A unique opportunity for therapeutic intervention

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The early observations by Dr Otto Warburg revealed that fundamentally metabolic differences exist between malignant tumor cells and adjacent normal cells. Many studies have further reported the relationship between altered cellular metabolism and therapeutic outcomes. These observations suggest that targeting the peculiar metabolic pathways in cancer might be an effective strategy for cancer therapy. In recent years, investigations have accelerated into how altered cellular metabolism promotes tumor survival and growth. This review highlights the current concepts of altered metabolism in cancer and the molecular targets involved in glycolysis, mitochondria and glutamine metabolism and discusses future perspective of cellular metabolism-based cancer treatment.

Introduction

The tendency of cancer research during the past decades has been focused on the gain of function of oncogenes and loss of function of tumor-suppressor genes. Although this paradigm provided major insights into cancer biology, it is now clear that there are additional factors contributing to carcinogenesis that cannot be ignored. Hanahan and Weinberg recently updated their review published in 2000 on the hallmarks of cancer by incorporating the remarkable progress in cancer research that was made over the last decade, one of which is "reprogramming of energy metabolism".1 Cancer metabolism has been arousing wide interest, not only because the large number of oncogenes and tumor-suppressor genes involved, but also because it reveals novel therapeutic targets for inhibiting tumor growth. Here, we summarize recent developments and therapeutic targets in the area of cancer metabolism and discuss the future direction of metabolism-targeted cancer therapy.

The Warburg Effect and Cancer

The earliest and best-known cancer metabolic anomaly is described by Otto Warburg in the 1920s. Dr Warburg reported that tumors showed unusually high rates of glucose uptake and

Correspondence to: Jin-Ming Yang and Yi Zhang; Email: juy16@psu.edu and zhangyi@suda.edu.cn Submitted: 10/17/12; Accepted: 11/18/12 http://dx.doi.org/10.4161/cbt.22958 lactate production compared with normal tissues, even in the presence of oxygen (Warburg effect).² In fact, the enhanced tumor uptake of 2-deoxy-2(18F)-fluoro-D-glucose in positron emission tomography (PET) scans is now exploited in the clinic for diagnostic purposes.³ Glycolysis produces ATP with lower efficiency, but at a faster rate than oxidative phosphorylation. This faster rate of ATP production is thought to aid the rapid proliferation of cancer cells. In addition to providing ATP, the high glycolytic rate may favor the growth of cancer cells through increasing biosynthesis of important molecules such as lipids, nucleotides, NADPH and amino acids.⁴ Other metabolic products of glycolysis, such as lactate and H⁺, cause a consistent acidification of the extracellular environment⁵ and favor cancer invasion.⁶ Because of the similarity, abnormally high aerobic glycolysis is now considered a hallmark of cancer. It is therefore greatly important to understand the molecular pathways that regulate aerobic glycolysis, to exploit the altered cancer metabolism for cancer therapy.

Glycolytic Pathways as Cancer Therapeutic Targets

In the standard model, glucose is first uptaken into cells through glucose transporters (GLUTs), after which it is phosphorylated by hexokinaseII (HKII) to become glucose-6-phosphate. Glucose-6-phosphate is then converted to fructose-6-phosphate by glucose-6-phosphate isomerase to proceed into glycolysis or it can be shunted into the pentose phosphate pathway (PPP) by glucose-6-phosphate dehydrogenase (G6PDH) to participate in nucleotide synthesis. The oxidative branch of PPP also produces NADPH, which is used in combating oxidative stress.7 Phosphofructokinase-1 (PFK-1), located at downstream of HKII in glycolysis, converts fructose-6-phosphate to fructose-1,6-bisphosphate. Fructose-1,6-bisphosphate is then converted either to glyceraldehyde-3-phosphate to proceed into glycolysis or converted to dihydroxyacetone phosphate which is critical for lipid synthesis. In glycolysis, glyceraldehyde-3-phosphate is then converted to glycerate-2-phosphate by G3PDH and glycerate-2-phosphate is converted to phosphoenol pyruvate by enolase. Pyruvate kinase (PK) catalyzes an ATP-producing step of glycolysis in which phosphoenol pyruvate is converted into pyruvate with the release of ATP. Further down the glycolysis pathway, lactate dehydrogenase-A (LDH-A) converts pyruvate to lactate and generates NAD⁺ from NADH. NAD⁺ is necessary for continued flux through glycolysis (Fig. 1).

REVIEW





The increased dependence of cancer cells on glycolysis offers a number of attractive potential therapeutic targets for selectively killing cancer cells. Unfortunately, both cancer cells and normal proliferating cells display the Warburg effect. It is thus challenging to specifically target cancer metabolism, while sparing normal tissues. Several candidates such as GLUT1, HKII, PK, LDH-A and G6PD are currently the focus of interest for targeting, as they are overexpressed in cancer.

