Table of Contents

Volume 12  Number 1  January 2022

Time-Fractal in Living Objects
A. Szasz.................................................................1

Contribution of $^{99m}$Tc-DPD Scintigraphy in the Diagnosis of Cardiac Amyloidosis in Black Africans
D. M. Salif, N. Boucar, B. E. H. A. Lamine, D. Ousseynou,
G. Kalidou, T. Gora, N. Oumar, M. Mamadou.................................................................27

Time-Fractal Modulation—Possible Modulation Effects in Human Therapy
A. Szasz.................................................................38
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Time-Fractal in Living Objects

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Abstract
Homeostasis creates self-organized synchrony of the body’s reactions, and despite the energetically open system with intensive external and internal interactions, it is robustly stable. Importantly the self-organized system has scaling behaviors in its allometry, internal structures, and dynamic processes. The system works stochastically. Deterministic reductionism has validity only by the great average of the probabilistic processes. The system’s dynamics have a characteristic distribution of signals, which may be characterized by their frequency distribution, creating a particular “noise” $1/f$ of the power density. The stochastic processes produce resonances pumped by various noise spectra. The chemical processes are mostly driven by enzymatic reactions, which also have noise-dependent resonant optimizing. The resonance frequencies are as many as many enzymatic reactions exist in the target.

Keywords
Homeostasis, Self-Organizing, Feedbacks, Complexity, Resonance, Stochastic Processes, $1/f$ Noise, Dissipation, Enzymatic Reactions

1. Introduction
All parts of the biosystems are energetically open. The micro and macro environment have a decisional influence on their processes. The system exchanges energy and information with its environment. According to a well-defined balance, the processes are dynamic and interconnected with each other, the homeostasis [1]. This dynamic stability is self-organized [2], and despite the intensive interactions, it is robustly stable at large order of magnitudes [3]. The dynamic stability is regulated and controlled by the homeostatic feedback mechanisms [4], keeping the balance between promoters and suppressors in the complete system [5]. The living network is undoubtedly not a simple addition of its parts [6]. It forms a complex structure [7]. Theoretical biology faces a severe
challenge of complexity [8].

Regardless of its living of lifeless state build forms, the natural structures are far from the possibility to describe them in the frame of Euclidean geometry with straight lines and circles. The natural structures are self-organized and mostly form fractal structures [9]. The fractal geometry in life makes it possible to categorize the living species by their allometric comparison [10] comparison of complex morphogenetic differences [11]. This type of universality of the complex feedback mechanisms controls the dynamic equilibrium maintaining the homeostasis [12]. Fractal models represent an excellent approach to explaining the living processes’ structural development [13], even for the genetic code structure [14].

The genetic code construction uses Kronecker products (KP) of matrices with binary numbers. The construction of KP sequences the same template and so represents fractals too [15] [16]. The generated nucleotide sequences characteristic of various living systems form a fractal pattern. An extension of KP construction introduces blocks and a multifractal approach [17], which fits the living complexity [18].

The fractal description is suitable for extending the dynamic physiological processes and analyzing the fractal properties in time [19]. The time-fractal studies are based on the research of the structure of various signals [20] [21]. The dynamism of the energetically open living systems dominantly involves self-organizing processes allowing their fractal description [22]. The time fractals reflect the complex space-time approach developed a new discipline, fractal physiology [23] [24], expressing the collectivity of the processes [25].

The modulation of the external bioelectromagnetic signals has well-explained principles. The carrier frequency helps in the selection mechanisms, while its modulation supports homeostasis by its time fractal (1/f) frequency distribution [26]. The modulation could have multiple effects locally and systematically. The local force for the homeostatic control acts as a further selection factor regarding the lost control of the tumorous cells. Furthermore, the modulation forces the healthy dynamical order providing a compulsory process for apoptosis of the out-of-control cells. HRV may characterize the homeostasis [27], presenting the complexity of the system.

The well applied time-fractal current flow may activate the structural fractals in the living systems, and the personal fractal structure could modify the time-fractal pattern, too [28]. The fundamentally nonlinear physiological system dynamics work on the edge of chaos, a border of order and disorder showing a constant dynamic interplay between these states [29]. The challenge of the homeostatic equilibrium is the apparent chaos. The chaos looks complete randomness only. However, the chaos in biosystems results from the stochastic self-organizing and the energetically open system, which directly and permanently interacts with the environment. Its structural and temporal structure is fractal, which appears in the fundamental arrangements of the self-similar building and dynamism of the
energy exchanges internally and externally. The living processes are complex. They are in self-organized criticality (SOC) [30], which is formulated, as the “life at the edge of chaos” [31]. This chaos is the realization of a well-organized stochastic (probabilistic) system [32]. The chaos is only an ostensible complete disorder [33].

2. Methods

2.1. Fluctuations

An organism has a finite number of possible states. These states could be characterized in terms of operational quality utilizing a diagnostic parameter (signals). All signals have an average in time, and the signals fluctuate around this value in a controlled band. The random fluctuation sets various states (microstates) of the body, which exist only briefly and appear as fluctuation. The temporal fluctuation is regarded as a noise of the signal. The noise of living processes usually does not fluctuate randomly. The homeostatic control of the body regulates them. The minimal number of diagnostic signals is defined by the quasi-independent, weakly overlapping regulation intervals. The number of these quasi-independent diagnostic signals does not change during the system meets the conditions of the healthy dynamical equilibrium, the homeostasis. The average values, the fluctuation band, and the distribution of the frequencies may vary, depending on age and adaptation to changing environmental conditions. These quantities are called macroscopic diagnostic determinants and the status vector with \( D_i \) diagnostic states:

\[
D_i = D_i(X,Y) \quad (i = 1,2,\cdots,n)
\]

where \( X \) and \( Y \) are the signals of the system and outside environment, respectively. Due to the short time realized microstates, the number of diagnostic states is significantly less than the number of its determinant signals \( D_i \), consequently, the microstates appear as statistical statements. The same homeostatic macrostate has a wide variety of microstates that change rapidly over time, fluctuating around the averages. The probability that the microstate falls in the interval \((X, X + dX)\) at time \( t \), i.e., the probability density \( w(X,t) \) with:

\[
P_X(X < \xi \leq X + dX) = w(X,t) dX
\]

Consequently \( D_i \) is given by \( w(X,t) \) it is a stochastic determinant which primarily we characterize with its average (mean value)

\[
\langle D_i \rangle \equiv \int_X D_i(X,Y) w(X,t) dX \quad (i = 1,2,\cdots,n)
\]

and its variance

\[
\sigma_{n_i} = \left\langle (D_i - \langle D_i \rangle)^2 \right\rangle \quad (i = 1,2,\cdots,n)
\]

where \( \langle \rangle \) denotes the average of the values. The failure of the dynamic equilibrium when \( |D_i - \langle D_i \rangle| \) is larger than a predetermined threshold with a limiting value \( (l_{D_i}) \). According to the Chebyshev theorem [34] the probability that
\[ |D_i - \langle D_i \rangle| > l_{D_i} \] (so the system is out from the healthy homeostasis) is:

\[
P_{\text{fail}} \left( |D_i - \langle D_i \rangle| > l_{D_i} \right) \leq \frac{\sigma_{D_i}^2}{l_{D_i}^2} = \frac{(D_i - \langle D_i \rangle)^2}{l_{D_i}^2} \quad (5)
\]

In a healthy state the \( P_{\text{fail}} \) is small. The \( \langle D_i \rangle \) average characterizes this state. The conventional diagnostics controls \( \langle D_i \rangle \) values only, regarding the patient healthy when the fluctuations \( f_{D_i} = D_i - \langle D_i \rangle \) remain within a tolerance band \( l_{D_i} \). However, the fluctuation carries essential information about the microstates. Changes in the regulative processes could drastically modify the fluctuation of the signal without changing its average value. Study the noise spectrum may predict modifications of the regulative feedbacks, so it has diagnostic value.

The living, dynamic equilibrium is well-regulated but in a probabilistic way. The time-dependent processes realize the observed signal with a probability, as the actual exposition from the possibilities of the fluctuations of the measured signal.

The vital principle is the feedback mechanism, which controls the balance within a predetermined range around the reference value. It is usually well modeled with fuzzy logic, an approach to counting “degrees of truth” rather than the usual “true or false” decisions [35]. This logic governs homeostatic equilibria in all ranges of space and time in living systems. This uncertain value is undoubtedly in a controlled reference interval, were strongly interconnected negative feedback loops regulate the balance in the micro and macro ranges, forming the system’s dynamic stability.

These phenomena request a stochastic approach (probability of events dependent on time) instead of conventional thinking based on deterministic changes [36]. Deterministic reductionism can mislead the research. The homeostasis is often ignored and used as a static framework for effects [37]. The stochastic approach is fundamental in biological dynamism [38]. The dynamic homeostatic equilibrium keeps the system in a stable but constantly changing state.

### 2.2. Stochastic and Deterministic Approach

A model calculation of tumor growth shows the strength of the stochastic approach. In a simple example, the growth of a tumor can be described deterministically. The deterministic change of tumor mass (\( \Delta M_i \)) by observation time (\( \Delta t \)) is proportional with its actual mass (\( M_i \)):

\[
\Delta M_i (t) = k M_i (t) \Delta t
\]

where \( k \) is a constant. A well-known exponential solution uses the mass of the tumor at the start of its observation (\( M_o \)):

\[
\frac{dM_i (t)}{dt} = k M_i (t) \Rightarrow M_i (t) = M_o e^{kt} \quad (7)
\]

In a deterministic way, the prognostic task of oncology would be simple regarding exponential growth. However, the process is stochastic, requesting the
step-by-step analysis of the development of the tumor. We follow the additional or disappearing individual cells producing the mass growth. The probability $P_{M_t}$ to add a cell to the tumor at $t$ time during $\Delta t$ interval is proportional with $kM_t(t)\Delta t$, as we assumed initially been in (6). Then the probability equation with the added and eliminated cells in time interval $\Delta t$ is:

$$P_{M_t}(t+\Delta t) = P_{M_t}(t) + k(M_t-1)\Delta t P_{M_{t-1}}(t) - kM_t\Delta t P_{M_t}(t)$$

(8)

It depends on the added cells to the tumor from the previous time interval ($P_{M_t}(t+\Delta t)$) and the eliminated cells in the actual time ($-kM_t\Delta t P_{M_t}(t)$) considering the process in one step before ($k(M_t-1)\Delta t P_{M_{t-1}}(t)$). In a differential equation form:

$$\frac{dP_{M_t}(t)}{dt} = k(M_t-1)P_{M_{t-1}}(t) - kM_tP_{M_t}(t)$$

(9)

When we start from a single cell ($P_{M_t}(0)=1$ if $M_0=1$, and $P_{M_t}(0)=0$ in every other case), the solution of (9) at $M_t \geq M_0$ cases:

$$P_{M_t}(t) = \left(\frac{M_t-1}{M_t-M_0}\right) e^{-kM_0 t} \left(1-e^{-k t}\right)^{M_t-M_0}$$

(10)

Compare (7) and (10) how they are different! The deterministic approach (7) is continuous in time, running in real values, while the stochastic, probability-based approach (10) jumps on integers, building up the tumor-mass step by step. The deterministic equation gives a fixed result, while the stochastic shows “only” probability. It is interesting to see that the deterministic result is the particular case of the stochastic one, the deterministic $P_{M_t}(t) = \langle M_t(t) \rangle$ condition does not depend of the actual number of steps. Consequently, the averaging of the stochastic probability results provide the deterministic solution:

$$P_{M_t}(t) = \langle M_t(t) \rangle = \sum_{M_t=M_0} M_t P_{M_t}(t) = M_0 e^{k t}$$

(11)

2.3. The Fluctuation Phenomena

The signals follow the living, dynamic interactions, the molecular changes, and the chemical and physical excitations give a structured noise. The power spectral density of a signal ($S(f)$), is the power of the noise (fluctuation) per unit of bandwidth. Define the work of the $x(t)$ stochastic process:

$$W := \int_{-\infty}^{\infty} x^2(t) dt$$

(12)

The (12) with the Parseval’s formula may be evaluated

$$W = \int_{-\infty}^{\infty} x^2(t) dt = \int_{-\infty}^{\infty} S(f) df$$

(13)

where $S(f)$ is the spectral power density in any random stationary case. The Fourier transform of $x(t)$ stochastic process is the primary step to study the phenomena [39],

$$X(f) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{\infty} x(t) e^{-j2\pi ft} dt := F\{x(t)\}$$

(14)
where the spectral density function \( S(f) \) is:

\[
S(f) = \frac{|X(f)|^2}{2\pi}
\]  

(15)

The even function of the frequency, i.e., \( S(f) = S(-f) \).

The \( S(f) \) gives the intensity of noise as a function of spatial frequency, measured in \( W/Hz = J \), characterizing the stochastic signal with the \( f \) frequency.

The most straightforward complex noise follows normal (Gaussian) distribution (the amplitudes have normal distribution), and its power function \( S(f) \) is self-similar through many orders of magnitudes. In this simple case, the \( S(f) \):

\[
S(f) = \frac{A}{f^\alpha}
\]  

(16)

The \( \alpha \) exponent in (16) formally refers to optics, noted as the “color” of the noise. The white-noise is flat (\( \alpha = 0 \)), the pink-noise has \( \alpha = 1 \), and other colors are described by various other numbers up to \( \alpha = 2 \), the brown-noise. So, the \( S(f) \) of pink-noise inversely depends on \( f \) frequency, noted as 1/\( f \) noise. The 1/\( f \) noise carries the self-similar structure of living processes having a time-fractal covering the life’s dynamism [40] [41]. The dynamical fractal structure of living systems marks the self-organizing both in geometric and time structures and dynamically regulates the living matter [42], defines time-fractal structure in stochastic way of the living systems [43], a 1/\( f \) fluctuation. The physiological control shows 1/\( f \) spectrum [44]. One of the most studied such spectra is the heart rate variability (HRV).

This 1/\( f \) noise has a particular behavior. Each octave interval (halving or doubling in frequency) carries an equal amount of noise energy. The living system makes special signal processing due to its self-organized symmetry, so it transforms the white noise to pink [45], forming the most common signal in biological systems [46].

Stochastic signals additionally to \( S(f) \) are usually characterized by their autocorrelation function \( R_{xx}(t_1, t_2) \). The autocorrelation measures how the signal correlates with a delayed copy of itself in the function of time-lag \( (\tau = t_2 - t_1) \), measuring the signal in \( t_1 \) and subsequent \( t_2 \) in \( X \) position. The autocorrelation evaluation is a mathematical tool for finding repeating patterns, looking for periodicity in the signal. It allows identifying the existence of the biological chain processes. The \( S(f) \) and \( R_{xx}(t_1, t_2) \) functions are not independent, they could be converted to each other by Fourier transformation. Measuring the power density \( S(f) \) of a signal is easier than its autocorrelation, so usually the studies concentrate on the power density function.

3. Results

3.1. White Noise

All frequencies in the entire interval have the same \( A \) amplitude in the white
noise spectrum:

\[ S(f) = A \propto \frac{1}{f^\alpha} \]  

(17)

i.e., from (16), \( \alpha = 0 \). Consequently, the autocorrelation function is completely uncorrelated:

\[ R_{xx}(\tau) = \frac{1}{2\pi} \int_{-\infty}^{\infty} S(f) \cos(2\pi f \tau) \, df = \int_{0}^{\infty} S(f) \cos(2\pi f \tau) \, df = A \frac{1}{2} \cos(2\pi f \tau) \, df = A \frac{1}{2} \delta(\tau) \]

(18)

The band constraint in a limited interval, up to \( f_{\text{max}} \) upper-frequency limit affects a longer-term correlation:

\[ R_{xx}(\tau) = \frac{1}{2\pi} \int_{-\infty}^{\infty} S(f) \cos(2\pi f \tau) \, df = \int_{0}^{f_{\text{max}}} S(f) \cos(2\pi f \tau) \, df = A \frac{1}{2} \sin(2\pi f_{\text{max}} \tau) \]

(19)

For example, the completely flat \( S(f) \) limited to the frequency-band \([-10 - 10] \) has well-defined autocorrelation Figure 1:

The correlation function oscillates, so the correlation length does not monotonically decrease in band-limited white noise.

