

Tumour Metabolism: An Emerging Therapeutic Target for Cancer Treatment

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Malignant cells are highly proliferative in nature and display invasion and metastatic properties. Hence, they need rapid energy and huge amount of macromolecules for their development and progression. They fulfil their energy and macromolecular demands either by speeding up/down or switching on /off the various metabolic pathways. Therefore, altered tumor cell metabolism is considered as one of the emerging hallmark features of cancer cells. Most of the cancerous cells usually show dependency on glycolysis, glutaminolysis and lipid biosynthetic pathways.

Several studies indicate an indispensable role of glycolytic pathway in the progression of cancer via supplying rapid ATP, precursors for macromolecular synthesis [1,2]. Moreover, the high glycolytic rate plays a central role in the development of acidified tumor microenvironment (TME) via generating lactate as an end product [3]. Further, the intracellular lactate produced through glycolytic pathway is transported outside with the help of monocarboxylate transporter which leads to acidification of TME. The acidified tumor microenvironment induces immunosuppression, apoptosis of surrounding normal cells and invasion & metastasis via activating the matrix-metalloproteases [3,4]. Due to these benefits, malignant cells perform glycolysis even in the presence of sufficient oxygen, a phenomenon first time observed by Nobel laureate Otto Heinrich Warburg and thus called as 'Warburg's effect' and popularly known as 'aerobic glycolysis'. Cancerous cells maintain a high rate of glycolysis through enhancing the expression of various glycolysis regulatory molecules, namely glucose transporter, hexokinase, phosphofructokinase, pyruvate kinase, lactate

dehydrogenase pyruvate dehydrogenase kinase, and monocarboxylate transporter [2]. Recently, several glycolytic regulatory molecules are being targeted either by their inhibitors or by respective siRNAs in cancers of various origins, for the development of safer and effective targeted therapeutic strategies for the treatment of cancer patients.

The role of glutaminolysis has also been observed in various malignancies and most of the cancerous cells are found glutamine addicted and show dependency on glutamine metabolism to fulfil their bioenergetics and biosynthetic requirements [5,6]. The reason behind showing their interest in glutamine is due to its easy availability and ability to supply various precursors and intermediate molecules for bioenergetics and macromolecular synthesis pathways. Interestingly, in mammals, the level of glutamine is high in blood and muscles among the all amino acids. Cancer cells utilize glutamine for the production of numerous metabolites, nonessential amino acids, antioxidants for the removal of reactive oxygen species, and nitrogenous bases and fatty acids for the synthesis of nucleic acid and lipids, respectively [5,6]. Therefore, glutamine metabolism is emerging as a novel therapeutic target for the treatment of numerous cancers.

Reports also suggest a crucial role of lipid metabolism enzymes in tumorigenesis and tumor progression [7-9]. This could be due to following facts: i) an important component of plasma membrane, ii) one of the key energy source during the nutrient deprivation and iii) crucial regulators of various signaling pathways [7,8]. Therefore, due to above discussed benefits; cancer cells modulate

their lipid metabolic pathways according to their requirement during the course their development and progression. Further, the augmented expression/activity of lipid metabolism enzymes, such as ATP citrate lyase (ACLY), acetyl-CoA carboxylase (ACC), fatty acid synthase (FASN), and carnitine palmitoyltransferase 1 (CPT1) is found in malignancies of various origins [8,10]. Thus, cancer-specific lipid metabolism enzymes may be exploited as diagnostic markers or therapeutic targets for developing the diagnostic tools or designing the targeted therapeutic protocols against cancers.

Therefore, reprogrammed tumor cell metabolism plays a crucial role in supporting the unhindered proliferation of cancer cells via accelerating the pathways involved in the generation of biosynthetic materials, rapid energy, and membrane biogenesis. Hence, targeting of tumor metabolism could help in designing the effective and promising therapeutic strategies for the treatment of cancer patients.

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