Probing Biomechanical Noise in Neuroblastoma Cells using the Quartz Crystal Microbalance

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SUMMARY:

Quantifying cellular behavior by motility and morphology changes are increasingly important in formulating an understanding of fundamental physiological phenomena and disease mechanisms such as cancer. However, cells are complex biological units which often respond to external environmental factors by manifesting subtle responses that may be difficult to quantify using conventional biophysical measurements. This paper describes the adaptation of the Quartz crystal microbalance (QCM) to the quantitative study of neuroblastoma cells under environmental stress wherein the frequency stability of the device can be co-related to changes in cellular state. It is seen that when neuroblastoma cells are seeded on to the sensor surface, a distinct variation in the pattern of frequency fluctuations is recorded, which can be attributed to the biomechanical noise originating from the cells. By employing time domain analysis of the resulting frequency fluctuations we study cellular motility and differentiate between different cell states induced by applied external heat stress for the SH-SY5Y neuroblastoma cell line. The changes in the frequency fluctuation data are correlated to phenotypical physical response recorded using optical microscopy under identical conditions of environmental stress. Our study shows that this technique can provide new insight into the physical behavior of cells by providing a basis to quantify the associated biomechanical noise paving the way for its use in monitoring cell activity, and intrinsic motility and morphology changes, as well as those resulting from the action of drugs, toxins and environmental stress.

INTRODUCTION

Depending on the cell type and environment, cells often undergo continuous morphological and motility changes. Cytoplasm and sub-cellular components self-organize to change shape and enable cells to move around. These processes, though superficially simple, are highly complex and represent nature’s unique solutions to challenges posed by mechanics, thermodynamics and biochemistry. Diverse biological processes ranging from embryogenesis, fetal and organ development to immune response against inflammation and wound healing involve movement or migration of cells[1]. Cellular motility also plays a crucial role in many diseases including in tumor invasion and spread (metastasis) of cancer. Therefore understanding and targeting cell motility is a topical strategy in reducing morbidity and fatality in cancer treatment[2]. Cellular morphology is studied intensively because it gives a fair idea of the conditions being experienced by the cells as well as of the various biological processes being executed by the cells[3].

Attachment-detachment to the underlying substrate and extra-cellular matrix (ECM) takes place through focal adhesion points which involve binding of integrin mediators to ECM molecules leading to the formation of focal complexes[4]. Integrin receptors are connected to cytoskeleton through actin networks which influence reorganization of cytoskeleton, hence, the cell structure itself[5]. These processes have biomechanical consequences and can be used to model the behavior of cells. Assuming environmental factors as input, cells (both individually and collectively) represent physical dynamic systems which respond to the input (stimuli) by manifesting their motility and morphological characteristics. These biomechanical manifestations are often stochastic and, thus, require a tool for sensing and interpreting these variations. Techniques based on optical microscopy[6–8], micromanipulation[9], modified AFM[10] and optical traps[11] have been used to understand the mechanics of focal adhesion[12], however, analysis of stochastic aspects of cellular motility and morphology is missing. Thus, there is a need of de-
development of a technique that could be simple, non-invasive and real-time. In this paper, we report one such technique which is based on performing frequency fluctuations analysis on Quartz Crystal Microbalance (QCM) data.

In the past few decades, the Quartz Crystal Microbalance (QCM)[33] has emerged as an important tool for highly sensitive, label-free, gravimetric measurements of physical processes involving physical adsorption and desorption of a variety of analytes. The QCM can be mechanically abstracted as a single degree-of-freedom spring-mass-damper system such that any attachment/detachment of analyte on the crystal manifests in the form of a change in resonant response that is read out using external electronics. This property of QCM has been extensively used to perform measurements on biological systems in liquid environments. In the presence of liquid, acoustic oscillations of QCM couple to the surrounding liquid media and the viscoelastic properties of the media at the interface[14] have predictable effects on the frequency \( f \) and dissipation \( D \) characteristics of QCM. A range of biological systems have been probed using the QCM including the physi/chemisorption of proteins and lipids[15], binding and reaction kinetics of biomolecules[16], cell attachment-detachment and spreading[17–20] etc. Descriptive models [21,22] have been proposed to correlate adsorption-desorption and binding studies using QCM but for complex systems such as cells, a quantitative description using conventional analysis \( (f-D) \) techniques is often inadequate. When such systems are influenced by actively driven dynamic and non-equilibrium processes like motility and morphological perturbations, fluctuations in the QCM response as a function of time are seen. There is, therefore, the possibility of utilizing the resulting recorded fluctuations in response as a means of characterizing such dynamic and complex biological systems.