3.2. The 1/f Noise

A stationary random process has an indefinite duration. To introduce a modified density spectrum, consider a finite segment of the random process \( x(t) \) of duration \( 2T \), defined by:

\[ x(t) = \int_0^{2T} S(f) \cos(2\pi ft) \, df \]

\[ X(t) = \int_0^{2T} S(f) \cos(2\pi ft) \, df \]

\[ R_{xx}(\tau) = \frac{1}{2\pi} \int_{-\infty}^{\infty} S(f) \cos(2\pi f \tau) \, df = \int_0^{f_{\text{max}}} S(f) \cos(2\pi f \tau) \, df = A \frac{1}{2} \sin(2\pi f_{\text{max}} \tau) \]

Figure 1. The correlation function \( R_{xx}(\tau) \) of band-limited white noise than the \( S(f) = f^0 = \cos f \).
According to (14), the Fourier transform of \( x_T(t) \) has the form of
\[
X(f,T) = \frac{1}{\sqrt{2\pi}} \int_{-T}^{T} x_T(t) e^{-ijft} dt
\]

(21)

The Fourier transform of the function \( x(at) \), where \( a \) is an arbitrary complex number, and \( f \) is the frequency:
\[
F\{x(at)\} = \frac{1}{a} X\left(\frac{f}{a}\right)
\]

(22)

Use (21) and (22) we get:
\[
F\{x_T(at)\} = \frac{1}{a} X\left(\frac{f}{a},T\right)
\]

(23)

Using Parseval’s formula and (15):
\[
\lim_{T \to \infty} \frac{1}{2T} \int_{-T}^{T} x^2(t) dt = \int_{-\infty}^{\infty} S(f) df
\]
\[
S(f) = \frac{1}{2\pi} \lim_{T \to \infty} \left| X(f,T) \right|^2
\]

(24)

The living processes are basically self-similar, so it is convenient to define the self-similarity of a stochastic process. A stochastic process is said to be self-similar if the effective power of the stochastic process representation \( x(t) \) equals the effective power of the representation \( x(at) \) defined over time scale \([at]\), for every a positive scalar, i.e.
\[
\lim_{T \to \infty} \frac{1}{2T} \int_{-T}^{T} x^2(t) dt = \lim_{T \to \infty} \frac{1}{2T} \int_{-T}^{T} x^2(at) d(at)
\]

(25)

And so from (22) and (20), we get
\[
a \int_{-\infty}^{\infty} \frac{1}{a^2} S\left(\frac{f}{a}\right) df = \int_{-\infty}^{\infty} S(f) df
\]

(26)

Also, for the power spectral density function, the functional equation may be expressed:
\[
S\left(\frac{f}{a}\right) = aS(f)
\]

(27)

for every positive scalar \( a \) and every scalar \( f \). To solve this equation, we assume that \( f > 0 \) and set for \( a \) the value \( a = f \):
\[
S(f) = \frac{S(1)}{f}
\]

(28)

On the other hand, if \( f < 0 \) then \( f = -|f| \), and
\[
\frac{1}{a} S\left(\frac{f}{a}\right) = \frac{1}{a} S\left(\frac{-|f|}{a}\right) = S(f)
\]

(29)

Let us set for \( a \) the value \( a = |f| \) and take into account that the power density function is even, so we obtain the 1/f spectrum, or “pink-noise”:
The autocorrelation function of \( S(f) \propto \frac{1}{f} \) pink noise with Fourier transformation has a singular result:

\[
R_{xx}(\tau) = \frac{1}{2\pi} \int_{-\infty}^{\infty} S(f) \cos(2\pi f \tau) \, df
\]

\[
= \int_{0}^{\infty} \frac{1}{f} \cos(2\pi f \tau) \, df
\]

\[
= \int_{0}^{\infty} \cos(2\pi f \tau) \frac{d(2\pi f \tau)}{2\pi f \tau}
\]

follows the \( Ci(x) \) function:

\[
Ci(x) = -\int_{x}^{\infty} \frac{\cos(2\pi f \tau)}{2\pi f \tau} \frac{d\tau}{\tau}
\]

Due to \( Ci(\infty) = 0 \), the autocorrelation of 1/f noise in long time-lag is zero

Figure 2.

By the ergodic hypothesis [47], the autocorrelation function of a stationary random process \( x(t) \) can be defined as

\[
R_{xx}(\tau) = \lim_{T \to \infty} \frac{1}{2T} \int_{-T}^{T} x(t)x(t+\tau) \, dt \quad \{R_{xx}(\tau) = R_{xx}(-\tau)\}
\]

where \( \tau \) is the time-lag. The relation between autocorrelation function and the power density spectrum can be expressed by the Fourier transform of the autocorrelation function (Wiener-Khinchine theorem), namely:

\[
R_{xx}(f) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{\infty} R_{xx}(\tau) e^{-2\pi i f \tau} \, d\tau
\]

\[
R_{xx}(\tau) = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{\infty} R_{xx}(f) e^{2\pi i f \tau} \, df
\]
From these (considering [36] and [48]), we may conclude

\[ R_{\alpha}(\tau) = \int_{-\infty}^{\infty} S(f) e^{i2\pi f \tau} df = \int_{-\infty}^{\infty} \frac{S(1)}{|f|} e^{i2\pi f \tau} df = \frac{\sqrt{2\pi} S(1)}{|\tau|} \]  

(35)

Assuming the lower cutoff frequency \( f_{\text{min}} \), the function of such an approximate 1/f noise correlation from (31)

\[ \phi(\tau) = \int_{f_{\text{min}}}^{\infty} \frac{1}{f} \cos(2\pi f \tau) df = \int_{f_{\text{min}}}^{\infty} \frac{\cos(2\pi f \tau)}{2\pi f \tau} d(2\pi f \tau) = -Ci(2\pi f_{\text{min}} \tau) \]  

(36)

The procedure is also shown in Figure 2.

It can be seen from the figure that here too, there is a problem with the introduction of the correlation length since the correlation function oscillates.

In the case where the lower cutoff frequency is minimal, the argument of the \( Ci \)-function is small even at significant offset times. Then the correlation function is as shown in Figure 3.

It appears that this case can be approximated by the sum of white noise and a virtually constant correlation function. More precisely, the can be asymptotically approximated by

\[ \varphi(\tau) = -\left(\gamma + \ln(2\pi f_{\text{0}} \tau)\right) \]  

(37)

with a function where \( \gamma \approx 0.5772 \) is the Euler-Mascheroni constant.

The autocorrelation function of \( 1/f^\alpha \) (\( \alpha \neq 0 \) and \( \alpha \neq 1 \)):

\[ R_{\alpha\alpha}(\tau) = \int_{0}^{\infty} \frac{1}{f^\alpha} \cos(2\pi f \tau) df = \frac{1}{\tau^{1-\alpha}} \frac{\sqrt{\pi}}{\sqrt{2} \sin\left(\frac{\alpha\pi}{2}\right) \Gamma(1-\alpha)} \]  

(38)

Note that colored noises do not fit the white and pink noises, so the basic noises have no common expression.

The pink noise cannot be described with the classical apparatus of nonequilibrium thermodynamics. Macroscopic fluctuation characterizes the

Figure 3. Autocorrelation function 1/f noise for very low cutoff frequency (negative, integral cosine function, \(-Ci(x)\)).
thermodynamic processes. The range of space in which the fluctuation occurs is not uniform concerning the fluctuating quantity \( s \) but is thermodynamically in equilibrium at all points. The latter means that the exchange of extensive amounts characteristic of fluctuation between spatial domains during the relaxation period of equilibrium is negligible. A further feature of thermodynamic fluctuations is that the fluctuation persists for a finite time and that the rate of change of each \( a_i \) \( i = 1, 2, \cdots, n \) extensive can be expressed in terms of the extensive amounts involved in the fluctuation, i.e.

\[
\frac{da_i}{dt} = f(a_1, a_2, \cdots, a_n) \quad (i = 1, 2, \cdots, n)
\]  \( 39 \)

Let be an extensive one whose relaxation time is much longer than the others. Then the fluctuation can be described by this single extensive one. When (39) is linear and returns to the equilibrium position of the system, then the equation is a one-sided fluctuation process, completely deterministic, with no noise in it:

\[
\frac{da}{dt} = -\lambda a
\]  \( 40 \)

Solving (41):

\[
a(t) = a(0)e^{-\lambda t}
\]  \( 41 \)

Then the correlation function is:

\[
R_{aa}(\tau) = \langle a(t)a(0) \rangle = \langle a(0) \rangle^2 e^{-\lambda|\tau|}
\]  \( 42 \)

and its power spectrum:

\[
S(\omega) = \int_{-\infty}^{\infty} R_{aa}(\tau)e^{-i\omega\tau}d\tau = \langle a(0) \rangle^2 \frac{\lambda}{\lambda^2 + \omega^2}
\]  \( 43 \)

For stochasticity, the necessary noise appears in the fluctuation and spectrum for the whole, but the considerations lead to (43) are deterministic. Therefore, it is assumed that this deterministic signal is repeated randomly, forming a noise of a series of randomly repeated deterministic signals. Introducing a white noise function into the deterministic equation (like is in the Langevin equation) applies the amplitudes of the white noise spectrum that corresponds to the noise spectrum given by the deterministic random fluctuation and accordingly with the correlation function too. This is white noise \( \langle \frac{1}{\omega} \rangle \) for small \( \omega \) values, while Brown noise \( \langle \frac{1}{\omega^2} \rangle \) for large values.

In the case of pink-noise, these considerations do not work. The Fourier transform connects the \( S(f) \) power function and the \( R_\omega \) autocorrelation function:

\[ S(i\omega) \leftrightarrow R_{aa}(\tau) \Rightarrow \frac{1}{|b|} S\left(\frac{i\omega}{b}\right) \leftrightarrow R_{aa}(b\tau) \]  \( 44 \)

Because
$$S(i\omega) = \frac{1}{|\omega|}$$  \hspace{1cm} (45)$$

because of this

$$\frac{1}{|b|} S\left(\frac{i\omega}{b}\right) = \frac{1}{|\omega|}$$  \hspace{1cm} (46)$$

so it follows that

$$R_{\omega\omega}(\tau) = R_{\omega\omega}(b\tau)$$  \hspace{1cm} (47)$$

The correlation function is constant in this case, so the pink noise correlated in the same way for each shift, so there can be no thermodynamic fluctuation!

Starting with such randomized deterministic fluctuations, we get equivalents to form of (40), like:

$$\frac{da}{dt} = -\lambda a = -\frac{1}{\tau} a$$  \hspace{1cm} (48)$$

In this case, instead of (41), we get the following spectrum:

$$S(i\omega) = \int_{-\infty}^{\infty} f_{\omega}(\tau) e^{-i\omega \tau} d\tau = [a(0)]^2 \frac{\tau}{1+(i\omega\tau)^2}$$  \hspace{1cm} (49)$$

Assuming that the temporal correlation length probability density function is lognormal, the resulting noise spectrum is: $1/f^{\alpha}$. It is the same as the originally white-noise pumped stochastic case. It is confusing, of course, that this process started from deterministic distribution, but it was overcome by assuming that there is a random series of such deterministic fluctuations.

Two stochastic processes can be considered equivalent if their noise spectrum is the same. Based on this, we introduce a stochastic excitation term $q(t)$ to (48):

$$\frac{da}{dt} = -\frac{1}{\tau} a + q(t)$$  \hspace{1cm} (50)$$

The $q(t)$ spectrum is chosen of the signal resulting from the solution of the equation is equal to the power spectrum of the fluctuation (49). This can always be done. To prove this, Fourier transforms Equation (50), then we get that

$$\left( i\omega - \frac{1}{\tau} \right) a = q(\omega) \rightarrow a(\omega) = \frac{\tau}{1+(i\omega\tau)} q(\omega)$$  \hspace{1cm} (51)$$

Hence the power spectrum

$$S(\omega) = \frac{\tau^2}{1+(\omega\tau)^2} \left| q(\omega) \right|^2$$  \hspace{1cm} (52)$$

The following choice leads to the desired result:

$$q(\omega) = a(0) \frac{\sqrt{\tau}}{\sqrt{\tau}}$$  \hspace{1cm} (53)$$

Consequently, if $q(t)$ is a white noise with $a(0) \frac{\sqrt{\tau}}{\sqrt{\tau}}$ amplitude, then the noise
spectrum of the signal is the same as the noise spectrum of the fluctuation.

### 3.3. Orstein-Uhlenbeck Process

The power spectrum of a random series of such deterministic fluctuations differs from the white-noise pumped Langevin solution only in a proportionality factor. We approach the fluctuation by decomposing it into the sum of quasi-periodic stochastic processes of different statistically independent time scales. The quasi-periodic stochastic processes with different time scales also have different frequency scales. All such component processes are assumed to be statistically similar. Note the increase of a stochastic $X(t)$ process $X(t + dt) - X(t)$ without memory with $\Theta$-function:

$$ X(t + dt) - X(t) = \Theta[X(t), t, dt]. $$

(54)

Assume that $\Theta[X(t), t, dt]$ is a smooth function of the $X, t, dt$ variables and that $X(t)$ is continuous:

$$ \lim_{dt \to 0} X(t + dt) = X(t). $$

(55)

The approach that the observed noise by the emission of subsequent process-chains in statistical mechanics, the Markov process [49] describes the chain reaction, which is used in biology too [50]. The Markovian recursive successive building the $X(t + dt)$, while the function $X(t)$ from where it was derived depends only from $t$ in memory-less construction, using:

$$ \Theta[X(t), t, dt] = \sum_{i=1}^{n} X \left( t + \frac{dt}{n} \right) - X \left( t + (i - 1) \frac{dt}{n} \right) $$

$$ = \sum_{i=1}^{n} \Theta \left[ X \left( t + (i - 1) \frac{dt}{n} \right), t + (i - 1) \frac{dt}{n}, t \right] $$

(56)

Since $dt$ can be chosen to be arbitrarily small, the $t_{i-1} = t + (i - 1) \frac{dt}{n}$ can be placed in any proximity of the $t$ times by choosing $n$ large enough. Exploiting the continuity, in this case:

$$ t_{i-1} \to t, \quad X(t_{i-1}) = X(t) $$

$$ \Theta[X(t), t, \frac{dr}{n}] = \sum_{i=1}^{n} \Theta \left[ X(t), t, \frac{dr}{n} \right] $$

(57)

Here, the $\Theta \left[ X(t), t, \frac{dr}{n} \right]$ terms can be considered as representations of the $\Theta \left[ X(t), t, \frac{dr}{n} \right]$ variable that is statistically independent due to being the memory free of the process. Since $n$ is arbitrarily large, it follows from the central limit theorem that $\Theta \left[ X(t), t, dt \right]$ is the sum of $n$ statistically independent $\Theta \left[ X(t), t, \frac{dr}{n} \right]$ probability variables. Hence, this probability variable distributes normally. The following properties follow from the property of normally distributed random variables:
where \( \Theta \) notes the mean, and \( \sigma \) is the standard deviation. Solving function equations

\[
\left\langle \Theta \left[ X(t), t, d\tau \right] \right\rangle = A \left[ X(t), t \right] dt
\]

\[
\left\langle \left\langle \Theta \left[ X(t), t, d\tau \right] \right\rangle \right\rangle = D \left[ X(t), t \right] dt
\]

where \( A \) and \( D \) are smooth functions of \( X \) and \( t \), and \( D > 0 \). Considering the normality of (55) and (60):

\[
X(t+dr) - X(t) = \Theta \left[ X(t), t, d\tau \right]
\]

\[
\begin{align*}
&= N \left[ A(X,t)dr, D(X,t)dr \right] \\
&= A(X,t)dr + D^2 N(0,1)dr^2
\end{align*}
\]

where \( N(0,1) \) is the unit standard deviation squared normal distribution stochastic process with zero means. Turning to a differential equation, we get the following nonlinear generalized Langevin equation

\[
\frac{dX}{dt} = A(X,t) + \frac{1}{2} D^2 (X,t)\Gamma(t)
\]

where \( \Gamma(t) = \lim_{dr \to 0} N(0,dr^{-1}) \)

In the Gillespie sense [51], the stochastic process is self-similar, resolved to a sum of statistically independent terms normally distributed within the studied interval. Consider the simplest of the self-similar stochastic processes in (61):

\[
\frac{dX}{dt} = -\frac{1}{\tau} X + \frac{1}{\tau} D^2 \Gamma(t)
\]

where \( \tau \) is the time constant of the process.

The describes an Ornstein-Uhlenbeck process (OUP), which is stochastic and follows a normal (Gaussian) distribution. The OUP is homogeneous in time. Its homogeneity in time allows the OUP to describe it simply with the stochastic interaction of an energy source and the connected energy-consuming system Figure 4, allowing linear transformations of space and time variables [52].

The central value is exponentially decreasing, and a white noise drives it. The exponential decay should be uniformly distributed rather than lognormal, the maximum entropy belongs to \( 1/f \), and then the equation and distribution of the distribution should lead to \( 1/f \).

If we use a lognormal distribution in the interval [53], modifying (63) by

\[
D = \frac{D_0}{\sqrt{\tau}} [54]:
\]
Figure 4. The simplest relation of the energy source (reservoir, mechanical, electronic, etc.) and the linear consumer (energy-sink mechanical electronic, etc.)

\[
\frac{dX}{dt} = -\frac{1}{\tau} X + \frac{D^2}{\sqrt{\tau}} \Gamma(t)
\]  

(64)

Thus, the power spectrum of this is distributed by the lognormal of the time domain, asymptotically $1/\tau$. The equation describes the noise of a system excited by white noise consisting of an energy store (e.g., mass, rotating mass, capacitor, inductance) and a linear attenuation (e.g., fluid resistance, ohmic resistance). The power spectrum of the process:

\[
S(\omega, \tau) = \frac{D^2 \tau^2}{1 + (\omega \tau)^2}
\]

(65)

Here $\tau$ is the time constant of the system, which can also be considered the natural time scale of the stochastic process. Let’s define

\[
\lambda = \frac{1}{\tau}
\]

(66)
a frequency scale at which we want to characterize stochastic processes. Let $G(\lambda) d\lambda$ be the number of stochastic processes in the frequency interval $(\lambda, \lambda + d\lambda)$, then the energy spectrum of the stochastic processes in the interval between the frequency scales $(\lambda_2, \lambda_1)$:

\[
S(\omega, \lambda_1, \lambda_2) = \int_{\lambda_1}^{\lambda_2} \frac{D \cdot G(\lambda)}{\lambda^2 + \omega^2} d\lambda
\]

(67)

If the distribution is uniform, that is, if,

\[
G(\lambda) d\lambda = \frac{d\lambda}{\lambda_2 - \lambda_1}
\]

(68)

then we get that

\[
S(f, \lambda_1, \lambda_2) = \int_{\lambda_1}^{\lambda_2} \frac{D \cdot G(\lambda)}{\lambda^2 + \omega^2} d\lambda =
\begin{cases}
D & \text{if } 0 < \omega \ll \lambda_1 \ll \lambda_2 \\
\frac{D\pi}{2\omega(\lambda_2 - \lambda_1)} & \text{if } \lambda_1 \ll \omega \ll \lambda_2 \\
\frac{D}{\omega^2} & \text{if } \lambda_1 \ll \lambda_2 \ll \omega
\end{cases}
\]

(69)
a well-known result gives white noise in the first interval, pink in the second, and brown (Wiener noise) in the third.

