





Detection of change in cancer breast tissues from fractal indicators:

A brief introduction

ANR MISTIC

Journées Textures à Vannes October, 10 & 11 2024

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^{*} Computational Modeling, Analysis of Imagery and Numerical Experiments

Tissue density fluctuations in normal vs. cancerous breasts

Overall mammographic density:

 \implies important risk factor for breast cancer radiological assessment

Tissue density fluctuations in normal vs. cancerous breasts

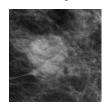
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Local fluctuations: self-similar textures ⇒ fractal analysis for

- classification of mammogram density (Caldwell et al., 1990, Phys. Med. Biol.)
- lesion detectability in mammograms (Burgess et al., 2001, Med. Biol.)
- assessment of breast cancer risk (Heine et al., 2002, Acad. Radiol.)

Mammogram



Tissue density fluctuations in normal vs. cancerous breasts

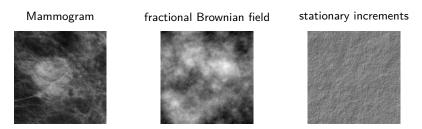
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Fractional Brownian fields: characterized by their local roughness



Motivations and goals

Breast microenvironment plays a crucial role in tumorigenesis:

- ullet structure integrity preserved \Longrightarrow lesions are suppressed
- ullet structure lost by tissue disruption \Longrightarrow tumor is promoted

Tumor vs. healthy not only in the tumor but also in its surrounding tissue

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- tissue disruption
- loss of homeostasis in breast tissue microenvironment
- bilateral asymmetry

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Main idea: quantify density fluctuations through the Hust exponent estimated in multifractal formalism based on 2D Wavelet Transform Modulus Maxima

 \Longrightarrow risk assessment and tumorous breasts detection without seeing a tumor

fBf of Hurst exponent $H \in [0,1]$ denoted $\{B_H(x), x \in \mathbb{R}^2\}$

- Gaussian field with zero-mean
- and for some $\sigma^2 > 0$, correlation function writing

$$\mathbb{E}\left[B_{H}(\boldsymbol{x})B_{H}(\boldsymbol{y})\right] = \frac{\sigma^{2}}{2}\left(\|\boldsymbol{x}\|^{2H} + \|\boldsymbol{y}\|^{2H} - \|\boldsymbol{x} - \boldsymbol{y}\|^{2H}\right)$$

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Stationary increments

$$\forall h \in \mathbb{R}^{2}, \quad \mathbb{E}\left[(B_{H}(x+h) - B_{H}(x))(B_{H}(y+h) - B_{H}(y))\right]$$
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$$= \|\mathbf{x} - \mathbf{y}\|^{2(H-1)}2H(2H - 1)\|\mathbf{h}\|^{2} + o\left(\|\mathbf{h}\|^{2}\right)$$

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- H < 1/2: anti-correlated
- H = 1/2: uncorrelated \Longrightarrow disruption
- H > 1/2: long-range correlated

Self-similarity

$$\forall \boldsymbol{h} \in \mathbb{R}^2, \lambda > 0, \quad B_H(\boldsymbol{x} + \lambda \boldsymbol{h}) - B_H(\boldsymbol{x}) \stackrel{(law)}{\simeq} \lambda^H(B_H(\boldsymbol{x} + \boldsymbol{h}) - B_H(\boldsymbol{x}))$$

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Local regularity: same roughness everywhere $h(x) \equiv H \Longrightarrow$ monofractal signature

The larger the Hurst exponent H, the smoother the texture.

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Singularity spectrum: $\mathcal{D}(h)$ Haussdorff dimension of $\{x \in \mathbb{R}^2, h(x) = h\}$

$$\mathcal{D}(h) = \begin{cases} 2 & h = H \\ -\infty & h \neq H \end{cases}$$

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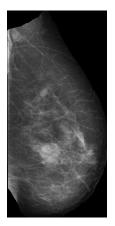
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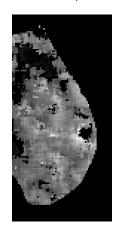
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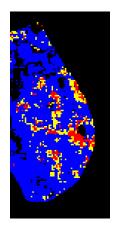
 \implies estimation of $h, \mathcal{D}(h)$: multifractal formalism based on wavelet transform

CompuMAINE local mammogram analysis (Marin et al., 2017, Phys. Med. Biol.)

