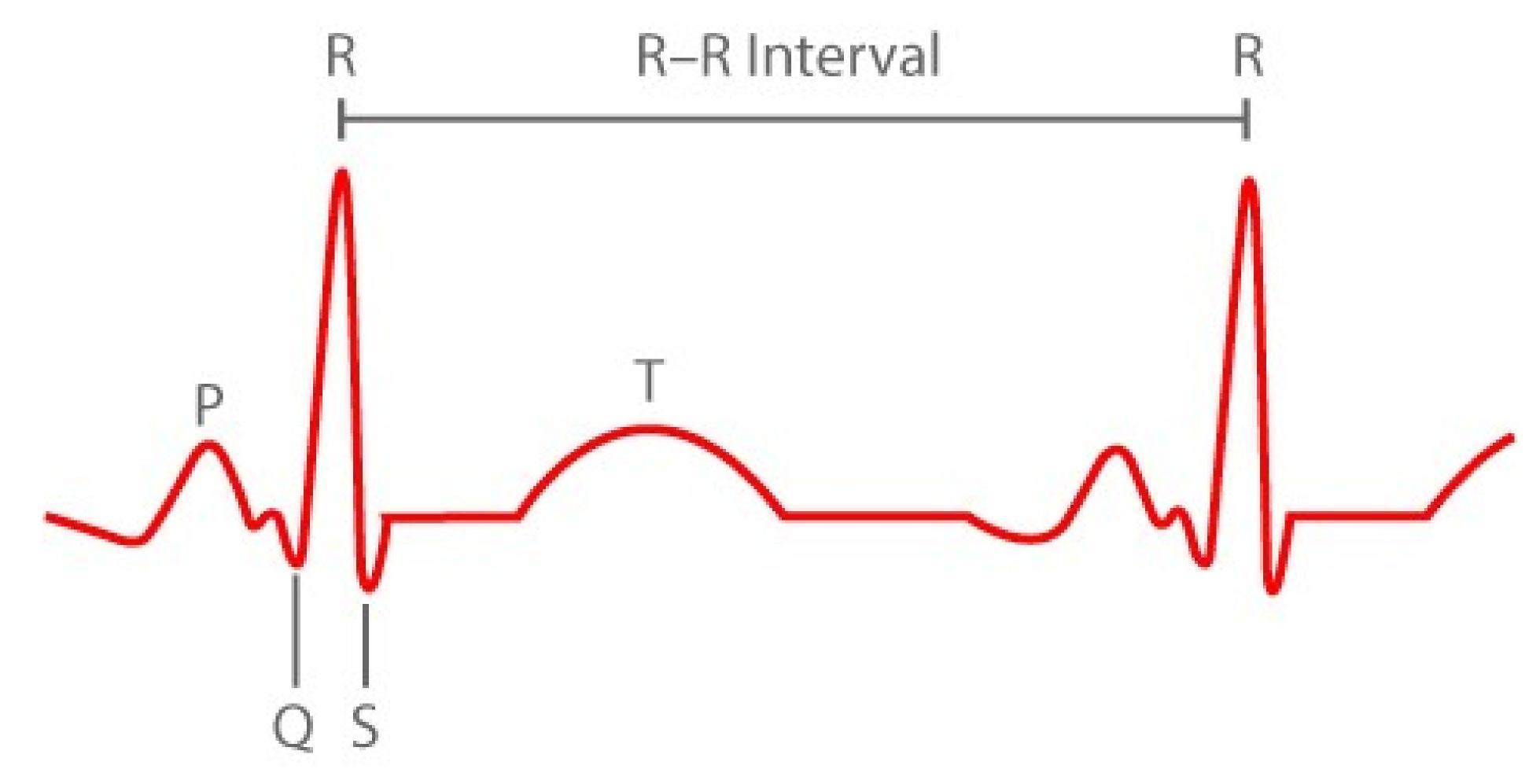
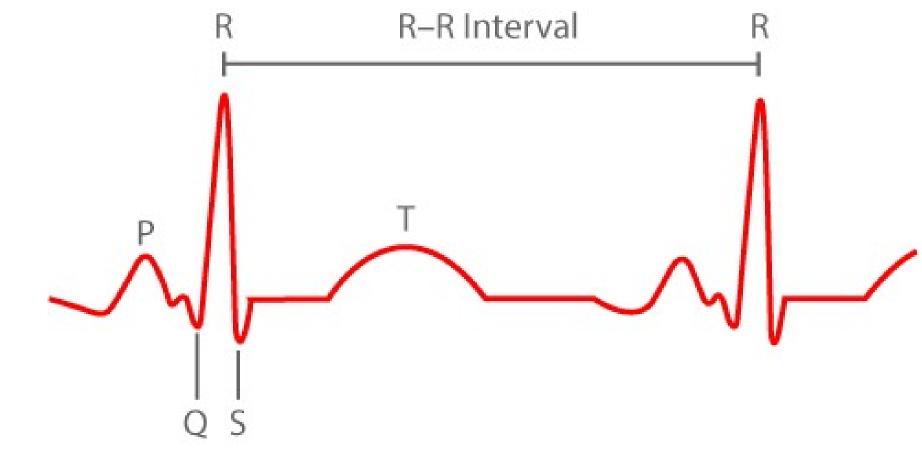
Heart rate variability (HRV)



