

## Lecture 17: Wavelets

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Recall

- ▶ One sample sign test, Wilcoxon signed rank test, large-sample approximation, median, Hodges-Lehman estimator, distribution-free confidence interval.
- ▶ Jackknife for bias and standard error of an estimator.
- ▶ Bootstrap samples, bootstrap replicates.
- ▶ Bootstrap standard error of an estimator.
- ▶ Bootstrap percentile confidence interval.
- ▶ Hypothesis testing with the bootstrap (one-sample problem.)
- ▶ Assessing the error in bootstrap estimates.
- ▶ Example: inference on ratio of heart attack rates in the aspirin-intake group to the placebo group.
- ▶ The exhaustive bootstrap distribution.

- ▶ Discrete data problems (one-sample, two-sample proportion tests, test of homogeneity, test of independence).
- ▶ Two-sample problems (location problem - equal variance, unequal variance, exact test or Monte Carlo, large-sample approximation, H-L estimator, dispersion problem, general distribution).
- ▶ Permutation tests (permutation test for continuous data, different test statistic, accuracy of permutation tests).
- ▶ Permutation tests (discrete data problems, exchangeability.)
- ▶ Rank-based correlation analysis (Kendall and Spearman correlation coefficients.)
- ▶ Rank-based regression (straight line, multiple linear regression, statistical inference about the unknown parameters, nonparametric procedures - does not depend on the distribution of error term.)
- ▶ Smoothing (density estimation, bias-variance trade-off, curse of dimensionality)
- ▶ Nonparametric regression (Local averaging, local regression, kernel smoothing, local polynomial, penalized regression)

- ▶ Cross-validation, Variance Estimation, Confidence Bands, Bootstrap Confidence Bands.

# Wavelets

## Spatially inhomogeneous functions

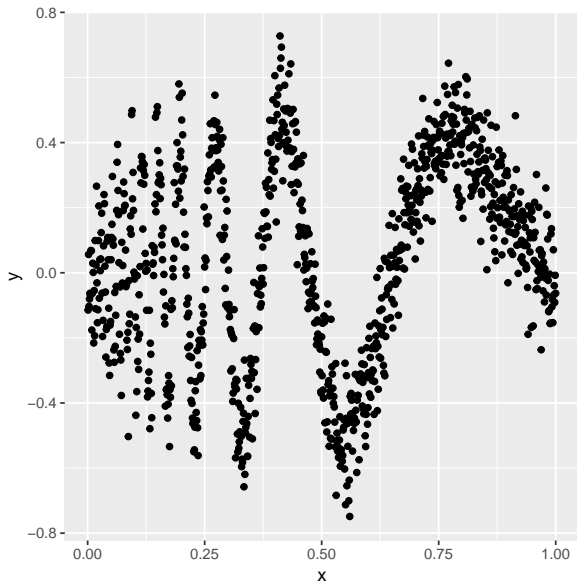
# Example

► Doppler function

```
library(ggplot2)
r = function(x){
  sqrt(x*(1-x))*sin(2.1*pi/(x+.05))
}
ep = rnorm(1000)
y = r(seq(1, 1000, by = 1)/1000) + .1 * ep
df = data.frame(x = seq(1, 1000, by = 1)/1000, y = y)
ggplot(df) +
  geom_point(aes(x = x, y = y))
```



# Example



## Example

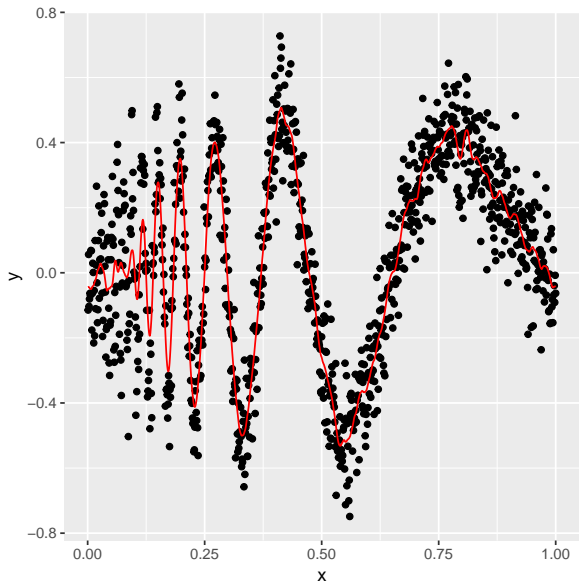
- ▶ Doppler function is spatially inhomogeneous (smoothness varies over  $x$ ).
- ▶ Estimate by local linear regression

```
library(np)
doppler.npreg <- npreg(bws=.005,
  txdat=df$x,
  tydat=df$y,
  ckertype="epanechnikov")

doppler.npreg.fit = data.frame(x = df$x,
  y = df$y,
  kernel.fit = fitted(doppler.npreg))

p = ggplot(doppler.npreg.fit) +
  geom_point(aes(x = x, y = y)) +
  geom_line(aes(x = x, y= kernel.fit), color = "red")
```