GLUT1 appears to be the predominant glucose transporter in many types of cancer.^{8,9} Inhibition of glucose transporter by phloretin can preferentially sensitize cancer cells to chemotherapeutic drugs¹⁰ (Table 1).

HKII is a key glycolytic enzyme catalyzing the first step of glycolysis. Wolf and colleagues recently found that malignant gliomas show elevated HKII expression. Depletion of HKII in malignant glioma cells stimulates the production of ROS and increases cell death.¹¹ Another reason for considering HKII as a therapeutic target is that this enzyme is reversibly bound to the mitochondrial membrane protein named porin, which not only plays a role in maintaining HKII activity to regulate glycolysis, but also prevents cytochrome c release during the intrinsic apoptosis cascade.¹² Some drugs such as 3-bromopyruvate (3-BrPA) and clotrimazole can induce the detachment of HKII from

mitochondria, abolish the activity of mitochondrial bound HKII and induce cancer cell death by a dramatic decrease in the level of ATP¹³ (**Table 1**).

PK converts phosphoenol pyruvate to pyruvate with release of ATP. PK has two splice variants: PKM1and PKM2. PKM1 is found in many normal cells, whereas tumors express higher levels of PKM2.14 Due to the high expression and activity of PKM2, this enzyme is also considered as an attractive target for cancer therapy. However, a recent study found that inhibition of PKM2 activity by cellular oxidative stress promotes accumulation of glycolytic intermediates and feeds other biosynthetic pathways (such as PPP pathway), resulting in tumor progression.¹⁵ On the other hand, Goldberg and colleagues demonstrated that knockdown of PKM2 expression by RNAi increases apoptosis in vitro and causes tumor regression in vivo.16 These paradoxical results may be explained by different cellular responses to varying degrees of hypoxia. Under moderate hypoxia, PKM2 is oxidized, leading to inhibition of its activity, promotion of PPP pathway and enhanced generation of cellular NADPH. This prevents accumulation of aberrantly high levels of ROS and oxidative cellular damage. Under severe hypoxia, oxidative phosphorylation is limited and cancer cells become dependent on PKM2 activity for ATP generation. Thus, PKM2 inhibitors

Table 1. Summary table of selective potential drugs/compounds targeting cancer metabolism pathways and mode of action

Drug/compound	Molecular target	Mode of action
Phloretin	GLUT1	Blocks glucose uptake
Fasentin	GLUT1	Blocks glucose uptake
2-Deoxyglucose	Hexokinase	Blocks glycolytic flux
3-Bromopyruvate	Hexokinase	Blocks glycolytic flux
Clotrimazole	Hexokinase	Blocks glycolytic flux
Iodoacetate	G3PDH	Blocks glycolytic flux
Fluoride	Enolase	Blocks glycolytic flux
FX11	Lactate dehydrogenase	Blocks metabolic flux
Dichloroacetate	Pyruvate dehydrogenase kinase	Promotes oxidative phosphorylation
PKM2 inhibitor	Pyruvate kinase 2	Prevents ATP production in hypoxia cancer cells
PKM2 activator	Pyruvate kinase 2	Promotes oxidative cellular damage in oxygenated cancer cells
Buthionine S'R'-sulfoximine	G6PD	Inhibits oxidative PPP pathway
6-Aminonicotinamide	G6PD	Inhibits oxidative PPP pathway
Chloroquine	Autophagy	Cuts off stromal fuel supply
Metformin	Energy sensing pathways(AMPK)	Inhibits lipolysis and oxidative phosphorylation
C93	FASN	Inhibits fatty acid synthesis
FAS31	FASN	Inhibits fatty acid synthesis
Rotenone	Mitochondrial respiration complex I	Inhibits oxidative phosphorylation
α-TOS	Mitochondrial respiration complex II	Genarates reactive oxygen species
Benzylisothiocyanate	Mitochondrial respiration complex III	Genarates reactive oxygen species
Oligomycin	Mitochondrial respiration complex IV	Inhibits ATP synthesis
Resveratrol	Mitochondrial respiration complex IV	Inhibits ATP synthesis
L-asparaginase	Glutamine uptake	Depletes blood glutamine level
L-γ-glutamyl- <i>p</i> -nitroanilide (GPNA)	SLC1A5 (a target of Myc)	Inhibits glutamine uptake and inhibits mTOR activation
968	Glutaminase	Inhibits Rho GTPases
Aminoxyacetic acid (AOA)	Glutamine aminotransferase	Inhibits transamination

prevent ATP production in severely hypoxic cancer cells, whereas PKM2 activators promote oxidative cellular damage in moderately hypoxic cancer cells. The therapeutic benefits of PKM2 inhibitors and activators are therefore context- and tumor type-dependent. PKM2 inhibitors may be therapeutically effective in relatively limited-oxygenated tissues such as brain, whereas PKM2 activators may be therapeutically beneficial in relatively well-oxygenated tissues such as lung.

