        vertex_count_top = addFaceToModel(V, FV, V_top, FV_top, f, old_vertex_count_top)
    elseif(isOnBottom(FV[f], V, imageDx, imageDy, imageDz))
        vertex_count_bottom = addFaceToModel(V, FV, V_bottom, FV_bottom, f, old_vertex_count_bot)
    elseif(isOnFront(FV[f], V, imageDx, imageDy, imageDz))
        vertex_count_front = addFaceToModel(V, FV, V_front, FV_front, f, old_vertex_count_front)
    elseif(isOnBack(FV[f], V, imageDx, imageDy, imageDz))
        vertex_count_back = addFaceToModel(V, FV, V_back, FV_back, f, old_vertex_count_back)
    else
        vertex_count_model = addFaceToModel(V, FV, V_model, FV_model, f, old_vertex_count_model)
    end
end

# Removing double vertices
return [removeDoubleVerticesAndFaces(V_model, FV_model, 0)],
[removeDoubleVerticesAndFaces(V_left, FV_left, 0)],
[removeDoubleVerticesAndFaces(V_right, FV_right, 0)],
[removeDoubleVerticesAndFaces(V_top, FV_top, 0)],
[removeDoubleVerticesAndFaces(V_bottom, FV_bottom, 0)],
[removeDoubleVerticesAndFaces(V_front, FV_front, 0)],
[removeDoubleVerticesAndFaces(V_back, FV_back, 0)]
end
end
◇

```



## Model2Obj

```
"src/Model2Obj.jl" 48≡
module Model2Obj

import LARUtils

using Logging

export writeToObj, mergeObj, mergeObjParallel

function writeToObj(V, FV, outputFilename)
    """
    Take a LAR model and write it on obj file

    V: array containing vertices coordinates
    FV: array containing faces
    outputFilename: prefix for the output files
    """

    if (length(V) != 0)
        outputVtx = string(outputFilename, "_vtx.stl")
        outputFaces = string(outputFilename, "_faces.stl")

        fileVertex = open(outputVtx, "w")
        fileFaces = open(outputFaces, "w")

        for v in V
            write(fileVertex, "v ")
            write(fileVertex, string(v[1], " "))
            write(fileVertex, string(v[2], " "))
            write(fileVertex, string(v[3], "\n"))
        end

        for f in FV

            write(fileFaces, "f ")
            write(fileFaces, string(f[1], " "))
            write(fileFaces, string(f[2], " "))
            write(fileFaces, string(f[3], "\n"))
        end

        close(fileVertex)
        close(fileFaces)

    end
end
```

```

end

function mergeObj(modelDirectory)
    """
    Merge stl files in a single obj file

    modelDirectory: directory containing models
    """

    files = readdir(modelDirectory)
    vertices_files = files[find(s -> contains(s,string("_vtx.stl")), files)]
    faces_files = files[find(s -> contains(s,string("_faces.stl")), files)]
    obj_file = open(string(modelDirectory, "/", "model.obj"), "w") # Output file

    vertices_counts = Array{Int64, length(vertices_files)}
    number_of_vertices = 0
    for i in 1:length(vertices_files)
        vtx_file = vertices_files[i]
        f = open(string(modelDirectory, "/", vtx_file))

        # Writing vertices on the obj file
        for ln in eachline(f)
            write(obj_file, ln)
            number_of_vertices += 1
        end
        # Saving number of vertices
        vertices_counts[i] = number_of_vertices
        close(f)
    end

    for i in 1 : length(faces_files)
        faces_file = faces_files[i]
        f = open(string(modelDirectory, "/", faces_file))
        for ln in eachline(f)
            splitted = split(ln)
            write(obj_file, "f ")
            if i > 1
                write(obj_file, string(parse(splitted[2]) + vertices_counts[i - 1], " "))
                write(obj_file, string(parse(splitted[3]) + vertices_counts[i - 1], " "))
                write(obj_file, string(parse(splitted[4]) + vertices_counts[i - 1]))
            else
                write(obj_file, string(splitted[2], " "))
                write(obj_file, string(splitted[3], " "))
                write(obj_file, splitted[4])
            end
        end
        write(obj_file, "\n")
    end
end

```

```

        end
        close(f)
    end
    close(obj_file)