# Example



# Example

- ▶ Doppler function fit using local linear regression.
  - ▶ Effective degrees of freedom 166.
  - ▶ Fitted function is very wiggly.
  - ▶ If we smooth more, right-hand side of the fit would look better at the cost of missing structure near  $x = 0$ .

## Introduction

- ▶ Construct basis functions that are
  - ▶ multiscale.
  - ▶ spatially/ locally adaptive.
- ▶ Find sparse set of coefficients for a given basis.

- ▶ Function  $f$  belongs to a class of functions  $\mathcal{F}$  possessing more general characteristics, such as a certain level of smoothness.
- ▶ Estimate  $f$  by representing the function in another domain.
- ▶ Use an orthogonal series representation of the function  $f$ .
- ▶ Estimating a set of scalar coefficients that represent  $f$  in the orthogonal series domain.
- ▶ Tool: Wavelets
  - ▶ ability to estimate both global and local features in the underlying function

Sparseness



- ▶ **W 2006** Chapter 9
- ▶ A function  $f = \sum_j \beta_j \phi_j$  is sparse in a basis  $\phi_1, \phi_2, \dots$  if most of the  $\beta_j$ 's are zero.
- ▶ Sparseness generalizes smoothness: smooth functions are sparse but there are also non smooth functions that are sparse.
- ▶ Sparseness is not captured by  $L_2$  norm.
  - ▶ Example  $\mathbf{a} = (1, 0, \dots, 0)$  and  $\mathbf{b} = (1/\sqrt{n}, 1/\sqrt{n}, \dots, 1/\sqrt{n})$ .
  - ▶  $\mathbf{a}$  is sparse.
  - ▶  $L_2$  norms are  $\|\mathbf{a}\|_2 = \|\mathbf{b}\|_2 = 1$ .
  - ▶  $L_1$  norms are  $\|\mathbf{a}\|_1 = 1$  and  $\|\mathbf{b}\|_1 = \sqrt{n}$ .

# Wavelets

- ▶ Data: There are  $n$  pairs of observations  $(x_1, Y_1), (x_2, Y_2), \dots, (x_n, Y_n)$ .
- ▶ Assumptions
  - ▶  $Y_i = f(x_i) + \epsilon_i$ .
  - ▶  $\epsilon_i$  are IID.
  - ▶  $\int f^2 < \infty$  and  $f$  is defined on a close interval  $[a, b]$ . For simplicity, we will consider  $[a, b] = [0, 1]$ .

Wavelet representation of a function

# Basis functions

- ▶  $\psi = \{\psi_1, \psi_2, \dots\}$  is called a basis for a class of functions  $\mathcal{F}$ .  
Then, for  $f \in \mathcal{F}$ ,

$$f(x) = \sum_{i=1}^{\infty} \theta_i \psi_i(x).$$

- ▶  $\theta_i$ - scalar constants/coefficients
- ▶ Basis functions are orthogonal if  $\langle \psi_i, \psi_j \rangle = 0$  for  $i \neq j$ .
- ▶ If basis functions are orthonormal, they are orthogonal and  $\langle \psi_i, \psi_i \rangle = 1$ .
- ▶ How do we construct basis functions  $\psi_i$ 's ?

# Basis functions

- ▶ If  $\psi$  is a wavelet function, then the collection of functions

$$\Psi = \{\psi_{ij}; j, k \text{ integers}\},$$

where

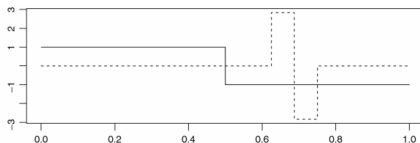
$$\psi_{ij} = 2^{j/2} \psi(2^j x - k),$$

forms a basis for square integrable functions.

- ▶  $\Psi$  is a collection of translation (shift) and dilation (scaling) of  $\psi$ .
  - ▶  $\psi$  can be defined in any range of real line.
  - ▶  $\int \psi = 1$ 
    - ▶ value of  $\psi$  is near 0 except over a small range.