It can be hypothesized that movement of cells on the surface can result in biomechanical noise that can couple to response of the QCM and be conspicuously evidenced in terms of enhanced levels of fluctuations in the recorded QCM response. Sapper et al[23] previously reported that the effect of biomechanical noise on frequency fluctuations in a QCM is several orders of magnitude higher than that contributed by other noise sources. By calculating the power spectral density (PSD) of the QCM frequency fluctuations in the case of attached mammalian cells, the authors were able to qualitatively associate frequency fluctuation data with cell motility. In a separate study, Tarantola et al[24] studied motility of different cancer cell lines using the same technique. In both the studies, outcome of the fluctuation analysis was compared with a similar analysis on the Electric Cell-Substrate Impedance Sensing (ECIS) data. In another study, periodic contractions of rat cardiac myocytes was seen coupled to the frequency data[25].

In the present paper, mammalian cancer cells are studied using a time-domain fluctuation analysis technique (Allan Deviation measurements) applied to the measured QCM response. The case study of neuronal cells (differentiated human neuroblastoma cell line) is considered where their interactions with the sensor surface are monitored, while subjecting the cells to external heat stress which eventually leads to apoptosis. This study provides insight into the mechanical response of the particular type of cancer cells due to the change in local temperature. By combining separate optical studies with QCM data, it is possible to correlate the observed changes in single cell state and morphology as well as total surface coverage of the cells to the QCM frequency fluctuation data. Thus, the paper establishes the basis for the use of time-domain QCM frequency fluctuation analysis for deciphering and monitoring physical behavior of cells and cell-substrate interactions and the changes induced due to environmental stress. The results indicate that this technique can be expanded to other case studies on cellular systems and the effect of varying environmental conditions on their mechanical behavior.

MATERIALS AND METHODS

A. QCM Set-up

AT-cut plano-plano thickness shear mode Cr/Au coated 5 MHz quartz crystals, 1 inch in diameter (see schematic in Fig. 1A) are used. Crystals were fitted into a dedicated Kynar® flow-cell, part of the standard QCM200® instrument from Stanford Research Systems (SRS). The flow-cell volume is approximately 150 μl. Crystal electrodes are isolated from the flow media and electrical connections to the interface electronics are achieved through spring-loaded contact pins. During experiments, the series resonance frequency and crystal resistance is logged and then later processed off-line. The gate-time (sampling time) of the instrument was set to 1 s. The temperature of the flow-cell was controlled by a heater with the whole set-up being placed inside an environmental chamber with fully humidified atmosphere at 37°C and 5% CO₂ during the length of the experiments on the live cells. The environmental chamber was placed on a vibration isolation table and the attached electrical cables, connecting the crystal to the measurement set-up (outside the chamber), were fitted with foam dampers at points of contact with the chamber to prevent external vibrations arising from the humidifier or other sources from impacting measurement.

B. Cell-culture and crystal chip preparation

Experiments were carried out using SH-SY5Y cells, a human neuroblastoma cell line, commonly used to study neuronal function and behaviour[26]. Cells were adherent with neurite outgrowth. Moreover, the cells tended to grow in clusters resulting in patchy surface coverage of the culture dish. Cells were cultured at 37°C in humidified atmosphere in a medium containing 87% Dulbecco’s Modified Eagle’s Medium (GIBCO), 10% heat inactivated fetal bovine serum (PAA), 2% B-27® (GIBCO) and 1% antibiotic-antimycotic (GIBCO). The addition of 10 nM staurosporine (Sigma) to the culture medium triggered cell differentiation into a neuroblast-like cell type, which
arrests proliferation and results in a stable population[27]. Thereafter, a required number of cells were reseeded on autoclaved, poly-L-lysine (PLL, Sigma) coated quartz crystals placed inside a 35 mm culture dish (Greiner). Prior to the seeding of the cells the quartz crystals were incubated with aqueous PLL (0.02 m/v %) solution for two hours at 37°C to facilitate adherence of cells to the surface. A schematic of a cell attached to a PLL-coated quartz crystal is shown in Fig 1B.

