# Package 'CTRing'

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Title Density Profiles of Wood from CT Scan Images
Version 0.1.0
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<b>Description</b> Computerized tomography (CT) can be used to assess certain wood properties when wood disks or logs are scanned. Wood density profiles (i.e. variations of wood density from pith to bark) can yield important information used for studies in forest resource assessment, wood quality and dendrochronology studies. The first step consists in transforming grey values from the scan images to density values. The packages then proposes a unique method to automatically locate the pith by combining an adapted Hough Transform method and a one-dimensional edge detector. Tree ring profiles (average ring density, early wood and latewood density, ring width and percent latewood for each ring) are then obtained.
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Add ring to pith to bark profile from CT scan image

# Description

addRingFromImage

Add ring to pith to bark profile from CT scan image

# Usage

```
addRingFromImage(n = 1, densProfile, im)
```

#### **Arguments**

n Number of rings to adddensProfile Density profileim Density matrix

# Value

Corrected density profile with new ring(s) added and blue bar in plot of added ring

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#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")</pre>
dcm <- readDICOM(file_path)</pre>
hdr_df <- dcm$hdr[[1]]</pre>
image_info <- getImageInfo(hdr = hdr_df)</pre>
im <- imageToMatrix(dcm$img)</pre>
im_8bit <- xBitTo8Bit(im, image_info$grayScale)</pre>
image_info <- getImageInfo(hdr = hdr_df)</pre>
im_dens <- grayToDensity(im_8bit)</pre>
pith_coord <- detect_pith(im_dens,</pre>
                             n_segments = 12,
                             pixel = TRUE,
                             toPlot = FALSE)
endPath <- c(472, 284)
densPath <- extractProfile(im_dens,</pre>
                              image_info,
                              pith_coord,
                              endPath,
                              k = 2, r = 5,
                              threshold = 0.002)
newPath2 <- addRingFromImage(n = 1, densPath, im_dens)</pre>
```

addRingFromProfile

Add ring to pith to bark profile from profile plot

#### **Description**

Add ring to pith to bark profile from profile plot

#### Usage

```
addRingFromProfile(n = 1, densProfile)
```

# **Arguments**

n Number of rings to add densProfile Density profile

#### Value

Corrected density profile with new ring(s) added and blue bar in plot of added ring

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#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)
plotProfile(path)
newPath <- addRingFromProfile(n = 1, path)</pre>
```

addYears

Add years to series

#### **Description**

Add years to series

#### Usage

```
addYears(lastYear, densProfile)
```

# **Arguments**

lastYear Last year of series densProfile Density profile

#### Value

Density profile with years

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)</pre>
```

calcAvgDens 5

```
im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)

endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)

path <- addYears(2021, path)

Calculate average wood, earlywood and latewood density for every ring</pre>
```

# Description

Calculate average wood, earlywood and latewood density for every ring

#### Usage

```
calcAvgDens(densProfile)
```

#### **Arguments**

densProfile Density profile

#### Value

List with several vectors

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image</pre>
```

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```
path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)

pathEwLw <- getEwLw(path)
plotProfile(pathEwLw)
path_avgDens <- calcAvgDens(pathEwLw)
names(path_avgDens)</pre>
```

checkProfile

Verify position of ring transitions of a density profile

# **Description**

Verify position of ring transitions of a density profile

#### Usage

```
checkProfile(profile_with_borders, totRings)
```

#### **Arguments**

# Value

xRing profile with corrected ring location

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deleteRingFromImage

Add ring to pith to bark profile from CT scan image

# Description

Add ring to pith to bark profile from CT scan image

#### Usage

```
deleteRingFromImage(n = 1, densProfile, im)
```

#### **Arguments**

n Number of rings to remove

densProfile Density profile im Density matrix

#### Value

Corrected density profile with ring(s) removed and red bar in plot of deleted ring

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
densPath <- extractProfile(im_dens, image_info, pith_coord,</pre>
```

deleteRingFromProfile

deleteRingFromProfile Delete ring from a pith to bark profile

#### **Description**

Delete ring from a pith to bark profile

#### Usage

```
deleteRingFromProfile(n = 1, densProfile)
```

#### **Arguments**

```
n Number of rings to remove densProfile Density profile
```

#### Value

Corrected density profile with ring(s) removed and red bar in plot of deleted ring