# Some examples for wavelets

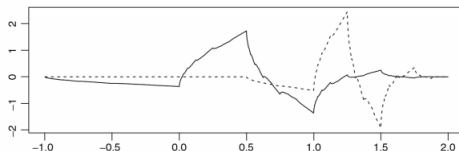
- Haar wavelets (1910)



Source: Hollander, Wolfe, and Chicken (2013)

# Some examples for wavelets

- Daubechies wavelets (1992)



Source: Hollander, Wolfe, and Chicken (2013)

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# Multiresolution analysis (MRA)

- ▶ Carefully construct wavelet function  $\psi$ .
- ▶ MRA: interpretation of the wavelet representation of  $f$  in terms of location and scale.
- ▶ Translation and dilation of  $\psi$  gives

$$f(x) = \sum_{j \in \mathcal{Z}} \sum_{k \in \mathcal{Z}} \theta_{jk} \psi(x),$$

where  $\mathcal{Z}$  is a set of integers.

- ▶ scale - frequency.
- ▶ For fixed  $j$ ,  $k$  represents the behavior of  $f$  at resolution scale  $j$  and a particular location.
- ▶ function  $f$  at differing resolution (scale, frequency) levels  $j$  and locations  $k$  - MRA.

# Multiresolution analysis (MRA)

- ▶ Cumulative approximation of  $f$  using  $j < J$ ,

$$f_J(x) = \sum_{j < J} \sum_{k \in \mathbb{Z}} \theta_{jk} \psi(x).$$

- ▶  $J$  increases -  $f_J$  models smaller scales (higher frequency) of  $f$  - changes occur in the small interval of  $x$ .
  - ▶  $J$  decreases -  $f_J$  models larger scale (lower frequency) behavior of  $f$ .
- ▶ A complete representation of  $f$  is the limit of  $f_J$ .

# Multiresolution analysis (MRA)

- ▶ Write  $f_J(x)$  as follows:

$$f_J(x) = \sum_{k \in \mathbb{Z}} \xi_{j0} \phi_{j0k}(x) + \sum_{j0 \leq j < J} \sum_{k \in \mathbb{Z}} \theta_{jk} \psi_{jk}(x),$$

where  $f_{j0} = \sum_{k \in \mathbb{Z}} \xi_{j0} \phi_{j0k}(x)$ .

- ▶ Add second term to  $f_{j0}$  allows for modeling higher scale-frequency behavior of  $f$ .
- ▶  $f_{j0}$  approximation at the smooth resolution level.
- ▶ Each of the remaining resolution level series is a “detail” level.
- ▶  $\phi$  - scaling function (Father wavelet).
- ▶  $\psi$  - wavelet function (Mother wavelet).

# MRA Using the Haar Wavelet (Example)

- ▶ Approximate  $f(x) = x, x \in (0, 1)$ .
- ▶ Define Haar wavelet function

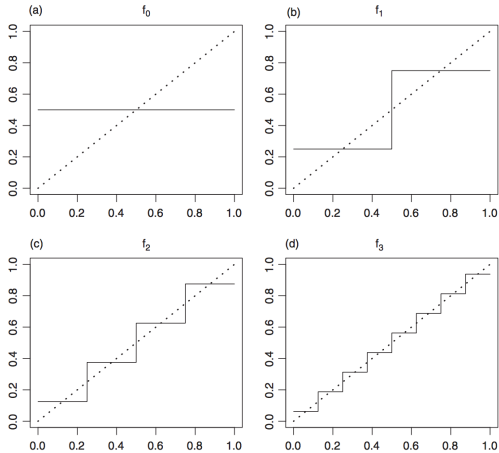
$$\psi(x) = \begin{cases} 1 & x \in [0, 1/2), \\ -1 & x \in [1/2, 1), \end{cases} \quad (1)$$

and

$$\phi(x) = 1, x \in [0, 1]. \quad (2)$$

- ▶ Haar wavelet allows exact determination of the wavelet coefficients  $\theta_{jk}$ .

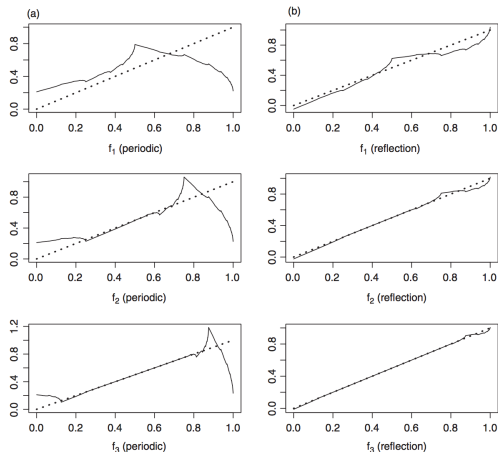
► Source **HWC**



**Figure 13.2** Cumulative approximations up to resolution levels  $J = -1, 0, 1, 2$  from Example 13.1 using the Haar wavelet. The underlying function is  $f(x) = x$ , shown with a dotted line.

