

# Nantes✓ Université



Detection of change in cancer breast tissues from fractal indicators:

A brief introduction

## ANR MISTIC

Journées Textures à Vannes

October, 10 & 11 2024

Barbara Pascal

Laboratoire des Sciences du Numérique de Nantes: B. Pascal

Laboratoire de Physique ENSL: P. Abry, B. Audit, L. Davy, N. Pustelnik

CompuMAINE:\* K. Batchelder, A. Khalil, B. G. White

\* 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

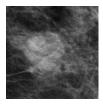
#### **Overall mammographic density:**

 $\implies$  important risk factor for breast cancer radiological assessment

Local fluctuations: self-similar textures  $\Longrightarrow$  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

#### Overall mammographic density:

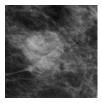
 $\Longrightarrow$  important risk factor for breast cancer radiological assessment

Local fluctuations: self-similar textures  $\Longrightarrow$  fractal analysis for

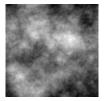
- classification of mammogram density (Caldwell et al., 1990, Phys. Med. Biol.)
- lesion detectability in mammograms (Burgess et al., 2001, Med. Biol.)
- assessment of breast cancer risk (Heine et al., 2002, Acad. Radiol.)

#### Fractional Brownian fields: characterized by their local roughness

Mammogram



fractional Brownian field



stationary increments



# Motivations and goals

Breast microenvironment plays a crucial role in tumorigenesis:

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

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

# Motivations and goals

Breast microenvironment plays a crucial role in tumorigenesis:

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

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

Pioneer work: Marin et al., 2017, Med. Phys. quantitatively and objectively assessed

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

via wavelet-based mammogram local analysis.

# Motivations and goals

Breast microenvironment plays a crucial role in tumorigenesis:

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

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

Pioneer work: Marin et al., 2017, Med. Phys. quantitatively and objectively assessed

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

via wavelet-based mammogram local analysis.

Main idea: quantify density fluctuations through the Hust exponent estimated in

multifractal formalism based on 2D Wavelet Transform Modulus Maxima

 $\implies$  risk assessment and tumorous breasts detection without seeing a tumor

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

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

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

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

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

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

Stationary increments

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

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

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

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

Stationary increments

$$\begin{aligned} \forall \boldsymbol{h} \in \mathbb{R}^2, \quad \mathbb{E}\left[ (B_H(\boldsymbol{x} + \boldsymbol{h}) - B_H(\boldsymbol{x}))(B_H(\boldsymbol{y} + \boldsymbol{h}) - B_H(\boldsymbol{y})) \right] \\ &= \|\boldsymbol{x} + \boldsymbol{h} - \boldsymbol{y}\|^{2H} + \|\boldsymbol{x} - \boldsymbol{h} - \boldsymbol{y}\|^{2H} - 2\|\boldsymbol{x} - \boldsymbol{y}\|^{2H} \end{aligned}$$
  
For  $\|\boldsymbol{h}\| \ll \|\boldsymbol{x} - \boldsymbol{y}\|, \qquad \mathbb{E}\left[ (B_H(\boldsymbol{x} + \boldsymbol{h}) - B_H(\boldsymbol{x}))(B_H(\boldsymbol{y} + \boldsymbol{h}) - B_H(\boldsymbol{y})) \right] \\ &= \|\boldsymbol{x} - \boldsymbol{y}\|^{2(H-1)} 2H(2H-1)\|\boldsymbol{h}\|^2 + o\left(\|\boldsymbol{h}\|^2\right) \end{aligned}$ 

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

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

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

Stationary increments

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

For  $\|\boldsymbol{h}\| \ll \|\boldsymbol{x} - \boldsymbol{y}\|$ ,  $\mathbb{E}[(B_H(\boldsymbol{x} + \boldsymbol{h}) - B_H(\boldsymbol{x}))(B_H(\boldsymbol{y} + \boldsymbol{h}) - B_H(\boldsymbol{y}))]$ =  $\|\boldsymbol{x} - \boldsymbol{y}\|^{2(H-1)}2H(2H-1)\|\boldsymbol{h}\|^2 + o(\|\boldsymbol{h}\|^2)$ 

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

#### Self-similarity

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

#### Self-similarity

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

**Local regularity:** same roughness everywhere  $h(x) \equiv H \implies \text{monofractal signature}$ 

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

#### Self-similarity

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

**Local regularity:** same roughness everywhere  $h(x) \equiv H \implies$  monofractal signature The larger the Hurst exponent H, the smoother the texture.