    # Removing all tmp files
    for vtx_file in vertices_files
        rm(string(modelDirectory, "/", vtx_file))
    end

    for fcs_file in faces_files
        rm(string(modelDirectory, "/", fcs_file))
    end

end

function assignTasks(startInd, endInd, taskArray)
    """
    This function choose the first files to merge
    creating a tree where number of processes is maximized

    startInd: starting index for array subdivision
    endInd: end index for array subdivision
    taskArray: array containing indices of files to merge for first
    """
    if (endInd - startInd == 2)
        push!(taskArray, startInd)
    elseif (endInd - startInd < 2)
        if (endInd % 4 != 0 && startInd != endInd)
            # Stop recursion on this branch
            push!(taskArray, startInd)
        end
        # Stop recursion doing nothing
    else
        assignTasks(startInd, startInd + trunc((endInd - startInd) / 2), taskArray)
        assignTasks(startInd + trunc((endInd - startInd) / 2) + 1, endInd, taskArray)
    end
end

function mergeVerticesFiles(file1, file2, startOffset)
    """
    Support function for merging two vertices files.
    Returns the number of vertices of the merged file

    file1: path of the first file
    file2: path of the second file
    """

```

```

startOffset: starting face offset for second file
"""

f1 = open(file1, "a")

f2 = open(file2)
debug("Merging ", file2)
number_of_vertices = startOffset
for ln in eachline(f2)
    write(f1, ln)
    number_of_vertices += 1
end
close(f2)

close(f1)

return number_of_vertices
end

function mergeFacesFiles(file1, file2, facesOffset)
    """
    Support function for merging two faces files

    file1: path of the first file
    file2: path of the second file
    facesOffset: offset for faces
    """

    f1 = open(file1, "a")

    f2 = open(file2)
    for ln in eachline(f2)
        splitted = split(ln)
        write(f1, "f ")
        write(f1, string(parse(splitted[2]) + facesOffset, " "))
        write(f1, string(parse(splitted[3]) + facesOffset, " "))
        write(f1, string(parse(splitted[4]) + facesOffset, "\n"))
    end
    close(f2)

    close(f1)
end

function mergeObjProcesses(fileArray, facesOffset = Nothing)
    """

```

Merge files on a single process

fileArray: Array containing files that will be merged

facesOffset (optional): if merging faces files, this array contains

offsets for every file

"""

```
if(contains(fileArray[1], string("_vtx.stl")))
  # Merging vertices files
  offsets = Array(Int, 0)
  push!(offsets, countlines(fileArray[1]))
  vertices_count = mergeVerticesFiles(fileArray[1], fileArray[2], countlines(fileArray[1]))
  rm(fileArray[2]) # Removing merged file
  push!(offsets, vertices_count)
  for i in 3: length(fileArray)
    vertices_count = mergeVerticesFiles(fileArray[1], fileArray[i], vertices_count)
    rm(fileArray[i]) # Removing merged file
    push!(offsets, vertices_count)
  end
  return offsets
else
  # Merging faces files
  mergeFacesFiles(fileArray[1], fileArray[2], facesOffset[1])
  rm(fileArray[2]) # Removing merged file
  for i in 3 : length(fileArray)
    mergeFacesFiles(fileArray[1], fileArray[i], facesOffset[i - 1])
    rm(fileArray[i]) # Removing merged file
  end
end
end
```

function mergeObjHelper(vertices\_files, faces\_files)

"""

Support function for mergeObj. It takes vertices and faces files  
and execute a single merging step

vertices\_files: Array containing vertices files

faces\_files: Array containing faces files

"""

numberOfImages = length(vertices\_files)

taskArray = Array{Int, 0}

assignTasks(1, numberOfImages, taskArray)

# Now taskArray contains first files to merge

numberOfVertices = Array{Int, 0}

tasks = Array{RemoteRef, 0}

```

for i in 1 : length(taskArray) - 1
    task = @spawn mergeObjProcesses(vertices_files[taskArray[i] : (taskArray[i + 1] - 1)])
    push!(tasks, task)
    #append!(numberOfVertices, mergeObjProcesses(vertices_files[taskArray[i] : (taskArray[i + 1] - 1)])
end

# Merging last vertices files
task = @spawn mergeObjProcesses(vertices_files[taskArray[length(taskArray)] : end])
push!(tasks, task)
#append!(numberOfVertices, mergeObjProcesses(vertices_files[taskArray[length(taskArray)] : end])

for task in tasks
    append!(numberOfVertices, fetch(task))
end

debug("NumberOfVertices = ", numberOfVertices)

# Merging faces files
tasks = Array{RemoteRef, 0}()
for i in 1 : length(taskArray) - 1

    task = @spawn mergeObjProcesses(faces_files[taskArray[i] : (taskArray[i + 1] - 1)],
                                    numberOfVertices[taskArray[i] : (taskArray[i + 1] - 1)])
    push!(tasks, task)

    #mergeObjProcesses(faces_files[taskArray[i] : (taskArray[i + 1] - 1)],
    #                  numberOfVertices[taskArray[i] : (taskArray[i + 1] - 1)])
end

