



ROLE OF MITOCHONDRIA IN CANCER

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ABSTRACT

Mitochondria is an important organelle of the body. The role of mitochondria in the body is to produce energy in the form of ATP. Mitochondria have their DNA that participates in coding certain enzymes which involve in oxidative phosphorylation. Mutation of any other genetic variation in mitochondrial DNA leads to dysfunction of mitochondrial DNA (mt DNA) which contributes to many types of cancer. Observations made during the last few decades revealed that a variety of disorders including cancer and different neurodegenerative disorders like Parkinson's disease, Alzheimer's disease, Huntington's disease and amyotrophic lateral sclerosis, have a mitochondrial origin. Disturbance of mitochondrial critical functions, such as ATP production, calcium buffering capacity, and excessive production of reactive oxygen species (ROS), can be potentially implicated in disease pathogenesis. In this article we explained the role of mitochondria behind the adenosine triphosphate production and also simultaneous production of (ROS) which causes mtDNA damage, leads to many types of cancers. In this article, we also explained what strategy we may use to minimize the production of ROS or free radical to reduce oxidative damage and another way which may be used to destroy dysfunctional mitochondria and accomplish healthy copy of the genome.

Key words: Mitochondria, mitochondrial DNA (mt DNA), Reactive oxygen species (ROS), Cancer

INTRODUCTION

There are very essential changes in cell physiology that normally might dictate malignant transformation: self-sufficiency in growth signals, sustained angiogenesis, insensitivity to growth-inhibitory (antigrowth) signals, limitless replicative potential, evasion of apoptosis and tissue invasion and metastasis (Hanahan and Weinberg 2000). Additionally, much evidence demonstrates that one of the unique qualities of tumor cells, they dependent on glycolysis for adenosine triphosphate (ATP) generation. Cancer cells generate their ATP through glycolysis still in aerobic conditions, Otto Warburg found this phenomenon in 1926 (Gogvadze, Zhivotovsky et al. 2010). There was an association between glycolytic ATP generation and violence of the tumor cells (Simonnet, Alazard et al. 2002). The universal characteristics of malignant cells are 'aerobic glycolysis' that was presumed by Warburg and he also proposed that impaired mitochondrial metabolism leads to cancer. Warburg hypothesized that cancer cells could be removed through the inhibition of mitochondrial oxphos (e.g. by average doses of ionizing radiation), which would reduce the action of these organelles lower than threshold level critical for cell survival and mitochondria in normal cells would yet be able to generate ATP. But further research challenged this and shows that

tumor cell's mitochondria do respire and generate ATP energy (Weinhouse 1976). Currently, for the visualization of tumors by positron emission tomography, the extensive glucose utilization by malignant cells is broadly used to highlight the significance of Warburg's observation. Nowadays the finding of tumor suppressor genes, finding of oncogenes, and other recent research in tumor biology shifted the attention of cancer researchers away from studies of energy metabolism to other areas. Mitochondria activities are not restricted to only ATP generation but it also produces ROS which participates in many physiological processes. But excessive ROS production might be harmful to the cell. In the process of apoptosis, mitochondria are essential for the control of intracellular Ca²⁺ homeostasis. Obviously, all these functions of mitochondria are crucial for tumor cell physiology, development, growth and survival. This review is dedicated to the important role of mitochondrial impairment in cancer.

MITOCHONDRIA

Mitochondria is a unique organelle in many other organelles because it has a unique property to produce energy in the form of ATP no other organelle has this type of property so mitochondria are called the powerhouse of the body. Cells have a high need for energy and they depend on mitochondria for it. Mitochondria metabolize oxygen to generate high amounts of energy. In contrast, mitochondria of tumor cells are defective in their ability to metabolize (O₂) oxygen (Wallace 2012).

PROCESS OF ENERGY PRODUCTION

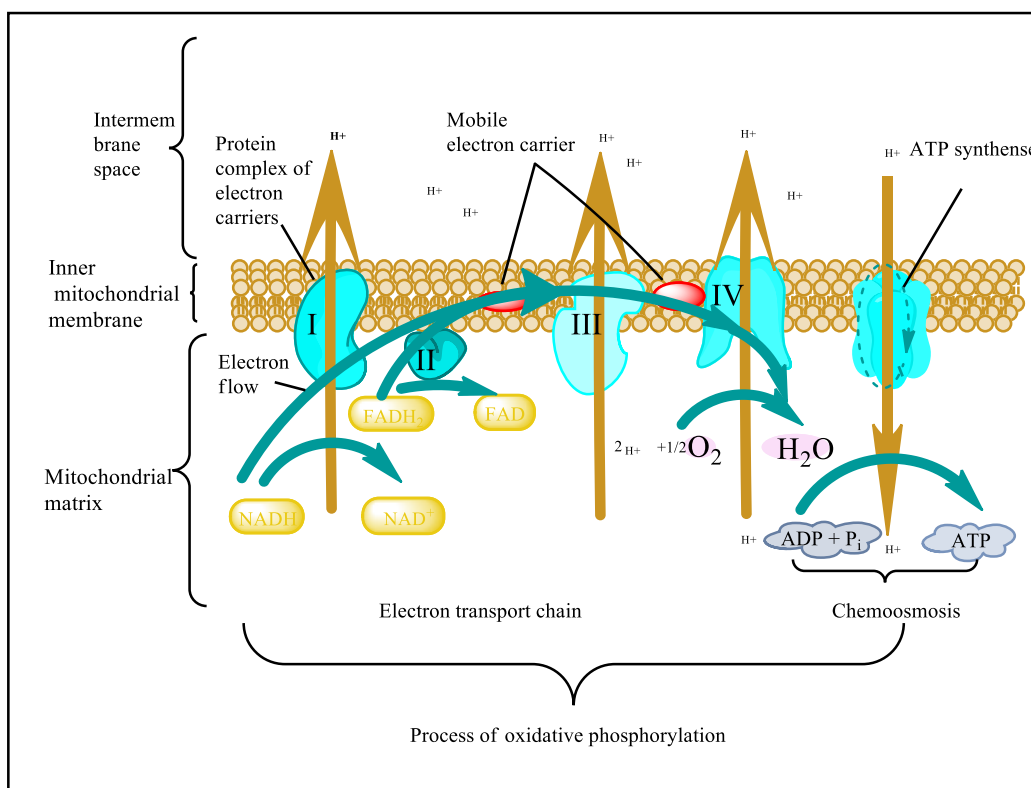
Normally, mitochondria complete their task of generating energy, by using three important foodstuffs which are Fatty acids, Carbohydrates, and Protein. Mitochondria uses 95% of oxygen which we breathe to oxidize foodstuffs in the respiratory chain where food is burned. By using a series of enzymes encoded in their DNA. The cells themselves can break down glucose without an oxygen requirement, called glycolysis. This process occurs in the cytoplasm of the cell outside the mitochondria. The breakdown of food occurs mainly in three steps. First, fuel breakdown occurs in cytoplasm then breakdown food is oxidized in Krebs cycle and at last by using four enzyme complexes or carriers (I, II, III, and IV) in electron transport chain Figure.1 (ETC) or chemically called oxidative phosphorylation (oxphos). In this process electrons (hydrogen ions) extracted from food are transferred from one complex to another complex in a redox process to a final stage IV where energy (ATP) is produced (Solaini, Sgarbi et al. 2011).