# MRA Using the D2 wavelet

## ► Source **HWC**



**Figure 13.4** Approximations up to resolution levels  $j = 0, 1, 2$  from Example 13.1 using the D2 wavelet. The left panels use periodic boundary handling, the right panels use reflection. The underlying function is  $f(x) = x$ , shown with a dotted line.

# MRA Using the D2 wavelet

- ▶ To avoid boundary issues using D2
  - ▶ Specify using reflection at the boundaries, rather than periodicity.
  - ▶ increase the number of indices  $k$  that must be considered at each resolution level  $j$ .

# Discrete wavelet transform

- ▶ Cascade algorithm provides MRA (Mallat 1989).
- ▶ Some restrictions
  - ▶  $J = \log_2(n)$ .
  - ▶ The number of resolution levels in the wavelet series is truncated both above and below in practice, resulting in  $J - j_0 + 1$  series, each representing a resolution level.
- ▶ Commands in R that make use of the DWT are `dwt`, `idwt`, and `mra` in package `waveslim` (Whitcher (2010)).



## Discrete wavelet transform (Example)

►  $y_i = x_i = (i - 1) / n, i = 1, 2, \dots, n.$

```
n = 2^12  
xi = (seq(1, n, by = 1) - 1)/n  
yi = xi  
library(waveslim)
```

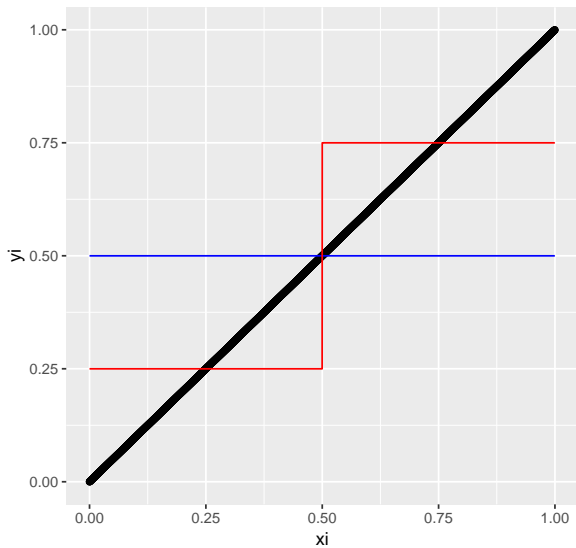
- ▶ Haar basis.
- ▶ Number of resolution levels  $J = 12$ .
- ▶ Decompose the sample data  $y$ .

```
dwt.fit = mra(yi, method="dwt", wf="haar", J=12)
```

- ▶ Output is a list of 13 vectors.
- ▶ The first vector is the change necessary to go from the approximation  $f_{12}$  to  $f_{13}$ - approximation at the highest detail resolution level.
- ▶ The next to last vector is  $f_1 - f_0$ .
- ▶ The final, thirteenth vector is the smooth approximation  $f_0$ .
- ▶ Summing the thirteenth vector and the twelfth vector results in  $f_1$ .
- ▶  $f_{13} - f_{12}, f_{12} - f_{11}, \dots, f_1 - f_0, f_0$ .

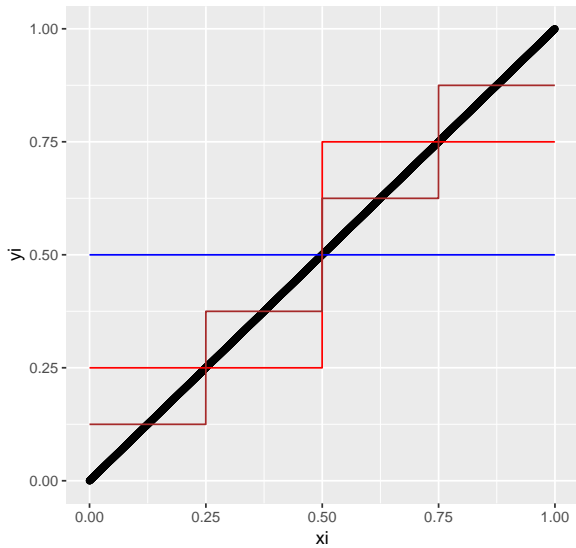
```
f0 = dwt.fit[[13]]
f1 = dwt.fit[[13]]+dwt.fit[[12]]
df = data.frame(x = xi, y = yi,
  f0=f0, f1 = f1)
p1 = ggplot() +
  geom_point(data = df ,
    aes(x = xi, y = yi))+
  geom_line(data = df ,
    aes(x = xi, y = f0), color = "blue") +
  geom_line(data = df,
    aes(x = xi, y = f1), color = "red")
```