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Source: https://www.myithlete.com/what-is-hrv

Linear methods – time domain



• mean RR

$$\overline{RR} = \frac{1}{N} \sum_{i=1}^{N} RR_{i}$$

RR variation

$$RR_{variation} = \frac{\left(RR_{max} - RR_{min}\right) \times 100}{RR_{mean}}$$

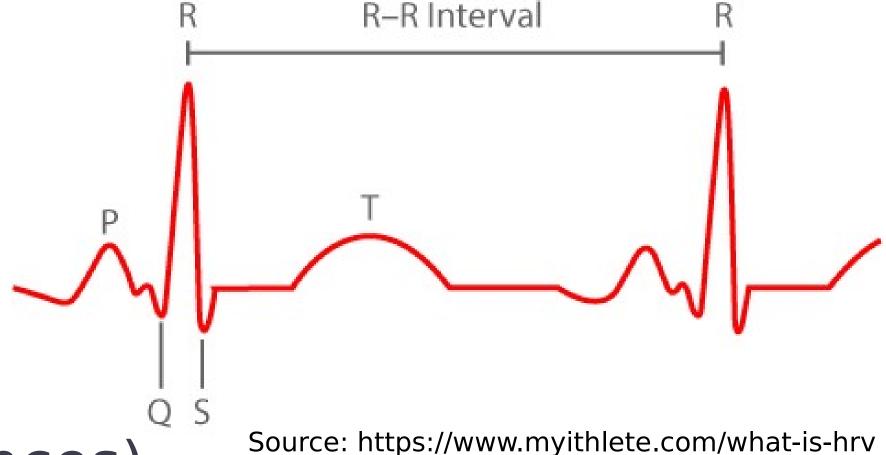
• std RR

$$std RR = \sqrt{\frac{1}{N-1} \sum_{i=1}^{N} \left(\overline{RR} - RR_i \right)^2}$$

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Linear methods – time domain



• RMSSD (root mean square of successive differences)