- H < 1/2 monofractal anti-correlated: fatty tissues (healthy)
- H > 1/2 monofractal long-range correlated: dense tissues (healthy)
- $H \simeq 1/2$ monofractal uncorrelated: disrupted tissues (tumorous)







Dataset: University of South Florida, Digital Database for Screening Mammography

- Mediolateral oblique views only;
- 43 normal, 49 cancer, 35 benign;
- for benign and cancer microcalcification only, masses excluded;

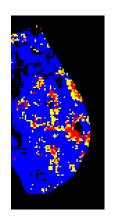


Image sliding-window analysis:

- squared 360 \times 360-pixel window
- with 32-pixel horizontal and vertical shifts
- \Longrightarrow analysis of all 360 \times 360-pixel overlapping patches

Example: mammogram of size 4459×2155 pixels

4457 patches \iff 4457 measures of the roughness H

Metric: number of yellow patches

 $H \sim 1/2 \Longrightarrow$ disrupted tissues

Q.: Is the quantity of disrupted tissues, $H \simeq 1/2$, indicative of a tumorous breast?

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If at least 20 samples, law of S_x well approximated by a Gaussian with

$$\mu = n_x n_y/2; \quad \sigma^2 = n_x n_y (n_x + n_y + 1)/2.$$

If $|S_x - \mu|/\sigma > 1.96$, **H0** is rejected with confidence level $\alpha = 0.05$.

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Tumorous breasts have more disrupted tissues compared to normal breasts:

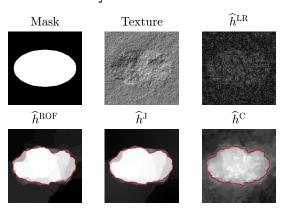
<u>normal vs. cancer:</u> $P \sim 0.0423$, <u>normal vs. benign:</u> $P \sim 0.0009$.

Fractal features piecewise constant estimation from leaders

Pascal et al., 2020, Ann. Telecommun.; Pascal et al., 2021, Appl. Comput. Harmon. Anal.; Pascal et al., 2021, J. Math. Imaging Vis.

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$$\left(\widehat{\boldsymbol{h}}, \widehat{\boldsymbol{v}}\right) (\boldsymbol{\mathcal{L}}; \lambda, \alpha) = \underset{\boldsymbol{h}, \boldsymbol{v}}{\operatorname{argmin}} \sum_{\boldsymbol{h}, \boldsymbol{v}} \left\| \log \boldsymbol{\mathcal{L}}_{\boldsymbol{a}, \cdot} - \log(\boldsymbol{a}) \boldsymbol{h} - \boldsymbol{v} \right\|^2 + \lambda \mathcal{Q}(\boldsymbol{\mathsf{D}} \boldsymbol{h}, \boldsymbol{\mathsf{D}} \boldsymbol{v}; \alpha)$$



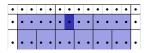
 \Longrightarrow estimation of the local regularity, i.e., roughness, at the **pixel** level

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Wavelet leaders: $\mathcal{L}_{a,n}$ at scale a and pixel \underline{n} supremum of wavelet coefficients

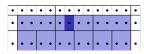
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For a grid of pixels $\Omega\subset\mathbb{R}^2$, scaling exponent au(q) accessible through

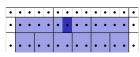
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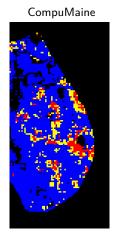
linear regression to estimate H for all 360 \times 360-pixel overlapping patches

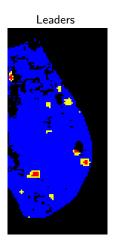
Wavelet leader coefficients (Wendt et al., 2009, Sig. Process.)