In order to study cell motility, the number of cells seeded on the quartz crystal should result in a relatively sparse coverage to minimize the influence of cell compression and contact inhibition[28]. Therefore, two extreme cell concentrations were studied to determine cell behavior and motility. Approximately 5,700 cells per cm² and 51,700 cells per cm² were seeded onto the quartz crystal containing cell culture dishes. These seeding densities resulted in a surface coverage of ~3.6% (surface coverage i) and ~30.7% (surface coverage ii), respectively, which was measured by quantifying the cell population over a relatively large area (~0.5 cm²) of the quartz crystal using optical images to remove the influence of local heterogeneity due to cell colonies/clusters. Experiments for intermediate surface coverages, 13.0% (surface coverage iii) and 22.5% (surface coverage iv) along with control experiments were also performed. Cells were allowed to attach to the substrate over 24hrs. Thereafter, cell attachment and viability was checked using a light microscope.

Crystals demonstrating good attachment of cells on the surface were then chosen for individual experiments. These crystals were placed inside the pre-warmed flow-cell which was filled with cell culture medium carefully avoiding introduction of air bubbles, which can interfere with the QCM readings. The flow-cell was sterilized with 70% ethanol prior to cell experiments and thereafter kept under sterile cell culture conditions to exclude contamination of the cells throughout the experiments. The flow-cell was then placed inside the environmental chamber, connections to electronics were made, and data recording with the QCM was initiated after an initial stabilization period of ~10 min. During QCM measurements there was no flow of the medium to preclude flow-related perturbations.

![Figure 1](image_url)

**Figure 1.** (A) Schematic of a quartz crystal Cr/Au top-surface electrode (yellow) and (B) Cross-sectional schematic view of a SH-SY5Y human neuroblastoma cell adhering to the poly-L-lysine (PLL) coated crystal. (Relative thicknesses of different components are not to scale.)

C. **Hyperthermia induced, time-lapse live cell experiments**

Changes in cellular morphology and motility were studied in SH-SY5Y cells seeded onto a 35 mm glass-bottom dish (Mattek, USA) with a surface coverage of ~30.7%. Hyperthermia was induced by gradual temperature increase from 37 °C to 45 °C in an environmental chamber (humidified atmosphere with 5% CO2) of a Leica microscope allowing for live cell, time-resolved phase-contrast imaging using a 40x objective. Time-lapse recording was started simultaneously with the initiation of temperature increase. The temperature was measured in the air chamber and a final temperature of 45 °C was recorded after 50 minutes. Time-lapse video recording was carried out capturing one frame every 20 seconds. For cell tracking, image sequences were pre-processed using threshold adaptation, background subtraction, followed by using an un-sharpening mask and binarization in ImageJ (NHI, USA)[29]. For every frame, boundaries were drawn using the marking tool in ImageJ and shape parameters like centroid, area and circularity were obtained. Centroid positions were superimposed on the optical frames to visualize the track of selected cells and further processing on the data was performed to calculate distance travelled, mean squared displacement and speed.

**RESULTS AND DISCUSSION**

Since QCM is a surface sensitive technique, it can be applied to study various phenomena associated with cell-surface interactions including factors governing motility and changes in their morphology. The cells studied in this work do not undergo cell division but nevertheless change their shape both as a result of movement and environmental stress[28,30,31]. Cellular motility is known to depend on many environmental factors including cell culture medium, surface properties[32,33], the cell line itself, the surface density of cells as well as temperature [34,35]. In order to determine the ability of QCM to measure the effect of temperature increase on live cells, heat stress experiments were carried out.