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

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

#### Self-similarity

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

**Local regularity:** same roughness everywhere  $h(x) \equiv H \implies$  monofractal signature The larger the Hurst exponent H, the smoother the texture.

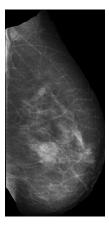
**Singularity spectrum:**  $\mathcal{D}(h)$  Haussdorff dimension of  $\{x \in \mathbb{R}^2, h(x) = h\}$ 

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

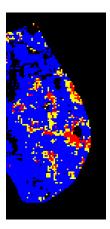
 $\implies$  estimation of  $h, \mathcal{D}(h)$ : multifractal formalism based on wavelet transform

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)

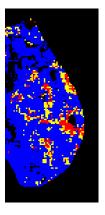






Dataset: University of South Florida, Digital Database for Screening Mammography

- 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  $4459 \times 2155$  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?

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

Wilcoxon rank test a.k.a. Wilcoxon-Mann-Whitney

Independent sets of real numbers X and Y, of cardinalities  $n_x$  and  $n_y$  respectively

**H0**:  $\mathbb{P}(X > Y) = \mathbb{P}(Y > X)$ 

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

Wilcoxon rank test a.k.a. Wilcoxon-Mann-Whitney

Independent sets of real numbers X and Y, of cardinalities  $n_x$  and  $n_y$  respectively

$$H0: \mathbb{P}(X > Y) = \mathbb{P}(Y > X)$$

(i) order elements of  $X \cup Y$  to form an increasing sequence;

(ii) assign to each element in  $X \cup Y$  its rank in the sequence;

(iii) sum the ranks of elements in X: variable  $S_x$ .

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

Wilcoxon rank test a.k.a. Wilcoxon-Mann-Whitney

Independent sets of real numbers X and Y, of cardinalities  $n_x$  and  $n_y$  respectively

**H0**: 
$$\mathbb{P}(X > Y) = \mathbb{P}(Y > X)$$

(i) order elements of  $X \cup Y$  to form an increasing sequence;

(ii) assign to each element in  $X \cup Y$  its *rank* in the sequence;

(iii) sum the ranks of elements in X: variable  $S_x$ .

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  $\alpha = 0.05$ .

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

Wilcoxon rank test a.k.a. Wilcoxon-Mann-Whitney

Independent sets of real numbers X and Y, of cardinalities  $n_x$  and  $n_y$  respectively

**H0**: 
$$\mathbb{P}(X > Y) = \mathbb{P}(Y > X)$$

(i) order elements of  $X \cup Y$  to form an increasing sequence;

(ii) assign to each element in  $X \cup Y$  its *rank* in the sequence;

(iii) sum the ranks of elements in X: variable  $S_x$ .

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

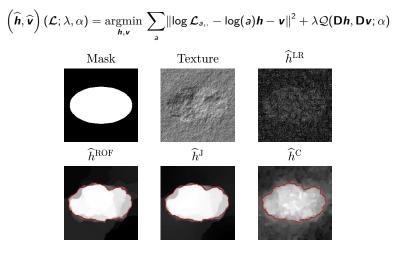
$$\mu = n_x n_y/2; \quad \sigma^2 = n_x n_y (n_x + n_y + 1)/2.$$

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

**Tumorous** breasts have more disrupted tissues compared to normal breasts: <u>normal vs. cancer</u>:  $P \sim 0.0423$ , normal vs. benign:  $P \sim 0.0009$ .