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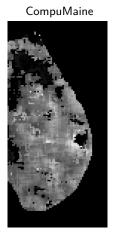
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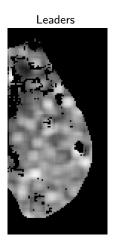




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2D Wavelet Transform: $\{f(x), x \in \mathbb{R}^2\}$ 2D-field

Smoothing function
$$\varphi(\mathbf{x}) \Longrightarrow$$
 wavelets $\psi_1(\mathbf{x}) = \partial_{x_1} \varphi(x_1, x_2), \ \psi_2(\mathbf{x}) = \partial_{x_2} \varphi(x_1, x_2)$

$$\mathbf{T}_{\psi}[f](\boldsymbol{b},a) = \begin{pmatrix} a^{-2} \int \psi_1 \left(a^{-1}(\boldsymbol{x} - \boldsymbol{b}) \right) f(\boldsymbol{x}) \, \mathrm{d}\boldsymbol{x} \\ a^{-2} \int \psi_2 \left(a^{-1}(\boldsymbol{x} - \boldsymbol{b}) \right) f(\boldsymbol{x}) \, \mathrm{d}\boldsymbol{x} \end{pmatrix} \stackrel{\text{(complex)}}{=} \mathbf{M}_{\psi}[f](\boldsymbol{b},a) \exp\left(\mathrm{i}\mathbf{A}_{\psi}[f](\boldsymbol{b},a)\right)$$

Example: Gaussian and Mexican hat smoothing functions

$$\varphi_{\mathsf{Gauss}}(\textbf{\textit{x}}) = \exp(-\|\textbf{\textit{x}}\|^2/2); \quad \varphi_{\mathsf{Mex}}(\textbf{\textit{x}}) = (2 - \|\textbf{\textit{x}}\|^2) \exp(-\|\textbf{\textit{x}}\|^2/2)$$

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Wavelet Transform Modulus Maxima

$$\{(m{b},a)\in\mathbb{R}^2, imes\mathbb{R}_+^* \quad \mathbf{M}_{m{\psi}}[f](m{b},a) ext{ locally maximal in direction } \mathbf{A}_{m{\psi}}[f](m{b},a)\}$$

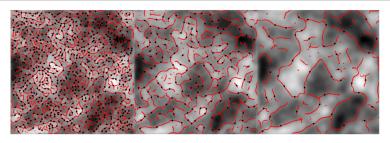


Figure 4.2: The maxima chains are shown for scales $a=2^1\sigma_w$ (left), $a=2^2\sigma_w$ (middle), and $a=2^3\sigma_w$ (right) (where $\sigma_w=7$ pixels) overlaid onto a 2D fBm image with H=0.5. The local maxima along \mathcal{M}_ψ (WTMMM) are shown through small filled black dots.

Source: Basel G. White

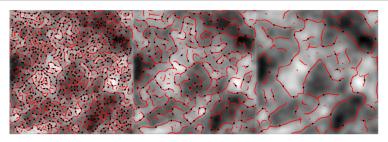


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Wavelet Transform space-scale skeleton: $\mathcal{L}(a)$

lines formed by WTMM maxima across scales

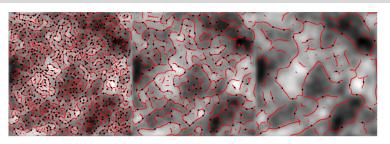


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Wavelet Transform space-scale skeleton: $\mathcal{L}(a)$

lines formed by WTMM maxima across scales

If a maxima line $\mathcal{L}_{x_0}(a)$ is pointing toward a singularity x_0 as $a \to 0^+$, then

$$\mathbf{M}_{\psi}[f](\mathcal{L}_{\mathbf{x}_0}(a)) \sim a^{h(\mathbf{x}_0)}, \quad a \to 0^+$$

provided that the wavelet has $n_{\psi} > h(x_0)$ vanishing moments.