When the relaxation rate is uniform in an interval \([f_1,f_2]\) and the applied amplitude doesn’t change. Hence the spectrum of OUP, \(S(f) = \frac{1}{f^\alpha}\) has three well distinguishable frequency parts Figure 5.

3.4. Importance of the Self-Similarity

The \(\tau_s\) the time constant of the system in (65) generates the stochastic signal. The \(\tau_s\) can be considered as the natural time scale of the stochastic process that characterizes the two-point correlation function of the stochastic process. Indeed, the two-point correlation function from (65) shows the degree of correlation decreases exponentially with \(\tau\) time constant:

\[
\phi_{ss}(\theta) = F^{-1}[S(\omega, \tau_s)] = F^{-1}\left[\frac{D_0 \tau_s}{1 + (\omega \tau_s)^2}\right] = D_0 e^{-\frac{\omega^2}{\tau_s}}
\] (70)

This feature of \(\tau_s\) is the temporal correlation length.

The complexity of the system involves a \(G(\tau_s)\) number of statistically independent stochastic processes in the temporal correlation length interval \((\tau_s, \tau_s + d\tau_s)\), then the resulting energy spectrum of the stochastic processes in the \((0,\infty)\) interval is:

\[
S(\omega) = \int_0^\infty D_0 \tau_s G(\tau_s) \frac{d\tau_s}{1 + (\omega \tau_s)^2}
\] (71)

when the distribution is scale variant, i.e.:

\[
G(\tau_s)\, d\tau_s = \frac{d\tau_s}{\tau_s}
\] (72)

form, then using Equation (70) a

Figure 5. The power density function is divided into three distinguishable parts in Ornstein-Uhlenbeck process. The \([f_1,f_2]\) interval, when the probability of realization of the \(f\) frequencies are equal.
improper integrated, we get the desired result:

$$S(\omega) = \int_0^\infty \frac{D_0 \tau_s G(\tau_s)}{1 + (\tau_s/\omega)} \, \text{d}\tau_s = D_0 \int_0^\infty \frac{\tau_s}{1 + (\tau_s/\omega)} \, \text{d}\tau = \frac{D_0 \pi}{2 \omega} \propto \frac{1}{f}$$  \hspace{1cm} (74)

The scale invariance means that the probability scale is independent,

$$G(\tau_s) \, \text{d}\tau_s = G(\alpha \tau_s) \, \text{d}\alpha \tau_s \Rightarrow \frac{\text{d}\alpha}{\alpha} = \frac{\text{d}\tau}{\tau_s}$$  \hspace{1cm} (75)

In the case where only self-similarity is required, e.g., as a function of density. That is

$$G(\alpha \tau_s) = \alpha^\beta G(\tau_s)$$  \hspace{1cm} (76)

then we get that

$$G(\tau_s) = \tau_s^\beta$$  \hspace{1cm} (77)

In this case \( \beta = -1 \), it provides 1/f noise. If we require only self-similarity, we get from (71) and (77) that the noise spectrum of signals in the interval \((0, \infty)\) is:

$$S(\omega) = \int_0^\infty \frac{D_0 \tau_s G(\tau_s)}{1 + (\tau_s/\omega)} \, \text{d}\tau_s = \int_0^\infty \frac{\tau_s}{1 + (\tau_s/\omega)} \, \text{d}\tau_s$$  \hspace{1cm} (78)

Due to the physical image, the integrated a

$$S(\omega) = \int_0^\infty \frac{D_0 \tau_s^{\beta+1}}{1 + (\tau_s/\omega)} \, \text{d}\tau_s = \frac{D_0}{\omega^{\beta+2}} \int_0^\infty \frac{(\omega \tau_s)^{\beta+1}}{1 + (\tau_s/\omega)} \, \text{d}(\omega \tau_s)$$  \hspace{1cm} (79)

to shape.

The integral is generally unpredictable. Fortunately, in the case of interest to us, if \(0 < \beta < 2\) the impropriety integral can be given in the closed-form:

$$\int_0^\infty \frac{(\omega \tau_s)^{\beta+1}}{1 + (\tau_s/\omega)} \, \text{d}(\omega \tau_s) = \frac{\pi}{2 \sin \left(\frac{(\beta + 2)\pi}{2}\right)} = A$$  \hspace{1cm} (80)

This gives (79) that

$$S(\omega) = \frac{D_0}{\omega^{\beta+2}} \int_0^\infty \frac{(\omega \tau_s)^{\beta+1}}{1 + (\tau_s/\omega)} \, \text{d}(\omega \tau_s) = \frac{D_0 A}{\omega^{\beta+2}}$$  \hspace{1cm} (81)

The self-similar distribution function is thus the condition a shaped power spectrum:

$$S(\omega) \propto \frac{1}{\omega^\beta}$$  \hspace{1cm} (82)

The above considerations can be generalized to a large extent.
Namely, if instead of \( D = \frac{D_0}{\sqrt{\tau}} \) in (64) use
\[
D = \frac{D_0}{\tau}
\]  
(83)

We start from the stochastic process described by the equation, using normally distributed white noise as before in (62). Then the power spectrum will be:
\[
S(\omega, \tau) = \frac{D_0 \tau^{2-\gamma}}{1+(\omega \tau)^2}
\]  
(84)

If we require only self-similarity, we get from (84) and (59) the noise spectrum of signals in the interval \((0, \infty)\):
\[
S(\omega) = \int_0^\infty D_0 \tau^{2-\gamma} G(\tau) d\tau = \int_0^\infty \frac{D_0 \tau^{\beta-\gamma+2}}{1+(\omega \tau)^2} d\tau
\]  
(85)

Due to the physical image, the integral is arranged into a form:
\[
S(\omega) = \int_0^\infty D_0 \tau^{\beta-\gamma+2} G(\tau) d\tau = \int_0^\infty \frac{D_0 \omega^{\beta-\gamma+3}}{1+(\omega \tau)^2} d(\omega \tau)
\]  
(86)

In the case of interest to us, if the \( 0 < \beta - \gamma + 3 < 2 \), the impropriety integral can be given again in closed form:
\[
\int_0^\infty (\omega \tau)^{\beta-\gamma+2} d(\omega \tau) = \frac{\pi}{2 \sin \left(\frac{(\beta - \gamma + 3)\pi}{2}\right)} = A
\]  
(87)

which gives from (78):
\[
S(\omega) = \frac{D_0}{\omega^{\beta-\gamma+3}} \int_0^\infty (\omega \tau)^{\beta-\gamma+2} d(\omega \tau) = \frac{D_0 A}{\omega^{\beta-\gamma+3}}
\]  
(88)

The self-similarity is again desired the power spectrum:
\[
S(\omega) \propto \frac{1}{\omega^d}
\]  
(89)

This result concludes to an important note: the self-similarity is a more fundamental feature of the noise than its \(1/f\) shape. Support this we derive instead of the \(1/f^d\) the noise spectrum from thermodynamic fluctuations, [55].

### 3.5. Energy Dissipation

Considering that the quantum theory of the dissipative systems is not adequately worked out, we stay within the range of the classical theory. We suppose that the pieces of information necessary for the communication are carried by the analog signals describing the physicochemical state of the individual cells. Furthermore, we are going to suppose that the self-similar Markov processes can represent the state of coaching biological subsystems. Gillespie could show that from this assumption, the equation describing the dynamics of processes can be concluded. This is the generalized Langevin equation [56]:

\[\]
\[
\frac{dX_i}{dt} = A_i(X_j,t) + D_i^{\frac{1}{2}}(X_j,t)\Gamma(t), \quad (i = 0, 1, 2, \cdots, N - 1)
\]  
(90)

where

\[
\Gamma(t) = \lim_{dt \to 0} N \left\{ 0, \frac{1}{dt} \right\}
\]  
(91)

is the white-noise with zero mean value, infinite dispersion, and normal distribution. Let us decompose the \( A_i(X_j,t) \) function into three parts:

\[
A_i(X_j,t) = f_i(t) + A_i(X_j) + \sum_{k=0}^{N-1} c_{ik} X_k
\]  
(92)

where the \( c_{ik} \) elements form a cyclic matrix.

\[
C = \begin{bmatrix}
c_0 & c_1 & \cdots & c_{N-1} \\
c_{N-1} & c_0 & \cdots & \vdots \\
\vdots & \vdots & \ddots & \vdots \\
c_1 & c_2 & \cdots & c_0
\end{bmatrix}
\]  
(93)

\( A_i(X_j) \) can be nonlinear and the \( f_i(t) \) is the time function generated by the internal active processes of the cell. It is reasonable to assume that \( A_i(X_j) \) is identical for each cell, and at the same way, we may suppose that \( D_i \) is constant for each cell. This latter can be justified because each cell is to be found in the same heat conditions. We did not assumed any confinement for the \( f_i(t) \) function. The proposed equation is the generalization of the model of the coupled damped oscillators, which showed [57] that the stochastic resonance is included in the forms of motion. We are going to examine a case where the social signal has low amplitude; therefore, the nonlinear members can be neglected. Then (91):

\[
\frac{dX_i}{dt} = f_i(t) + \sum_{k=0}^{N-1} c_{ik} X_k + D_i^{\frac{1}{2}}\Psi(t), \quad (i = 0, 2, \cdots, N - 1)
\]  
(94)

### 3.6. Cellular Communication in a Noisy Environment

The effective field strength of thermal noise was first calculated by Weaver and Astumian [58]. The Weaver & Astumian model (W-A model) assumed changes in the field strength result from fluctuations of space charges on both sides of the cellular membrane and further showed a thermal noise limit at low frequencies. Kaune [59] revisited the W-A model and showed that the field strengths typical of thermal noise converge to zero at low frequencies. Therefore, the W-A model does not describe this region appropriately. However, thermal noise in Kaune’s model [19] is assumed to be synchronized (coherent) over the entire cell membrane. This assumption is called the coherence condition. Unfortunately, thermal noise is unlikely to be coherent over a large structure such as a cell. Therefore, the calculation that followed is limited to a highly unlikely special case. Kaune set all noise-generators to be equipotential based on the coherence condition by assuming parallel connectivity and the equivalent electrical circuit. As the coherence condition does not hold in the general case, the equipotential assumption also does not hold in the general case. We generalized the problem...
and developed a solution [60]. Our results proved when there are only zero-mode currents present. The limit does not exist. However, at non-zero currents, the thermal noise does limit the efficacy of electromagnetic effects in low frequencies. The zero mode is the action by central symmetry for all individual cells instead of the translation symmetry of the usually applied outside field effects.

The topological construction is an essential factor of the cellular organization, [61], irrespective it is alive or not. The cellular structure, because of some topological reasons, develops preferring special coordination arrangements [62] and could arrange a self-organized collectivity [63] [64]. It was discovered that the division tendency is very low in the cell population, small in number [65]. For the start of a significant cell division, a critical cell density is necessary. This was later observed on a self-synchronization of chemical oscillators [66]. The topological importance was assumed in living cellular cultures also, [67], declaring that not the cell density but the position (coordination number) of cells related to each other determines what is favorable or not favorable from the point of view of division. This hypothesis was later justified experimentally [68].

The cells in developed multicellular living objects are grouped into organs to perform certain tasks in a network together. This network extends inside the cells and has suitable connection points outside the cell wall, ensuring with this to involve the cellular mechanisms in the tasks of the network. The cytoskeleton of the cells provides the basic cellular information-transfers intracellularly. The internal cytoskeleton network has transmembrane bridges (e.g., adherent connections, junctions) connecting the matrix structure on the outer side of the cell through the polar protein molecules [69]. The network develops by polymerization [70], where the water structures of aqueous electrolyte arrange the extracellular matrix partially. For example, the formed “intercellular filaments” in epithelial tissues implements the mechanical coupling of individual cells [71] [72]. Ordered water creates efficient proton conduction mechanisms [73] that disordered water does not have. The hydrogen bridges transport the protons, which is crucial in living systems [74]. This high-speed and low dissipation of the transport propagation is based on Grotthuss-mechanism [75].

The healthy cells are under the control of others in the network (“social” signaling [76], a collective action). Social information should spread within the body without loss of information. However, the environment is noisy, and the living information exchange faces this challenge. Now, we are going to prove that among the modes belonging to the eigenvectors of the matrix (9 3) of equation (91), there are modes of zero noise spectrum. It is well known that any cyclic matrix can be diagonalized by the transformation matrix [77], that is

\[
\begin{pmatrix}
1 & 1 & \cdots & 1 & \cdots & 1 \\
1 & a & \cdots & a^i & \cdots & a^{N-1} \\
1 & a^i & \cdots & a^{i^2} & \cdots & a^{i(N-1)} \\
\vdots & \vdots & \ddots & \vdots & \ddots & \vdots \\
1 & a^{N-1} & \cdots & a^{(N-1)i} & \cdots & a^{(N-1)i^2}
\end{pmatrix}
\]

\[
T = \frac{1}{\sqrt{N}}
\]
where \( a = e^{\frac{i2\pi t}{N}} \). Applying this transformation to the Equation (94), we obtain:

\[
\frac{dx_i}{dt} = \lambda_i x_i + f_i(t) + \Gamma_i(t) \quad (i = 0, \ldots, N - 1).
\]

(96)

Here the new coordinates and the eigenvalues of the cyclic matrix are

\[
x_i = \frac{1}{\sqrt{N}} \sum_{k=0}^{N-1} a^{-ik} x_k', \quad \lambda_i = D^2 \frac{1}{\sqrt{N}} \sum_{k=0}^{N-1} a^{-ik} \Gamma(t),
\]

(97)

\[
\lambda_j = \sum_{k=0}^{N-1} a^{jk} c_k, \quad f_m(t) = \frac{1}{\sqrt{N}} \sum_{k=0}^{N-1} a^{-ik} f_k(t), \quad (j = 0, \ldots, N - 1)
\]

Let us consider any one of the new noise components for which \( k \neq 0 \) (non-zero order component). Let us take the Fourier to transform thereof and consider that the amplitudes are unitary in the white-noise spectrum. Then we get that

\[
\Gamma_m(t) = D^2 \frac{1}{\sqrt{N}} \sum_{k=0}^{N-1} a^{-ik} \Gamma(t)
\]

(98)

On the other hand, we know that

\[
\sum_{k=0}^{N-1} a^{-ik} = 0
\]

(100)

In consequence, every non-zero order mode is noiseless because:

\[
\Gamma_m(t) = 0, \quad k \neq 0
\]

(101)

So the zero-order noises are not only limitless by thermal noises, but the signal exchange in such a way is noiseless.

4. Conclusion

The stochastic processes drive the homeostatic harmony, synchronizes the processes by environmental noises, while the system performs the important internal signal communications noiselessly. The dynamic stochastic living systems involve characteristic resonances. Particular resonant frequencies differentiate and describe the various enzymatic processes.

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Conflicts of Interest

The author declares no conflicts of interest regarding the publication of this paper.

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Contribution of $^{99m}$Tc-DPD Scintigraphy in the Diagnosis of Cardiac Amyloidosis in Black Africans

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Abstract

Cardiac amyloidosis presents a picture of hypertrophic cardiomyopathy with heart failure with preserved ejection fraction. It is largely underdiagnosed, especially in black Africans, and therefore falls under the category of heart disease classified as idiopathic. Light chain amyloidosis (AL) is mainly found in Caucasian subjects and the mutant variant of transthyretin (TTRm) in negroid subjects. Numerous studies have shown that ATTRm was found predominantly in black American and black British patients. In African countries the entity of idiopathic heart failure is quite important because of lack of diagnosis, ETT, MRI and immuno-histochemistry are expensive or not available. We can probably assume that the proportion of cardiac amyloidosis is quite important in black Africans. The question is if $^{99m}$Tc-DPD really easy to perform, can probably help to investigate in the nuclear medicine department in Africa. No large-scale study has been able to demonstrate the prevalence or not of cardiac amyloidosis in black-African subjects and by extension reduce this nosological entity of idiopathic heart disease. The $^{99m}$Tc-DPD scintigraphy using Perrugini’s visual sore allows localization and classification of amyloid damage. The mechanism of binding of $^{99m}$Tc-DPD to amyloid fibril depositions is not well known, its binding to TTR-type (mutated or wild type) amyloidosis is greater than the AL variant. In the diagnostic algorithm, endomyocardial biopsy is the gold standard but remains invasive, ETT with the strain allows a presumptive diagnosis and remains an operative examination dependent and is not reproducible. Cardiac MRI allows some localization of amyloid deposits but still remains less sensitive than scintigraphy. In addition, performing the whole-body MRI is very restrictive (time, antenna change and cost). The aim of this literature review was to show the superiority of...
99mTc-DPD scintigraphy compared to other diagnostic modalities and to consider its use given its simplicity when it comes to usage in Sub-Saharan Africa to diagnose the disease. Cardiac amyloidosis and by extension reduce the number of cases of heart disease classified as idiopathic and thus allow early and appropriate management.

Keywords
Amyloidosis, TTRm, AL, 99mTc-DPD, Bone Scintigraphy

1. Introduction

Amyloidosis is a heterogeneous group of diseases linked to the extracellular accumulation of insoluble fibrillar proteins: amyloid fibrils [1]. These can be hereditary or acquired diseases. They can be localized or systemic, asymptomatic or, on the contrary, have a redoubted prognosis. The clinical manifestations result from the location and type of these amyloid deposits, as well as from the toxicity of certain soluble precursors (for example immunoglobulins). These are relatively rare (underdiagnosed) pathologies, which can pose diagnostic and therapeutic problems. About twenty proteins have already been identified as potential amyloid precursors, thus determining the type of amyloidosis. Two precursors are responsible for the three types of cardiac amyloidosis: immunoglobulin light chains, involved in AL amyloidosis; mutated transthyretin (TTRm) involved in hereditary amyloidosis and wild type or wild type TTR (TTRwt) involved in senile amyloidosis. The epidemiology of amyloidosis is poorly understood around the world. Amyloidosis is described to be rare in sub-Saharan Africa [2]. Amyloidosis is mimicking a heart failure with preserved ejection fraction (HFpEF). Indeed, if we exclude the African autopsy series [3] [4] [5], less than a hundred cases of amyloidosis have been reported. The prevalence and incidence of amyloidosis in sub-Saharan Africa are likely to be underestimated and probably can be added in the “idiopathic heart failure”. In addition, the specific characteristics of the black African subject are little studied [6].