## 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. → Journées ANR Mistic, April 2023



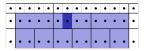
 $\Longrightarrow$  estimation of the local regularity, i.e., roughness, at the pixel level

But first: assess that the wavelet leaders formalism agrees with WTMM on patches

But first: assess that the wavelet leaders formalism agrees with WTMM on patches

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

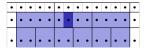
- at all finer scales  $a' \leq a$
- in a spatial neighborhood



But first: assess that the wavelet leaders formalism agrees with WTMM on patches

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

- at all finer scales  $a' \leq a$
- in a spatial neighborhood



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

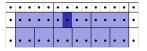
$$\frac{1}{|\Omega|}\sum_{\underline{n}\in\Omega}\mathcal{L}_{a,\underline{n}}^{q}=F_{q}a^{\tau(q)},\quad a\to 0^{+}$$

homogeneous monofractal texture of Hurst exponent  $H \Longrightarrow \tau(q) = qH$ 

But first: assess that the wavelet leaders formalism agrees with WTMM on patches

Wavelet leaders:  $\mathcal{L}_{a,n}$  at scale *a* and pixel <u>*n*</u> supremum of wavelet coefficients

- at all finer scales  $a' \leq a$
- in a spatial neighborhood



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

$$\frac{1}{|\Omega|}\sum_{\underline{n}\in\Omega}\mathcal{L}_{a,\underline{n}}^{q}=F_{q}a^{\tau(q)},\quad a\to 0^{+}$$

homogeneous monofractal texture of Hurst exponent  $H \Longrightarrow \tau(q) = qH$ 

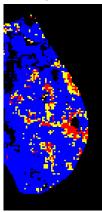
linear regression to estimate H for all  $360 \times 360$ -pixel overlapping patches

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)

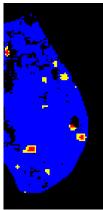
Wavelet leader coefficients (Wendt et al., 2009, Sig. Process.)

- *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)



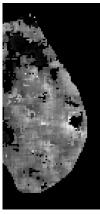
CompuMaine

Leaders



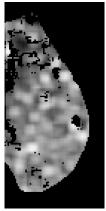
Wavelet leader coefficients (Wendt et al., 2009, Sig. Process.)

- *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)



CompuMaine

Leaders



#### Multifractal analysis of mamographic microenvironment

Kestener et al., 2001; Marin et al., 2017; Gerasimova-Chechkina et al., 2021

#### Multifractal analysis of mamographic microenvironment

Kestener et al., 2001; Marin et al., 2017; Gerasimova-Chechkina et al., 2021

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

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

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

Example: Gaussian and Mexican hat smoothing functions

$$\varphi_{\text{Gauss}}(\boldsymbol{x}) = \exp(-\|\boldsymbol{x}\|^2/2); \quad \varphi_{\text{Mex}}(\boldsymbol{x}) = (2 - \|\boldsymbol{x}\|^2)\exp(-\|\boldsymbol{x}\|^2/2)$$

#### Multifractal analysis of mamographic microenvironment

Kestener et al., 2001; Marin et al., 2017; Gerasimova-Chechkina et al., 2021

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

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

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

Example: Gaussian and Mexican hat smoothing functions

$$\varphi_{\text{Gauss}}(\mathbf{x}) = \exp(-\|\mathbf{x}\|^2/2); \quad \varphi_{\text{Mex}}(\mathbf{x}) = (2 - \|\mathbf{x}\|^2)\exp(-\|\mathbf{x}\|^2/2)$$

#### Wavelet Transform Modulus Maxima

 $\{(\boldsymbol{b},\boldsymbol{a})\in\mathbb{R}^2,\times\mathbb{R}^*_+\quad\mathsf{M}_\psi[f](\boldsymbol{b},\boldsymbol{a})\text{ locally maximal in direction }\mathsf{A}_\psi[f](\boldsymbol{b},\boldsymbol{a})\}$ 