Wavelet representation with resolution  $J = 0$  and  $1$



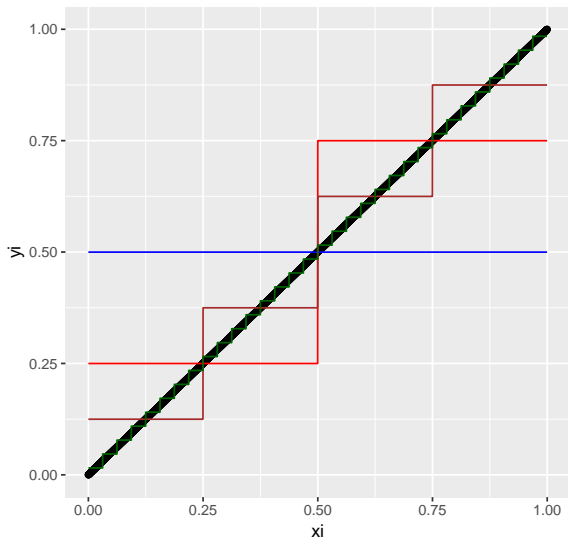
```
f2 = dwt.fit[[13]]+dwt.fit[[12]] + dwt.fit[[11]]
df = data.frame(x = xi, y = yi,
  f0 = f0, f1 = f1, f2 = f2)
p2 = ggplot() +
  geom_point(data = df ,
    aes(x = xi, y = yi)) +
  geom_line(data = df,
    aes(x = xi, y = f0), color = "blue")+
  geom_line(data = df,
    aes(x = xi, y = f1), color = "red")+
  geom_line(data = df,
    aes(x = xi, y = f2), color = "brown")
```

Wavelet representation with resolution  $J = 0, 1, 2$



```
f5 = dwt.fit[[13]]+dwt.fit[[12]] + dwt.fit[[11]] + dwt.fit
df = data.frame(x = xi, y = yi,
  f0 = f0, f1 = f1, f2 = f2, f5 = f5)
p5 = ggplot() +
  geom_point(data = df ,
    aes(x = xi, y = yi)) +
  geom_line(data = df,
    aes(x = xi, y = f0), color = "blue")+
  geom_line(data = df,
    aes(x = xi, y = f1), color = "red")+
  geom_line(data = df,
    aes(x = xi, y = f2), color = "brown")+
  geom_line(data = df,
    aes(x = xi, y = f5), color = "darkgreen")
```

Wavelet representation with increasing resolution  $J = 0, 1, 2$





- ▶ What if we choose  $J$  is less than 12 for this example?
- ▶ Set  $J = 3$
- ▶  $j_0 > 0$ , for example, when  $J = 3$ ,  $j_0 = 9$ .

```
dwt.fit.J3 = mra(yi, method="dwt", wf="haar", J=3)
```

```
length(dwt.fit.J3)
```

```
## [1] 4
```

```
f9 = dwt.fit.J3[[4]] # f0
```

```
f10 = dwt.fit.J3[[4]] + dwt.fit.J3[[3]] # f1
```

```
f11 = dwt.fit.J3[[4]] + dwt.fit.J3[[3]] + dwt.fit.J3[[2]] #
```

```
df = data.frame(x = xi, y = yi,
```

```
  f0 = f9, f1 = f10, f2 = f11)
```

```
p.J3 = ggplot() +
```

```
  geom_point(data = df ,
```

```
    aes(x = xi, y = yi)) +
```

```
  geom_line(data = df,
```