A. **Heat stress experiments**

Cell covered QCM crystals were prepared as detailed earlier (Materials and Methods) and loaded into the flow chamber. After the recordings on the QCM crystal stabilized, the temperature of the heater was ramped up from 37 °C to 45 °C. This change in temperature induces stress resulting in changes in the cell shape and texture indicating apoptotic processes and ultimately resulting in apoptosis (as shown in the later sections). Data was recorded beyond the time after the temperature monitor (attached to the flow cell) reached the final set temperature to allow for equilibration and compensate for any thermal lag. After every experiment, the crystals were viewed under the microscope to confirm that the cells had undergone apoptosis triggered by the heating process and formed typical spherical shapes alongside with detachment from the surface. Frequency data for two such experiments are shown in Fig. 2(A). As the temperature was increased, both crystals showed a gradual positive frequency shift (ΔF) indicating a change in mass loading (Δm) on the surface. Frequency shift for the larger cell population (~250 Hz) is greater than the one observed in the case of the smaller cell population (~50 Hz) as expected roughly from
the Sauerbrey equation \( \Delta F = -2F_0 \frac{\Delta m}{A\sqrt{\rho_0 \mu_q}} \), where \( F_0, A, \rho_0, \mu_q \) are the unloaded crystal resonance frequency, active crystal area i.e. area constrained by the electrodes, density of quartz and shear modulus of quartz respectively. It should be noted that the Sauerbrey equation has limited validity in this case and a more elaborate model is necessary for quantitative correlation to the experiment due to the complex nature of the physical system under investigation. For a complex surface involving multiple layers (PLL, cell medium) in a viscous medium, the penetration depth \( \delta = \sqrt{\frac{\eta_L}{\pi F_0 \rho_L}} \) where \( F_0, \eta, \rho_L \) are resonance frequency of the crystal, dynamic viscosity and density of the liquid layer at the interface respectively provides information on the dimensional extent of the physical region probed. For a crystal of 5 MHz frequency operating in shear contact with a liquid at room temperature, the penetration depth is about 250 nm. While accurate prediction of changes based on frequency data alone is challenging due to the complex nature of the interfacial surface, the changes could be qualitatively interpreted due to rupture of strong adhesion bonds of cells, as they progress towards apoptosis, which are slowly replaced by loose physisorption, both for the cells which remain on the surface and those that sink after complete detachment. This observation is consistent with previous experiments on studying cell apoptosis using the QCM[36].

B. Frequency fluctuation analysis

Frequency and dissipation readings provide valuable information on changes occurring at the crystal-liquid interface and have proven to be especially useful in ligand binding experiments. While cell adhesion, spreading, detachment and morphological changes in presence of an analyte have been previously studied by QCM using the frequency-dissipation approach[18-20,37] dynamic changes or fluctuations in these parameters have been given little attention. Fluctuations in QCM parameters have been shown to correlate with the subtle changes in mass and viscous loading at the interface[23,24]. Cell-substrate interactions probed using the QCM involve constant attachment and detachment at focal adhesion points, morphological changes and cell-cell interactions which can result in perturbations in viscoelastic loading on the QCM.

Based on QCM frequency measurements, three zones are defined on the basis of variability in environmental and interfacial conditions. Zone 1 is defined as the time period starting from the point where data recording is commenced and the system is held at 37°C. This zone corresponds to the case where cells are viable and kept in ideal culture conditions. Zone 2 is defined as the zone starting immediately after the temperature is increased and up to the point where all cells are in the late stages of apoptosis due to hyperthermia. Zone 3 is defined as the region where the environment has reached a plateau phase following cell death due to the heat shock treatment in Zone 2. The frequency fluctuation data for both experiments is plotted in Fig. 2(B)-(C). It is clear, even by visual examination, that the amplitude of frequency fluctuations is reduced significantly for Zone 3 compared to Zone 1 for higher cell coverage. An increased amplitude of frequency fluctuations indicates large scale activity on the surface, which is to be expected when the cells are healthy and mostly stable in clusters (colonies) compared to when they are largely static (apoptotic). However, to establish

![Figure 2](image_url)
more quantitative insight into these observations and
cases for low cell coverage, time-domain frequency stabili-
ty analysis is employed.

C. Time-domain fluctuation analysis

Two sample Allan Deviation

Frequency fluctuations can be characterized either by
performing frequency domain analysis (to calculate power
spectral density (PSD)) of the data or by analyzing the
Allan variance in time domain. Though inter-convertible,
Allan variance is chosen over PSD as it is more convenient
to interpret in the terms of our experimental set-up
which, due to use of a frequency counter, outputs ob-
erved frequency as a time series. A quantitative measure
of frequency stability is established by analyzing the Allan
variance (see Supplementary Information) for the data
sets corresponding to the various cases of cell coverage in
Zones 1-3. The Allan variance is chosen over standard M-
Sample variance as it is convergent for most types of noise
processes and can help distinguish between different
noise mechanisms as well. An overlapped Allan
ance, \( \sigma^2_{\tau}(\tau) \), for \( M \) fractional frequency values, is
employed to provide higher statistical confidence (see Sup-
plementary Information).