## Multifractal framework: Wavelet Transform Modulus Maxima

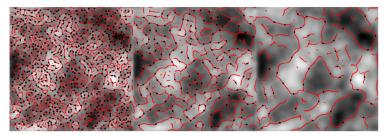


Figure 4.2: The maxima chains are shown for scales  $a = 2^{1}\sigma_{w}$  (left),  $a = 2^{2}\sigma_{w}$  (middle), and  $a = 2^{3}\sigma_{w}$  (right) (where  $\sigma_{w} = 7$  pixels) overlaid onto a 2D fBm image with H = 0.5. The local maxima along  $\mathcal{M}_{\psi}$  (WTMMM) are shown through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

Source: Basel G. White

## Multifractal framework: Wavelet Transform Modulus Maxima

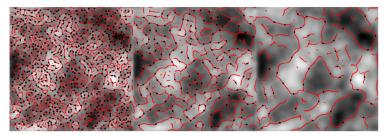


Figure 4.2: The maxima chains are shown for scales  $a = 2^{1}\sigma_{w}$  (left),  $a = 2^{2}\sigma_{w}$  (middle), and  $a = 2^{3}\sigma_{w}$  (right) (where  $\sigma_{w} = 7$  pixels) overlaid onto a 2D fBm image with H = 0.5. The local maxima along  $\mathcal{M}_{\psi}$  (WTMMM) are shown through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

Source: Basel G. White

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

lines formed by WTMM maxima across scales

## Multifractal framework: Wavelet Transform Modulus Maxima

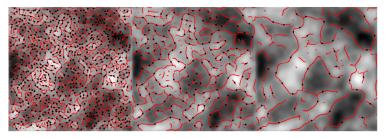


Figure 4.2: The maxima chains are shown for scales  $a = 2^{1}\sigma_{w}$  (left),  $a = 2^{2}\sigma_{w}$  (middle), and  $a = 2^{3}\sigma_{w}$  (right) (where  $\sigma_{w} = 7$  pixels) overlaid onto a 2D fBm image with H = 0.5. The local maxima along  $\mathcal{M}_{\psi}$  (WTMMM) are shown through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

Source: Basel G. White

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

lines formed by WTMM maxima across scales

If a maxima line  $\mathcal{L}_{\mathbf{x}_0}(a)$  is pointing toward a singularity  $\mathbf{x}_0$  as  $a \to 0^+$ , then

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

provided that the wavelet has  $n_{\psi} > h(\mathbf{x}_0)$  vanishing moments.

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

$$\mathcal{Z}(q, a) = \sum_{\ell \in \mathfrak{L}(a)} \left( \sup_{(b, a') \in \ell, a' \leq a} \mathsf{M}_{\psi}[f](b, a') 
ight)^q$$

q: statistical order moment

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

$$\mathcal{Z}(q, a) = \sum_{\ell \in \mathfrak{L}(a)} \left( \sup_{(b, a') \in \ell, a' \leq a} \mathsf{M}_{\psi}[f](b, a') 
ight)^q$$

q: statistical order moment

Roughness, quantified by Hölder exponent, characterized by  $\tau(q)$  spectrum

$$\mathcal{Z}({m q},{m a})\sim {m a}^{ au({m q})}, \quad {m a}
ightarrow 0^+$$

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

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

$$\mathcal{Z}(q, a) = \sum_{\ell \in \mathfrak{L}(a)} \left( \sup_{(b, a') \in \ell, a' \leq a} \mathsf{M}_{\psi}[f](b, a') 
ight)^q$$

q: statistical order moment

Roughness, quantified by Hölder exponent, characterized by  $\tau(q)$  spectrum

$$\mathcal{Z}({m q},{m a})\sim {m a}^{ au({m q})}, \quad {m a} o 0^+$$

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

**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  $\tau$ )

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

 $\Longrightarrow$  Mean quantities in a canonical ensemble with Boltzmann weights

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

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

 $\Longrightarrow$  Mean quantities in a canonical ensemble with Boltzmann weights