This is the first stage of fuel breakdown, fatty acids, carbohydrates, and proteins from food are broken down into their individual molecular components and used for energy production. Oxygen (O₂) is used by mitochondria as fuel to oxidize foods in the oxphos chain (Alberts, Johnson et al. 2002). Several important molecules are needed to run the Krebs cycle, like all the enzymes and other substrates. In the process of food digestion first glucose (from sugar), enters into the cell using some special molecule in the membrane called "glucose transporters". Glucose converts into pyruvate in multi-steps through the process of glycolysis. Pyruvate then transported from the cytosol into the mitochondria. Pyruvate then converted into acetyl COA, by a soluble multienzyme pyruvate dehydrogenase complex. In this energy-rich form, the molecule is fully oxidized to carbon dioxide (CO₂) by reactions of the citric acid cycle or Krebs cycle (Lodish, Berk et al. 2000).

It is very necessary to continue the breakdown process of the citric acid cycle inside the mitochondria. Carbon and hydrogen, the best fuels, are released during the process of the Krebs cycle. These carbon molecules are used to make more CO₂ that together with hydrogen

ions (H^+) is take up by NAD and FAD. The CO_2 is a form of toxic waste in the body. This is quickly removed by red blood cells and hydrogen ions along with their partner electrons reach the terminal respiratory enzyme complex cytochrome oxidase to combine with oxygen to make water (H_2O). The electrons obtained in the oxidation of acetyl COA are a couple with oxaloacetate to make citric acid. Then citric acid gets oxidized to succinic acid, then succinate is converted to fumaric acid than malic acid and back to oxaloacetate. The electrons released during the oxidation are then transferred into the third stage of the “electron transport chain” with four major respiratory enzyme complexes that are-

- A. Complex I NADH dehydrogenase
- B. Complex II succinate dehydrogenase
- C. Complex III cytochrome C – reductase
- D. Complex IV Cytochrome oxidase a/a 3



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Figure.1 Mitochondrial energy metabolism.

This process is based on oxidation-reduction potential. Electrons are transported across the inner mitochondrial membrane to the final cytochrome oxidase complex a/a3 (complex IV). From here it passes them to molecular oxygen and combines with protons, generating energy in the form of ATP. Then the synthesis of ATP is done by complex V. The mitochondrial enzyme ATP synthase and energy stocked can be transported to cells where it is needed. This ATP is utilized in the various purpose of the cell such as growth, development, and differentiation.

Another mechanism that occurs in the mitochondria is fatty acid oxidation. The enzymes of the beta-oxidation pathway are also present in the mitochondrial matrix. Some enzymes of the gluconeogenesis and urea cycle are found in the mitochondrial matrix. Cytosolic NAD which is required for the substrate-level phosphorylation step in glycolysis is regenerate by mitochondria. Mitochondria are also maintained the intracellular homeostasis of inorganic ions such as phosphate and calcium. Many studies demonstrate that mitochondria play an important role in the series of intracellular events that lead to apoptosis (Zamzami, Susin et al. 1996).

MITOCHONDRIAL LINK WITH CANCER

Now a day researchers are interested to develop anti-cancer drug targeting to mitochondria because in many studies researchers found that mitochondria have a link with cancer. In tumor cells, many metabolic pathways altered which occurs only in mitochondria. Metabolic alterations that are related to mitochondrial function have been documented in tumor cells are increased rate of gluconeogenesis (Lundholm, Edstrom et al. 1982), increased lactic acid production and reduced pyruvate oxidation (Mazurek, Boschek et al. 1997), increased glutaminolytic activity (Mazurek, Boschek et al. 1997), and reduced fatty acid oxidation (Ockner, Kaikaus et al. 1993).

MITOCHONDRIAL DNA

Normally all mammalian cells have about 1000 mitochondria and up to 10,000 copies of mitochondrial DNA (mtDNA). The mitochondrial genome is double-helical a 16.6 kb, closed-circular molecule. Two rRNAs, 22 tRNAs, and 13 polypeptides are encoded by mitochondria (Figure.2 (Baysal 2006)). Mitochondria have seven, one, three, and five subunits of respiratory enzyme Complex I NADH dehydrogenase, Complex III cytochrome C – reductase, Complex IV Cytochrome oxidase a/a 3, and complex V respectively. Other mitochondrial proteins, among those participating in the replication, transcription, and translation of mitochondrial DNA, are encoded by nuclear genes, then a specific transport system targeted to these proteins to mitochondrion by a specific transport system (Schatz 1996). Mitochondrial genes show maternal inheritance means that they are inherited from the female, this is probably because the number of mtDNA copies in the egg is normally 103-fold higher than that present in the sperm (Song, Ballard et al. 2014).