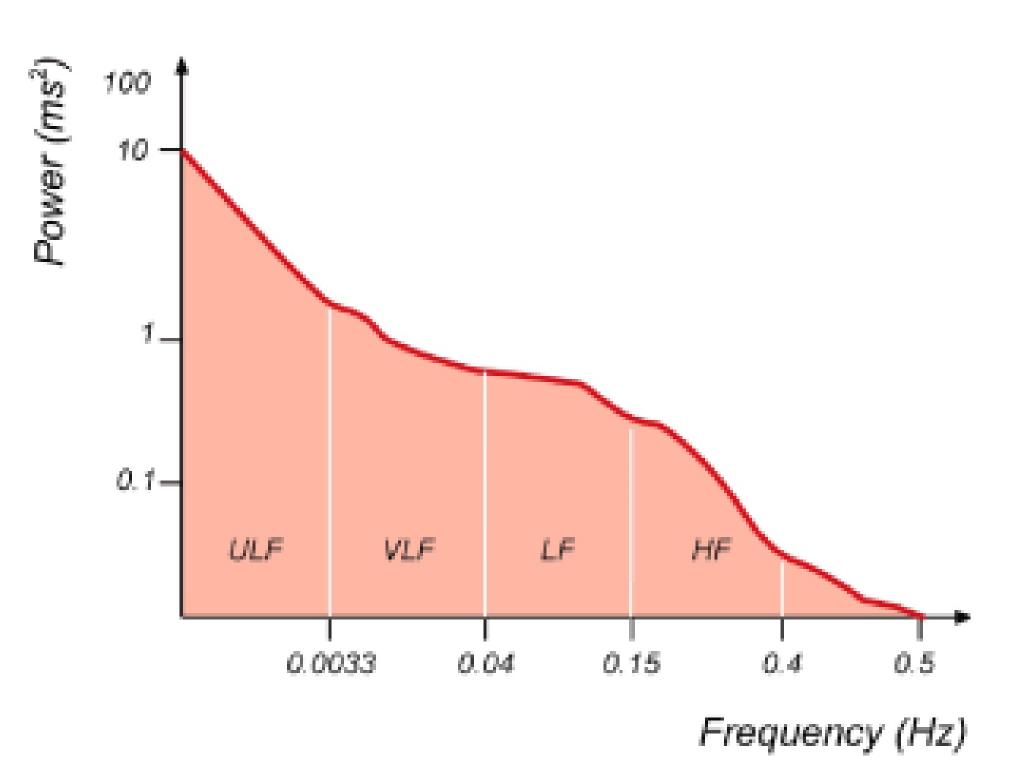
$$RMSSD = \sqrt{\frac{1}{N-1} \sum_{i=1}^{N} \left(RR_{i+1} - RR_i \right)^2}$$

successive RR interval > 20 or 50 ms

$$P(|RR_{i+1}-RR_i|>50 ms)$$

Linear methods – frequency domain

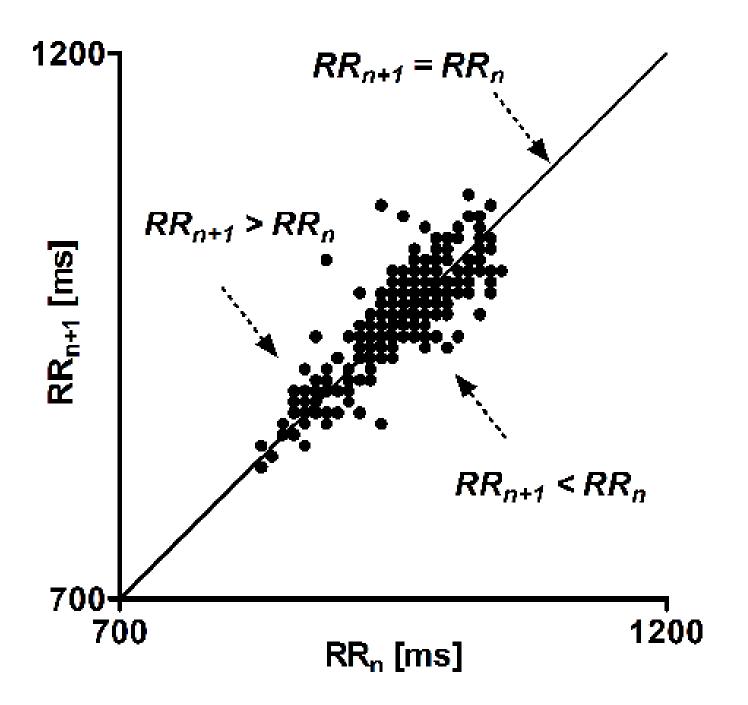
- total power (TP);
- ultra-low frequency (ULF) (<0,003Hz);
- very low frequency (VLF) (0,003-0,04Hz);
- low frequency (LF) (0,04-0,15Hz);
- high frequency (HF) (0,15-0,4Hz);



Source: Yilmaz M, Kayancicek H, Cekici Y, Heart rate variability: Highlights from hidden signals, Journal of Integrative Cardiology, January 2018, 5:265

Nonlinear method - Poincare plot

- Visual pattern whereby the shape of the plot is categorized into functional classes that indicate the degree of heart failure;
- Can be evaluated quantitatively through the computation of the SD indexes of the plot.