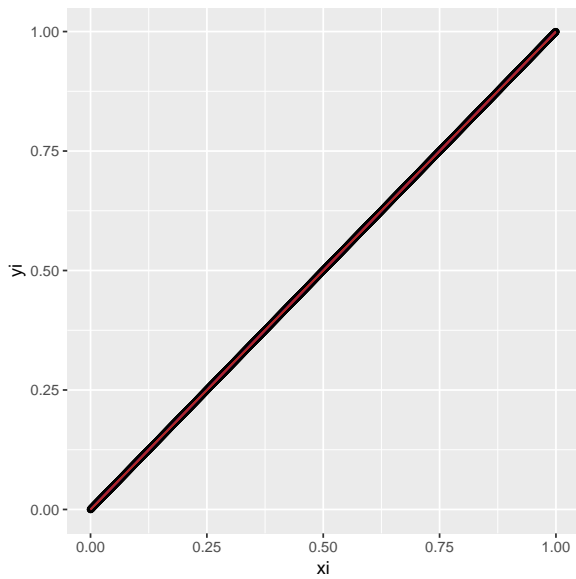
```
    aes(x = xi, y = f0), color = "blue")+
```

```
  geom_line(data = df,
```

```
    aes(x = xi, y = f1), color = "red")+
```

```
  geom_line(data = df,
```

```
    aes(x = xi, y = f2), color = "brown")
```

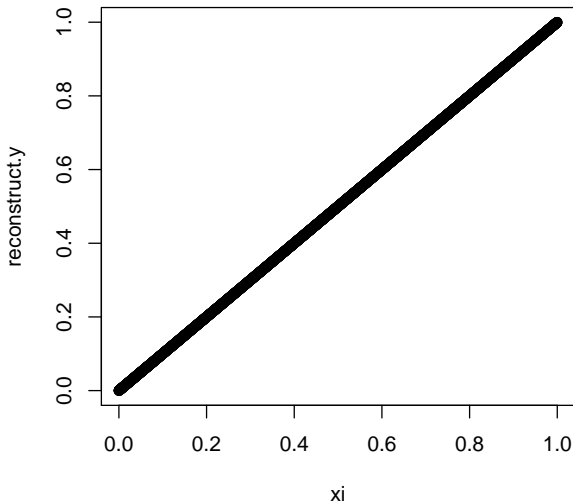


- ▶ dwt determines the wavelet coefficients at each resolution level.
  - ▶ n.levels - resolution levels to determine.
- ▶ Read Page 637 for more detail.

```
y.dwt <- dwt(yi, wf="haar", n.levels=12)
```

- The resulting R list of coefficients may be used to reconstruct the original vector of sampled data  $y$ .

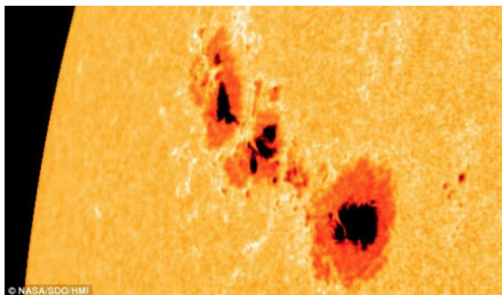
```
reconstruct.y = idwt(y.dwt)  
plot(xi, reconstruct.y)
```



## Wavelet Thresholding

- ▶ We saw how a function  $f$  may be represented with a wavelet basis.
  - ▶ DWT, a sample of length  $n$  from  $f$  may be decomposed into  $n$  wavelet coefficients making up a single smooth approximation and up to  $J = \log_2(n)$  detail resolution levels.
- ▶ Sparsity - the ability of wavelets to represent a function by concentrating or compressing the information about  $f$  into a few large magnitude coefficients and many small magnitude coefficients.
- ▶ Compression (thresholding) is applied to the wavelet coefficients of a sampled function  $f$  prior to its reconstruction.
- ▶ Thresholding - provides a significant level of data reduction for the problem.

# Sparsity of the Wavelet Representation



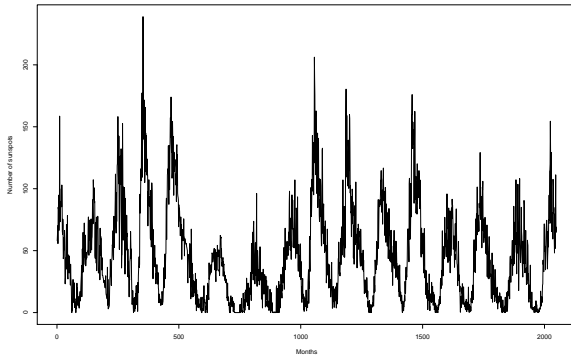
- ▶ **HWC** Example 13.3
- ▶ Monthly sunspot numbers from 1749 to 1983.
- ▶ Sunspots are temporary phenomena on the photosphere of the sun that appear visibly as dark spots compared to surrounding regions.