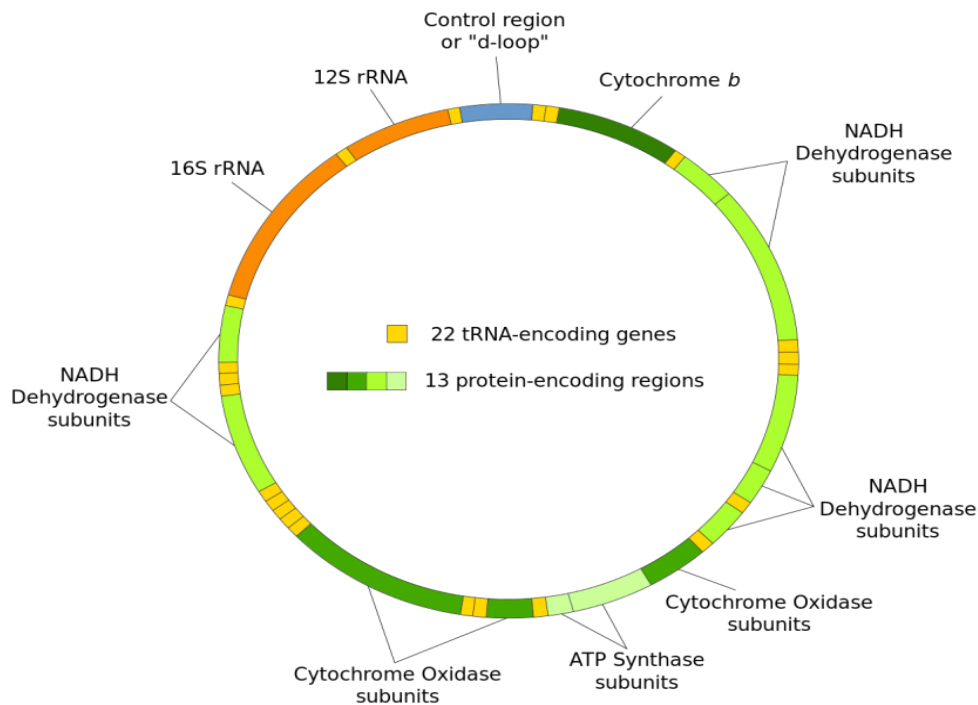


Figure.2 Structure of mitochondrial DNA

By derivative work: Shanel (talk) Mitochondrial DNA de.svg: translation by Knopfkind; layout by jhc (Mitochondrial DNA de.svg) [GFDL (<http://www.gnu.org/copyleft/fdl.html>) or CC-BY-SA-3.0 (<http://creativecommons.org/licenses/by-sa/3.0/>)], via Wikimedia Commons

MITOCHONDRIAL DNA IS MORE VULNERABLE TO MUTATION THEN NUCLEAR DNA

Mammalian mitochondrial DNA (mtDNA) represents approximately less than 1% of total cellular DNA. Its gene products are required for normal cellular function. Nuclear DNA has defensive histones and introns but mtDNA does not have defensive histones and intron are vulnerable to ROS generated by oxidative phosphorylation. Additionally, the replication of mtDNA is highly error prone (Kunkel and Loeb 1981, Shay and Werbin 1987, Singh 2004).

The mutations in mtDNA are about tenfold greater compared to nuclear DNA (Grossman and Shoubridge 1996). Normally all mitochondria have identical DNA sequences into a single cell are called homoplasmic, when differences in mtDNA sequences into the same cell, happens it is called heteroplasmy. This may happen due to de novo or “somatic” mutations in cells. When a minimum threshold of heteroplasmy has been reached mitochondrial pathologies arise. This threshold value is different for different tissue types and it is based on the energy status of a cell.

MITOCHONDRIA AND CANCER

ALTERATIONS OCCURS AT METABOLIC LEVEL

The specifically mitochondrial bioenergetic function associated with metabolic aberrations has been observed in cancer cells. These comprise differences between normal and tumor cells regarding, rate of electron transport and anion transport, preference for respiratory substrates, and also the capacity to accumulate and retain calcium (Pedersen 1978). There are activities of some enzymes participating in the procedure of oxidative phosphorylation that is

decreased in malignant versus normal cells. For example, in hepatocellular carcinoma, the measured maximal velocity of ATPase activity in isolated mitochondria (Pedersen and Morris 1974) and submitochondrial particles (Capuano, Guerrieri et al. 1997) is lower than in normal liver cells. In cellular homogenate and the mitochondrial samples from cultured human carcinoma cell lines, the activity of cytochrome c oxidase is less than that calculated in the control epithelial cell line (Modica-Napolitano and Singh 2002). Decreased activity of mitochondrial cytochrome c oxidase has been documented in biopsies of human colonic adenocarcinoma compared to normal colon mucosa (Sun, Sepkowitz et al. 1981), and in cultured rat HC252, hepatoma cells compared with non-neoplastic liver cells (Sun and Cederbaum 1980). In certain hepatoma compared to normal liver cells the adenine nucleotide exchange purpose of adenine nucleotide translocase (ANT) and the reactivity of (ANT) enzyme to bongkreikic acid (Chan and Barbour 1983) are also decreased in mitochondria. The membrane potential of mitochondrial is higher in carcinoma cells than in normal epithelial cells (Johnson, Walsh et al. 1981, Summerhayes, Lampidis et al. 1982). It is very important to note that despite the large number of metabolic alterations identified, undoubtedly none of these is common to all tumor cells because of the number of differences in the molecular constitution of the mitochondrial inner membrane among the normal and tumor cells have also been found. Alterations in the level of gene expression of cytochrome c oxidase also decrease this enzyme's specific activity in tumor cells. In colonic biopsies of carcinoma versus normal mucosa samples collected from humans, the mean level of expression of the mtDNA encoded subunit COX III was shown to be minimum (Heerdt, Halsey et al. 1990). In addition alterations in gene expression between normal and tumor cells comprise the anti-apoptotic oncogenes encoding Bcl-XL and Bcl-2, and genes encoding the PBR-linked protein Prax-1 and the peripheral benzodiazepin receptor (PBR), and mitochondrial creatine kinase (Kroemer 1997, Reed 1997). The lower expression of BAX a pro apoptotic, inner mitochondrial membrane protein also found in cancer cell lines (Nishikawa, Oshitani et al. 2005).

ALTERATION OCCURS AT GENETIC LEVEL CHANGES

The mitochondrial genome, known to be a mutational "hot spot" in human cancer particularly the displacement loop (or D-loop) region. The D-loop is a triple-stranded and non-coding region of mtDNA. In humans, the 16,569 base pairs of mitochondrial DNA encode for only 37 genes (Anderson, Bankier et al. 1981) and houses cis-regulatory elements that are required for replication and transcription. Somatic mutations in the region of the D-loop region have been documented in uterine serous carcinoma (Pejovic, Ladner et al. 2004), hepatocellular carcinoma (Tamori, Nishiguchi et al. 2004), melanoma (Takeuchi, Fujimoto et al. 2004), gastric cancer (Zhao, Yang et al. 2005), colorectal cancer (Lievre, Chapusot et al. 2005), breast cancer (Tan, Bai et al. 2002), and ovarian cancer (Liu, Shi et al. 2001). These studies suggest that mutations in the D-loop region of mitochondrial DNA maybe take part in the carcinogenesis of human cancers.