Source: Guzik P., Piskorski J., Krauze T., Correlation between the Poincare Plot and conventional Heart Rate Variability Parameters Assessed during Paced Breathing, The Journal of Physiological Science, March 2007, 57(1):63-71

Time irreversibility/asymmetry in HRV

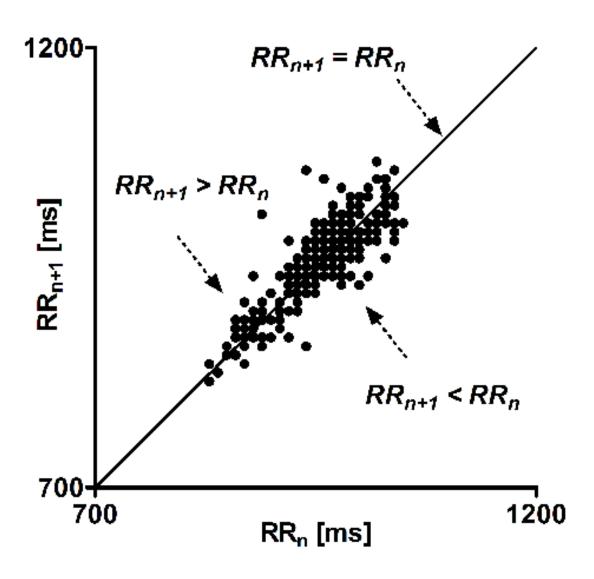
Guzik index (GI)

$$D_i = \frac{\left| RR_i - RR_{i+1} \right|}{\sqrt{2}}$$

$$GI = \frac{\sum_{i=1}^{C(P_i^+)} \left(D_i^+\right)^2}{\sum_{i=1}^{N-1} \left(D_i^-\right)^2} \times 100\%$$

Porta index (PI)

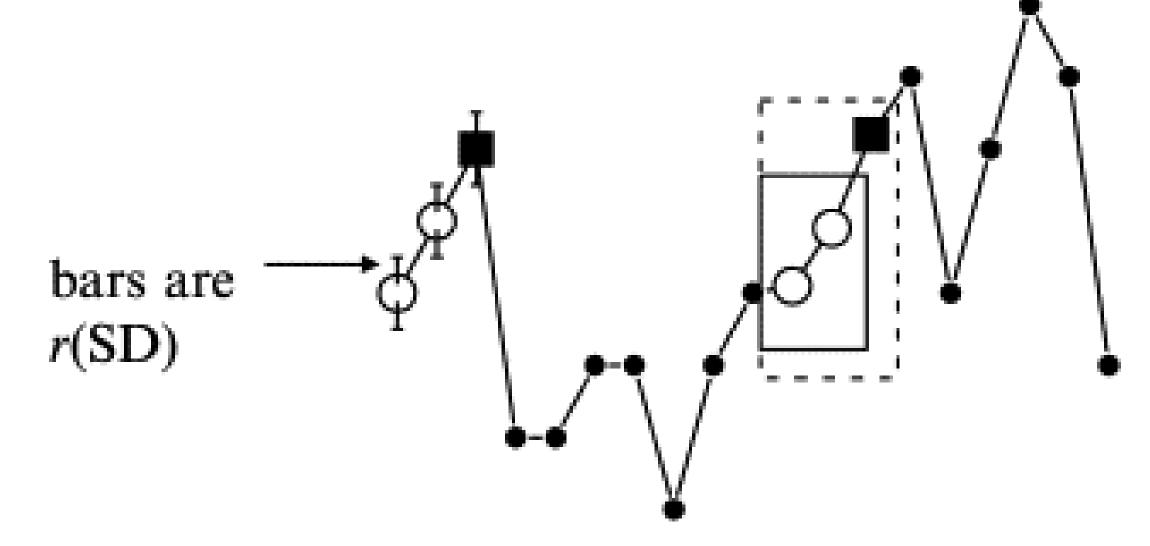
$$PI = \frac{C(P_i^-)}{C(P_i^+) + C(P_i^-)} \times 100\%$$



Source: Guzik P., Piskorski J., Krauze T., Correlation between the Poincare Plot and conventional Heart Rate Variability Parameters Assessed during Paced Breathing, The Journal of Physiological Science, March 2007, 57(1):63-71