#Merging last faces files
task = @spawn mergeObjProcesses(faces_files[taskArray[length(taskArray)] : end],
                                numberOfVertices[taskArray[length(taskArray)] : end])

push!(tasks, task)
#mergeObjProcesses(faces_files[taskArray[length(taskArray)] : end],
#                  numberOfVertices[taskArray[length(taskArray)] : end])

for task in tasks
    wait(task)
end

end

function mergeObjParallel(modelDirectory)
    """

```

Merge stl files in a single obj file using a parallel approach. Files will be recursively merged two by two generating a tree where number of processes for every step is maximized

Actually use of this function is discouraged. In fact speedup is influenced by disk speed. It could work on particular systems with parallel accesses on disks

```

modelDirectory: directory containing models
"""

files = readdir(modelDirectory)

# Appending directory path to every file
files = map((s) -> string(modelDirectory, "/", s), files)

# While we have more than one vtx file and one faces file
while(length(files) != 2)
  vertices_files = files[find(s -> contains(s,string("_vtx.stl")), files)]
  faces_files = files[find(s -> contains(s,string("_faces.stl")), files)]

  # Merging files
  mergeObjHelper(vertices_files, faces_files)

  files = readdir(modelDirectory)
  files = map((s) -> string(modelDirectory, "/", s), files)
end

mergeVerticesFiles(files[2], files[1], 0)
mv(files[2], string(modelDirectory, "/model.obj"))
rm(files[1])

end

function mergeAndRemoveDuplicates(firstPath, secondPath)
  """
  Merge two boundary files removing common faces between
  them

  firstPath, secondPath: Prefix of paths to merge
  """

  firstPathV = string(firstPath, "_vtx.stl")
  firstPathFV = string(firstPath, "_faces.stl")

  secondPathV = string(secondPath, "_vtx.stl")

```

```

secondPathFV = string(secondPath, "_faces.stl")

if(isfile(firstPathV) && isfile(secondPathV))

    V = Array(Array{Int}, 0)
    FV = Array(Array{Int}, 0)

    offset = 0

    # First of all open files and retrieve LAR models

    f1_V = open(firstPathV)
    f1_FV = open(firstPathFV)

    for ln in eachline(f1_V)
        splitted = split(ln)
        push!(V, [parse(splitted[2]), parse(splitted[3]), parse(splitted[4])])
        offset += 1
    end

    for ln in eachline(f1_FV)
        splitted = split(ln)
        push!(FV, [parse(splitted[2]), parse(splitted[3]), parse(splitted[4])])
    end

    close(f1_V)
    close(f1_FV)

    f2_V = open(secondPathV)
    f2_FV = open(secondPathFV)

    for ln in eachline(f2_V)
        splitted = split(ln)
        push!(V, [parse(splitted[2]), parse(splitted[3]), parse(splitted[4])])
    end

    for ln in eachline(f2_FV)
        splitted = split(ln)
        push!(FV, [parse(splitted[2]) + offset, parse(splitted[3]) + offset, parse(splitted[4])])
    end

    close(f2_V)
    close(f2_FV)

    V_final, FV_final = LARUtils.removeVerticesAndFacesFromBoundaries(V, FV)

```



```

    # Writing model to file
    rm(firstPathV)
    rm(firstPathFV)
    rm(secondPathV)
    rm(secondPathFV)
    writeToObj(V_final, FV_final, firstPath)
end
end

function mergeBoundariesProcess(modelDirectory, startImage, endImage,
                                imageDx, imageDy,
                                imageWidth, imageHeight)

    """"
    Helper function for mergeBoundaries.
    It is executed on different processes

    modelDirectory: Directory containing model files
    startImage: Block start image
    endImage: Block end image
    imageDx, imageDy: x and y sizes of the grid
    imageWidth, imageHeight: Width and Height of the image
    """"

    for xBlock in 0:(imageHeight / imageDx - 1)
        for yBlock in 0:(imageWidth / imageDy - 1)

            # Merging right Boundary
            firstPath = string(modelDirectory, "/right_output_", xBlock, "-", yBlock, "_", startImage)
            secondPath = string(modelDirectory, "/left_output_", xBlock, "-", yBlock + 1, "_", startImage)
            mergeAndRemoveDuplicates(firstPath, secondPath)

            # Merging top boundary
            firstPath = string(modelDirectory, "/top_output_", xBlock, "-", yBlock, "_", startImage)
            secondPath = string(modelDirectory, "/bottom_output_", xBlock, "-", yBlock, "_", endImage)
            mergeAndRemoveDuplicates(firstPath, secondPath)

            # Merging front boundary
            firstPath = string(modelDirectory, "/front_output_", xBlock, "-", yBlock, "_", startImage)
            secondPath = string(modelDirectory, "/back_output_", xBlock + 1, "-", yBlock, "_", startImage)
            mergeAndRemoveDuplicates(firstPath, secondPath)
        end
    end
end

function mergeBoundaries(modelDirectory,
                          imageHeight, imageWidth, imageDepth,
                          imageDx, imageDy, imageDz)

```