Apart from D-loop region mutations, some other mutations like deletions, point mutations, insertions, and duplications are also found in the mitochondrial genome. These mutations have been documented in a variety of human cancers such as thyroid, ovarian, liver, lung, salivary, colon, and prostate cancer (Clayton and Vinograd 1967, Boultonwood, Fidler et al. 1996). For example, insertion of a 40 bp sequence confined in the COX I gene found to be particular for renal cell oncocyoma (Welter, Dooley et al. 1989). and a deletion mutation resulting in the loss of mitochondrial DNA into the NADH dehydrogenase subunit III is related to renal carcinoma (FIGURE.3) (Selvanayagam and Rajaraman 1996). In renal cell carcinoma, the co-occurrence of somatic and germ-line mitochondrial DNA mutations has

also been reported (Sangkhathat, Kusafuka et al. 2005). In human cancer, tumor-specific changes in mitochondrial DNA copy number have been documented. For example, the mtDNA content was found to be increased in primary tumors of head and neck squamous cell carcinoma (Canter, Kallianpur et al. 2005) and the histopathologic rank was found to be an increase in premalignant and malignant head and neck lesions (Mandelker, Yamashita et al. 2005). Mitochondrial DNA copy number was revealed to increase conversely, it was also reported that mitochondrial DNA content is decreased in 80% of breast tumors compared to normal controls cells. Additionally, one current study exploring mitochondrial DNA copy number, in the number of cancers, both increases, and decreases in mitochondrial DNA content associate to controls were discovered for each type of cancer (Liu, Shi et al. 2001). Hence, the actual mitochondrial DNA copy number in specific cancers might depend upon the particular site of mutation, which means that mutations caused in the D-loop region, which regulate mitochondrial DNA replication, would be predicted to result in lowering its copy number. Conversely, mitochondrial DNA mutations in genes encoding oxphos proteins might be probable to increase in mitochondrial DNA copy number. It has been hypothesized that this response might happen due to compensatory response to mitochondrial dysfunction (Kim, Lee et al. 2004).

Tissue	D-Loop	12 S RNA	16 S RNA	ND1	ND2	ND3	ND4	ND4L	ND5	ND6	COX 1	COX 2	COX 3	COX 4	ATP 6	ATP 8
Breast	↑	↑		↑	↑								↑	↑	↑	
Ovary	↑	↑												↑		
Bladder/ Esophagus/ Head/Neck /Hepatic/ Lung	↑															
Colon		↑	↑	↑				↑	↑		↑	↑	↑	↑		
Gastric	↑			↑					↑		↑					
Pancreas	↑	↑	↑	↑	↑		↑	↑	↑	↑	↑	↑	↑	↑	↑	↑
Thyroid	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑	↑
Prostate	↑		↑				↑	↑								
ALL									↑			↑			↑	
MDS												↑	↑	↑		↑

FIGURE. 3. mtDNA mutations found in different types of tumors investigated, Arrows point out the coding and non-coding (D-loop) regions of mtDNA mutated in tumors. (Modica-Napolitano, Kulawiec et al. 2007)

MITOCHONDRIAL BIOENERGETICS IN TUMOR CELLS

PRODUCTION OF ENERGY

It is very necessary to generate a huge amount of ATP for energy and also for the synthesis of biomolecules required for the cells such as nucleotides, lipids, and proteins, required for cancer cell's rapid proliferation.

RATE OF GLYCOLYSIS INCREASED IN TUMOR CELLS

Cancer cells use glucose for many reasons (Kroemer and Pouyssegur 2008). First, it enables cells to use the extracellular nutrient, glucose to generate an adequate amount of ATP. The total yield of ATP per glucose molecule consumed is low. However, the study suggests that if the process of glycolysis is high enough, the total percentage of ATP generated from glycolysis can go greater than that of that generated from oxphos (Guppy, Greiner et al. 1993).

Second, in the anaerobic glycolysis, tumor cells may tune themselves to conditions of various oxygen concentrations (because of variable hemodynamics of distant blood vessels) that would be very toxic for cells that depend on oxphos to synthesize cellular ATP (Pouyssegur, Dayan et al. 2006).

In third, the process of anaerobic glycolysis, tumor cells produce lactic acid which creates a favorable acidic condition, essential for tumor cell invasion (Swietach, Vaughan-Jones et al. 2007). Fourth, and very necessary, for biosynthetic pathways, tumor cells utilize intermediates of the glycolytic pathway. Hence, tumors can metabolize glucose via the pentose phosphate pathway (PPP) to generate NADPH that can take part in the synthesis of fatty acid and together with ribose 5-phosphate, to synthesis of nucleotide from de novo pathway (DeBerardinis, Lum et al. 2008). Long-chain fatty acids synthesize from acetyl-CoA, malonyl-CoA and NADPH by fatty acid synthase, its activity is up regulated in a lot of tumor cells and contrary to normal cells, de novo fatty acid synthesis also take place at higher rates in cancer cells (Kuhajda 2000). Besides, tumor cells utilize intermediates of the glycolytic pathway for synthesis of glycogen, lipid, and alanine and malate (Gatenby and Gillies 2004).

WARBERG EFFECT

Cancer cells consume glucose at a higher rate and produce lactate, not carbon dioxide (CO₂) Otto Warburg has been the first to observed (Warburg 1956). This observation suggests that tumors use less of the highly efficient oxphos and use glycolysis for ATP production. This shift happens even when, there was adequate oxygen available to provide mitochondrial energy metabolism and so, this event has been known as "aerobic glycolysis" or the "Warburg effect". While the enhanced rate of glycolysis is a main characteristic of tumorigenicity. Global ATP concentration and adenylate energy charge change a little bit in cancer cells versus normal cells have shown by the number of bioenergetics measurements (Brizel, Schroeder et al. 2001). It is suggesting that mitochondrial metabolism is not essentially shut down. Otto Warburg first hypothesized that an increased rate of glycolysis under normal oxygen conditions arose from a deficiency in the mechanism of oxidative phosphorylation in mitochondria (Warburg 1956). Many studies have shown that alteration in

the mitochondrial energy metabolism in tumor cells as the number of mitochondria decreases, changes in mitochondrial ultra structure, and composition of the oxphosprotein in the respiratory chain activity and occurrence of somatic mutations mtDNA (Polyak, Li et al. 1998, Isidoro, Casado et al. 2005). Interestingly, it has been described in the AMPK pathway mediate the up regulation of glycolysis and the related attenuation in the oxphos in two human cancer cell lines (Wu, Neilson et al. 2007).