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

Roughness: robust local regularity estimation

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

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

 $\implies$  Mean quantities in a canonical ensemble with Boltzmann weights

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

Roughness: robust local regularity estimation

$$egin{aligned} h(q, \mathbf{a}) &= \sum_{\ell \in \mathfrak{L}(\mathbf{a})} \ln \left( \mathrm{W}_{\psi}[f](q, \ell, \mathbf{a}) 
ight) \mathrm{W}_{\psi}[f](q, \ell, \mathbf{a}), \ h(q) &= rac{\mathrm{d} au}{\mathrm{d}q} = \lim_{\mathbf{a} o 0^+} rac{h(q, \mathbf{a})}{\ln \mathbf{a}} \end{aligned}$$

Singularity spectrum:

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

**Roughness:**  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; **Singularity spectrum:**  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

- The larger the patch, the larger the range of q values, the better the estimate;
- but risk of confusing average of several monofractal signatures and multifractal.
- $\Longrightarrow$  estimation on overlapping patches of size 360  $\times$  360 pixels with 32-pixel shift

**Roughness:**  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; **Singularity spectrum:**  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

- The larger the patch, the larger the range of q values, the better the estimate;
- but risk of confusing average of several monofractal signatures and multifractal.
- $\Longrightarrow$  estimation on overlapping patches of size 360  $\times$  360 pixels with 32-pixel shift

#### Image sliding window analysis

- 1. Check that the central  $256 \times 256$  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  $\times$  360 pixels, i.e.,  $15.5 \times 15.5 \text{ mm}$ 

roughness: 
$$h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$$
; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

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

For each patch of 360  $\times$  360 pixels, i.e.,  $15.5 \times 15.5 \text{ mm}$ 

roughness:  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

 $\implies$  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

$$\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, \$$
with  $\ \sigma_w = 7$  pixels

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

For each patch of 360  $\times$  360 pixels, i.e.,  $15.5 \times 15.5 \text{ mm}$ 

roughness:  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

 $\implies$  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

$$\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, \$$
with  $\ \sigma_w = 7$  pixels

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

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

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

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

For each patch of 360  $\times$  360 pixels, i.e.,  $15.5 \times 15.5 \text{ mm}$ 

roughness:  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

 $\implies$  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

$$\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, \$$
with  $\ \sigma_w = 7$  pixels

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

• 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  $a \rightarrow 0^+$ 

For each patch of 360  $\times$  360 pixels, i.e.,  $15.5 \times 15.5 \text{ mm}$ 

roughness:  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

 $\implies$  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

$$\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, \$$
with  $\ \sigma_w = 7$  pixels

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

• 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

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

roughness:  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

 $\implies$  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

$$\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, \$$
with  $\ \sigma_w = 7$  pixels

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  $a_{\max}$ 

 $sd_w < 0.06$ 

 $\implies$  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

17/24

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

roughness:  $h(q) = \lim_{a \to 0^+} \frac{h(q, a)}{\ln a}$ ; singularity spectrum:  $\mathcal{D}(q, a) = \lim_{a \to 0^+} \frac{\mathcal{D}(q, a)}{\ln a}$ 

 $\implies$  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

$$\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, \$$
with  $\ \sigma_w = 7$  pixels

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  $a_{\max}$ 

 $\langle R_w^2 \rangle > 0.96$ 

 $\implies$  ensures self-similarity

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

17/24

For **each** patch of  $360 \times 360$  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_w} = 0.0, 0.1, \dots, 2.1, \ , \ \log_2 \frac{a_{\max}}{\sigma_w} = 2.0, 2.1, \dots, 4.1, \$$
with  $\ \sigma_w = 7$  pixels

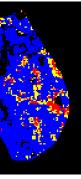
and select  $[a_{\min}, a_{\max}]$  if and only if

- -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<sub>w</sub> < 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**.

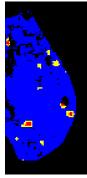
Wavelet leader coefficients (Wendt et al., 2009, Sig. Process.)

- *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)



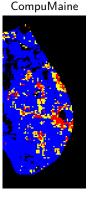
CompuMaine

fixed scales

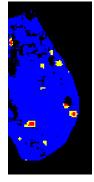


Wavelet leader coefficients (Wendt et al., 2009, Sig. Process.)