Plots showing overlapped Allan deviation as a function
of averaging time show a power-law dependence as a
function of averaging time (\( \tau \)) with characteristic slopes
\( \mu(\sigma_{\tau}(\tau) \approx \tau^{\mu/2}) \) in such plots and thus can be readily iden-
tified. White noise (\( f^0 \)) describes \( \mu = -1 \), Flicker noise
(\( 1/f \)) describes \( \mu = 0 \), Random walk frequency modulated
noise (\( 1/f^2 \)) describes \( \mu = 1 \), while frequency drift
describes \( \mu = 2 \) [38,39]. While the frequency fluctuation
data shown in Fig. 2 is linear drift compensated, there is
no pre-processing of the frequency fluctuation data used
to calculate overlapped Allan deviation. Further, error
estimates for 1-sigma confidence intervals are calculated
for all the three zones. Fig. 3(a) shows the sigma-tau plot
for the high surface coverage cell experiment where the
difference between the Allan deviation values (fluctua-
tion levels) for three different zones is very clear. This agrees
well with the frequency fluctuations observed in the three
zones. Further, the deviation plot for each zone is consid-
ered separately and slopes between every pair of adjacent
points are obtained using linear fitting. Relevant plots are
shown in Fig. 3 (A)-b,c,d. Similarly, Fig. 3(B) represents
sigma-tau plots for the low cell surface coverage. In this case,
for short averaging times, the Allan deviation values
are very close to each other while they diverge a bit for
longer averaging times.

![Figure 3.](image)

(A) (B)

Figure 3. (A)-(B) Log-log overlapped Allan-deviation plots
for the larger and smaller surface coverage cases respectively.
From (A)-(a) it is quite evident that the fluctuation levels for
all averaging times (\( \tau \)) are highest for Zone 2, intermediate
for Zone 1 while levels are lowest for Zone 3. This indicates
when cells are alive, due to their activity; fluctuation levels
are higher as compared to when cells are dead. Also, when
cells are in a stressed state higher fluctuation levels are ob-
served. (A)-(b-d)- Individual linear fitting curves (black, con-
tinuous and slightly offset for clarity) for three zones. Simi-
larly, from (B)-(a) it is evident that while fluctuation levels
for shorter averaging times are higher for both Zone 1 and
Zone 2 relative to Zone 3 but between Zone 1 and Zone 2,
fluctuation levels are only slightly higher for Zone 2. Linear
fitting (b-d) indicate contribution from different power law
noises as in the former case.

Interpretation of log-log overlapped Allan deviation plots

Individual deviation values for each time factor is calcu-
lated and can be used to compare the fluctuation levels
between different zones. In Table 1, deviation values for
shortest averaging time (is) are used for representation.
In the case where cells are cultured on the QCM, it is
quite evident that deviation values are the maximum for
Zone 2, are the least for Zone 3 and lie in-between these
two values for Zone 1. The relative difference between
deviation values is clear between high and low surface
coverage of cells. In all the cases though, for Zone 3, fluc-
tuation values are \( \approx 10^{-8} \) which approximately corre-
sponds to the baseline fluctuation level. To determine the
effect of heating on the fluctuation levels, two control experiments, without any cells, were performed with, first, just water and, secondly, with the cell culture medium. All other conditions were kept the same. Log-log plots of deviation values for the control experiments are shown in Fig 4 (C)-(D). Clearly there is very little change in the fluctuation levels when the temperature is increased implying that the observed noise levels (Fig 4A-B) are generated due to the presence of the cells on the surface. Experiments with intermediate cell coverage of 13.0% (surface coverage iii) and 22.5% (surface coverage iv) were also performed. Both the experiments agree with the trends in the frequency fluctuation data observed with the high and low surface coverage experiments. For averaging factors $\tau = 2 \, s$, $4 \, s$, a similar table can be formulated to provide similar trends (see Supplementary Information). For larger averaging times ($\geq 8 \, s$), the plots start to intersect pointing towards the fact that values must be interpreted with care. To understand the contribution of different noise sources at various time scales, we can separate the contribution of device noise from the biomechanical noise emerging from the activity of the cells themselves. Some of the principal device noise could be evidenced as thermal noise or flicker noise with noise contributions from the crystal and the amplifier, along with sensor drift due to changes in environmental conditions. Certain types of physical processes associated with cellular activity such as attachment-detachment of individual focal contacts, morphological changes in response to external environmental conditions and internal cellular processes, migration of cells on the surface, cluster dynamics of cells, fluctuations in the viscoelasticity of the fluid due to cell activity and external environmental perturbations are expected to contribute to the measured frequency fluctuations. Each of the above processes are expected to operate on different time scales, for example, attachment-detachment between extracellular matrix and cells are much faster ($t_{off}$ for spontaneous dissociation of a closed bond at a focal adhesion site can range from fraction of seconds to tens of seconds[40]) while motility and cluster dynamics can be relatively slower. As shape changes are sensitive to external conditions, any fast external processes are going to influence the cell morphology at a similar rate. A plot of Allan deviation slopes for different averaging times can be found in the Supplementary Information. Nevertheless, in order to correlate and verify predictions from the time-domain frequency analysis with cell behavior optical imaging experiments were carried out.