Though, these studies show no reliable pattern and do not explain the hypothesis of Warburg. Furthermore, several studies have demonstrated that mitochondrial energy metabolism is not changed in all cancer cells (Zu and Guppy 2004). Consequently, several mechanisms in cancer cells for increased glycolysis have been reported. Nowadays further research in molecular biology and biochemical proof representing those oncogenes most likely take part in the mechanisms which run the switch to anaerobic glycolysis.

HIF-1 ACTIVATION AND SUPPRESSION OF MITOCHONDRIAL ACTIVITY

HIF-1 α , which is stabilized and functional at lower oxygen consumption, increases the rate of glycolysis by increasing the glycolytic enzymes and glucose transporters (Semenza, Jiang et al. 1996). Recent studies demonstrate that HIF-1 suppresses mitochondrial function in cancer cells, suggesting that it altered the reciprocal relationship between the process of glycolysis and oxidative phosphorylation.

The switch among two processes, glycolysis and oxphos is regulated by the corresponding activities of two important enzymes, pyruvate dehydrogenase (PDH) and lactate dehydrogenase (LDH). Pyruvate dehydrogenase kinase 1 (PDK1) regulated the activity of pyruvate dehydrogenase (PDH). HIF-1 was found to stimulate Pyruvate dehydrogenase kinase 1 (PDK1) and in that way, it inactivates pyruvate dehydrogenase (PDH) the enzyme help in the conversion of pyruvate into acetyl-CoA as a result, it suppresses the citric acid cycle (Kim, Tchernyshyov et al. 2006, Papandreou, Cairns et al. 2006) and the process of mitochondrial respiration (Kim, Tchernyshyov et al. 2006). The enzyme lactate dehydrogenase A is encoded by the gene which stimulates by the HIF-1. This enzyme converts pyruvate into lactate (Semenza, Jiang et al. 1996). This result would further diminish the use of pyruvate by mitochondria, inhibiting the mitochondrial respiration. HIF-1 can also alter the expression of cytochrome oxidase, under hypoxic conditions; the composition of COX subunit is altered to optimize its activity. The expression of the COX4-2 subunit is enhanced, whereas the COX4-1 subunit, which optimizes COX activity under normal aerobic conditions, is degraded by a mitochondrial protease known as LON (Fukuda, Zhang et al. 2007).

Consequently, some other mechanisms for the increased glycolysis in tumor cells have been recognized and molecular and biochemical confirmation signifying that oncogenes underlie the process that drives the switch to anaerobic glycolysis particularly by stabilization of HIF-1 α (Almeida, Moncada et al. 2004, Elstrom, Bauer et al. 2004). Therefore, oncogenes, for example, Ras or Myc have been trigger glycolysis by induction of glucose transporters and glycolytic enzymes phosphofructokinase or hexokinase (Dang and Semenza 1999). Fascinatingly, it has been looked like that the above displayed cancer-gene idea is completely in tune with the Warburg effect but, most likely, with oncogenes as the result cause metabolic alterations (Ramanathan, Wang et al. 2005). In contrast to oncogenes, the induction of the tumor suppressor gene p53 cause diminishes the level of glycolysis and a raise in oxidative phosphorylation (Bensaad, Tsuruta et al. 2006). Therefore, the loss of p53, the tumor suppressor can enhance the glycolytic flux in cancer cells (Levine and Oren

2009). Several studies have shown that ROS are involved in the induction of HIF and stabilize it under hypoxic conditions (Brunelle, Bell et al. 2005, Guzy, Hoyos et al. 2005). Fascinatingly, ROS also appears to act downstream of some oncogenes to stabilize HIF under the condition of normoxia, leading to activation of HIF and the support tumorigenesis (Gao, Zhang et al. 2007). Recently, it has been shown that some mitochondrial proteins can be tumor suppressors whose modification indirectly induce anaerobic glycolysis, four genes are concerned with it, SDHD and SDHC, and the fumarase gene FH, the succinate dehydrogenase (SDH) genes (Gottlieb and Tomlinson 2005).

ROS DAMAGE

Oxygen is very essential for the survival of the living organism and forms the basis of cellular respiration. But oxygen (O_2) in the form of free radical serves as a molecule that causes oxidative stress (Stamati, Mudera et al. 2011). When free radicals produced in a large amount to the human body these state is caused by a deleterious effect on the life cycle of living cells (Bensaad, Tsuruta et al. 2006). So this oxidative stress is the origin of many age-related diseases such as Alzheimer's and Parkinson's, cancer and aging. When the ratio of oxidative stress generators like free radicals and the antioxidants get imbalance, then cell suffers from oxidative cellular stress (Liguori, Russo et al. 2018). Reactive oxygen species (ROS) include Singlet oxygen, hydroxyl radicals, superoxide, hydrogen peroxide, hydroperoxyl radicals, and ozone, etc (Halliwell and Cross 1994). The hypoxic environment of proliferating tumor cells provides a facility for ROS production. However, in hypoxia conditions, when electron transport complexes are in the reduced state, the ROS level can also be enhanced (Guzy and Schumacker 2006). Consequently, under hypoxic circumstances and in particular, after normalization of oxygen (O_2) supply, production of ROS in cancer cells can be enhanced to level that might leads to damage fundamental biomolecules such as mitochondrial DNA, lipid and protein. This might trigger a dangerous cycle (hypoxia, ROS generation, mitochondrial DNA (mtDNA) mutations, failure of the mitochondrial respiratory chain, etc), this series of events damaging mitochondrial structure and function, causing a shift to glycolytic ATP generation.

ROS DETOXIFICATION IN CELL

Under normal physiological conditions of cells, the intracellular levels of ROS are continuously maintained by some special types of enzymes known as an antioxidant enzyme like superoxide dismutase, catalase, Peroxiredoxins, etc. Some other non-enzymetic molecules also serve as a ROS scavenger like glutathione, flavonoids, and vitamins A, C, and E, to prevent cells from damage caused by ROS (Nita and Grzybowski 2016). Superoxide dismutases (SODs) are a type of metalloenzymes that catalyze the dismutation of superoxide anion to oxygen (O_2) and hydrogen peroxide (H_2O_2). Superoxide dismutases utilize metal ions such as copper (Cu^{2+}), zinc (Zn^{2+}), manganese (Mn^{2+}), or iron (Fe^{2+}) as cofactors (Younis 2018). The various types of SOD enzymes are present in various compartments of the cell and are highly specific in controlling associated biological processes (Copin, Gasche et al. 2000). The decomposition of H_2O_2 to water and oxygen facilitates by Catalase. The localization of catalase in most eukaryotes is in the peroxisomes and cytosol (Glorieux and Calderon 2017). These enzymes play a major role in the detoxification of ROS.