Partition function: for a set $\mathfrak{L}(a)$ of maxima lines

$$\mathcal{Z}(q, a) = \sum_{\ell \in \mathfrak{L}(a)} \left(\sup_{(oldsymbol{b}, a') \in \ell, a' \leq a} oldsymbol{\mathsf{M}}_{\psi}[f](oldsymbol{b}, a')
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Roughness, quantified by Hölder exponent, characterized by $\tau(q)$ spectrum

$$Z(q,a)\sim a^{\tau(q)},\quad a\to 0^+$$

For 2D fractional Brownian field: $\tau(q) = qH - 2$ is **linear**.

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Singularity spectrum: $\mathcal{D}(h)$ Haussdorff dimension of $\{x \in \mathbb{R}^2, \ h(x) = h\}$

$$\mathcal{D}(h) = \min_{q} (qh - \tau(q))$$
 (Legendre transform of τ)

Numerically: unstable estimation of $\tau(q)$ and $\mathcal{D}(q)$

⇒ Mean quantities in a canonical ensemble with Boltzmann weights

$$W_{\psi}[f](q,\ell,a) = \frac{\left| \sup_{(\boldsymbol{b},a') \in \ell, a' \leq a} \mathbf{M}_{\psi}[f](\boldsymbol{b},a') \right|^{q}}{\mathcal{Z}(q,a)}$$

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Roughness: robust local regularity estimation

$$\begin{split} h(q,a) &= \sum_{\ell \in \mathfrak{L}(a)} \ln \left(\mathbf{W}_{\psi}[f](q,\ell,a) \right) \mathbf{W}_{\psi}[f](q,\ell,a), \\ h(q) &= \frac{\mathrm{d}\tau}{\mathrm{d}q} = \lim_{a \to 0^+} \frac{h(q,a)}{\ln a} \end{split}$$

Numerically: unstable estimation of $\tau(q)$ and $\mathcal{D}(q)$

⇒ Mean quantities in a canonical ensemble with Boltzmann weights

$$\mathrm{W}_{\psi}[f](q,\ell,a) = rac{\left| \sup_{(oldsymbol{b},a')\in\ell,a'\leq a} oldsymbol{\mathsf{M}}_{\psi}[f](oldsymbol{b},a')
ight|^q}{\mathcal{Z}(q,a)}$$

Roughness: robust local regularity estimation

$$h(q, a) = \sum_{\ell \in \mathfrak{L}(a)} \ln \left(W_{\psi}[f](q, \ell, a) \right) W_{\psi}[f](q, \ell, a),$$

$$h(q) = \frac{\mathrm{d}\tau}{\mathrm{d}q} = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$$

Singularity spectrum:

$$\mathcal{D}(q,a) = \sum_{\ell \in \mathfrak{L}(a)} \ln \left(\mathrm{W}_{\psi}[f](q,\ell,a) \right) \mathrm{W}_{\psi}[f](q,\ell,a),$$
 $\mathcal{D}(q) = \lim_{a o 0^+} rac{\mathcal{D}(q,a)}{\ln a}$

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Image sliding window analysis

- 1. Check that the central 256×256 pixels are contained in the mask;
- 2. if so, compute the Wavelet Transform for 50 scales, from a=7 to 120 pixels;
- 3. extract the space-scale skeleton from the central 256 $\times\,256$ pixels;
- 4. compute h(q, a) and $\mathcal{D}(q, a)$ from the partition function $\mathcal{Z}(q, a)$;
- 5. linear regressions h(q, a) vs. $\log_2(a)$ and $\mathcal{D}(q, a)$ vs. $\log_2(a)$:

how to choose the range of scales $[a_{min}, a_{max}]$?