Sample Entropy

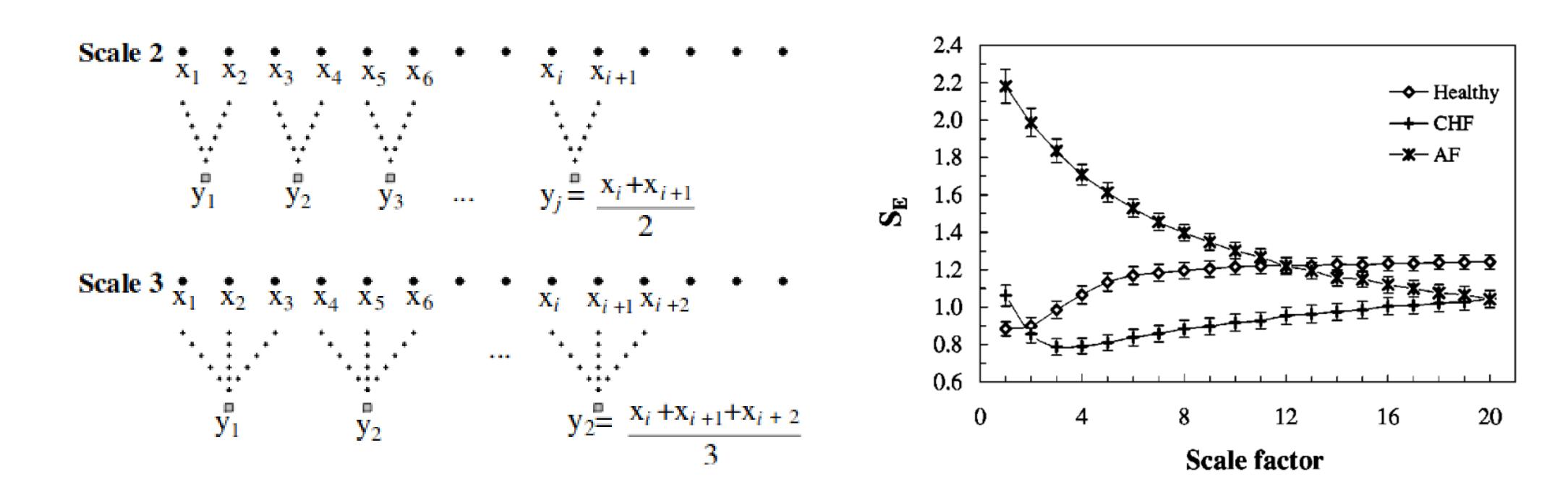
$$SampEn(m,r,N) = -\ln\left[\frac{A^{m}(r)}{B^{m}(r)}\right]$$



- \blacksquare A_i = number of matches of length m+1 with ith template
- $OB_i = number of matches of length m with ith template$

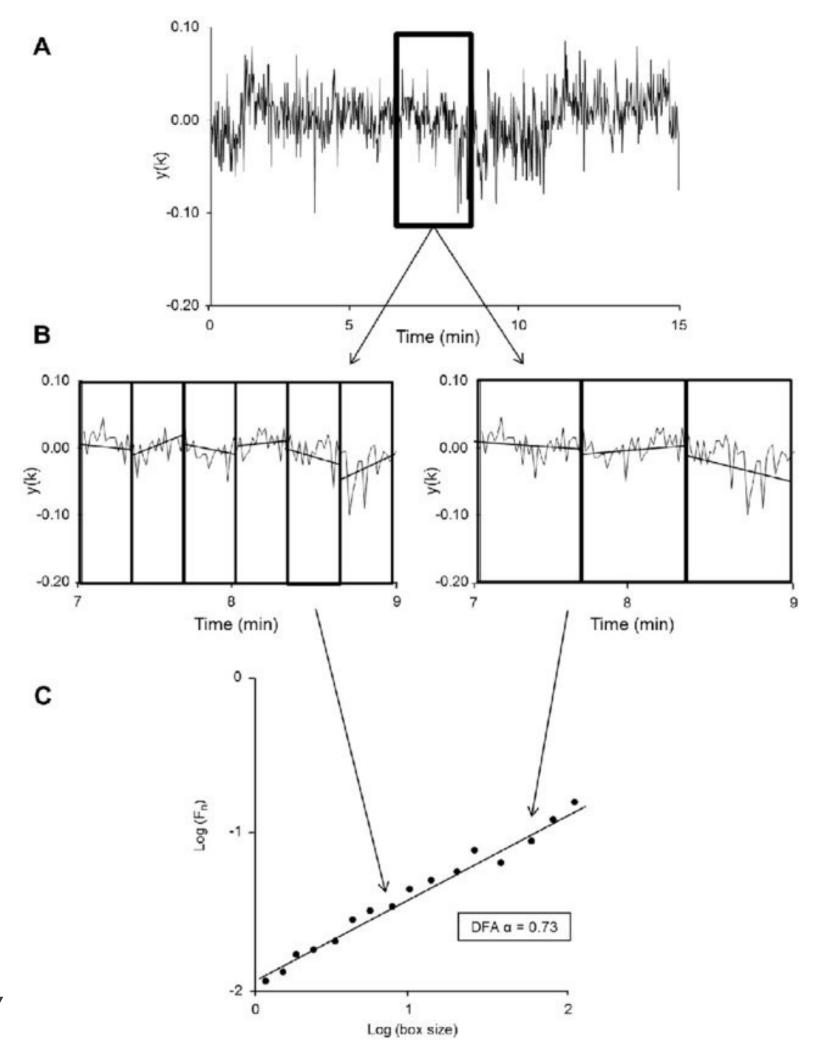
SampEn =
$$-\log ((\Sigma A_i)/(\Sigma B_i)) = -\log A/B$$