In cancer cells increased levels of ROS can result from mitochondrial impaired, enhanced cellular receptor signaling, high metabolic activity, oncogene action, increased activity of oxidases, peroxisome action, thymidine phosphorylase and lipoxygenases or during cross-talk between infiltrating immune cells (Storz 2005).

MAJOR SIGNALING PATHWAYS IN CANCER REGULATED BY ROS

In various types of cancer a number of ROS-sensitive signaling pathways are elevated where they participate in cell growth, proliferation, differentiation, glucose metabolism, cell survival, and inflammation process (Kumari, Badana et al. 2018). ROS, mainly (H_2O_2), can serve as a second messengers in cellular signaling (Di Marzo, Chisci et al. 2018). H_2O_2 controls the protein activity by reversible oxidation of its targets involving receptor tyrosine kinases, proteintyrosine kinases, protein tyrosine phosphatases, and transcription factors (Truong and Carroll 2013).

ROS-MEDIATED CONTROL OF THE MAPK/ERK1/2 PATHWAY

In cancer, the MAPK (mitogen-activated protein kinase)/Erk1/2 (extracellular-regulated kinase 1/2) pathway is activated by growth factors and K-ras. It was functionally associated with enhanced cell growth (Khavari and Rinn, 2007; Roberts and Der, 2007). For instance, in human breast cancer cells, Erk1/2 activated through (H_2O_2) generated as a by product through the metabolism of estrogen, enhances cell proliferation (Aggarwal, Tuli et al. 2019).

For instance, Ras is known as an upstream activator for Erk1/2. It can be activated directly through oxidative modification at its cysteine 118 residue, leading to the inhibition of GDP/GTP exchange (Mitchell, Hobbs et al. 2013). It was shown that increased Erk1/2 activity in ovarian cancer cells in the presence of the elevated concentration of endogenous ROS consequences from sustained ubiquitination and loss of endogenous MKP3 (mitogen-activated protein kinase phosphatase 3), a phosphatase that negatively-regulates Erk1/2 activity (Chan, Liu et al. 2008). In Addition to its effects on cell growth, it was also shown in multiple cancers types like ovarian cancer, breast cancer, melanoma, and leukemia that the activation of Erk1/2 through ROS anchorage-independent growth, motility and increases cell survival. While a Participation of ROS-activated Erk1/2 signalling in cell growth is well established (Roberts and Der 2007). The ability of ROS to control cancer cell survival appears to be cell-type definite (Rygiel, Mertens et al. 2008). For instance, MCF-7 and MDA-MB-435 breast cancer cells when treated with ROS inhibitor or scavengers or that support apoptosis and cell adhesion by targeting Erk1/2 or its upstream kinase MEK (mitogen-activated protein kinase) (Ostrakhovitch and Cherian 2005).

OXIDATIVE STRESS REGULATES THE PI3K/AKT PATHWAY

Akt or protein kinase B; PKB mediates cell survival by phosphorylation and inactivation of its substrates such as the pro-apoptotic proteins Bad, Bax, Bim, or FOXO transcription factors that take part in apoptosis which leads to cell death (Bonni, Brunet et al. 1999, Zhang, Tang et al. 2011). In breast cancer, ROS production during estrogen metabolism or many other potential mammary carcinogens was shown to activate the PI3K/Akt signaling pathway (Park, Na et al. 2009, Zhang, Tang et al. 2011). Hydrogen peroxide (H_2O_2) produced by epithelial growth factor (EGF) in human ovarian cancer cells activates Akt and p70 S6K1, a substrate of Akt that regulates protein synthesis process (Liu, Hu et al. 2006). Furthermore, the inhibition of ROS by ROS scavenger in the human pancreatic tumor cell line Panc-1 reduced the levels of phosphorylated (active) Akt and induced apoptosis (Subramani, Gonzalez et al. 2016). The activity Of AKT is firmly controlled by a signaling cascade that includes the kinases PDK-1 (3'-phosphoinositide-dependent kinase-1), mTOR and PI3K as well as the phosphatase PTEN (phosphatase and tensin homolog deleted on chromosome 10). PDK-1 and mTOR control Akt activating phosphorylations at S473 and T308, whereas PI3K generates phosphatidylinositol-3,4,5-triphosphate (PIP3), which serves as a membrane anchor (Storz and Toker 2002). PTEN serves as a negative regulator of the PIP3 level and thus decreases Akt activity (Ogg and Ruvkun., 1998; Sun et al., 1999). Study shows that

treating cells with exogenous hydrogen peroxide leads to activation of Akt and PDK-1 (Higaki, Mikami et al. 2008). It is observed that PTEN is reversibly inactivated by (H_2O_2) (Lee, Yang et al. 2002). Loss of PTEN increases basal levels of hydrogen peroxide (H_2O_2) and superoxide due to a reduction in the expression of several antioxidant enzymes such as copper/zinc superoxide dismutase and peroxiredoxins (Basseres and Baldwin 2006). This reveals a constant Akt activation through increase ROS production due to depletion of PTEN but also activation of its upstream kinases mediated by oxidative stress.