Table 1. Two sample overlapped Allan deviation values at $\tau = 1 \, s$ for different zones and varying surface coverage

<table>
<thead>
<tr>
<th>Experiment</th>
<th>Zone 1 ($10^{-9}$)</th>
<th>Zone 2 ($10^{-9}$)</th>
<th>Zone 3 ($10^{-9}$)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Surface coverage i (30.7%)</td>
<td>2.248</td>
<td>116.2</td>
<td>1.122</td>
</tr>
<tr>
<td>Surface coverage ii (3.6%)</td>
<td>1.172</td>
<td>1.310</td>
<td>0.899</td>
</tr>
<tr>
<td>Surface coverage iii (13.05%)</td>
<td>2.31</td>
<td>3.2</td>
<td>1.024</td>
</tr>
<tr>
<td>Surface coverage iv (22.5%)</td>
<td>1.67</td>
<td>99.186</td>
<td>1.612</td>
</tr>
<tr>
<td>Control-DI Water</td>
<td>0.861</td>
<td>0.865</td>
<td>0.742</td>
</tr>
<tr>
<td>Control-Cell Culture</td>
<td>0.975</td>
<td>0.888</td>
<td>0.742</td>
</tr>
</tbody>
</table>
D. Correlating frequency fluctuation data with optically recorded cellular mechanical behavior

Cellular changes for increased temperatures

As discussed before, cellular motility depends on many environmental factors such as temperature, cell culture medium[34] and substrate properties such as rigidity, microstructure[41] and coating[33], as well as being dependent on the cell type or cell line and the seeding density. In particular, it has been shown that cellular confluency influences cellular behavior and cell morphology as higher cell densities compromise the individual cell area impacting on cell morphology and motility[42]. Many intracellular processes are involved in cell migration. In migration direction, a leading edge is formed by extending flat membrane processes composed of an inner actin crisscross network called lamellar ruffling. This is accompanied by the cell-substrate interactions in order to form adhesion points at the leading processes. Then, the cell body, including cytoplasm and organelles, is shifted while the trailing cell rear is retracted. Thereby, temperature affects cell motility and related characteristics such as lamellar ruffling protrusion as shown by Shelden et al [31] for human neuroblastoma cells. Lamellar ruffling has been observed to increase for temperatures up to 40°C (mild hyperthermia) but declines for further temperature increase. Furthermore, Hartmann-Petersen et al[34] have studied cellular process domains as well as mean cell speed and area in murine fibroblasts under temperature conditions of 37 to 40°C and demonstrated an increase of characteristic values proposing the assumption that cells at slightly higher temperatures are more spread out and motile.

Similar observations have been made for the cell motility of 3T3 cells showing a peak between 36.5 to 38.5°C, with strongly declining motility towards lower or higher temperatures[43]. It is commonly known that prolonged treatment of cells at hyperthermic temperatures compromises cell viability and effectively triggers apoptosis. Treatment of cells with temperatures as high as 43°C for at least one hour has been shown by Luchetti et al[44] to already trigger apoptosis in human neuroblastoma cells. This is accompanied by changes in cell morphology such as blebbing, nuclear swelling, chromatin margination and changes in the cell shape eventually followed by cell rounding. Cytoskeletal proteins undergo aggregation and
rearrangements leading to the final breakdown of microtubules. Additionally, cellular membrane characteristics are altered leading to cellular adhesiveness impairment [45, 46].