Multiscale Entropy



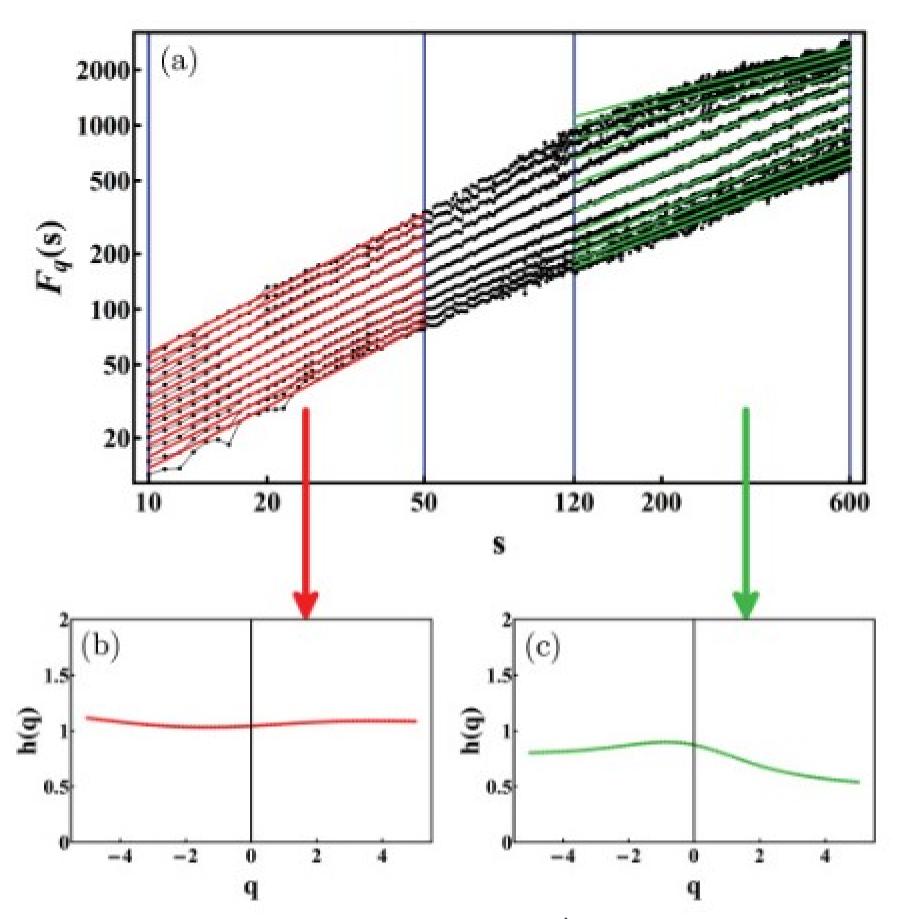
Source: Costa M.; Golderberg G.; Peng C.-K. Multiscale entropy analysis of complex physiologic time series. Physical Review Letters, 89, 2002

Multiscale Multifractal Analysis (MMA)