REGULATION OF THE IKK/NF- κ B PATHWAY BY ROS

In various cancer types, the transcription factor known as NF- κ B is uncoupled from its usual state of regulation and demonstrates increased activity (Biswas, Shi et al. 2004). Recent studies have recognized a pivotal role for NF- κ B in the regulation of cell cycle and proliferation, tumor cell survival, cellular adhesion, and development of drug resistance in cancer cells during therapy (Ahmed, Cao et al. 2006, Bourguignon, Spevak et al. 2009). NF- κ B serves as a redox-regulated sensor for oxidative stress (Li and Karin 1999) and low doses of (H_2O_2) activated it (Schreck, Albersmann et al. 1992). In its inactive state, NF- κ B is inflexibly bound to I κ B, its inhibitor that sequesters the transcription factor in the cytosol (Beg, Finco et al. 1993, Bonizzi, Bebién et al. 2004). The canonical activation of NF- κ B is mediated by the NF- κ B-inducing kinase (NIK) and the I κ B kinase (IKK) complex, consisting of IKK α , IKK β , and NEMO. Upon its activation by cytokines for example TNF α or IL-1, NIK phosphorylates and activates its downstream targets, the kinases IKK α and IKK β (Bergmann, Hart et al. 1998). Active IKKs phosphorylate I κ B and this process leads to its subsequent ubiquitination and proteasomal degradation (Singh, Darnay et al. 1996, Ghoda, Lin et al. 1997). Degradation of I κ B translocates NF- κ B to the nucleus, where it acts as a transcription factor to provoke the expression of anti-inflammatory genes and anti-apoptotic genes (Liu, Zhang et al. 2017). Oxidative stress activates NF- κ B by distinct signaling pathways (Janssen-Heininger, Poynter et al. 2000).

INVOLVEMENT OF MITOCHONDRIA IN CELL DEATH

In both normal and cancer cells, mitochondria not only participate in cellular ATP production but also play vital life-supporting roles in cellular homeostasis as well as ROS production, calcium signaling, and amino acid production (cataplerosis). ATP formation and cataplerosis are synchronized during the shared Krebs cycle in the mitochondria. To preserve the integrity of mitochondria due to cataplerosis, the entry of biosynthesis intermediates (anaplerosis) is therefore required. Glutamine was found to be an essential factor in this (Kovacevic and McGivan 1983). In contrast to normal cells, in cancer cells, glutamine becomes a major substrate for the development and proliferation of cells (Bustamante and Pedersen 1977).

Mitochondria also play a central role in the intrinsic pathway of apoptosis. In response to several different stimuli that result in the permeabilization of the outer mitochondrial membrane (OMM), mitochondria release the number of soluble proteins from their intermembrane space to the cytosol that can activate cellular apoptotic programs (Saelens, Festjens et al. 2004). These critical and lethal functions are connected to a few important components of the mitochondrial bioenergetic metabolism, like cytochrome c, cardiolipin, ROS production, and membrane permeability.

A soluble nuclear-encoded protein of mitochondria known as Cytochrome c works as a single electron shuttle from the complex III to the complex IV of the electron transport chain. Though when cytochrome c is released into the cytosol, from mitochondria it takes part in the construction of the apoptosome and the progression programmed cell death (Wang 2001).

Normally most of the cytochrome c lies within the cristae, which are supposed to form an obstacle against its diffusion into the intermembrane space of mitochondria. In this context, the release of cytochrome c through mitochondria would require two main steps during the process of apoptosis. The first step needs an early remodeling of cristae configuration (Sun, Williams et al. 2007) which would need an expansion of the narrow tubular cristae junction, building the cytochrome cristae pool releasable after its relocation in the intermembrane space (Scorrano, Ashiya et al. 2002). Studies show that cristae remodeling is connected with the disassembly of the Optic Atrophy 1 (OPA1) oligomer (Frezza, Cipolat et al. 2006). This oligomer is essential for an adequate amount of cytochrome c to be released for the process of the cascade of apoptosis (Yamaguchi, Lartigue et al. 2008). This hypothesis is also supported by information that a definite threshold of released cytochrome c must be passed to activate caspases and induce apoptosis (Clayton, Clark et al. 2005). The second step is which need for cytochrome c release is Bcl-2 family proteins that regulate the oligomerization of pro-apoptotic protein Bax/Bak in the outer membrane of mitochondria, firmly regulate the movement of cytochrome c across the outer membrane (Kluck, Bossy-Wetzel et al. 1997). This results in construction of apoptosis-induced channel (Dejean, Ryu et al. 2010).

Moreover, in this way of cytochrome c release, the mitochondrial permeability transition pore (mPTP) is another way that plays a critical role in cell death following necrosis and ischemic injury (Halestrap 2009). mPTP mostly assembled at the contact sites of mitochondrial inner and outer membranes, it opens a large pore with low ion selectivity that enhances the permeability of the inner mitochondrial membrane (Kinnally, Peixoto et al. 2011). The important activators of mPTP opening are the mitochondrial ROS, calcium, and phosphate (Halestrap 2009). At the time of apoptotic cell death, mPTP might participate in the amplification loop that essential for cytochrome c exciting calcium mobilization through the endoplasmic reticulum (Boehning, Patterson et al. 2003). Therefore as apoptosis is stimulated, the construction of mitochondrial apoptosis-induced channel releases cytochrome c that might connect to and ease the calcium-dependent inactivation of inositol (Hanahan and Weinberg 2000) triphosphate receptors, allow to leave calcium through the endoplasmic reticulum. Calcium would then stimulate the mPTP, as a result, mitochondrial swelling occurs, this leads to disruption of the outer mitochondrial membrane which in turn leads to cytochrome c release from intermembrane space (Boehning, Patterson et al. 2003).

MITOCHONDRIA AS A CLINICAL MARKER FOR CANCER

The homoplasmic nature of mtDNA and its abundance make it an attractive molecular marker for cancer (Kirches 2017). Undoubtedly, it has been revealed that the presence of mutant mtDNA in cancer cells to be 220 times more than the mutated nuclear DNA marker (Chatterjee, Dasgupta et al. 2011). Mutant mtDNA is simply detectable in blood, urine, and saliva samples collected from patients with, bladder, neck, head, and lung cancers. mtDNA mutations have been used as clonal markers in breast cancer (Sultana, Rahman et al. 2012) and hepatocellular carcinoma (Nomoto, Yamashita et al. 2002). It is expected that 1,000 different proteins comprise mitochondria. A new approach in proteomic technologies has made possible the quantitative analysis of protein expressed in mitochondria. The mitochondrial proteomic database has newly been documented by the National Institutes of Standards and Technology (Gulcicek, Colangelo et al. 2005). The help of mitochondrial protein profiles through new research efforts in normal and cancer cells will lead to the identification of markers for a clinical finding of cancer that may provide an understanding of how differential protein expression might affect the proportion of the disease (Cho 2007). A recent study showed that mitochondrial biomarkers also help in the prediction of tumor progression and poor overall survival in gastric cancers (Sotgia and Lisanti 2017).