The definitive diagnosis of amyloidosis is histological. However, the cardiac biopsy is not without risk of hemorrhage, increased by the fragility of the vessels due to the disease. Cardiac amyloidosis is an infiltrative cardiomyopathy mimicking hypertensive and hypertrophic cardiomyopathies. Raising awareness and advances in imaging over the past two decades have improved the diagnosis of cardiac amyloidosis, an increasingly recognized cause of heart failure for elderly people [7].

In addition, it has been reported that black subjects presented heart failure at a much younger age and for different causes compared to the Caucasian subjects [8]. Physical examination, ECG, echocardiography, and MRI can help in the diagnosis but cannot distinguish between the different types of cardiac amyloidosis. With good sensitivity, those examinations facilitate the diagnosis of amyloidosis with cardiac involvement and thus provide assistance in the differential
diagnosis of hypertrophic cardiomyopathies.

The objective of this systematic review of the literature was to show the place of nuclear imaging and more particularly of bone scintigraphy in the diagnosis of cardiac amyloidosis, which is a significant cause of idiopathic heart failure. Its diagnosis, facilitated by DPD bone scintigraphy, would allow a better assessment of the prevalence of cardiac amyloidosis in this heterogeneous group of idiopathic heart failure.

2. Methodology

A comprehensive literature search was performed using the Medline database to identify relevant articles. We used various search algorithms based on a combination of the following terms: amyloidosis, cardiac, MRI, ETT, bone scintigraphy, amyloidosis in black African. No language restrictions were applied. All articles reporting data on the diagnostic performance of 99mTc-DPD or 99mTc-PYP scintigraphy were considered relevant and studies reporting data about patient involve in cardiac amyloidosis and particularly in black population.

We have thus analyzed the scientific publications found to first show the real problem of diagnosis of cardiac amyloidosis, then to describe the scintigraphic aspects of cardiac amyloidosis and finally to better appreciate the place of bone scintigraphy with DPD compared to other modalities of imaging in diagnosis.

3. Results and Discussion

3.1. Problem of Diagnosis

In current practice, the diagnosis is made based on a series of arguments in favor of cardiac amyloidosis. These include the clinical history of the disease, additional tests such as ETT, MRI, and extracardiac biopsies. These investigations do not make it possible to distinguish AL amyloidosis from ATTR. This is why the key examination to make the diagnosis with certainty is based on the endo-myocardial biopsy. Given the cardiovascular comorbidities of patients with suspected cardiac amyloidosis, the biopsy is not systematic, and the amyloidosis typing done by immunohistochemistry or immunofluorescence has certain limitations. The positivity of an anti-light chain antibody labeling (κ or λ) will sign AL amyloidosis. The positive labeling of transthyretin (TTR) indicates cardiac transthyretin amyloidosis [9]. Despite these performances, this examination does not make it possible to distinguish ATTRm amyloidosis from ATTRwt. The sensitivity of detection in AL amyloidosis is reduced in the event of low affinity of the antibodies for certain free chains. Some anti-TTR antibodies also frequently lack specificity and can lead to false positives. Subtyping is difficult, even in experimental laboratories, due to the background noise associated with serum contamination and epitope loss correlated with crosslinking of proteins after formalin fixation. Mass spectrometry is the gold standard for amylose typing with an accuracy of 98%. It is performed after laser microdissection of the sample, followed by formalin fixation and paraffin embedding before spectrometric
However, it is still not readily available and remains expensive. In developing countries, non-invasive examinations such as $^{99m}$Tc-DPD or $^{99m}$Tc-HMDP whole body scintigraphy may be an alternative.

The prevalence of cardiac amyloidosis is unknown and underestimated worldwide and particularly in Sub-Saharan Africa. Indeed, the causes of heart failure with preserved ejection fraction associated with myocardial hypertrophy are rarely investigated amongst black subjects.

Taking into account all the TTR mutations (~100), familial amyloidosis is disseminated in 30 countries including 15 countries where the identification of the Val30Met mutation (the valine in position 30 is replaced by methionine) has been reported, with the reservation that molecular biology techniques allowing the identification of mutations are not available in all countries. The average age of subjects with symptomatic TTR amyloidosis is 35.3 years (18 to 83 years) [10]. The most frequently identified mutation is the Val122Ile mutation (or isoleucine 122, substitution of isoleucine for valine at position 122) found in approximately 4% of African Americans regardless of location. This mutation is also common in people from West Africa. A study by Jacobson found 66 alleles of Val122Ile in DNA samples from 1688 patients [11]. However, it is likely that this mutation is a grossly underdiagnosed cause of heart failure in African and Caribbean populations, the evidence of which is based on a subgroup analysis performed on an African American population participating in the trial “Beta-Blocker Evaluation in Survival Trial (BEST)”. In addition, the prevalence and incidence of amyloidosis in sub-Saharan Africa are underestimated as the specific characteristics of the black African subject are little studied [6] and it has been reported that subjects of African origin present with heart failure at one level at younger age and for different reasons than Caucasians [7]. In fact, Cardiac amyloidosis is a significant cause of heart failure, studies have shown relatively high penetrance in African American and black British [12]. It has also been proven that mutated ATTR is more frequent in black subjects than in white subjects [13]. These facts could lead one to believe in a strong penetrance among the Negro-African subjects. All the more so since no study has assessed the prevalence of idiopathic heart disease in black Africa.

“Shah KB” and “al have” shown that cardiac amyloidosis with mutated transthyretin affects black subjects much more compared to white subjects who develop more senile cardiac amyloidosis [13]. Myocardial fixation on scintigraphy with di-phosphonates (bone tracers) is more frequent and intense in transthyretin amyloidosis (senile or hereditary) than in AL amyloidosis. The place of scintigraphy therefore appears essential in the decision-making strategy for establishing the diagnosis.

### 3.2. Cardiac Amyloidosis Scintigraphic Aspects

The $^{99m}$Tc-DPD scintigraphy using Perrugini’s visual score allows localization and classification of amyloid damage. In fact, myocardial fixation during whole body scintigraphy was found and made it possible to establish the Perugini score in favor of amyloidosis. This score uses a visual metric to establish 4 grades:
• Grade 0: absence of significant binding of the radio-tracer in projection of the cardiac area.
• Grade 1: discreet fixation of the radio-tracer of the cardiac area, inferior to the fixation of the rib grid.
• Grade 2: intense fixation of the cardiac area, superior to fixation of the rib grid.
• Grade 3: very intense fixation of the cardiac area with weak physiological fixation of the tracer by the rest of the skeleton.

Figure 1 illustrates the different grades of Perugini on scintigraphy [14]. Indeed, grade 0 does not demonstrate cardiac amyloidosis. On the other hand, a grade 1 of the Perugini score is positive and increases the sensitivity of the diagnosis of ATTR, but its specificity is poor. For other types of cardiac amyloidosis such as AL in particular, the authors found a grade 1 and more rarely a grade 2 or 3 [14].

In the etiological diagnosis, scintigraphy also has a role to play. Indeed, if its interpretation takes into account the biological results, the determination of the free light chains in the blood and in the urine, it undoubtedly makes it possible to differentiate AL amyloidosis from transthyretin amyloidosis. From the above, some authors concluded that cardiac amyloidosis scintigraphy is therefore used to confirm the diagnosis of cardiac transthyretin amyloidosis, which has a different prognosis from light chain amyloidosis. Claudio Rapezzi et al. compared the overall survival among three subtypes of cardiac amyloidosis AL, mutated ATTR and wild-type ATTR, the unadjusted two-year survival was 63% versus 98% and 100% in the mutated and wild type ATTRs, respectively [15]. The prognosis is generally more favorable in ATTR than in AL amyloidosis, although very significant differences are observed depending on the mutations. In a study by Connors & al. in patients with predominantly cardiac amyloidosis, survival was significantly better in the mutated ATTR group (n = 30) compared to the
AL group (n = 31): 27 vs. 5 months. The group consisted of 156 African American patients with amyloidosis; the most frequent mutation, V122I, was present in 36 patients, i.e. 23.7% [16]. The clinical presentation of the mutated Val122Ile ATTR frequently mimics that of wild-type (senile) ATTR with often isolated cardiac involvement, preceding carpal tunnel syndrome in some cases. Very rarely, neurological damage is found [17].

3.3. Place of Bone Scintigraphy at DPD Compared to Other Imaging Modalities

- Quantitative comparison between ETT and $^{99m}$Tc-DPD scintigraphy

In a rare study relating to the degree of cardiac amyloid deposition as assessed by $^{99m}$Tc-DPD scintographies with BNP values and echocardiographic data in patients with familial amyloidosis linked to TTR, twenty-eight patients (9 men, 19 women) of mean age 50 ± 14 years underwent, on the same day, the following examinations: BNP assay, 2D Doppler echocardiography and a $^{99m}$Tc-DPD scintigraphy. At the time of inclusion, all patients were Class I or II according to the New York Heart Association (NYHA) classification and without a clinical history of heart disease. In addition, 14 normal asymptomatic control subjects (5 men, 9 women), with a mean age of 48 ± 9 years also without a history of heart disease were recruited and had undergone 2D Doppler and stress echocardiography [18].

$^{99m}$Tc-DPD SPECT demonstrated uptake in 14 of 28 patients (50%). Cardiac uptake was mild in 5 patients (mild AC group) and severe in 9 patients (severe AC group). No patient showed moderate cardiac uptake.

At $^{99m}$Tc-DPD, the study of the ratios shows the extent of cardiac amyloid deposition because it is well correlated with the values of BNP.

Morphologic echocardiography shows a decrease in longitudinal LV strain indicating an increase in amyloid deposition, while circumferential and radial deformities increase with severe cardiac deposition as a compensatory phenomenon. These results show that when the amyloid deposit increased, an aggravation of the longitudinal strain occurred. It has been observed that trans-thoracic cardiac ultrasound is more of an entry point into the process of looking for amyloidosis [18].

The usefulness of ETT in diagnostic and etiological confirmation remains less compared to that of $^{99m}$Tc-DPD scintigraphy where the binding of $^{99m}$Tc-DPD to the amyloid deposit seems more specific to the ATTR type than to the AL type [18, 19, 20] but this binding does not seem to be able to differentiate ATTRwt from ATTRm. The elevation of BNP values only in patients with severe amyloid deposition confirms that a slight amyloid deposition is detected earlier by $^{99m}$Tc-DPD scintigraphy. From the above, it is clear that echocardiography makes it possible to detect cardiac amyloidosis but belatedly compared to the $^{99m}$Tc-DPD scintigraphy, especially since the echocardiography remains a dependent operator examination and does not allow visualization of an extracardiac extension unlike bone scintigraphy.

- Quantitative comparison between MRI and $^{99m}$Tc-DPD scintigraphy.

Other authors have also compared $^{99m}$Tc-DPD scintigraphy and MRI in the
assessment of cardiac involvement in patients with familial amyloid transthyretin polyneuropathy. For this, a study was carried out in 18 patients who received 99mTc-DPD imaging and MRI with late enhancement with gadolinium [21].

Whole body scintigraphy and SPECT revealed cardiac hyperfixation to 99mTc-DPD in 10 of 18 patients (56%) and no hyperfixation in the others. Cardiac hyperfixation was inferior to bone fixation in 5 out of 10 patients and superior to bone fixation in the remainder. MRI with late enhancement with gadolinium was positive in 8 out of 18 patients (44%). These authors specified that all the patients with late myocardial enhancement with gadolinium also presented cardiac hyperfixation of 99mTc-DPD, whereas no late myocardial enhancement with gadolinium of positive areas was found in the 8 patients without cardiac uptake of 99mTc-DPD Fig 2. All patients with positive MRI results were symptomatic (four patients with polyneuropathy, three with carpal tunnel syndrome and one patient with heart failure), while all patients with negative MRI results were asymptomatic. The LV patterns of late enhancement with gadolinium were circumferential subendocardial in one patient (the only one with signs of heart failure), focal in six patients, and diffuse in one patient. **Figure 2 and Figure 3**

![Figure 2](image-url)

**Figure 2.** Asymptomatic 50-year-old man with positive results of genetic testing for transthyretin familial amyloid polyneuropathy (Glu89Gln). (A)-(C), Technetium-99m-diphosphonate SPECT images in short-axis (A), horizontal long-axis (B), and vertical long-axis (C) views show radiotracer uptake (arrowheads) involving entire left ventricle. Left ventricle uptake was lower than that in bone. R = rib, S = sternum. (D)-(F), MRI with late gadolinium enhancement in short-axis (D), horizontal long-axis (E), and vertical long-axis (F) views show no cardiac enhancement.
Figure 3. 35-year-old man with somatic polyneuropathy and positive results of genetic testing for transthyretin familial amyloid polyneuropathy (Thr49Ala). (A)-(C), Technetium-99m-diphosphonate SPECT images in short-axis (A), horizontal long-axis (B), and vertical long-axis (C) views show radiotracer uptake (arrowheads) involving entire left ventricle. Left ventricle uptake was higher than that in bone. Also right ventricle (A), right atrium (B), and left atrium (C) show radiotracer uptake. S = sternum. (D)-(F), MRI with late gadolinium enhancement in short-axis (D), horizontal long-axis (E), and vertical long-axis (F) views show focal enhancement of left ventricle only (arrows) involving basal inferoseptal and inferior segments. Also right ventricle (arrowheads, (D)), right atrium (arrowheads, (E)), and left atrium (arrowheads, (F)) show late gadolinium enhancement.

illustrate the results of the MRI [21].

Amongst nine patients, the hyperfixation of $^{99m}$Tc-DPD affected the entire LV (17 segments in each patient). In only two out of these nine patients, the authors observed full LV involvement (one patient with a late gadolinium diffuse enhancement model and one patient with a subendocardial circumferential late gadolinium enhancement model). In the remaining seven patients, they noted less segment involvement than scintigraphy (Figure 3). Hyperfixation of $^{99m}$Tc-DPD involved RV in 8 patients and both atria in 5 patients. The burden of cardiac amyloid infiltration is considerably underestimated by visual analysis of MRI with late enhancement with gadolinium compared to imaging with $^{99m}$Tc-DPD. This is more emphasized since the $^{99m}$Tc-DPD allows a complete mapping of extracardiac uptake where full-body MRI is difficult to perform [21].

Since $^{99m}$Tc-DPD scintigraphy is superior to MRI and echo, it remains the reference imaging modality for the diagnosis of cardiac amyloidosis. It is reproducible, inexpensive, and is now widely available in black African countries. It uses a very accessible radiopharmaceutical protocol with the very practical Mo-
lybdenum generator. In fact, $^{99m}$Tc-DPD is accessible in all nuclear medicine departments in the same way as the HMDP, which is most frequently used as a bone tracer in the assessment of the extension of osteolytic cancers. This scintigraphic indication is timely because the recognition of the amyloid nature of heart disease has a direct therapeutic impact. This is because symptomatic drug treatment cannot be strictly superimposed on that for usual heart failure. Some drug classes are used with caution according to Bodez et al. [22]. In cardiac amyloidosis, the decrease in myocardial compliance leads to a decrease in the telediastolic volume of the LV. Diuretics should therefore be used with caution as they reduce the preload. These two mechanisms combined can lead to symptomatic arterial hypotension or even true hemodynamic failures. For the same reason, ACE inhibitors and angiotensin II receptor antagonists (ARA2) may be hemodynamically poorly tolerated, particularly in AL amyloidosis. Tachycardia is the predominant mechanism of adaptation to reduce ejection volume in advanced forms. The use of negative chronotropic drugs such as β blockers should therefore be avoided. These are frequently used in other causes of altered ejection fraction heart failure to increase the diastolic filling phase and thereby improve LVEF.

In a single-center study on 20 patients with ATTR with a wild subtype (10), mutant (10), the authors found a positive scintigraphy (in the sense of Perugini) at $^{99m}$Tc-PYP [23]. On average, a year and a half later, a new scintigraphy performed under the same conditions with a semi-quantitative analysis (H/CL ratio) [23] did not find any significant modification of the fixation despite the clinical progression of the disease, according to serum biomarkers, echocardiographic or NYHA score [23].

Admittedly, $^{99m}$Tc-DPD cannot allow monitoring of the evolution of the pathology but will make it possible to have a first series on the incidence of cardiac amyloidosis in subjects with african origins.

4. Conclusion

This literature review has highlighted the problem of diagnosing cardiac amyloidosis in the world and more particularly in Sub-Saharan Africa. $^{99m}$Tc-DPD can make a difference on all these heart failures so frequent in black African classified as idiopathic. DPD scintigraphy is a superior contribution to MRI and ultrasound, it remains a reproducible and inexpensive examination. In addition, it is widely available in black African countries and the supply of technetium for DPD labeling is relatively easy. Hence the interest in improving the awareness of practitioners on the contribution of scintigraphy to DPD for the diagnosis and adequate management of cardiac amyloidosis. $^{99m}$Tc-DPD can be used to have an idea of the prevalence of Cardiac amyloidosis in black African people.