**Observing a sample cell**

The aim of this section is to correlate the frequency fluctuation analysis with cellular changes and behavior in terms of motility, substrate adhesiveness and morphology resulting from hyperthermia. Therefore, cells were exposed to increasing temperature under conditions comparable to those used during the QCM experiment and optical images were obtained using a phase-contrast microscope to extract information on morphology and motility.

Fig 5 (A)-(H) shows eight selected time points for the same cell sample exemplifying the drastic cellular changes due to hyperthermia. The position of the same cell which moves on the surface and changes in morphology is marked with a star/asterisk (*). Significantly, the size of the nucleus increases on increasing the temperature and thereafter, the nucleus shrinks (Fig 5 (E)-(F)) until distinction from the cytoplasm is not possible any more (Fig 5 (G)-(H)). Furthermore, pigmentation of the nucleus becomes uneven showing darker spots as visible in Fig 5 (C)-(H). Nuclear swelling and chromatin aggregation have been observed in the context of apoptosis induced hyperthermia [43, 44]. In the same images, similar abnormal coloration of the cytoplasmic region is observed that is likely triggered by the aggregation and rearrangement of cytoskeletal proteins caused by the heat treatment. Several other changes in the morphology can be seen including reduced branching of neurites and blebbing (bubble-shaped protrusions) as highlighted by the white arrows in Fig 5 (F)-(G). The cell size sharply decreases after prolonged heat treatment as seen in Fig 5 (E)-(H) with gradual rounding the cell shape indicating apoptosis and ultimate detachment from the surface indicated by halo around each cell.

**Figure 5.** (A)-(H) Phase-contrast optical images of cells seeded on a glass surface. Temperature is gradually increased from 37°C to 45°C over a period of 150 min during which cells change morphology, ultimately shrink in size and become round which is characteristic of apoptosis.

**Cell tracking analysis**

For further quantification of cellular movement during heat stress treatment, we applied cell tracking analysis. Fig 6 (A), (B) and (C) represent the frames recorded at the beginning, during and end of the optical imaging experiment, respectively. Individual cell movement tracks are labeled and colored in the frames. One clear observation from cell tracking is that the motion of cells changes from being directed (elongating and reaching out in a particular direction) with nearly linear displacement track towards localized, confined movement. This change occurs simultaneously with the change in cell shapes outlined in the previous section.

To identify the movement of cells during the heat stress stage of the experiment, mean-square displacement (MSD) for each cell is calculated using the following expression for two dimensional motion:

\[ \text{MSD}(\Delta t) = \langle (\Delta r)^2 \rangle = \langle (x(t) - x(t + \Delta t))^2 + (y(t) - y(t + \Delta t))^2 \rangle \]

where, \(x(t), y(t)\) are spatial position loci of a cell. By plotting the MSD vs \(\Delta t\), the motion can be characterized as either confined random walk (Fig 6 (D)) using MSD(\(\Delta t\)) = 4D\(\Delta t^a\) or as a superposition of directed motion and diffusion using MSD(\(t\Delta t\)) = 4D\(\Delta t + (v\Delta t)^2\) where, D is the diffusion constant and \(\alpha\)-\(\varsigma\). During the first 16 mins of the time-lapse experiment, the temperature was not observed to increase more than 0.2°C and allows us to assume that cells are exposed to normal culture conditions. Cell motility was characterized during this phase to make following heat-induced changes apparent and comparable. Roughly, two groups of cells were identified, one with faster motility and directed motion (\(v\sim0.38 \mu m/min\)) with a diffusion constant of D=2 \(\times\) \(10^{-2}\) \(\mu m^2/s\) for both. These observed motilities take cell-cell interactions into account which slows down the overall speed and causes observed differences between the groups. The observed motility of cells along with the constant morphological readjustments within and around clusters continuously varies the viscoelastic loading of the QCM and contributes to the fluctuations in resonant frequency observed. Even when cells are observed under healthy and stable conditions, there is constant motility contributing to the variations in resonance frequency. Thus, as cells are stressed, their motility increases initially and then drops. Further, during the final part of the experiment, the remaining small and round cell bodies appear to describe random motion. To quantitatively characterize this cellular motion in order to find out whether this motion could contribute to the noise levels, images from 120 frames between 150th and 190th minute were used. Fitting the tracking data to the above equations revealed that cells have a diffusion constant of \(-10^{-3} \sim -10^{-2}\) indicative of Brownian motion (Fig 6 (E)). This confirms the hypothesis of cellular detachment as the temperature increases. Thus, following the final stages of apoptosis, cells are eventually
either loosely attached or completely detached. The detached cell body is located above the penetration depth region, which accounts for the reduced fluctuations observed in Zone 3 in the QCM experiments.