MITOCHONDRIA AS A TARGET IN TUMOR CELLS FOR CANCER THERAPY

Even if the main contribution of mitochondrial energy metabolism remains controversial in cancer cells, the participation of mitochondria in cancer cell proliferation persists. For instance, the down regulation of some mitochondrial protein linked with disease detection (Cuezva, Krajewska et al. 2002, Isidoro, Casado et al. 2005). For that reason, mitochondria appear to be a promising therapeutic target for cancer treatment. It is essential to target pathways that found to differ between tumor cells and normal cells since the energy metabolism of tumors and the host cells rely on the same essential pathways for ATP. Amongst the potential therapeutic pathway targets in mitochondria that could put off-tumor progression was glutaminase or aminotransferases, enzymes that permit glutamine to trigger the Krebs cycle to make sure anaplerosis. Certainly, inhibiting aminotransferases by aminoxyacetic acid inhibits xenograft tumor growth of MDA-MB-231 breast cancer cells (Thornburg, Nelson et al. 2008). Fatty acid synthesis which takes place in mitochondria is another important target for the treatment of cancer cells. Hence, blocking the fatty acid oxidation by etomoxir seems to provoke cell death in human glioblastoma cells by diminishing NADH production, increase ROS generation, and reduce the generation of ATP (Pike, Smift et al. 2011). Additionally, the growth of human ovarian cancer cells in SCID mice was noticeably diminished by blocking the activity of fatty acid synthase (Wang 2001). Dichloroacetate indirectly triggers mitochondrial ATP formation by blocking pyruvate dehydrogenase kinase (PDK) and therefore activating pyruvate dehydrogenase (PDH) (Gudi, Bowker-Kinley et al. 1995). Dichloroacetate has been reducing in vivo and in vitro tumor cell growth (Chen, Cairns et al. 2009). In recent times, it was examined that blocking of MCT-1 results in the interruption of the metabolic symbiosis between oxygenated tumor cells and hypoxia by preventing the ability of aerobic cells to utilize lactate for oxidative phosphorylation and enforce them to utilize glucose instead of lactate. As a result, hypoxic tumor cells are deprived of glucose and died. This growth retardation of tumor cell and make available the rest of the cell (positioned at the neighborhood of blood vessels) susceptible to radiotherapy (Sonveaux, Vegran et al. 2008). To date, therapy targeting the intrinsic apoptosis pathway of mitochondria is one of the most exciting areas of cancer research and its treatment. In this point of view, mPTP is also a becoming target of selection. It was shown that lonidamine, a putative adenine nucleotide translocator (ANT), one of the major components of (mPTP) ligand, applies a cytostatic effect on tumor growth and survival in Phase II clinical study (Oudard, Carpentier et al. 2003). Furthermore, clodronate that serves as

a competitive ANT inhibitor has newly been seeming to get better the overall survival of patients suffering from primary breast cancer(Diel, Jaschke et al. 2008). To induce apoptosis in tumor cells, it is believable to therapeutically initiate mitochondrial outer membrane permeabilization (mPTP) through targeting mitochondrial membranes(Debatin, Poncet et al. 2002). The involvement of mitochondrial ROS in tumorigenic pathways demonstrates that the inhibition of ROS generation in mitochondria may be a useful strategy for cancer treatment. By using antioxidants such as quercetin, curcumin, vitamin C, and vitamin B, targeted particularly to mitochondria could signify a very fascinating approach toward cancer therapy(Mut-Salud, Alvarez et al. 2016, Forni, Facchiano et al. 2019).

The mtDNA also serves as a therapeutic target for cancer. Several discrete differences in mtDNA structure and function among normal cells and tumor cells provide the probable clinical use of mitochondria not only as markers for the diagnosis of cancer but also as targets for the new and site-specific anti-cancer drug for cancer treatment(Rowe, Weissig et al. 2001, Verschoor, Ungard et al. 2013).

Mitochondrial membrane potential plays an important role in the chemotherapeutic approach, which uses delocalized lipophilic cations (DLCs) that aggregate selectively in carcinoma cells in response to raised membrane potential. Some of these compounds have found to be effective in carcinoma cell killing *in vivo* and *in vitro*(Anderson, Wood et al. 1993). Some DLCs have been employed in photo chemotherapy (PCT), an investigational type of cancer treatment implicating light activation of a photo reactive drug or photosensitizer, that is selectively taken up or preserved by tumor cells (Weiss, Wong et al. 1987, Agostinis, Berg et al. 2011). The use of PCT as a form of treatment for neoplasms of the lung, brain, breast, or any other tissue available to light transmitted either by the body surface or internally through fiber-optic endoscopes. Cationic photosensitizers are principally capable as strong PCT agents. Similar to other DLCs, these compounds are aggregated by the cells into mitochondria in response to –inside negative trans membrane potentials and are hence, selectively accumulate in the mitochondria of tumor cells. By the use of localized photo irradiation, the photosensitizer can be transformed into a more reactive and toxic species, increasing theselective toxicity to tumor cells and providing a highly specific way to kill tumor cells without harm to normal cells of the body. Many efforts are being made to develop mitochondrially targeted drug and DNA delivery systems. A study has been demonstrated that by attaching mitochondriotropic residues to the surface of liposomes mitochondria-specific conventional liposomes can be accomplished (Ellerby, Arap et al. 1999). Another recent study suggests that targeting mitochondria in the therapy of cancer could offer a foundation for selective anti-cancer activity (Rozanov, Cheltsov et al. 2019).

The long-term therapeutic objective of this type of research in the field of cancer biology is to someday successfully create mitochondria-specific vehicles that will potentially deliver drugs or mtDNA into the organelle to eliminate dysfunctional mitochondria or restore mitochondria with functional copies of the genome.

CONCLUSION

Mitochondria play a critical role in most of the important processes of energy metabolism in the body. Thus impairment of mitochondrial function or dysfunction of mt DNA leads to many neurological disorders and cancer development. For all those reasons mitochondria serve as a promising target for cancer therapy. Furthermore, besides particularly target mitochondria, targeting the anaerobic glycolysis, for example, hexokinase and oncogenes will significantly contribute to these anti-cancer therapies. Studies have shown that all types of

cancer cells illustrate a high glycolytic flux but not all have diminished mitochondrial oxidative phosphorylation. For that reason, contrary to Warburg theory, not all tumor cells rely on glycolysis for ATP and the increased glycolysis is not the outcome of altered oxidative phosphorylation in all tumor cells. However, mitochondria stay in the middle of critical physiological processes in tumor cells and play a vital role in the progression of tumors.

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