Lactate overproduction is another important hallmark of advanced cancer. The first metabolic target for cancer therapy was LDH-A, an enzyme which converts pyruvate to lactate and converts NADH to NAD⁺. In addition to facilitating sustained glycolysis, the overproduction of lactate leads to an acidic tumor microenvironment associated with metastasis, tumor recurrence and poor survival.¹⁷ Inhibition of LDH-A by a small-molecule inhibitor (FX11 [3-dihydroxy-6-methyl-7-(phenylmethyl)-4-propylnaphthalene-1-carboxylic acid]) reduces ATP levels, induces significant oxidative stress and inhibits the progression of sizable human lymphoma and pancreatic cancer xenografts¹⁸ (Table 1).

In addition to the above mentioned candidate targets, there are other molecules in the glycolytic pathway that are potentially implicated in cancer therapy. Fox example, 2-deoxyglucose (2-DG), a non-metabolizable glucose analog, could improve the efficacy of ABT-263/737 (molecular antagonists of the Bcl-2 family) without harming healthy cells.¹⁹ Iodoacetate, an inhibitor of G3PDH, increases cellular necrosis in pancreatic cancer.²⁰ Fluoride, an inhibitor of enolase, has been shown to induce C6 glioma cell death.²¹ Dichloroacetate (DCA), an inhibitor of pyruvate dehydrogenase kinase 1 (PDK1), increases mitochondrial production of ROS in cancer cells by shifting metabolism from glycolysis to glucose oxidation.²² Given that NADPH is important in combating oxidative stress, inhibiting PPP could also be an attractive target. Buthionine S'R'-sulfoximine (BSO), a G6PDH inhibitor, has been shown to suppress the colony-forming efficiency of G6PDH-expressing cells.23 The combination of 2DG and 6-AN (6-aminonicotinamide, another promising inhibitor of G6PDH) has been shown to greatly decrease the glutathione(GSH) content of cancer cells and enhance radiationinduced damage in human glioma and squamous carcinoma cell lines²⁴ (Table 1).

Additionally, a number of oncogenes are known to be involved in the metabolic switch from oxidative phosphorylation (OXPHOS) to glycolysis in tumor cells, such as AMP-activated protein kinase(AMPK),²⁵ hypoxia-inducible factor 1 α (HIF-1 α),²⁶ Myc,²⁷ PI3K/Akt/mTOR²⁸ and rat sarcoma viral oncogene homolog (RAS).²⁹ Targeting of these oncogenes may selectively kill cancer cells by inhibiting glycolytic pathway (Fig. 1).



Figure 2. Mitochondrial oxidative phosphorylation in cancer cells. Some energy-rich metabolites (L-lactate, ketones and fatty acids shown in yellow) derived from the tumor stroma can be transferred to the adjacent cancer cells and used for energy production via mitochondrial oxidative phosphorylation.

In 2008, while sequencing 22 human glioblastomas, researchers found that five of these tumors harbored mutations in the isocitrate dehydrogenase 1 (IDH1) gene.³⁰ IDH is a key player in the tricarboxylic acid cycle or Krebs cycle and catalyzes the oxidative decarboxylation of isocitrate to produce α -ketoglutarate (α -KG). Mutations in the enzyme isocitrate dehydrogenase 1 (IDH1) have been recently identified in secondary brain glioblastomas (> 70%).³¹ One explanation for the role of mutant IDH1 in cancer was first described in 2009. Researchers found that mutant IDH1 could reduce levels of α -KG, but lead to hypoxia-inducible factor (HIF) upregulation, a condition often seen in tumors.³² Other researchers inserted mutant IDH1 into cells and found resulting high levels of a single metabolite, 2-hydroxyglutatarate (2HG). This metabolite was thought to potentially disable mitochondria and push the cancer cells toward anerobic glycolysis.33 The IDH1 pathway thus represents a novel therapeutic target for cancer, particularly cancers like brain glioblastomas or acute myeloid leukemia that most commonly harbor IDH1 mutations.