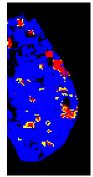
- *H* < 1/2 monofractal anti-correlated: fatty tissues (healthy)
- H > 1/2 monofractal long-range correlated: dense tissues (healthy)
- $H \simeq 1/2$  monofractal uncorrelated: disrupted tissues (tumorous)



fixed scales



adaptive scales



19/24

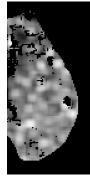
$$[a_{\min}, a_{\max}] = [2^3, 2^5] \quad [a_{\min}, a_{\max}] \subset [2^2, 2^8]$$

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

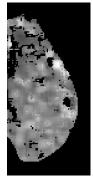
- *H* < 1/2 monofractal anti-correlated: fatty tissues (healthy)
- H > 1/2 monofractal long-range correlated: dense tissues (healthy)
- $H \simeq 1/2$  monofractal uncorrelated: disrupted tissues (tumorous)



#### fixed scales



adaptive scales



19/24

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

## Mammogram datasets

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

### Mammogram datasets

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:  $P \sim 0.0009$ .

Gerasimova-Chechkina et al., 2021, Front. Physiol.

**Russian:** Perm Regional Oncological Dispensary

81 cancer vs. 23 benign

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

Cancerous breasts have more disrupted tissues compared to breasts with benign lesions:

cancer vs. benign:  $P \sim 0.003$ 

### Mammogram datasets

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:  $P \sim 0.0009$ .

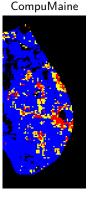
Gerasimova-Chechkina et al., 2021, Front. Physiol.  $\implies$  shared with us, with analyses Russian: Perm Regional Oncological Dispensary

81 cancer vs. 23 benign

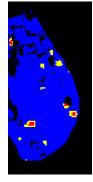
 $\implies$  digitally acquired mammograms: uncompressed 8-bit BMP images ([0, 255]) Cancerous breasts have more disrupted tissues compared to breasts with benign lesions: cancer vs. benign:  $P \sim 0.003$ 

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

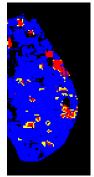
- *H* < 1/2 monofractal anti-correlated: fatty tissues (healthy)
- H > 1/2 monofractal long-range correlated: dense tissues (healthy)
- $H \simeq 1/2$  monofractal uncorrelated: disrupted tissues (tumorous)



fixed scales



adaptive scales



21/24

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

Gerasimova-Chechkina et al., 2021, Front. Physiol.  $\implies$  shared with us, with analyses Russian: Perm Regional Oncological Dispensary

81 cancer vs. 23 benign

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

Cancerous breasts have more disrupted tissues compared to breasts with benign lesions:

cancer vs. benign:  $P \sim 0.003$ 

Gerasimova-Chechkina et al., 2021, Front. Physiol.  $\implies$  shared with us, with analyses **Russian:** Perm Regional Oncological Dispensary

81 cancer vs. 23 benign

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

Cancerous breasts have more disrupted tissues compared to breasts with benign lesions:

cancer vs. benign:  $P \sim 0.003$ 

Wavelet leaders with

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

Gerasimova-Chechkina et al., 2021, Front. Physiol.  $\implies$  shared with us, with analyses **Russian:** Perm Regional Oncological Dispensary

81 cancer vs. 23 benign

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

Cancerous breasts have more disrupted tissues compared to breasts with benign lesions:

cancer vs. benign:  $P \sim 0.003$ 

Wavelet leaders with

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

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) cancer vs. benign  $P \sim 0.0030$ 

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

# 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) cancer vs. benign  $P \sim 0.0030$ 

 $\implies$  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

 $\implies$  increase the sensibility in the measurement of the quantity of disrupted tissues

### 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

 $\implies$  increase the sensibility in the measurement of the quantity of disrupted tissues

#### 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

### 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

 $\implies$  increase the sensibility in the measurement of the quantity of disrupted tissues

#### 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

#### Anisotropic Gaussian fields for mammogram modeling

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