Conflicts of Interest

The authors declare no conflicts of interest regarding the publication of this paper.
References


Time-Fractal Modulation—Possible Modulation Effects in Human Therapy

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Abstract

The malignant processes deviate from the healthy homeostatic control, and various “tricks” enable malignant cells to avoid the healthy regulation. Consequently, the malignant structures miss the apoptosis and proliferate without restriction, and without the formation of communication networks in the newly formed cells. The modulation supports the homeostatic control to rearrange the health regulation processes in various ways. The modulation acts with stochastic processes, using stochastic resonances for molecular excitations, supporting the regulative enzymatic processes. The number of stochastic resonant frequencies is as many as the number of enzymatic reactions. The malignant cells differ structurally and dynamically in their connections and interactions from their healthy host tissues. The radiofrequency carrier is modulated with an appropriate time-fractal (1/f) noise to select the autonomic cancer-cells, destroy them, or force the precancerous, semi-individual cells to participate in the networking connections. The modulation in this way limits the cellular autonomy of malignant cells and boosts the healthy control. The resonant energy triggers apoptotic processes and helps immunogenic actions deliver extracellular genetic information for antigen-presentation. The modulation is applied in clinical practice. The therapy (modulated electro-hyperthermia, mEHT) is intensively used in oncology in complementary applications and for palliative stages, and occasionally even as a monotherapy.

Keywords

Pink-Noise, Homeostasis, Amplitude-Modulation, Cellular-Communication, Antigen-Presentation, Stochastic Resonances, Resonance Frequencies, Cellular Networking

1. Introduction

Healthy homeostasis controls the dynamic balance in the organism, ensuring the
harmony of complex micro- and macro-interactions. Cancer destroys this harmony. The modulation goal is to find and force the malignant cells into apoptosis and to restore the healthy synchrony between cells and their communication.

The malignant processes are driven by the unicellular behavior and the individualism of the involved cells. The result is the breakdown of the integrity of the multicellular organization (which normally has a healthy networking structure). Cells with autonomy behavior have potential to better adapt to environmental changes. The transformation from the organized multicellular structure seen in healthy tissues to the structure seen in tumors is driven by the primitive transcriptional programs active in malignant cells [1]. The reorganization of the tumors structure supports the unicellular behavior and autonomy of the malignant cells, promoting the survival of the “colony” of malignant cells [2]. In an attempt to correct the abnormality, the healthy host initiates processes, such as angiogenesis, nerve healing, and numerous other supports, which instead provide essential conditions for the development of the malignancy. This regulation follows the general homeostatic control of the body. From the time that the malignancy appears in the tissue, cancer becomes a systemic disease. The dynamic control mechanisms of the healthy host are not able to repair the malignant lesion due to various reasons: genetic aberrations [3], mitochondrial dysfunction [4], and other intra-[5] and extracellular [6] hallmarks of cancer. Additional challenges which the host must face are the permanent uncontrolled stress on the system exerted by the malignancy [7], the recognition of the lesion as an unhealed wound [8], inflammation [9], and the blocking of apoptotic activity in malignant cells [10].

Cancer is an organizing (networking) disease, where the cells abandon the cooperative advantages [11]. The application of a tool that can help the homeostasis mechanisms to correct the cellular disorder of the malignant lesion and induce apoptosis, could allow for the cancerous cells, which are acting autonomously, to return to a healthy network by forcing cooperative harmony. Our objective is to show the possible effects of the modulated radiofrequency (RF) carrier, and its preclinical and clinical applications, in order to achieve this goal.

2. Methods—The Modulationphenomena

The concept of modulation applied in physiology is centered on the stochastic dynamics (time-dependent events) in the biosystems. The chosen frequency spectrum is devoted to promoting healthy controls and intended to suppress cancerous processes. The carrier frequency is in the radiofrequency (RF) range which delivers an audio range (<20 kHz) to the target. This method is well known and intensively applied by the various telecommunication networks (radio, TV, phone, GPS, etc.) to transfer information between the source and the distant targets.

The masking of the carrier can be achieved using different features of the RF wave, like its amplitude, frequency, or phase. The amplitude modulation was
historically the first form of modulation, and it is the easiest to decode the delivered information (demodulation). The resulting shape of the signal follows the shape of modulation (Figure 1); mirroring the modulation signal at both the positive and negative side of the delivery. This amplitude modulation is very vulnerable to environmental noises and absorbents.

2.1. Why Modulation and not Direct Excitations?

The question naturally arises: while the advantages of frequency modulation include better stability over distance, less vulnerability to interferences, better selectivity, why then do we not propose applying directly the modulation frequencies in order to transmit information in biological systems without carrier frequency?

The challenge is to deliver the low frequency to the body and the selected places. The low frequencies are blocked by the heterogenic isolating (capacitive) factors in the application. The adipose tissue in the layer at the skin, the various membranes, and isolation compartments block the low-frequency current because their electric impedance inversely depends on the frequency. This resistivity becomes too high in low frequencies, and no deep targeting of structures is possible. In order to overcome the adipose layer, invasive application could be introduced, but other isolations remain.

The proper solution for the direct effect would be the invasive application of low-frequency, electrodes in the tumor itself, where the low-frequency effect of isolators is negligible. However, the inserted electrodes develop a layer on the surface of electrodes in the tumor, which produces a complex impedance effect (Warburg impedance, $Z_w$) [12] and has an inverse-square-root dependence on the frequency ($f$). In a case of planar-electrode and sinusoidal supply, the Warburg impedance is [13]:

$$Z_w(f) = A \frac{(1-i)}{\sqrt{f}}$$

(1)

where $A$ is a constant and $i = \sqrt{-1}$. To avoid the Warburg impedance, a high

![Figure 1](image-url). The amplitude modulated carrier frequency. (a) The modulation with a low-frequency periodic signal, (b) the modulation with a non-periodic signal with frequency variants.
frequency is necessary. This noise could limit the accuracy of the measurements. However, in low frequencies, the large Warburg resistance also produces large electric noise because the electric noise of the resistance is proportional to the value of the actual resistivity [14].

The other advantage of the high frequency carrier is its impedance selection, which focuses the effects on the malignant cells, with the dispersion relation to attack the membrane rafts for exciting extrinsic signal pathways to apoptosis.

2.2. The Modulation Process

The electromagnetic interactions with real biological systems are faced with a heterogeneous, non-linear, and complexly regulated target. The \textit{in vitro} (cell lines), the \textit{ex-vivo} (tissue samples), the \textit{in-vivo} (animal experiments), and human applications differ from each other due to the heterogeneity and organization of the biological target. The active regulation processes in the targets, which are complex, could change the electrolytes in the tissue, varying the inhomogeneities, causing further complications in the evaluation of the interactions. The heterogeneity of the targets influences the results; however the structure and function of the cellular membranes have strong similarities, which allows for the use of some unified considerations.

The average isolation of the various membranes in the tissues is enormous $\approx 10^{13} \text{ } \Omega/m$ [15]. It is so large that it can keep $\approx 70$ mV at a distance of $\approx 7$ nm, which is equivalent to ten million volts at a distance of one meter ($=10^7 \text{ } V/m$), and its capacity is $\approx 10^{-2} \text{ } (\text{A} \cdot \text{s})/(\text{V} \cdot \text{m}^2) = \text{F/m}^2$.

The cellular membrane, regulating the ionic transport of selected ionic species in and out of the cells. The idea corresponds to the triode or transistor when the current that flows through the device is non-linearly regulated with an intermediate action (net of base), which could amplify the time-dependent signal. The membrane is also a non-linear element [16]. Applying an RF signal on the membrane, it increases non-linearly Figure 2. The rectification is not ideal because a small amount of the opposite current also exists.

Measurement of the non-temperature dependent rectification (non-linearity) is not simple because it is not measurable through the living object when the integrity of the cells is intact. The current traveling through the cell meets twice with the transmitted signal, first from outside to inside the cell, and the second meeting is in the opposite direction. The two effects eliminate each other. The lipid bilayer, together with the ionic exchanges, completes the rectification phenomena [17] [18].

We had shown [19] the white noise excited linear system with infinite freedom, and cyclic symmetry emits pink noise. It works like a special filter creating $1/f$ noise from the non-correlated white noise spectrum, which was measured [20].

The Fourier transformation of \( x(t) \) pink noise is:

\[
F(f) = \frac{A}{i2\pi\sqrt{|f|}}
\]  

(2)
Figure 2. The membranes of cells have enormously large polarization potential (≈70 mV/5 nm = 1.4 x 10^7 V/m), which non-linearity rectifies the RF current. Due to the non-linearity, the rectification increases the signal amplitude $A$ to $B$ (gain: $G = B/A$).

where $A$ is the amplitude. The noise power spectrum:

$$S(f) = F^*(f)F(f) = -i \frac{A}{2\pi \sqrt{|f|}} \cdot \frac{A}{2\pi \sqrt{|f|}} = \left( \frac{A}{2\pi} \right)^2 \frac{1}{|f|}$$

(3)

where the star is a sign of conjugation. Let be the signal function $v(t)$ of the carrier signal

$$v(t) = U_0 \cos 2\pi f_c t$$

(4)

where $f_c = 13.56$ MHz is the frequency of the carrier, $U_0$ is its amplitude (voltage in electric signal). The $f_c$ carrier frequency is in the overlapping range of $\beta/\delta$-dispersions. The Fourier transform of the modulated signal:

$$Y(f) = \frac{1}{2} F(f - f_c) + \frac{1}{2} F(f + f_c)$$

(5)

The Fourier transform of the pink noise modulated carrier is:

$$Y(f) = \frac{1}{2} \frac{AU_0}{i2\pi} \frac{1}{\sqrt{|f - f_c|}} + \frac{1}{2} \frac{AU_0}{i2\pi} \frac{1}{\sqrt{|f + f_c|}}$$

(6)

Physically, only positive frequencies can be realized, so the power spectrum is:

$$S(f) = -i \frac{1}{2} \frac{AU_0}{2\pi} \frac{1}{\sqrt{|f - f_c|}} \cdot \frac{1}{2} \frac{AU_0}{2\pi} \frac{1}{\sqrt{|f + f_c|}} = \left( \frac{AU_0}{4\pi} \right)^2 \frac{1}{|f - f_c|}$$

(7)

The power spectra of pink noise and pink noise embedded in the carrier are shown in Figure 3. The noise power spectrum shifts to the $\omega_0 = 2\pi f_0$ circular frequency of the periodic carrier.

2.2.1. Amplitude Modulation-Stochastic Resonance

Many chemical reactions have a coordinated subsequent chain, having a series of
reactions in a definite order. This set is called the Markov process, a chain-like stochastic reaction-line. When the sequence of the realized states is such, each step depends solely on the state realized in the previous event **Figure 4**.

The biological processes have well-organized and controlled Markovian chain reactions, avoiding the sudden single-step liberation of energy during the catabolism. The series of effects in the time dynamics are Markovian. A two-state Markov process in which we assume that the coefficients can be influenced by an external electric field are described as follows:

\[
\frac{d p_1}{dt} = -\alpha p_1 + \beta p_2 \\
\frac{d p_2}{dt} = \alpha p_1 - \beta p_2, \\
p_1 + p_2 = 1
\]

where \( p_i (i=1,2) \) is the probability of the actual state, and \( \alpha \) and \( \beta \) depend on the external electric field. This is the master equation of a Brownian particle bouncing back and forth in a potential well with two minima, excited by a force.

**Figure 3.** Power spectrum of pink-noise voltage amplitude modulated signal. The shifted pink-noise signal modifies the one-sided distribution towards symmetry around \( \omega_0 \).

**Figure 4.** The Markov process is a sequential series of stochastic processes with individual reaction probabilities.
of an external periodic field.

The cell works appropriately as long as the catalyst enzymes are not poisoned and, obviously, as long as the catalyst has something to catalyze. More specifically, although coenzymes provide a high degree of enzyme selectivity, there may be substances which have suppressor functions (enzyme poisons) that react in an opposite enzymatic way, or the coenzymes act to support the enzyme reaction (promoter function). On the other hand, when the cell actually does not have molecules to catalyze, the chemical reactions are terminated. We assume that the two states of the enzyme, \( A \) and \( B \), are stable but interchanging. The two states are the result of chemical reactions, with \( \alpha \) forward, and \( \beta \) backward reaction rate:

\[
A \overset{\alpha}{\underset{\beta}{\rightleftharpoons}} B
\]

The concentration of the enzyme by \([A]\) and \([B]\) in confirmation states \( A \) and \( B \). Then,

the kinetic equations of the reaction in (9):

\[
\frac{d[A]}{dt} = -\alpha[A] + \beta[B]
\]

\[
\frac{d[B]}{dt} = \alpha[A] - \beta[B]
\]

Enter the quantities in (10): \([T] = [A] + [B]\), \( p_1 = \frac{[A]}{[T]} \), \( p_2 = \frac{[B]}{[T]} \), the equation has the same form as the master Equation (8).

The study of a voltage-gated ion channel defines \( O \equiv A \), and \( C \equiv B \), where \( O \) is the open state and \( C \) is the closed state. In this way, the kinetic equation of the ion channel is again the master Equation (8).

To determine the mathematical form of transient probabilities, we examine the stationary state of (8). Then, \( \frac{dp_1}{dt} = 0 \) so

\[
\frac{p_1}{p_2} = \frac{\beta}{\alpha}
\]

(11)

The energy of the two states \( \pm \Delta E \) deviates from the reference level and assumes a Boltzmann distribution:

\[
\frac{p_1}{p_2} = \frac{\beta}{\alpha} = \frac{e^{\frac{\Delta E}{kT}}}{e^{\frac{-\Delta E}{kT}}}
\]

(12)

We obtain the Arrhenius law for reaction rates from (12):

\[
\alpha = Ce^{\frac{-\Delta E}{kT}}, \quad \beta = Ce^{\frac{\Delta E}{kT}}
\]

(13)

With this, from (8), we get a differential equation; e.g., for \( p_1 \)

\[
2\tau \frac{dp_1}{dr} = -\left( e^{\frac{\Delta E}{kT}} + e^{\frac{-\Delta E}{kT}} \right) p_1 + e^{\frac{\Delta E}{kT}}
\]

(14)
Apply Taylor series and stop at the first term, then

\[ \frac{dp_1}{dt} = -\frac{1}{\tau} p_1 - \frac{1}{2\tau} E kT \]

\[ 2\tau = \left( C e^{\frac{E_0}{kT}} \right)^{-1} \] (15)

When the field strength is a harmonic function of time, the energy depends on it linearly:

\[ \Delta E = a \sin \omega t \] (16)

From (15) and (16), we get

\[ \frac{dp_1}{dt} = -\frac{1}{\tau} p_1 + \frac{1}{2\tau} a \sin \omega t \] (17)

To find the form of the excitation, the stationary solution of the following inhomogeneous differential equation has to be determined:

\[ \frac{d}{dt} + \frac{1}{\tau} p = -\frac{1}{2\tau} a \sin \omega t \] (18)

which is

\[ \hat{p}(kT) = \frac{2a}{2kT} \frac{1}{\sqrt{\omega^2 + 1}} = \frac{a}{2kT} \sqrt{1 + \left( \frac{E_0}{2C \omega} \right)^2} \] (19)

The solution shows that the probability associated with such an excitation varies according to a harmonic function with time, regarding the noise intensity of the thermal background \( kT \). It follows a sharp maximum at a given frequency as a function of \( kT \) noise intensity Figure 5.

Formulate this result oppositely: this also means that there is a frequency where the amplitude is maximum for a given noise level. The stochastic resonance rules also such microscopic molecular structural changes like the protein folding phenomenon [21].

### 2.2.2. Demodulation Stochastic Resonance

The modulated carrier signal targets the selected malignant cells, and the cells rectify (demodulate) the received signal. The demodulation is basically a rectification process, which uses stochastic resonance [22]. The decomposition of (14):

\[ 2\tau \frac{dp_1}{dt} = -\left( 2 + \left( \frac{\Delta E}{kT} \right)^2 \right)p_1 + \frac{1}{2} \left( \frac{\Delta E}{kT} \right)^2 \]

\[ 2\tau = \left( C e^{\frac{E_0}{kT}} \right)^{-1} \] (20)

Assume the field strength of the modulated signal is a harmonic function of time and let the energy depend linearly on it

\[ \Delta E = a (1 + m \cos \Omega t) \sin \omega t \] (21)
where \( m < 1 \), the modulation depth \( \Omega \) is the modulator and the \( \omega \) carrier frequency.

Substituting this into our previous equation, we get that

\[
2\tau \frac{dp_t}{dt} = -\left( 2 + \left( \frac{a}{kT} \right)^2 \left( 1 + m \cos \Omega t \right)^2 \sin^2 \omega t \right) p_t \\
+ 1 - \frac{a}{kT} \left( 1 + m \cos \Omega t \right) \sin \omega t + \frac{1}{2} \left( \frac{a}{kT} \right)^2 \left( 1 + m \cos \Omega t \right)^2 \sin^2 \omega t,
\]

(22)

As a function of probability over time, there is a slow change and a rapid fluctuation. The first is from the modulation, and the second is from the carrier.

We are only interested in slow change now. To filter this out of the above equation, we average it over the time of the carrier period. In this case, the slow signal can be considered constant for the time functions, and the time average can be replaced by the fast one.

Under this rule, the slow change is

\[
2\tau \frac{dp_t}{dt} = -\left( 2 + \left( \frac{a}{\sqrt{2}kT} \right)^2 \left( 1 + m \cos \Omega t \right)^2 \right) p_t \\
+ 1 + \frac{1}{2} \left( \frac{a}{\sqrt{2}kT} \right)^2 \left( 1 + m \cos \Omega t \right)^2,
\]

(23)

2\tau = \left( C e^{\frac{E_0}{kT}} \right)^{-1}
equation describes.

Suppose the modulation depth is small. Then our above equation can be further simplified:

\[
2\tau \frac{dp}{dt} = -\left(2 + \left\{ \frac{a}{\sqrt{2kT}} \right\}^2 \left(1 + 2m\cos\Omega t\right) \right) p_j + 1
\]

\[
+ \frac{1}{2} \left( \frac{a}{\sqrt{2kT}} \right)^2 \left(1 + 2m\cos\Omega t\right)
\]

(24)

This is now a linear differential equation with variable coefficients.