- ▶ Sunspots correspond to concentrations of magnetic field flux that inhibit convection and result in reduced surface temperature compared to the surrounding photosphere.
- ▶ The original data has length 2820, but only the first 2048 are used here to make it a dyadic number.
- ▶ So the filtered data is monthly sunspot data from January 1749 through July 1919.

```
library(datasets)  
data(sunspots)
```

```
plot.ts(sunspots[1:2048],  
        ylab = "Number of sunspots",  
        xlab = "Months")
```



- ▶ The DWT is applied to this data resulting in 2048 coefficients.

```
dwt.sunspot = dwt(sunspots[1:2048], n.levels = 4, wf = "la8
```

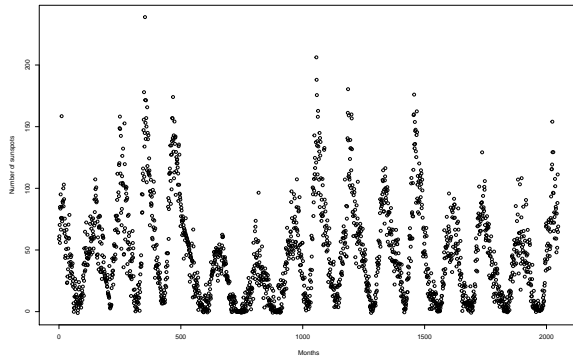
- ▶ These coefficients are sorted in magnitude and the smallest 50% (1024) are set to 0.
  - ▶ Reconstruction nearly indistinguishable from the original data.

```
dwt.sunspot.coeff = unlist(dwt.sunspot)
dwt.sunspot.coeff = sort(dwt.sunspot.coeff,
  decreasing = T)
val = as.numeric(quantile(dwt.sunspot.coeff,
  p = .5))
manual.thresholding = manual.thresh(dwt.sunspot,
  value = val)
```

- The inverse DWT is applied to this compressed (50% thresholding) set of coefficients, resulting in the reconstruction.

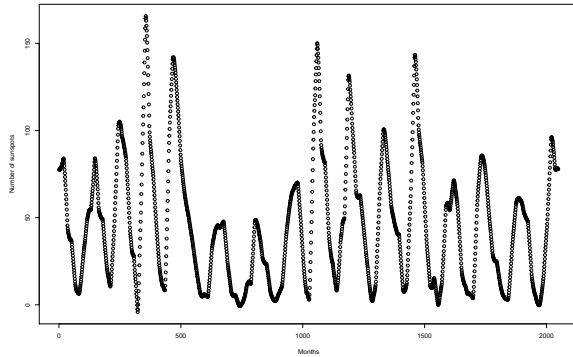
```
y.idwt.manual.thresholding = idwt(manual.thresholding)
plot(y.idwt.manual.thresholding,
     ylab = "Number of sunspots",
     xlab = "Months",
     main = "50% Thresholding")
```

50% Thresholding



- ▶ Set smallest 95% of the coefficients to 0 prior to reconstruction.
  - ▶ Reconstruction with the basic shape of the original data, but with the very localized variability mostly removed.

95% Thresholding



# Thresholding

- ▶ A drawback to compression - need to specify the amount of reduction.
- ▶ Thresholding specifies a data-driven compression.
- ▶ Many methods of thresholding are based on assuming that the errors are normally distributed.



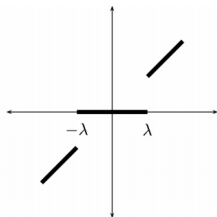
# Thresholding

- ▶ Let  $\theta$  is a coefficient estimated with the DWT and  $\lambda$  is a specified threshold value.
- ▶ Hard thresholding
  - ▶ sets a coefficient to 0 if it has small magnitude and leaves the coefficient unmodified otherwise.
- ▶ Soft thresholding
  - ▶ threshold sets small coefficients to 0 and shrinks the larger ones by  $\lambda$  toward 0.
- ▶ DWT operation may be represented as a matrix operator  $W$

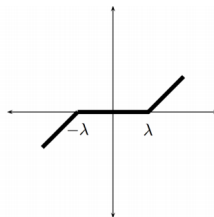
$$\tilde{\theta} = Wf + W\epsilon.$$

- ▶  $\theta = Wf$  represents the wavelet coefficients of the unobserved sampled function  $f$ .
- ▶  $\tilde{\epsilon} = W\epsilon$  represents the coefficients of the errors.