For each patch of 360×360 pixels, i.e., 15.5×15.5 mm

roughness:
$$h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$$
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The Autofit Methodology: imposing $\log_2 a_{\text{max}} - \log_2 a_{\text{min}} \ge 1$ explore

$$\log_2 \frac{a_{\min}}{\sigma_w} = 0.0, 0.1, \dots, 2.1, \;, \;\; \log_2 \frac{a_{\max}}{\sigma_w} = 2.0, 2.1, \dots, 4.1, \;\; \text{with} \;\; \sigma_w = 7 \;\; \text{pixels}$$

and select $[a_{\min}, a_{\max}]$ if and only if

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and select $[a_{min}, a_{max}]$ if and only if

• linear regression on h(q = 0, a) from a_{min} to a_{max} yields

$$-0.2 < \hat{h}(q=0) = \hat{H} < 1$$

- $-H \le -0.2$: high roughness \Longrightarrow abnormally high noise
- $H \ge 1$: low roughness, differentiable field \Longrightarrow artificially smooth

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• linear regression on $\mathcal{D}(q=0,a)$ from a_{\min} to a_{\max} yields

$$1.7 < \widehat{\mathcal{D}}(h(q=0)) < 2.5$$

for a monofractal field of Hurst exponent H, expected to be $\mathcal{D}(H)=2$ **but** finite size effect affect the maxima lines as $a \to 0^+$

For each patch of 360×360 pixels, i.e., 15.5×15.5 mm

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and select $[a_{\min}, a_{\max}]$ if and only if

ullet coefficient of determination of linear regression on h(q=0,a) from a_{\min} to a_{\max}

$$R^2 > 0.96$$

sufficiently linear to extract the Hurst exponent H

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and select $[a_{min}, a_{max}]$ if and only if

• weighted standard deviation across q of the $\widehat{h}(q)$ estimated from a_{\min} to a_{\max}

$$sd_w < 0.06$$

 \Longrightarrow excludes multifractal scaling

| q | -2 | -1.5 | -1 | -0.5 | -0.3 | -0.2 | -0.1 | 0 | 0.1 | 0.2 | 0.3 | 0.5 | 1 | 1.5 | 2 | 2.5 | 3 |
|---|-----|------|----|------|------|------|------|----|-----|-----|-----|-----|---|-----|---|-----|-----|
| W | 0.1 | 0.5 | 1 | 3 | 5 | 7 | 9 | 10 | 9 | 8 | 7 | 5 | 3 | 2 | 1 | 0.5 | 0.2 |

For each patch of 360 \times 360 pixels, i.e., 15.5 \times 15.5 mm

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and select $[a_{min}, a_{max}]$ if and only if

• weighted average of goodness of fit of $\widehat{h}(q)$ estimated from a_{\min} to a_{\max}

$$\langle R_w^2 \rangle > 0.96$$

⇒ ensures self-similarity

| - | 1 | -2 | -1.5 | -1 | -0.5 | -0.3 | -0.2 | -0.1 | 0 | 0.1 | 0.2 | 0.3 | 0.5 | 1 | 1.5 | 2 | 2.5 | 3 |
|---|---|-----|------|----|------|------|------|------|----|-----|-----|-----|-----|---|-----|---|-----|-----|
| V | / | 0.1 | 0.5 | 1 | 3 | 5 | 7 | 9 | 10 | 9 | 8 | 7 | 5 | 3 | 2 | 1 | 0.5 | 0.2 |

For each patch of 360×360 pixels:

$$\implies$$
 linear regressions $h(q,a)$ vs. $\log_2(a)$ and $\mathcal{D}(q,a)$ vs. $\log_2(a)$ across $[a_{\min},a_{\max}]$

The Autofit Methodology: imposing $\log_2 a_{\max} - \log_2 a_{\min} \ge 1$ explore 418 couples

$$\log_2 \frac{a_{\min}}{\sigma} = 0.0, 0.1, \dots, 2.1, , \log_2 \frac{a_{\max}}{\sigma} = 2.0, 2.1, \dots, 4.1, \text{ with } \sigma_w = 7 \text{ pixels}$$

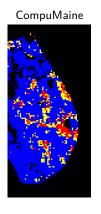
and select $[a_{\min}, a_{\max}]$ if and only if

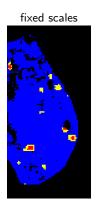
- -0.2 < h(q = 0) < 1: expected roughness
- $1.7 < \widehat{D} < 2.5$: expect 2
- $R^2 > 0.96$: accurate estimation of H
- $sd_w < 0.06$: monofractal scaling
- $\langle R_w^2 \rangle > 0.96$: h(q, a) sufficiently linear

 \implies If no scale range $[a_{\min}, a_{\max}]$ for which all conditions are satisfied: **no scaling**.