Although there is a general solution to this, we are not going anywhere with it, so we will apply the method of successive approximation to the solution.

Since we are looking for resonance, we only study the effect of the harmonic excitation function:

\[
2\tau \frac{dp}{dt} + \left[2 + \left( \frac{a}{\sqrt{2kT}} \right)^2 + \left( \frac{a}{kT} \right)^2 \right] m\cos\Omega t \]

\[
p = \left( \frac{a}{\sqrt{2kT}} \right)^2 m\cos\Omega t
\]

(25)

Introduce the

\[
F(p) := 2\tau \frac{dp}{dt} + \left[2 + \left( \frac{a}{\sqrt{2kT}} \right)^2 \right] p,
\]

\[
f(p) := p \left( \frac{a}{kT} \right)^2 m\cos\Omega t
\]

expressions that are continuous functions of \( p \).

Steps of successive approximation:

\[
F(p_j) = \left( \frac{a}{\sqrt{2kT}} \right)^2 m\cos\Omega t,
\]

\[
F(p_2) = f(p_1),
\]

\[
F(p_i) = f(p_{i-1})
\]

(27)

Now it's true that

\[
F(p_j) + F(p_2) + \cdots + F(p_1) = F(p_j + p_2 + \cdots + p_1)
\]

\[
= \left( \frac{a}{\sqrt{2kT}} \right)^2 m\cos\Omega t + f(p_j) + f(p_2) + \cdots + f(p_{i-1})
\]

\[
= \left( \frac{a}{\sqrt{2kT}} \right)^2 m\cos\Omega t + f(p_j + p_2 + \cdots + p_{i-1})
\]

(28)

If the procedure is convergent, then if \( i \to \infty \), the line proceeds to the solution, and due to the required continuity the above expression reduces to the simple equation:

\[
F(p) = \left( \frac{a}{\sqrt{2kT}} \right)^2 m\cos\Omega t + f(p)
\]

(29)

Now let's look at the first approximation! This gives a solution of a smooth diff equation.
The amplitude of the variable probability according to the harmonic function with modulating frequency can now be determined as follows:

\[
2\tau \frac{dp_{i}}{dt} + \left[ 2 + \left( \frac{a}{\sqrt{2kT}} \right)^2 \right] p_{i} = \left( \frac{a\sqrt{m}}{\sqrt{2kT}} \right)^2 \cos \Omega t
\]  

(30)

The amplitude of the variable probability according to the harmonic function with modulating frequency can now be determined as follows:

\[
\hat{p}_{i}(RT) = \left( \frac{a\sqrt{m}}{\sqrt{2kT}} \right)^2 \frac{1}{\sqrt{4\tau^2 \Omega^2 + \left[ 2 + \left( \frac{a}{\sqrt{2kT}} \right)^2 \right]^2}}
\]

\[
= \left( \frac{a\sqrt{m}}{\sqrt{2kT}} \right)^2 \frac{1}{\sqrt{\left[ 2 + \left( \frac{a}{\sqrt{2kT}} \right)^2 \right]^2 + \left( \frac{\epsilon_{0}}{C\tau} \Omega \right)^2}}
\]  

(31)

It is a resonance again Figure 6.

2.3. Effects of Modulation in Human Therapy

The expected actions mark out the following basic goals:

1) Supports the healthy network over the unhealthy network.
2) Selectively targets the cancer.
3) Supports the immune system, boosts the homeostatic harmony and chains of reactions.
4) Excites the selected molecules in selected cells for special molecular and immunogenic changes.

Figure 6. The amplitude of the demodulation stochastic probability depends on the noise intensity too (Calculated by PTC Math Cad).
Due to the complex interconnection of the living objects, these effects are overlapping and are supporting each other.

In the following sections, we focus on emphasizing the features of modulation for fighting against cancer, which is a disease of the dynamic equilibrium breaking the multicellular cooperative network into unicellular individualism of the autonomic “selfish” cells. The task is twofold:

1) find and destroy the malignant cells. The intention is to act with the local treatment systemically, extend the task to the entire body;

2) force cellular homeostasis harmony with a homeostatic synchronization signal.

2.3.1. Boost the Healthy Network

The role of each cell supports each other in the cellular network in a tissue, there must be an interaction between them to realize this. One such possibility is the cyclic interaction in which the thermal noise is eliminated and instead a noiseless mode of information exchange appears [23]. When an abnormality occurs in cellular functions the cyclic symmetry locally brakes. The networked communication between the cells develops disturbances and that became noisy resulting in a break in the communication between cells. The thermal noise at least locally overtakes the healthy harmony. This is a change in the physiological information and is accompanied by broken psychological harmony causing a malaise state.

The social inclination is encoded by evolution into DNA. It is capable of creating a form of movement in which only noiseless modes are realized. As a result, two effects are realized, the noiseless communication, and harmony of physiochemical processes with stress and hormone dependent psychochemical ones. This later is a brain-controlled systemic process, fixing the well-being.

1) One of the crucial steps to re-establish the cyclic symmetry is the synchrony between the cellular activities. The self-synchronization of the chemical processes was observed in the famous Belousov-Zhabotinsky (BZ) reaction [24]. The BZ reactions are far from thermal equilibrium and evolve chaotic features [25], providing a chemical model of self-organized, synchronized but non-equilibrium biological processes and noise-assisted ordering [26]. The self-synchronization of chemical oscillators is also a known phenomenon [27]. An interesting example of neurological harmony is the synchrony promoted by the music affecting neuronal interactions merging the γ, β, and θ EEG signals as electrical synchrony [28]. Recognition of the harmony in 1/f noise fluctuations in music [29], connected to the short-range autocorrelation of rhythms [30]. The synchrony between the living components and the connected, synchronized coupling in living organisms is essential, and the resonance is a highly effective way to achieve it [31]. When the living system functions normally, there are noiseless modes of internal communications. The environment is however noisy, and the living information exchange faces this challenge. This means that the system must work among various external noises and must be perfectly controlled, even in changing thermal white-noise resulting from temperature in a physiologic interval.
This ensures the dynamic equilibrium, the homeostasis.

2) The physiological state includes a dynamic chemically driven psychological state in the living organism corresponding to well-being. The relationship between physiological and conscious processes has puzzled many researchers, including Schrödinger [32]. Apparently, the physicochemical processes that take place in each cell are the same, but the healthy cells are under the control of others in the network (“social” signaling [33]) which is a collective action. Social information should spread within the body without loss of information. This accompanied psychological process corrects the well-being. The realization of stochastic resonance indicates that the states of well-being are discrete. Their change is like a “resonance”. Based on a finite number of distinguishable states of well-being, it is also likely that the number of modes of communication or modes of disintegration is finite.

The stored information passed down to the new cells through the individual DNA molecules regulates the organ’s function, but cannot be designated as controllers of the described self-organized communication. Despite their ability to store all the information about the body in the nucleic DNA, the cells, due to the organ collectivity, use only those pieces of information appropriate to their tasks in the organ. The intranuclear DNA is a blueprint for the production of the new cell. This condition provides amazingly high safety in the functioning of the organ. The “social” communication allows the organ to function even when a part of it is damaged. This is essentially Neumann’s concept of realizing a high-reliability system from low-reliability ones [34].

As the network becomes extensive, the communication signal goes over greater distances in an intensively noisy environment, so more and more information is being lost by distance. In the end, the cell cannot recover the received information. Beyond a limit, noise wins, no matter how good the noise-suppressing ability of the communication network of cells is. It can be assumed that the disintegration of the information network is a process similar to the phenomenon of stochastic resonance, i.e., the addition of a small amount of noise leads to the emergence of novel, primarily chaotic behavior. It is reasonable to assume that with stochastic resonance, that cell that lost its communication commits suicide, i.e., ends up with apoptosis.

The collective excitations comprise the non-local waves and activate the energy flow in the homeostatic networks. These excitations are mainly in the low-frequency range, and the expected frequency spectrum follows the natural 1/f fluctuations [35].

The self-similar self-organizing process is collective [36] and relates to the scale-independent phenomena [37]. Terminals of a circulatory system to supply an organ adjacent to the cells are equivalent and supply the cells with the same functions equally. The collectivity subordinates the individual needs to the groups and optimizes the energy distribution for the best survival with the lowest energy consumption. This energy-share works like some kind of democracy [38].
One of the fundamental ordering principles for healthy living subjects is a characteristic “democratic” energy supply in their organisms for optimal energy consumption. The “democratic” idea is more general than just the energy supply. It characterizes the information distribution, as well [39].

This “democracy” is regulated by general biophysical rules, governed by the competition for resources in micro- and macro-phenomena, having intensive interaction with their environment. The goal of the multicellular living object is forming cooperation, optimizing the energy intake. Due to its energetically openness, the individual living object needs internal cooperative distribution of the incoming energy. The survival of its parts determines the overall survival of the object, so the optimization of energy distribution is crucial.

In healthy multicellular systems, numerous mitochondria help the individual cellular functions. Even the otherwise similar cells may vary substantially in their mitochondrial content, together with their size and membrane potential [40]. The variation arises from uneven partitioning at cell division, producing extrinsic differences in energy demand and supplied cellular processes [41]. The cells which have considerable energy demand for their normal function have a higher number of mitochondria. For example, the mitochondria fill up ~20% of the entire cell volume of human liver cells, and they work cooperatively to serve the extreme hepatocellular ATP demand [42]. The uneven mitochondrial density in different kinds of cells shows the energy distribution problem between the cells. The cells use different energies depending on their function, so the “democratic” distribution of energy between all cells equally obviously does not work. However, another kind of “democracy” works: the cells have well-controlled and balanced energy parts depending on their function in the collective. The structure of the entire system optimizes the energy distribution. The energy balance realizes a variation of the transport network supplying the variations of the demand. This directly requires that the fractal structure of transport is not unified as structure in the system, but of the formed or homeostatic demands.

The balancing of autocracy and democracy characterizes the evolution when the “selfish gene” [43] attempting to dominate all processes, but the environment-dependent self-organization requests democratic decisions; otherwise, the process blocks the system’s complexity. The game with two actions to cooperate or defect well approaches the biological interactions [44]. The relevance of the strategies of such alternating games is sometimes more appropriate than synchronous games [45].

Moreover, the complex homeostatic control shows the balancing of feedback mechanisms. The negative-feedback interaction tunes to keep the actual state as close to the accurate value, determined by the self-organizing system. Nevertheless, the system is energetically open, which is the mandatory condition for life, so the positive feedback mechanisms force some reactions, determining the metabolic processes in both directions (catabolism and anabolism). This obligatory
constraint derives a certainly autocratic line when the probability of reactions drives the defined direction so the processes. Such necessary autocratic strategies can be beneficial for exerting control over asymmetric interactions [46]. The regulatory networks have many similarities, from microbes to humans [47]. The regulation of collaboration and cooperation massively boosts the democratic character with overall genomic complexity. The regulatory effects in complexity have a propensity to a partnership supporting the democratic structure, whereas others regulate primarily in isolation, in a more autocratic fashion [48]. The degree of collaboration forming autocratic, in opposition to democratic, behavior is a specific character of the complexity in the open living systems.

Considering the complexity, the compulsory synchronization attempt by external modulated electric forces looks like a promising strategy in the cases when the collective harmony (the homeostatic regulation) is lost.

2.3.2. Selecting Mechanisms

The biochemically masked, hidden tumor cells show biophysical peculiarities. This biophysical evidence is not enough to turn the immune attention on the tumors but could select them and force the selected cells to show their factual genetic errors.

The first recognizable feature is their enlarged metabolic rate. Due to the intensive proliferation, the energy demand of these cells exceeds the rate of their healthy hosts. The ion transports in their environment increase the conductivity as a result of the higher ionic concentrations. So the adequately chosen RF current has a higher current density in the tumor region, where the electric conductivity is higher than in the healthy host. The high current density in tumor allows for the use of the RF energy to act selectively on cancer and host tissues.

The impedance measurements on Erlich solid tumors [49] identifies the particular fractal structure of malignancy. As the above discussions explained, the malignant specialties dynamically lower the conductivity of the tumor and the definite percolative self-similarity, with a lower impedance than the healthy host [50].

The biophysical selectivity has another distinguishing phenomenon due to the cellular autonomy of cancer cells [51] [52]. The intensive proliferation forms the autonomic feature of the cancer cell. The cytoskeleton of the cells frequently collapses and re-polymerises due to frequent fission. The intercellular connection is connected to the mitotic spindle [53], so the dividing cell breaks its connections with the neighbors during the division process [54]. Autonomy is a perfect concept as a hallmark of malignant cells. However, this hallmark formulates a more uncomplicated situation than the complex process works in reality [5].

Nevertheless, one factor of this phenomenon offers a selection possibility, the weakening or complete loss of the networking connections of these cells to their neighbors. The loss of the cadherin intercellular links promotes tumor growth [55], and allows increased motility of the cells [56], and changes the microenvi-
ronment around it. The rearrangement of the structure in this region, due to the autonomic behavior, changes the dielectric permittivity, distinguishing these cells' changed impedance from the healthy cells [57]. The modified environment makes the cell recognizable by the RF-current, which flows through the selected tumor.

The third selective factor uses the modified structure of the tissue’s pattern, which is a usual experience of the pathologists to identify the stage and prognosis of the malignancy. The pattern changes the impedance for the RF-current, and also its reaction to the modulation of the carrier frequency fundamentally differs when it forces healthy harmony. The well-chosen modulation delivers such a time pattern, which corresponds with the homeostatic dynamism, and is synchronized with some basic phenomena. But, the cancer is out from this harmony, and the modulation drastically differs from their dynamic properties. There is a simple analogy to represent this process with the pushing of the swing in the playground. When the push is in harmony with the swing’s pendulum, it keeps moving with a small amount of energy; however, when the energizing push is out of harmony, considerable energy is lost, and the swing loses its kinetic energy. The applied modulation (synchronization) strategy has to be harmonized with the homeostatic regulation of individual systems.

2.3.3. Control the Autonomy of Cells

Distinguishing the networked, and individual cells was studied in detail by the Nobel Laureate A. Szent-Gyorgyi. He applied an etiological approach: how does the cellular collectivity disappear [53]? He focused on the pyruvate metabolic pathways for ATP production. He termed the cell which uses the pyruvate dominantly on fermentative way α-state, which characterized the start of cellular evolution when no free oxygen was available [58]. This developmental period was unicellular, with the simple operation of life and replicated as much as possible. These individual cells compete for their demands; acting autonomously, without any cooperative communication. The appearance of free oxygen changed the game of life and made it possible to develop cooperative multicellular units, called β-state. The energy production was taken by a “power plant” in the cell, the mitochondria. Intensive diversity of life was available. The cooperatively connected cells in β-state share the tasks of living processes, to optimize the efficacy of the life.

The evolution on the cellular level is explained by an interplay of α- and β-stages [53]. The regular cooperative system in β-stages interrupted by the reproduction process when the cell breaks its collective functions, became an individual (α-state) to produce its daughter cells, and after this, integrates with the new cells into the system forming β-states. So the α-state locally terminates the multicellular cooperative complexity. The cooperative living complexity easily and naturally transforms into a basic, dividing α-state. When the network is broken, the system becomes unstable, parallel with stable, independent cells in α-state form [59]. The cells in β-state are cooperative, energizing themselves with
oxidative metabolism provided by their mitochondria. Their division should be strictly controlled by the networking cells and dynamic processes.

Normal cellular division transforms locally the β-state to α-state, which is a normal process in the multicellular system, although it hurts the systemic collectivity in that local volume. Irregularity happens when the dividing cell remains in α-state, because the individual local conditions prefer autonomy, which ensures a high amount of energy intake to the cell. The long-time optimal collective demand overcame the individual interest of the cell. This state develops cancer which is regarded as “dismantling of multicellularity” [60].

The embryonic cellular organization could be a model of how the cancerous cluster can be stabilized. The “renegade” single-cell [61] is the cell born with a genetic deviation allowing its further development to remain individual. This behavior could be the starting step of forming a malignancy. The driving principle is the mass-dependent allometry of tumor mass [62]. The observed scaling follows from the linear dependence of the relative changes of the parameters. In general, when \( c \) is a constant and \( a \) and \( b \) are the parameters, then:

\[
\frac{db}{b} = c \frac{da}{a}
\]  

The (32) scaling is observed in the allometric comparison of the various living masses and their metabolic rates [63], where the basal metabolic rate of tumor (\( B_t \)) is proportional with \( 3/4 \)th of the \( M \) mass of the tumor: \( B_t \propto M^{3/4} \), so the metabolic rate for unit volume:

\[
\frac{B_t}{M} \propto M^{-3/4}
\]  

On the other the lifespan \( T \) of the tumor also has approximately the same dependence from the mass [64]:

\[
T \propto M^{1/4}
\]  

Note that the universality of the values of exponents has intensive debates [65] [66] [67]. However, there is a consensus that the exponent \( \alpha < 1 \). This also means that as the final mass increases, the metabolism of unit mass decreases, and at the same time, the mortality rate of the cells lowers. Compare the approximated lifespan of tumor (\( T_{\text{tumor}} \)) and the healthy part of the body (\( T_{\text{healthy}} \)), using the estimate in (34):

\[
\frac{T_{\text{tumor}}}{T_{\text{healthy}}} = \left( \frac{M_{\text{tumor}}}{M_{\text{healthy}}} \right)^\alpha
\]  

where \( \alpha < 1 \) (most frequently it is \( \alpha \approx 1/4 \)) and \( M_{\text{tumor}} \) and \( M_{\text{healthy}} \) are the mass of tumor and the healthy body part which is affected by the tumor, respectively. It is obvious, that \( M_{\text{tumor}} < M_{\text{healthy}} \), so from (35)

\[
\frac{T_{\text{tumor}}}{T_{\text{healthy}}} < 1
\]  

We assume that the regulators of complete homeostatic control (proliferation,
A. Szasz

morphogenesis, physiology, immune system, etc.) have an intervention time varying according to (34). As a result of (36), the regulation time of the tumor is shorter than the healthy reaction. Hence, the tumor reacts quicker than the healthy regulation system, so its growth escapes from the standard supervision. The consequence of this is that the regulators in the body are unable to control an activated cancerous cellular cluster.