```

"""
Merge boundaries files. For every cell of size
(imageDx, imageDy, imageDz) in the model grid,
it merges right faces with next left faces, top faces
with the next cell bottom faces, and front faces
with the next cell back faces

modelDirectory: directory containing models
imageHeight, imageWidth, imageDepth: images sizes
imageDx, imageDy, imageDz: sizes of cells grid
"""

beginImageStack = 0
endImage = beginImageStack

tasks = Array{RemoteRef, 0}
for zBlock in 0:(imageDepth / imageDz - 1)
    startImage = endImage
    endImage = startImage + imageDz
    task = @spawn mergeBoundariesProcess(modelDirectory, startImage, endImage,
                                         imageDx, imageDy,
                                         imageWidth, imageHeight)

    push!(tasks, task)
end

# Waiting for tasks
for task in tasks
    wait(task)
end
end
end
◇

```

## PngStack2Array3dJulia

```

"src/PngStack2Array3dJulia.jl" 57≡
module PngStack2Array3dJulia

    { modules import PngStack2Array3dJulia 11a }
    { Convert to png 14 }
    { Get image data 15 }
    { Centroid computation 16 }
    { Pixel transformation 20 }

```

end  
◇

## 10.1 Installing the library

## 11 Conclusions

### 11.1 Results

### 11.2 Further improvements

## References

- [CL13] CVD-Lab, *Linear Algebraic Representation*, Tech. Report 13-00, Roma Tre University, October 2013.
- [PDFJ15] Alberto Paoluzzi, Antonio DiCarlo, Francesco Furiani, and Miroslav Jirik, *CAD models from medical images using LAR*, Computer-Aided Design and Applications **13** (2015), To appear.
- [W3C] W3C, *Portable Network Graphics (PNG) Specification (Second Edition)*, Tech. report.

## A Utility functions

## B Tests

### Generation of the border matrix

```
"test/generateBorderMatrix.jl" 58≡
    push!(LOAD_PATH, "../..")
    import GenerateBorderMatrix
    import JSON
    using Base.Test

    function testComputeOriented3Border()
        """
        Test function for computeOriented3Border
        """
        boundaryMatrix = GenerateBorderMatrix.computeOriented3Border(2,2,2)
```



## Conversion of a png stack to a 3D array

```
"test/pngStack2Array3dJulia.jl" 59≡
    push!(LOAD_PATH, "../..")
    import PngStack2Array3dJulia
    using Base.Test

    function testGetImageData()
        """
        Test function for getImageData
        """

        width, height = PngStack2Array3dJulia.getImageData("images/0.png")

        @test width == 50
        @test height == 50

    end

    function testCalculateClusterCentroids()
        """
        Test function for calculateClusterCentroids
        """

        path = "images/"
        image = 0
        centroids = PngStack2Array3dJulia.calculateClusterCentroids(path, image, 2)

        expected = [0, 253]
        centroids = vec(reshape(centroids, 1, 2))

        @test sort(centroids) == expected
    end

    function testPngstack2array3d()
        """
        Test function for pngstack2array3d
        """

        path = "images/"
        minSlice = 0
        maxSlice = 4
        centroids = PngStack2Array3dJulia.calculateClusterCentroids(path, 0, 2)
        image3d = PngStack2Array3dJulia.pngstack2array3d(path, minSlice, maxSlice, centroids)
```

```

    @test size(image3d)[1] == 5
    @test size(image3d[1])[1] == 50
    @test size(image3d[1])[2] == 200

end

function executeAllTests()
    @time testCalculateClusterCentroids()
    @time testPngstack2array3d()
    @time testGetImageData()
    println("Tests completed.")
end

executeAllTests()

◇

```

## Test for LAR utilities

```

"test/LARUtils.jl" 61≡
push!(LOAD_PATH, "../..")
import LARUtils
using Base.Test

function testInd()
    """
    Test function for ind
    """

    nx = 2
    ny = 2

    @test LARUtils.ind(0, 0, 0, nx, ny) == 0
    @test LARUtils.ind(1, 1, 1, nx, ny) == 13
    @test LARUtils.ind(2, 5, 4, nx, ny) == 53
    @test LARUtils.ind(1, 1, 1, nx, ny) == 13
    @test LARUtils.ind(2, 7, 1, nx, ny) == 32
    @test LARUtils.ind(1, 0, 3, nx, ny) == 28
end

function executeAllTests()
    @time testInd()

```

```
    println("Tests completed.")  
end  
  
executeAllTests()  
  
◇
```