Wavelet leader coefficients (Wendt et al., 2009, Sig. Process.)

- H < 1/2 monofractal anti-correlated: fatty tissues (healthy)
- H > 1/2 monofractal long-range correlated: dense tissues (healthy)
- $H \simeq 1/2$ monofractal uncorrelated: disrupted tissues (tumorous)

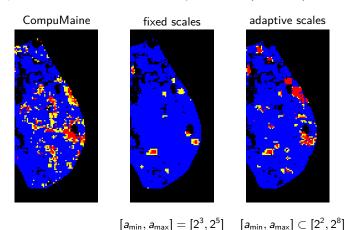




$$[a_{\min}, a_{\max}] = [2^3, 2^5]$$

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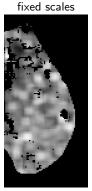
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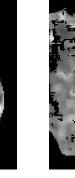


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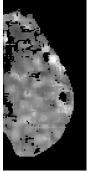
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 $[a_{\min}, a_{\max}] = [2^3, 2^5] \quad [a_{\min}, a_{\max}] \subset [2^2, 2^8]$



adaptive scales

Marin et al., 2017, Phys. Med. Biol.

DDSM: *University of South Florida*, Digital Database for Screening Mammography 43 normal vs. 49 cancer, 35 benign

 \implies digitized films: lossless LJPEG 12-bit images (pixel values: integers in [0, 4095])

Tumorous breasts have more disrupted tissues compared to normal breasts:

<u>normal vs. cancer:</u> $P \sim 0.0423$, normal vs. benign: $P \sim 0.0009$.

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Russian: Perm Regional Oncological Dispensary

81 cancer vs. 23 benign

 \implies digitally acquired mammograms: uncompressed 8-bit BMP images ([0, 255])

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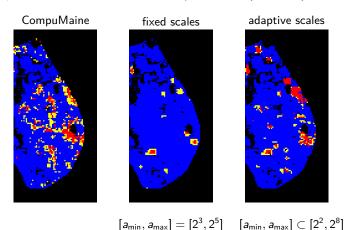
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Wavelet leaders with

- Daubechies wavelets with $n_{\Psi} = 2$ vanishing moments
- $\bullet~\sim$ scales selected by the CompuMaine autofit method, up to rounding errors

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cancer vs. benign: $P \sim 0.074$

Conclusions

Patch-wise fractal analysis of mammograms with WT modulus maxima method

- disrupted tissues, characterized by $H \sim 1/2$, indicate loss of homeostasis
- quantity of disrupted tissues discriminates between

```
(Marin et al., 2017) <u>tumorous vs. normal</u> P \sim 0.0006
(Gerasimova-Chechkina et al., 2021) <u>cancer vs. benign</u> P \sim 0.0030
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 \implies exploration of 418 couples of (a_{\min}, a_{\max}) for each patch and strict conditions

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 \implies exploration of 418 couples of (a_{\min}, a_{\max}) for each patch and strict conditions

Reproduction with wavelet leaders formalism on Russian dataset

- range of scales for each patch extracted from CompuMaine analyses,
- remains less informative: $P \sim 0.0740$

Perspectives

From patch-wise to pixel-wise fractal analysis

- using wavelet leaders framework,
- combined with variational methods,
- with PyTorch implementation to benefit from fast GPU computing,
- reduced number of hyperparameters & fine-tuned automatically

⇒ increase the sensibility in the measurement of the quantity of disrupted tissues

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Asymmetry in tissue disruption in cancerous cases

- assessed both in Marin et al., 2017 and Gerasimova-Chechkina et al., 2021,
- to be evaluated with (pixel-wise) wavelet leader fractal analysis

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Anisotropic Gaussian fields for mammogram modeling

- observed in Richard & Biermé, 2010
- many tools that have never been applied to mammogram yet!