Since the embryonic cell must be connected to the food network of the adult body, it must develop and increase in its direction by a known mechanism of vascularization. But in the adult body, it is also slow as it has adapted to the adult body. The process goes through the genetic instability of cancer [68] [69]. The p53 gene is the guardian of the genome. It ensures that the genome is stable for the long term. This requires that the p53 gene maintains the appropriate level of p53 protein in the intracellular space. A change in environmental conditions is a stress effect. When it persists for a long time, permanent stress is created. Inflammation and hypoxia are lasting stress effects, reducing the level of p53 protein in the cell. The weakened stability of the genome is observed by inflammation [70], and hypoxia [71], and so increased likelihood of mutations appears. The bystander effect produces precancerous cells in the neighborhood of the cells with intensive permanent stress and the consequent genetic instability [72]. The control needs re-regulation of the tumorous area locally and also systemically [73].

2.3.4. Excite Selected Molecules

The special autonomy and high metabolic activity of the malignant cells allow recognition, and the selection and attack of them. The energy provided with the carefully chosen RF current targets the selected cells and excites a few chemical changes in the selected malignant cells.

Electromagnetic selection of the malignant cells guides the energy delivery. The living targets are inherently heterogenic in thermal and electric properties, so the structural selection at the cellular level needs additional selection to target the chosen heterogeneity of the malignant cells. For this, we have to choose the carrier frequency, acting on different parts of the tissue structure. The applied RF current triggers various processes, which are often grouped in dispersion regions by frequency, defined by the main character of the bio-electromagnetic exchanges.

The β-dispersion (~1 kHz - ~100 MHz broad spectrum) is linked to the cellular structure of biological materials [74]. It occurs at the interface of membrane-electrolyte structures, using Maxwell-Wagner relaxation [75]. The effect is an interfacial polarization of the cell membranes [76], resulting in the charge distribution at the cellular of interfacial boundaries [77]. The time dependence of the charge accumulation causes the characteristics of the β-dispersion [78]. In an experiment with sacrificed haddock muscle [79], the β-dispersion increases after a few hours of the sacrifice of the animal, showing the mechanism more connected to the cells than to the intact tissues. The high end of β-dispersion

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differs from the main effects of the range, called $\beta$1-dispersion caused from the torque of biological macro-molecules representing significant dipole moments, like proteins, orienting these against the disordering force of the thermal background [80]. However, the large molecules have limited possibility to follow the applied frequency and to grow further. Due to the massive heterogeneity of the biological tissues, this dispersion could have multiple effects, depending on the excited molecules. The conformational change of the present polymers [81], and macro-molecular relaxation with the exchange of the ionic effect in the vicinity of them [82]. The relaxation of amino-acid chains [83] and the proton fluctuations [84] also could have a role, which has no significant importance [85].

The upper tail of the $\beta$-dispersion continues to the $\delta$-dispersion (~1 MHz - ~1 GHz), which represents another mechanism. The $\delta$-dispersion occurs at a distinguished frequency range above the main $\beta$-dispersion [86] and is connected to the dipolar moments of proteins and other large molecules (like cellular organelles, biopolymers) [87]. This second Maxwell-Wagner dispersion ($\delta$) is characteristic of suspended particles surrounded by a cell [88]. Proteins also cause these effects, protein-bound water and cell organelles such as mitochondria [89] [90].

Other relaxation processes like those of molecular side chains, bound water molecules, diffusion of charged molecules, and near membrane bonds, could be added to the $\delta$-dispersion.

In addition, the $\beta$/$\delta$-dispersion of the chosen carrier frequency orients the attack on the membrane reaction of the impedance selected cells [91] [92]. The optimally chosen carrier frequency is in the interval of intensive overlap of $\beta$ and $\delta$ dispersion and using in a practical point of view is an ISM-frequency 13.56 MHz. (The ISM spectrum is reserved internationally for industrial, scientific, and medical use.) The model calculation also shows the importance of the 13.56 MHz [93]. The chosen modulation enhances the selection. The RF energy dominantly absorbs on the clustered transmembrane proteins (rafts) [94], exciting their receptors [95]. The excitation destructs the malignant cells dominantly in an apoptotic way [96] Figure 7.

### 2.3.5. Support the Immune System

Homeostatic dynamic equilibrium is too complex for external constraints to be effective in repairing it. Tightly connected feedback mechanisms regulate the system, and the reaction of homeostatic control is against any simple constraints. An excellent example of this is the response to conventional hyperthermia, which aims to kill the tumor by the thermal effect of the absorbed energy. The heating is a valid objective, but unfortunately, homeostatic control mechanisms begin to correct this heating through various actions to maintain thermal balance. The most effective reaction is increased blood flow and perfusion, which aims to cool the heated lesion. However, this feedback carries a danger, as it increases the delivery of nutrients to the tumor and promotes metastasis by cell invasion into the bloodstream.
Figure 7. The extrinsic energy absorption excites the TRAIL death receptor and induces extrinsic, caspase-dependent signal transduction to apoptosis. The stress induces double intrinsic signals going on caspase-dependent and independent pathways.

Consequently, any winning strategy to influence the dynamic equilibrium must work in conjunction with homeostatic controls, utilizing natural processes and supporting the immune system to recognize and destroy malignant cells throughout the body. The preparation of the immune system could be a perfect target for oncology treatments, instead of targeting cancer’s main strength, its proliferation. In this case, the lack of adaptive immunity to tumors can be revised, and the malignancy can be attacked by the host system itself.

The malignancy lacks the homeostatic complexity, turning the structural and dynamic conditions out of the healthy homeostatic network. Consequently, the challenging general tasks are fighting against cancer and the immune surveillance to support healthy homeostasis. The situation is complicated because the tumor cells hide their genetic information from the immune surveillance. They are well masked to avoid any systemic action against them, even the opposite: their demands are met as the system attempts to correct the irregularities. These “cheats” mislead the regulative actions and start to support the malignant cluster. Due to this falsely presented tumor status, the regular immune system does not act against it. The primary task is to present the signs of danger and stimulate the immune action to target and destroy the dangerous, highly proliferation cells.

The critical point is the immunological recognition of the malignancy. The immune system needs recognizable signs to direct its actions. However, the highly adaptive hiding strategy of malignant cells protects them from being identified by immune cells. A practical possibility for cancer invasion is the in-
nate antitumor immune action of NK cells [97] [98]. Natural Killer (NK) cells do not need information through the host’s histocompatibility complex I (MHC-I) molecules and act in the absence of priming. The cytotoxic activity of NK potentially controls tumor growth [99]. To complicate the complement of the available positive effect of NK cells, it could also promote tumor progression and angiogenesis [100] by inducing Reactive Oxygen Species (ROS) dysfunction [101]. The NK-cell activation could be achieved with a low-frequency electric field [102] as well as by electroporation [103]. Intensive low-frequency components in the spectrum of the modulated treatment may trigger the NK activity, enriching the NK cells in the targeted, selected tumor [104].

As a standard in immune homeostatic regulation, all processes have promoters and suppressors, so in complexity, every effect maybe a friend or foe, which appears for NK cell activity as well [105].

The modulation also may effectively support the healthy additive immune effects with developing tumor-specific immune reactions. The challenge is how to present the genetic information of the malignancy in order to ignite the immune system. One of the possible solutions would be immunogenic cell death (ICD). The main point is the particular apoptosis starting with an external signal with the electric field, exciting the Trail R2 (DR5) death receptor with FADD-FAS complex, and the external signal through caspase (Cas) pathway goes to Cas8 and Cas3 finishing in apoptosis [106]. The external signal pathway connects to the internal mitochondrial apoptotic path as well, and forms Bax and cytochrome-c (point of no return) through Cas9, and Cas3 [107] finishing in apoptosis again [108]. The internally excited apoptosis also can follow the caspase-independent pathway through apoptosis-inducing factor (AIF) [104], so the apoptotic end can be reached by three different signal pathways. The XIAP protein could block the main external signal path, but the modulated RF blocks this XIAP activity with Septin4 [109], and with the same function, the SMAC/Diabolo [106].

All of the above processes develop damage-associated molecular patterns (DAMP) [106]; as a factor of the immunogenic cell death (ICD) [108]. The cancer cells, during its apoptosis, in addition to apoptotic bodies, liberates calreticulin (CRT, “eat me” signal), HMGB1 (“danger” signal), and heat-shock protein 70 (HSP70, “info” signal). These processes prepare antigen recognition producing antigen-presenting cells (APCs), which trigger the presence of helper (CD4+) and killer (CD8+) T-cells with explicit recognition of the malignant cells over the body [110]. This process allows the attack of distant micro- and macro-metastases in the system (abscopal effect), turning the local treatment to systemic [73]. Noteworthy, the addition of dendritic cell medication may boost the overall immune effects [112] [113], and also an independent immune-stimulator works in harmony with modulated treatment [110]. Figure 8 shows the schematic summary of the immune processes.

2.3.6. Use Non-Thermal Effects

The amplitude modulation (AM, <20 kHz) of the RF carrier frequency can
The modulated electromagnetic carrier selects the tumor cells and excites the membrane rafts inducing apoptosis. The special process is immunogenic, and the dying tumor-cell produces damage-associated molecular pattern with calreticulin (CRT), ATP, HMGB1 and HSP70; which generate antigen presentations in lymph-nodes and develops adaptive and innate immune reactions to attack the tumor.

Intensify the tumor-specific absorption as a part of the selection mechanism [114]. Despite that, by a theoretical approach of the little energy absorption [115], the membrane demodulation effect could cause damages in the cytosol [116], or trigger apoptotic signals and destroy the cell [106]. The result of the non-thermal processes is the change of the chemical or structural situation in the targeted (investigated) compound assembly [117]. The non-thermal processes form a new phase (structural or chemical), which is a phase transition. The transition does not use heat from the field but directly uses the work of the electric field for the actual changes by absorption. The energy from the applied electric field is absorbed by the electrons or any other particles in a compound. It could be done without temperature change, but the phase of the material alters and the entropy changes without adjusting the temperature.

This phenomenon, however, has two connections to the temperature:

1) We know that 100% energy efficacy does not exist (entropy law), so the “lost” energy (which is kept as internal energy of the system) produces heat as well.

2) We know that the changing temperature changes the conditions of the “non-thermal” processes, which are in this way sharply dependent on the temperature. In other words, the reaction rate changes in the simplest case by the Arrhenius law. This complex process, which contains the thermal dependence of
the “non-thermal” effect (transition effect), is described by Eyring transition state theory, which is proposed to describe the hyperthermia where this complexity appears. In Eyring’s description, the thermal part is the same as in the Arrhenius law. Still, it has a part, which is a structural dependent factor, which could change without any temperature change, describing the “non-thermal effect” in complex systems.

The DNA damage is non-thermal. It is a phase transition, which has to be fixed by oxygen or by blocking the repairing enzymes. It does not need a temperature change at all. However, we know that both the oxygen-related or enzymatic fixing reactions (which conserves the break as it is) are temperature-dependent. The oxygen and enzyme reactions rapidly change according to temperature, mostly with Arrhenius law. The radiotherapy result has two parts: a temperature-independent and a temperature-dependent part, in the same way as this model describes the modulated electro-hyperthermia (mEHT) effect. This is why the same dose (Gray) is proposed for both the radiation processes (radiotherapy: ionizing; and mEHT: non-ionizing radiation).

3. Results—Applications

The dynamic structures, performing random stationary stochastic self-organizing processes, have practical applications in recognizing the diseases [118] [119]. The self-similarity used modeling of cancer by fractals [120], described by a generalized model [121]; helping to evaluate the various images in oncology [122]. The collectivity of the organization of the biological systems could be monitored by the fractal concept [123]. The fractal geometry describes the pathological architecture of tumors and their growth mechanisms accompanying time-dependent processes [124], and prognostic value [125].

The pink-noise modulation applies the fractal knowledge to support the harmonization efforts of homeostatic regulation and induce stochastic resonance near the frequency of the carrier, to demodulate and excite the desired signal pathways. The intensive backing of the healthy influence on the cancer is mandatory because the tumors are out from the structural and dynamic coordination of the natural organizing control.

Technically a high frequency (RF) carrier delivers the well-chosen time pattern of the modulation, which transports the modulated signal to the body with a carefully fitted applicator [126]. The applicator is driven by a voltage signal formed, as shown in Figure 9.

The spectrum and the power density of the signal which drives the applicator are studied by decomposing the applicator voltage into a signal \( s(t) \) and a noise component \( n(t) \):

\[
\begin{align*}
  u(t) &= s(t) + n(t), \\
  s(t) &= U_0 \sin \omega_0 t, \\
  n(t) &= mz(t) U_0 \sin \Omega t 
\end{align*}
\]

To calculate the power spectrum, we need to construct the autocorrelation
Figure 9. Principle of amplitude modulation with pink noise. The Ω is the carrier circular frequency, \( U_0 \) the carrier amplitude, \( m \) is the modulation depth, and \( z(t) \) is the pink noise.

Function \( u(t) \):

\[
u(t) \otimes u(t + \tau) := \lim_{T \to \infty} \frac{1}{T} \int_{-T}^{T} u(t)u(t + \tau) \, dt
\]

where the sign \( \otimes \) notes the necessary time integration of the signal with self in \( \tau \) time-lag, and where we have taken into account a sufficiently sizeable averaging time. Considering the signal and noise components, we get:

\[
u(t) \otimes u(t + \tau) = s(t) \otimes s(t + \tau) + n(t) \otimes n(t + \tau)
\]

where the average of the function formed by the product of signal and noise approaches zero. First, the properties of the \( F \{ f(t) \} \) Fourier transformation of the correlation function has to be calculated:

\[
F \{ f(t) \otimes g(t + \tau) \} = F(\omega)G^*(\omega),
F \{ f(t) \} = F(\omega),
F \{ g(t) \} = G(\omega)
\]

where the * star notation is a sign of conjugation. The Fourier transform is the pink noise in this approximation, so:

\[
F \{ z(t) \} = Z(\Omega) = \frac{1}{i\sqrt{\Omega}}
\]

thus the Fourier transform of noise:

\[
F \{ n(t) \} := \frac{i}{2} mU_0 \left[ Z\left(\omega_0 - \Omega\right) \right] = \frac{i}{2} mU_0 \left[ \frac{1}{i\sqrt{\omega_0 - \Omega}} \right]
\]

consequently, the voltage power spectrum of the applicator:

\[
\Phi_{uv}(\omega_0) := F \{ u(t) \otimes u(t + \tau) \} \:
= \frac{m^2 U_0^2}{4} \delta(\omega_0 - \Omega) + \frac{1}{\left|\omega_0 - \Omega\right|}
\]

Figure 10 shows the modulated signal spectrum. According to the Wiener-Khinchin theorem [127], the square of the voltage amplitude in the frequency interval is:
The second term of (44) shows that the square of the voltage amplitude can be extremely large near the carrier frequency, where the term tends to be infinite when $\Delta \Omega \rightarrow 0$. It follows that the average potential:

$$\Delta U^2 = \int_{\omega_0 - \Delta \Omega}^{\omega_0 + \Delta \Omega} R_{\omega_0}(\Omega) d\Omega$$

$$= \int_{\omega_0 - \Delta \Omega}^{\omega_0 + \Delta \Omega} \left( U_0^2 \delta(\omega - \Omega) + \frac{m^2 U_0^2}{4} \frac{1}{|\omega - \Omega|} \right) d\Omega$$

(44)

$$= U_0^2 + \frac{m^2 U_0^2}{4} \ln \frac{\Delta \Omega}{\Delta \Delta \Omega}$$

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(44)

$$= U_0^2 + \frac{m^2 U_0^2}{4} \ln \frac{\Delta \Omega}{\Delta \Delta \Omega}$$

(45)

And the field strength is represented as:

$$\Delta E_{\text{eff}} = \frac{\Delta U_{\text{eff}}}{d} = \frac{1}{d} \sqrt{U_0^2 + \frac{m^2 U_0^2}{4} \ln \frac{\Delta \Omega}{\Delta \Delta \Omega}}$$

(46)

where $d$ is the distance between the electrodes. Of course, arbitrarily high performance cannot be physically realized. In practice, due to fast Fourier transformation, only a point spectrum is realized. Nevertheless, the field strength amplitude can be large enough, and so the modulation does not demand a high power for appropriate effect. In conclusion, the modulation effect transmits the low-frequency power spectrum of the pink noise near the 13.56 MHz carrier, highly efficiently, and forms two sidebands. The modulation results in very high amplitude field strength peaks near the carrier frequency.

### 3.1. Modulation for Adherent Bonds

The E-cadherin-β-catenin complex plays a crucial role in cellular adhesion [128].
The suppression of E-cadherin expression leads to dysfunction in cell-cell adhesion, which can cause local invasion following tumor development [53]. Loss of adhesion has been associated with increased invasiveness and metastasis of tumors [129]. On the other hand, the appropriate expression of E-cadherins has a vital role in tumor suppression [130]. The changes in cadherin expression may affect signal transduction leading to the cancer cells growing uncontrollably [131]. This challenge also appears in the bioelectric considerations [132] of cancer development. The junctions as intercellular connections, exchanging molecules in the same way as cadherins exchange molecules [133]. Importantly the adhesive connections have a vital role in tissue repairing [134].