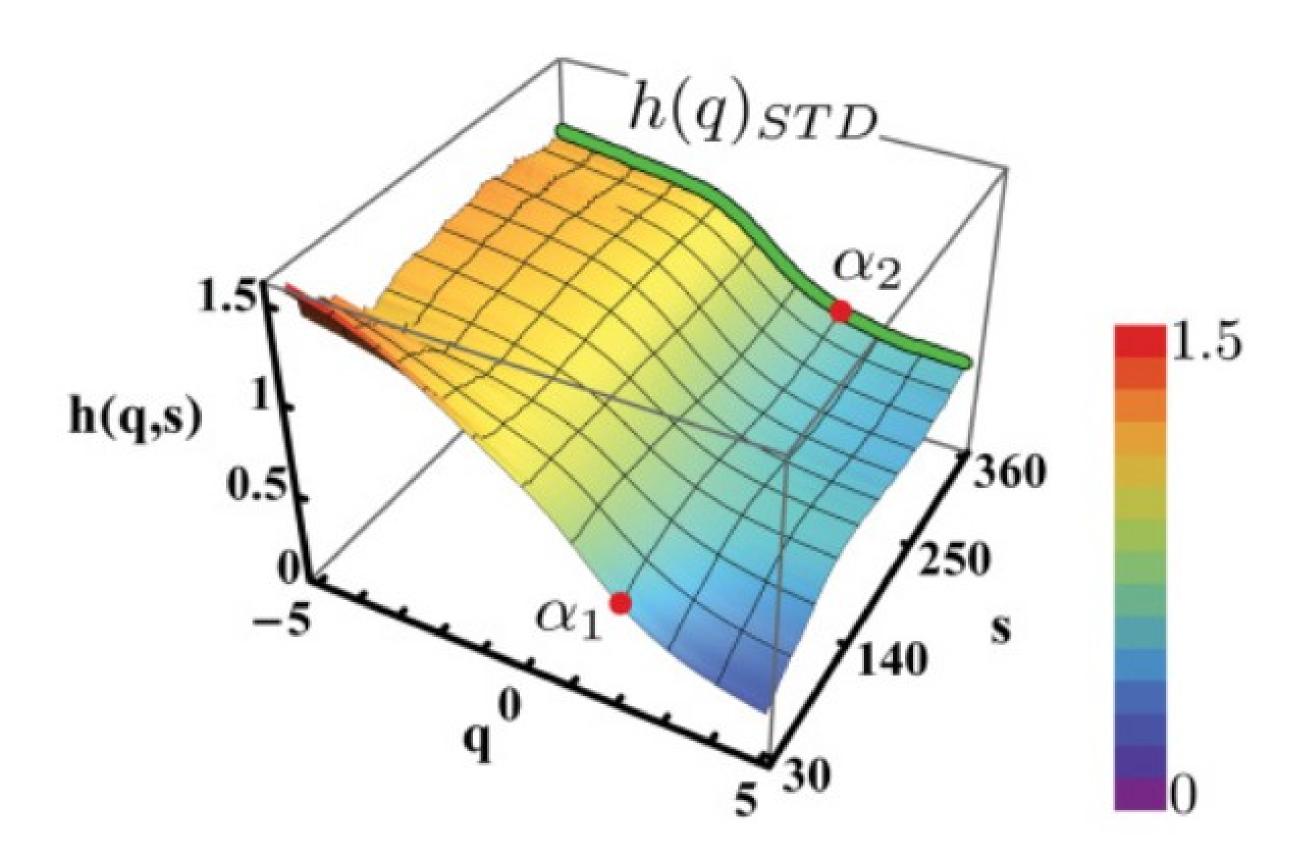
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Source: Costa M.; Golderberg G.; Peng C.-K. Multiscale entropy analysis of complex physiologic time series. Physical Review Letters, 89, 2002



Source: Gieraltowski J.; Żebrowski J.J.; Baranowski R. Multiscale multifractal analysis of heart rate variability recordings with a large number of occurrences of arrhythmia. Physical Review E, 85, 2012

Multiscale Multifractal Analysis (MMA)



Source: Gieraltowski J.; Żebrowski J.J.; Baranowski R. Multiscale multifractal analysis of heart rate variability recordings with a large number of occurrences of arrhythmia. Physical Review E, 85, 2012