Figure 6. (A)-(C) Phase-contrast optical images of SH-SY5Y cells at different time instances during the heating process. Colored lines represent the tracks of various cells undergoing heat induced stress. In (B) arrows are indicating the direction of motion of individual cells while in (C) they are indicating the final position of the cells. (D) Mean-square displacement plots of cells prior to the heat-stress phase of the experiment (first 16 minutes) indicating the presence of at least two categories of cells movements, fast and slow, while, a close fit to \( \text{MSD}(\Delta t) = 4D\Delta t + (v\Delta t)^2 \) is performed to extract diffusion coefficients and speed values (E) Similarly, MSD log-log plot for the last stage of the experiment (150th-190th minute). (F) Mean and standard deviation values for normalized change in area of the cells under consideration are plotted against time which peak almost half way through and then rapidly decrease until the cells become spherical. (G) Total surface coverage shows a similar trend.

Apart from the cell motility data, another observable factor that correlates well with the fluctuation analysis is the cellular contact area. To calculate the relative change in area, we normalize instantaneous contact area with respect to final contact area where sphericity is observed for each cell. The procedure is repeated for the whole
sample set and mean and standard deviation values were calculated for the recorded frames. From Fig 6 (F), it is quite evident that, cellular contact area reaches a maximum midway and this is also where maximum deviation values are observed. Depending on the immediate vicinity and state of the cells, some show a larger change in area while others do not. As the cells undergo apoptosis, we observe almost uniform contact area across the sample set with reduced deviation values. Cells attach to surfaces through focal adhesion points and therefore a change in contact area also is likely to result in the increase of these contact points, leading to larger perturbations per cell. Also as the area changes, viscoelastic loading within the penetration depth will also change. All this effectively changes the loading conditions of the QCM. It is clear from the calculations that the total surface coverage (Fig 6 (G)) increases by 34% first as the temperature increases and eventually decreases by 67% (relative to the initial surface coverage).

Both the motility and morphological data obtained from parallel optical experiments strongly correlate with the fluctuation analysis of QCM signals. The prediction from the fluctuation analysis that on heating an initial increase in fluctuations is a result of increase in activity as proved by the concomitant increase in motility and contact area (morphology change) observed in optical experiments. Similar observations of increased motility on slight increase of temperature before declining in another cell line[43] boost the validity of our time-domain fluctuation analysis. Overall the fluctuation analysis and its classification into the three different zones map directly to the cell state induced by thermal stress.

**CONCLUSION**

Understanding motility and cell behavior in response to stress is vital for many physiological processes and diseases. In this work we have studied the motility and cell behavior of living neuroblastoma cells and the effects of thermal stress on them by QCM. Time-domain frequency fluctuation analysis is employed to provide insight into the complex behavior of neuroblastoma cells and this data is optically correlated to changes in motility and morphology. The fluctuation analysis revealed three distinct types of behavior which included the prediction of an increased state of activity of cells following an increase of temperature before a decrease of activity on account of cell death. This study thus establishes the initial basis of the QCM and time-domain fluctuation analysis as a label-free bio-physical tool to quantify cellular activity and the effect of drugs, stress agents or environmental factors on cells. Furthermore, the experiments provide interesting insight into the nature of cell-substrate interactions under healthy conditions and in response to thermal stress. The high degree of correlation between the frequency fluctuation data and optically recorded mechanical behavior also provides the basis for future studies using the QCM as a means of inferring the nature of physical processes underlying cell-substrate interactions by quantifying the associated bio-mechanical noise.

**REFERENCES**


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**TABLE OF CONTENTS/ABSTRACT GRAPHICS**

![SH-SY5Y Cell QCM](image1)

![Zone 1 Zone 2 Zone 3](image2)

Cell death