Together with the collective and individual states, a third entity exists too: the pluripotent embryonic resting state (δ), stem cells that adapt themselves to their actual micro-environment [135]. Healthy cells are in a communicative network, maintaining the complexity of a well-controlled system, but the δ cells could form malignant transformations [133]. The malignant cells are mostly autonomic, having lost their “social” connections with other cells, and fighting for nutrients with every other cell, irrespective of their health status. The network state categorizes cells as connected or non-connected [136]. The non-connected cells also have a network but not by cell-cell bonds. The connections are formed by clusters of fighting cells trying to trigger the healthy neighbors to supply them (forced angiogenesis) [137].

The collective activity of cells emits pink noise, which is the noise of self-organization, and so it is the noise originated from the function of homeostasis. The essence of the pink noise of homeostasis is that the function of the organ-specific networked cells is regulated actively in their energy exchange in the harmony of the complex surveillance. Consequently, the variance of physiological signals remains constant over time.

The glycocalyx shell and cell membrane of a cancer cell are also different from those of a healthy cell. Because of this, the movement of proteins in the cell membrane by lateral diffusion is also more limited. Adherents are particular protein formulas that can move in the cell membrane by lateral diffusion. Contact between two cells can occur when two such proteins meet and chemically bond with each other by lateral diffusion Figure 11. The re-established intercellular connections following mEHT have been were experimentally demonstrated [107] [138].

The glycocalyx shell of cancer cells is known to be charged highly negatively. This results in repulsion, so the attraction, which is one of the preconditions for cell recognition and the start of the adhesion process, cannot prevail. Consequently, the bond-formation is hampered by the strong negative charge of the cancer cell glycocalyx shell, so the formation of intercellular bonds and networking in malignancy is unlikely.

The condition for the formation of cellular collectivity is that the cell must be fixed with adherent bonds and junctions, which is largely missing in malignant
tumors. The modulation/demodulation stochastic resonance of noise contains signals of very high amplitude, forcing the cell clusters to intercellular bonding, manifested in communication between cells, forming functional collectivity. The self-organizing signal transformed around a carrier produces large fields. The formed gradient promotes the adhesion bonding of neighboring cells. The pink-noise modulated electric field with a very high field strength which the amplitude is transformed near 13.56 MHz, helps build the bonds of adherent proteins.

**Figure 12.**

The pink noise modulation transformed around 13.56 MHz creates a field strength gradient towards the cell contact site. Attraction by a gradient of field strength towards the point of contact occurs between the cells. Proteins are normally dipolar with polarization vector $\mathbf{p}$. Therefore, a cataphoretic force $F = (\mathbf{p} \nabla)\mathbf{E}$ acts to move the proteins toward the point of contact. The gradient

![Diagram of cell-cell interaction](image1)

**Figure 11.** The movement of adherent proteins in the cell membrane promotes the reconnection of broken E-cadherin intercellular bonds.

![Diagram of electric field](image2)

**Figure 12.** Evolution of field strength for two contact cells. The sizeable potential gradient actively promotes cellular contacts.
generates a high cataphoretic force Figure 13 which promotes the contact of the cells. The process increases the likelihood of forming intercellular bonding by the connection of adherent proteins.

Due to the increased temperature in the contact point, the barrier between the protein electronic states could be easier overcome, so the moderate temperature helps the bonding. So it is expected that the thermal effect of mEHT helps the process. However, when the temperature is high, the break of bonds is more likely.

3.2. Modulation for HSP

The heat-shock proteins (HSPs) are chaperons to help in stabilizing the dynamic equilibrium of the cells. They are named according to their molecular weight. The first observations have connected these proteins to the heat-stress [139], but later their expression for many other stresses was discovered [140] and intensive research into their role in health and diseases began [141]. The extreme stress effect promote HSP translocation to the cellular membrane [142] and liberation to the extracellular matrix. The HSP70 has dual function [143]: inside the cell protects it (chaperone function) and outside it is a danger signal to trigger immune reactions [144]. This feature makes these proteins possible therapeutic targets in various diseases [145] [146], and especially in oncology [147].

We elaborate on a theory that pink noise mEHT is suitable for enhanced HSP expression in the membrane and extracellular electrolyte. There are two important arguments for triggering the effect:

1. The carrier frequency is chosen in the range of the β/δ-dispersion, acting on lipid/protein bonds in the cell membrane.

2. The existence of high power amplitudes produced by the pink noise shifted to the carrier-frequency location.

The selection ensures that we excite the intended targeted cells with the high power pink-noise fluctuation near the carrier frequency. The RF-current density is increased at the cell membrane due to the β/δ-dispersion. The carrier would shift the pink noise into the β/δ-dispersion range, and there we expose the membrane to the pink noise. Streamlines pass from one cell to another at the points of contact of the cells. Electric fields with very high field strengths and gradients are generated in these places Figure 14.

Figure 13. Formation of cataphoretic force. The inhomogeneous electric field turns the proteins in extracellular electrolyte.
The field gradients are shown in Figure 12 promote the dielectric forces and induces large SAR at the contact point and reconnects the transmembrane proteins as well as provides mild heating for the cells [94] [148].

The connection points have high local SAR. However, due to the high field strength gradient enhanced by pink noise, a dielectrophoretic force also occurs, which moves half of this high SAR location to the micro-domains of the membrane thermo-sensor. The micro-domains are exposed to increased stress due to high power amplitudes. On the one hand, the stress effect is heat stress due to the high SAR and, on the other hand, mechanical deformation stress due to the high field strength gradient.

The β/δ-dispersion is used to produce HSP, and the selected target triggers the production of HSP70 for immune stimuli [149]. The immunomodulation of extracellular HSP70 makes it a promising candidate for vaccination of cancerous diseases by presenting malignant cells’ antigens [150]. The danger signal of extracellular HSP is an essential mediator for intercellular signaling and transport [151]. The molecule is part of the damage-associated molecular pattern formed by immunogenic cell death, making a perfect way to produce CD8+ killer T-cells for a systemic fight against cancer.

3.3. Modulation for Cellular Communication

The biological transformation of normal (β) to cancerous (α) is a communication phase transition which is explained on the level of individual cells [136]. The starting point of Szent-Gyorgyi is an etiological approach: the disappearance of cellular collectivity [53]. The $\beta \rightarrow \alpha$ state change driven by changes of the micro-environment of the malignant cells [4].

The division naturally needs higher energy income than the regular life of the cells. The cytoskeleton polymerization during the cell division breaks the networking connections, allowing the free formation of the new cytoskeleton and intensifying support of the microenvironment of the dividing cells by higher diffusion in the free electrolyte. After finishing the division, the cells are reconnected, the network control activates again. The internal energy resources pro-
mote the appropriate, satisfactory amount of energy. Together with the environmental sources, the cells build up the daughter cells with the healthy cellular division [152]. However, the malignant cells have no reserves for the extreme energy demand to proliferate [153]. The usual healthy process does not work in malignant proliferation. The tumor cells do not return to the network. They remain as individuals and continue the proliferation. One explanation concentrates on the dielectric changes in the cell microenvironment [154], which is high in the cancerous process in vivid form [155]. Noisy communication can also cause cancer. This can occur when the relative arrangement of the cells does not allow for an interaction leading to cyclic symmetry, like requested [34]. From our perspective, this can be interpreted, meaning that if the symmetry is not cyclic, it is divisible. Otherwise, it is not, which supports the idea, that the cellular position has a vital role in favoring the malignant proliferation [156].

4. Discussion—Cancer Therapy

The strategy to eliminate the cancerous processes focuses on the weakest point of the tumor: individualism. The missing overall network makes them vulnerable, as they are alone in fighting the stress.

The malignant development misses the healthy homeostatic regulation. The systemic controls do not work. Their extreme energy demand drives their systemic effect. The cancerous process concentrates on intensive proliferation and uses healthy hosts to support it. This complex process misleads the healthy host, presenting their actions as normal energy-demand for healing the lesion like a wound (as was described shown 25 years ago [88]). Cancer presents itself as trying to heat, and the system cannot recognize the “cheating” due to the hidden identification signs. This shows that the entire malignant process is essentially a wound repair process [157]. The idea has been investigated more recently, with comparisons between the hallmarks of the two processes [158], demonstrating the role of this mechanism in the formation of metastases [159]. The bio-system falsely recognizes the tumor-like wound [160] and stimulates its environment to heal the irregularity (meaning to produce cells to heal).

The permanent reparation demand depletes the available stem cells, [161] [162], and is emphasizes a different repair pathway: the proto-oncogenes are activated, and malignant transformation could happen, inducing the clinical manifestation of cancer [163]. The malignant transformation of the wound results in the secretion of proto-oncogenes [164] [165] [166], and intensive capture and stimulation of the stem-cells from other places [167] [168].

Ignoring the healthy regulations and the collectivity of the host tissue has some structural and temporal signs to recognize it. The structure of the individual cells is more disordered than the networked healthy tissue, and the intensive individual metabolic demand changes the structure of extracellular electrolyte. The permanent cellular division periodically collapses the cell’s actual cytoskeleton and builds up new ones again, decomposed by the subsequent fission and
repolymerization. This changes the cellular form individually, and the active dynamism, which differs from the healthy regulated one, is expressed in the noise of the tumor processes. The noisy difference is measurable by electric impedance [169], detecting the “color” of the noise. As shown above, the “color” is determined by the self-similarity scaling exponent of power density by frequency. The measured exponent decreases from the near-1 (pink noise), depending on the weakened cellular interactions, and so the autonomy is field. The same change was observed in the autocorrelation [170], lowering the function value at 1/e, so the correlation was decreasing, which characterizes the trend towards autonomy. The weakness of the malignant process is this autonomy when the cells are alone to fight against any external stresses. So the weakness, which we have to use [171], is the missing network, the missing collective harmony. The individualism of the cells is similar to atavistic reverse development in the evolutionary meaning. The atavistic model could be used as a starting point, but this model does not consider all the crucial details (hallmarks) that keep the single-celled units of cancer development alive [172]. The wound healing support of the host tissue is missing from the atavistic model, and so it is not applicable. However, the proper noise application places the theranostic key in the hand of the therapist. The noise spectrum is measurable for diagnosis [171], and the pink noise as a homeostatic harmony character could make a forceful selected harmonization factor, which harmonizes the precancerous cells, rebinding their broken intercellular connections [107] [173], forcing back the healthy order. When the rebinding is impossible, the harmonizing force “softly” destroys the cells, exposing their genetic content for antigen presentation, and igniting the normal immune reaction against the malignancy. In other words, the mask falls off.

The practice of modulated electromagnetic applications connects the knowledge of hyperthermia with electromagnetic effects. The modulated electro-hyperthermia (mEHT, trade name oncothermia®) has used this method for years in a broad application for various tumors. Below we summarize the most crucial information-set about its processes and results, showing publications.

Numerous practical applications proved the feasibility of the mEHT treatments. Based on some principal considerations, the method uses the thermal and non-thermal effects [116] in a synergy. The non-thermal effects are connected to the cellular membranes [114] [115], emphasizing the physical potential of the amplitude modulation [174]. The strong connection between the thermal-factors and electric effects involves the similarities between their actions [175], and the strong synergy was indeed observed in vivo [176]. The absorbed power develops the temperature [177] which has a selective heating possibility [178] by electric inhomogeneities and depends on the technical solution of the coupling [179].

The temperature development is a complex process [180], and connected to the dosing in treatment practice [181] [182].
The extracellular action complexly interconnects the electric field with the thermal effects [183]. The thermal noise influences the electric field effect [22][184], but it does not block the intracellular processes [185]. The modulation technology has been described in oncology [186].

The RF energy is predominantly absorbed on the clustered transmembrane proteins (rafts) [94], exciting their receptors [94] [187] [188]. The excitation damages the malignant cells dominantly in an apoptotic way [105] [149]. The mEHT promoted extracellular HSP70 supports the apoptotic processes [108]. The membrane rafts are the key elements of energy absorption [189], which is an energy intake in the nano-range [148]. The whole mEHT process is immunogenic [190]. The method may be applied complementary to radiotherapy [191] [192]; an example is the resensitization of the radioresistant pancreatic cells for further treatment [193]. The complementary application of mEHT with chemotherapy has also been successful [194] [195].

A recent review summarizes the molecular biology background of the tumor damage mechanisms revealed in cancer models [196].

In the clinical application, the safety of the method was investigated in one of the most sensitive organs, the brain. The prospective safety study involved the dose-expansion for non-invasive transcranial therapy in 3rd and 4th line, advanced; relapsed high-grade gliomas and proved the safety even in the daily applications [197]. Efficacy studies validated the feasibility of this method for such a short survival, high-mortality stage of the disease, [198] [199] [200]. The special safety [201] and efficacy [202] studies in advanced non-small cell lung cancer patients justified the use of mEHT in this type of malignancy as well. The analyses of brain [203] and lung [204] studies support the economic advantage of mEHT. The applications in gastrointestinal tracts like colon, [205], rectum [206], liver [207], and pancreas [208] [209] demonstrated the versatile functions of the mEHT method. The relapsed, refractory, or progressive heavily treated ovarian cancer [210], and the advanced, recurrent cervix uteri [211] [212] were also successfully treated with complementary combinations of conventional chemoradiotherapy and mEHT, obtaining low toxicity and longer survival, with improved quality of life [213]. The applicability for sarcomas in complementary therapy protocols was also proven [214] [215]. The immunogenic influence in combination with other immune supports is also remarkable [216] and realizes tumor-specific immune reactions for distant metastases (abscopal effect) [217] [218]. These observations orient the treatment to tumor-directed immunotherapy [219] and could be a basis of the personalized medicine [220]. Remarkable application of mEHT for the peritoneal carcinomatosis with malignant ascites, [221], which extends the study focus to ascites cases.

Recently a review had summarized the clinical evidence of mEHT for the practicing oncologists [222], and an application guideline was published by an international collective [223].

Notably, most studies have single-arm protocols, which have inherent com-
lications in evaluating the mEHT method. The patients involved in these studies are in relapsed advanced stages when the conventional therapies are limited or have failed. Collecting a cohort for reference at this stage is exceptionally challenging and, in most cases, impossible [224]. However, the evaluation of single-arm trials has specialties, which could improve the level of evidence of the results. The simplest one is the comparison of the single-arm results from different studies with the same protocol. For example, the comparison of advanced glioblastoma (Figure 15) [225] [226] [227] [200] [203], and pancreas (Figure 16) [228] [229] [225] [230] [231], studies shows the significant difference of the common median from the historical or database control.

Other analysis of single-arm results uses the self-similarity of living processes [232] [233], building up the reference arm from the parameters of the active single-arm [234] [235].

5. Conclusion

The modulation of radiofrequency carriers causes stochastic resonances, which act mainly on enzymatic processes. The demodulation is also effectively performed by stochastic resonance, which allows the low-frequency modulation information to be delivered selectively to the cancerous cells. There are as many resonant frequencies as there are enzymatic reactions that exist.

The modulation intensively changes the paradigm of cancer treatment. It boosts the healthy network and selectively destroys the malignant cells. The cell-killing process is immunogenic, allowing for the release of genetic information

![Overall survivals](image-url)

Figure 15. The comparison of the single-arm mEHT studies with the database results for advanced, relapsed glioblastoma multiform. The different groups with different studies based on the same mEHT protocol well correspond with each other and significantly differ from the database.
in a molecular pattern (DAMP), which could form antigen presenting cells and develop tumor-specific immune reactions. The modulated signal is a complex mixture of non-thermal electron excitation for chemical reactions, and thermal boosting of the enzymatic reactions.

The complex homeostatic network has a cyclic symmetry which controls the balance and the apoptotic processes. The living cellular structures are open to the effects of energy, requiring energy input, followed by the output of waste. This organized transport system is not possible without direct cellular communication (no “social signal”). The cyclic topological order makes the noiseless inter- and intra-cellular communications possible despite the thermal noise background.

The malignant transformation breaks the organized transport system and seeks to build up new structures for the new demands. However, there are fundamental differences: the healthy construction is driven by the collective signal and seeks to optimize the energy-use for highest efficacy, while the malignant construction is only driven by the topology and physics of the mass number of individual, competing cells, irrespective of the efficacy of the energy conversion. The new structure modifies the cell-cell communications and arranges the malignant cellular microenvironment suitable for the autonomic actions. The topology in this meaning has significant diagnostic (pathologic) meaning: the more the pattern of tumor (networking “mesh” and form of the cells) resembles
the healthy cells, the less malignant the tumor is.

The modulation affects the malignant construction in various ways:
1) boosts the healthy network;
2) limits the autonomy of the cells;
3) selects the malignant cells and targets them with resonance energy-absorption;
4) excites selected molecules, influencing and promoting the enzymatic processes;
5) supports the immune system and triggers selective antitumor effects;
6) regenerates the lost intercellular adherent bonds, and with this limits the invasion of the malignant cells;
7) frees the HSP70 extracellularly to carry genetic information about the malignant cell to antigen-presenting cells;
8) supports the cellular communications and the synchrony of the cells.

The modulation signal is a time-fractal representing the systemic homeostatic fluctuations. It is self-similar, having $1/f$ noise of the power density. It might promote the establishing the lost cellular communications or stimulate apoptosis of the cells which remain individual.

The modulation principles to support the healthy processes have broad feasibility to treat the patients. The therapy is a complex, multi-component clinical process, where one of the factors is the modulated electro-hyperthermia (mEHT). The mEHT has proved its exceptional capability to treat late-stage, relapsed cancerous tumors, complementary to any other conventional therapies. In instances where there are no further treatment options available, mEHT could be applied as monotherapy, or in some cases, could desensitize tumors to the previously ineffective conventional treatment.

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**Conflicts of Interest**

The author declares no conflicts of interest regarding the publication of this paper.

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