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")</pre>
dcm <- readDICOM(file_path)</pre>
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)</pre>
im <- imageToMatrix(dcm$img)</pre>
im_8bit <- xBitTo8Bit(im, image_info$grayScale)</pre>
im_dens <- grayToDensity(im_8bit)</pre>
pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)</pre>
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image</pre>
densPath <- extractProfile(im_dens,</pre>
                              image_info,
                              pith_coord,
                              endPath,
                              k = 2, r = 5,
```

densityDataFrame 9

```
threshold = 0.002)
```

```
plotProfile(densPath)
newPath <- addRingFromProfile(n = 1, densPath)
oldPath <- deleteRingFromProfile(n = 1, newPath)</pre>
```

densityDataFrame

Convert to dataframe

#### **Description**

Convert to dataframe

# Usage

```
densityDataFrame(densProfile, sampleID = "NoID", addTransitionType = FALSE)
```

# Arguments

```
densProfile Density profile
sampleID Sample ID
addTransitionType
add transition type to dataframe
```

#### Value

Dataframe with cambial age, density, years, transition type

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)
pathEwLw <- getEwLw(path)
plotProfile(pathEwLw)</pre>
```

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```
path_avgDens <- calcAvgDens(pathEwLw)
densityDf <- densityDataFrame(pathEwLw)</pre>
```

 ${\tt detect\_pith}$ 

Automatically detect pith in a CT scan image

# Description

Automatically detect pith in a CT scan image

# Usage

```
detect_pith(
  im,
  toPlot = TRUE,
  n_segments = 25,
  flag = TRUE,
  x_0 = 0.5,
  y_0 = 0.5,
  n_run_max = 15,
  threshold = 0.1,
  pixel = TRUE
)
```

# Arguments

im	Matrix of the CT scan image
toPlot	Boolean to plot the location of the pith on the image
n_segments	Number of segements used to locate pith
flag	FALSE if pith location is known
x_0	Estimate of pith location in x
y_0	Estimate of pith location in y
n_run_max	Maximum number of iterations
threshold	Thershold value for identifying ring transition points
pixel	If TRUE, returns x,y coordinates in pixel numbers, else FALSE returns x,y coordinates in relative values of x and y

#### Value

x,y pith coordinates

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#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)</pre>
```

extractProfile

Get profile between two points of the CTScan image matrix

#### **Description**

Get profile between two points of the CTScan image matrix

#### Usage

```
extractProfile(
  im,
  imHeader,
  beginPath,
  endPath,
  r = 10,
  k = 2,
  threshold = 0.01
)
```

#### **Arguments**

im Density matrix imHeader image header

beginPath X,Y coordinates of the start point of the path endPath X,Y coordinates of the start point of the path

r Profile width

k Rolling window width, integer

threshold Threshold value between maximum and minimum density to establish change

of ring

# Value

Density profile

12 getEwLw

#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)</pre>
```

getEwLw

Establish the transition point from earlywood to latewood for a series of rings

#### **Description**

Establish the transition point from earlywood to latewood for a series of rings

# Usage

```
getEwLw(densProfile)
```

#### **Arguments**

densProfile Density profile

#### Value

xRingList with EW to LW transition points with transition type added (1: low number of points in ring; 2: inflexion point estimated by polynomial; 3: min or max are out of range; 4: inflexion point close to min or max; 5: convex-concave)

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)</pre>
```

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```
im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)

endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)

pathEwLw <- getEwLw(path)

densityDf <- densityDataFrame(path)</pre>
```

getImageInfo

Extract from header of CT scan image grayscale number of bits and pixel size

# Description

Extract from header of CT scan image grayscale number of bits and pixel size

# Usage

```
getImageInfo(hdr)
```

#### **Arguments**

hdr

Header dataframe

#### Value

List with grayscale values, and pixel size

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
getImageInfo(hdr = hdr_df)</pre>
```

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grayToDensity

Convert from 8bit gray scale to density

# Description

Convert from 8bit gray scale to density

# Usage

```
grayToDensity(im, a = -0.1321, b = 0.01834)
```

# Arguments

im Matrix of CT scan image in 8bit gray scalea Intercept of the calibration curveb Slope of the calibration curve