# Thresholding



Hard thresholding



Soft thresholding

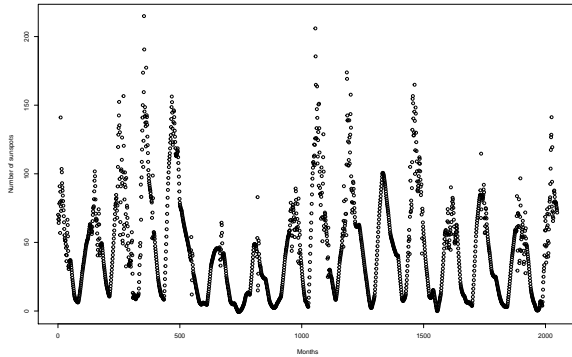
Source: Wasserman (2006)

# Thresholding - VisuShrink (Donoho and Johnstone (1994))

- ▶ Applying a single threshold  $\lambda$  (Donoho and Johnstone 1994).

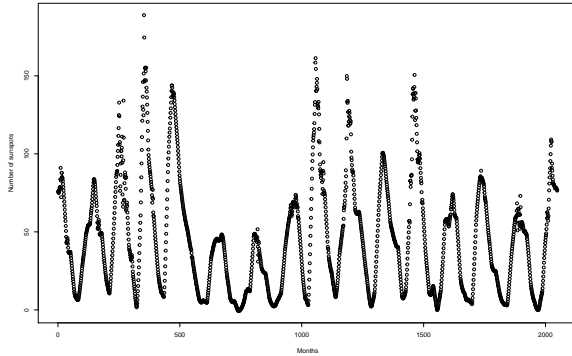
```
y = sunspots[1:2048]
y.dwt = dwt(sunspots[1:2048])
y.visuShrink = universal_thresh(y.dwt, hard = TRUE)
y.idwt.visuShrink = idwt(y.visuShrink)
plot(y.idwt.visuShrink,
     ylab = "Number of sunspots",
     xlab = "Months",
     main = "VisuShrink-Hard Thresholding")
```

VisuShrink-Hard Thresholding



```
y.visuShrink.soft = universal thresh(y.dwt, hard = FALSE)
y.idwt.visuShrink.soft = idwt(y.visuShrink.soft)
plot(y.idwt.visuShrink.soft,
      ylab = "Number of sunspots",
      xlab = "Months",
      main = "VisuShrink-Soft Thresholding")
```

VisuShrink-Soft Thresholding

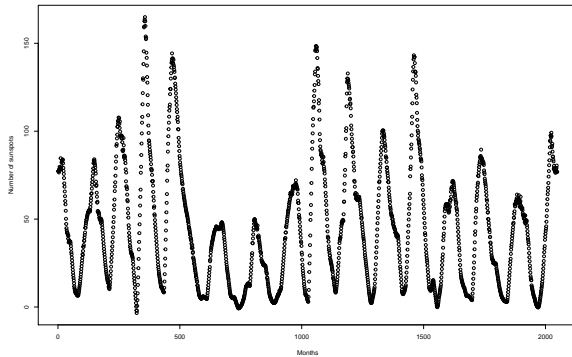


# Thresholding - SureShrink (Donoho and Johnstone (1995))

- ▶ Uses a different threshold at each resolution level of the wavelet decomposition of  $f$  (Donoho and Johnstone 1995).
- ▶ SureShrink is actually a hybrid threshold method
  - ▶ certain resolution levels can be too sparse.
  - ▶ revert SureShrink to using the universal threshold of VisuShrink at the resolution level in question.

```
y.sureshrink = hybrid.thresh(y.dwt, max.level = 4)
y.sureshrink.idwt = idwt(y.sureshrink)
plot(y.sureshrink.idwt,
     ylab = "Number of sunspots",
     xlab = "Months",
     main = "SureShrink-Soft Thresholding")
```

SureShrink-Soft Thresholding





## Other use of wavelets

- ▶ Nonparametric density estimation (Vidakovic (1999)).
- ▶ Use for understanding the properties of time series and random processes.

- ▶ Can do thresholding without strong distributional assumptions on the errors using cross-validation (Nason 1996).
- ▶ Practical, simultaneous confidence bands for wavelet estimators are not available (Wasserman 2006).
- ▶ Standard wavelet basis functions are not invariant to translation and rotations.
  - ▶ Recent work by (Mallat 2012) and (Bruna and Mallat 2013) extend wavelets to handle these kind of invariances.
  - ▶ Promising new direction for the theory of convolutional neural network.

# References for this lecture

**HWC** Chapter 13 (Wavelets)

**W** Chapter 9

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