#### Value

Matrix of density values

# **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
range(im_8bit)

im_dens <- grayToDensity(im_8bit)
range(im_dens)</pre>
```

imageToMatrix

Convert dicom image to matrix

#### **Description**

Convert dicom image to matrix

# Usage

```
imageToMatrix(img)
```

locatePathEnd 15

#### **Arguments**

img Dicom image

#### Value

Matrix of image

# **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
dim(im)
image(im)</pre>
```

locatePathEnd

Get coordinates of the end of the path on a CT scan image

# Description

Get coordinates of the end of the path on a CT scan image

#### Usage

```
locatePathEnd(im, pithCoord)
```

#### **Arguments**

im CT scan image

pithCoord X,Y coordinates of the pith

#### Value

Coordinates of the end of the path

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)
im <- imageToMatrix(dcm$img)</pre>
```

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```
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image</pre>
```

pithCoordinates

convert pith coordinates from pixels to length units

#### **Description**

convert pith coordinates from pixels to length units

#### Usage

```
pithCoordinates(pith_coord, pixel_size_x, pixel_size_y)
```

# **Arguments**

```
pith_coord Pith coordinates in pixels
pixel_size_x Pixel size in x
pixel_size_y Pixel size in y
```

#### Value

Pixel coordinates in length units

plotImageProfile

Plot scan image, profile path and ring limits

#### **Description**

Plot scan image, profile path and ring limits

# Usage

```
plotImageProfile(densProfile, im)
```

# Arguments

densProfile Density profile im Density matrix

plotProfile 17

#### Value

Plot

#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)
plotProfile(path)
plotImageProfile(path, im_dens)</pre>
```

plotProfile

Plot density profile

# Description

Plot density profile

#### Usage

```
plotProfile(densProfile)
```

#### **Arguments**

```
densProfile Density profile
```

#### Value

Figure

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#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)
plotProfile(path)</pre>
```

relToPixel

Change from relative to fixed pixel coordinate system

#### **Description**

Change from relative to fixed pixel coordinate system

# Usage

```
relToPixel(pith_coord, im)
```

#### **Arguments**

 $\begin{array}{ll} \text{pith\_coord} & \text{Pith coordinates in relative space } (x,\,y) \\ \\ \text{im} & \text{Density matrix} \end{array}$ 

#### Value

Pixel coordinates in number of pixels (x, y)

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removeLastYear

Remove the last year of a profile

#### **Description**

Remove the last year of a profile

#### Usage

```
removeLastYear(densProfile)
```

#### **Arguments**

```
densProfile Density profile
```

#### Value

Density profile with the last year removed

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
endPath <- c(472, 284) # manual
# not run - endPath <- locatePathEnd(im_dens, pith_coord) # using the image

path <- extractProfile(im_dens, image_info, pith_coord, endPath, k = 2, r = 5, threshold = 0.002)
path_last_year_2021 <- addYears(2021, path)
path_last_year_2020 <- removeLastYear(path_last_year_2021)</pre>
```

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verifyPith

Check if pith location is correct

#### **Description**

Check if pith location is correct

#### **Usage**

```
verifyPith(im, pith_coord)
```

# **Arguments**

im Density matrix of image

pith\_coord Pith coordinates

#### Value

Corrected pith coordinates

#### **Examples**

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
im_8bit <- xBitTo8Bit(im, image_info$grayScale)
im_dens <- grayToDensity(im_8bit)

pith_coord <- detect_pith(im_dens, n_segments = 12, pixel = TRUE, toPlot = FALSE)
pith_coord_checked <- verifyPith(im_dens, pith_coord)</pre>
```

xBitTo8Bit

Convert gray scale from measured bits to 8bit

#### **Description**

Convert gray scale from measured bits to 8bit

#### Usage

```
xBitTo8Bit(im, bits)
```

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# **Arguments**

im Matrix of values in x bits

bits Number of bits of the original gray scale

#### Value

Matrix of gray scale values in 8bits

```
library(oro.dicom)
file_path <- system.file("extdata", "disk.dcm", package = "CTRing")
dcm <- readDICOM(file_path)
hdr_df <- dcm$hdr[[1]]
image_info <- getImageInfo(hdr = hdr_df)

im <- imageToMatrix(dcm$img)
range(im)

im_8bit <- xBitTo8Bit(im, image_info$grayScale)
range(im_8bit)</pre>
```

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