Although the increased dependence of cancer cells on glycolysis offers an attractive therapeutic strategy in cancer therapy, some clinical trials have reported that utilization of glycolytic inhibitors may not be appropriate for certain types of tumors. Moreno-Sanchez and colleagues reported that some tumors located in the lung, mammary gland, cervix, ovaries and skin are not dependent on glycolysis, but are instead dependent on mitochondrial oxidative phosphorylation.³⁴ This suggests that glycolytic pathway-targeted strategies might only be useful in cancer cells with mitochondrial function defects or under hypoxia conditions when glycolysis is the major source of ATP.³⁵

Mitochondrial Oxidative Phosphorylation and Therapeutic Intervention

Mitochondria are important cellular organelles that fulfill a dual role in cell metabolism. During anabolic cell proliferation, they are the "power center" of the cell. However, under stressful circumstances, mitochondria control the intrinsic pathway of apoptosis by regulating the translocation of pro-apoptotic proteins to the cytosol.

Inspired by his initial observations, Warburg hypothesized that in tumors, mitochondrial respiration becomes defective, followed by an increasing reliance on glycolysis for ATP production.² However, this view has been disputed in recent years and may need a revision. Although glycolysis is the best-known metabolic phenotype of cancers, it is not a unique feature of all human cancers. In fact, in highly glycolytic tumor cells, defects in mitochondrial oxidative metabolism are not found.34 Fogal and colleagues reported that high levels of glycolysis, in the absence of adequate OXPHOS, may not be beneficial for breast cancer growth.36 Jessie and colleagues showed that introduction of activated Ras not only promotes tumorigenesis but also upregulates basal autophagy, leading to maintenance of mitochondrial

function and providing energy substrates required for the tricarboxylic acid (TCA) cycle during periods of nutrient limitation.³⁷ These studies suggest that under certain circumstances, cancer cells will depend on mitochondrial oxidative phosphorylation to meet with cellular ATP demands.

Recently, some studies have shown that hydrogen peroxide secreted by epithelial cancer cells can induce oxidative stress in adjacent tumor stroma. This oxidative stress induces autophagy, mitophagy and glycolysis in the tumor stroma.^{38,39} L-lactate derived from glycolytic tumor stroma can be transferred to the adjacent cancer cells and used for energy production via mitochondrial oxidative phosphorylation^{38,40} (Fig. 2). Similarly, Pierre reported that lactate generated by hypoxic tumor cells is a prominent substrate that fuels the oxidative metabolism of oxygenated tumor cells. These observations imply that glycolytic and oxidative tumor cells mutually regulate their access to energy metabolites.⁴¹ It has also been reported that starvation mobilizes lipid stores from adipocytes to form ketone bodies, which can also fuel the mitochondrial oxidative phosphorylation of adjacent epithelial cancer cells^{38,40,42} (Fig. 2). Tumor stroma metabolites such as lactate and ketone may thus promote tumor growth by acting as high-energy metabolites. In support of this view, a recent study by Nieman and colleagues demonstrated that lipolysis in adipocytes leads to the release of free fatty acids. Ovarian cancer cells showed increased free fatty acids uptake and β-oxidation in mitochondria, which promoted ovarian cancer metastasis and provided energy for rapid tumor growth (Fig. 2). This study suggests that metabolic coupling between adipocytes and cancer cells may also favor tumor growth and that this can

be blocked by lipid transport from adipocytes to cancer cells.⁴³ Thus, stromal-epithelial metabolic coupling and mitochondria are promising candidate targets for devising novel therapeutic intervention for cancer treatment and prevention. For instance, chloroquine inhibits autophagy and cuts off the stromal fuel supply, preventing high-energy metabolites moving from the stroma to cancer cells.⁴⁴ Metformin, an anti-diabetic reagent, inhibits both lipolysis in adipocytes and oxidative phosphorylation,⁴⁵ preventing cancer cells from using the energy-rich metabolites derived from the tumor stroma. The mitochondrial respiration complex I inhibitors, rotenone or bullatacin, induce apoptosis in a variety of cancer cell lines.^{46,47} The complex II inhibitor, α -TOS (α -tocopheryl succinate, vitamine E analogs), causes electron leakage and subsequent generation of reactive oxygen species (ROS), which are cytotoxic to tumor cells.⁴⁸ The complex III inhibitor, benzylisothiocyanate, generates ROS and induces apoptosis in breast cancer, liver and lung solid tumor.⁴⁹ The complex IV and ATP synthesis inhibitors, resveratrol or oligmycin, show antitumor actions in skin, neuroblastoma and lymphoblastoid cancer models^{47,50} (Table 1).

Target Glutamine Metabolism for Cancer Therapy

The unique role of glutamine in proliferating cells was first observed by Eagle in 1955.51 Eagle observed that many cell lines consumed 10-fold higher rates of glutamine than any other amino acids and these cell lines could not proliferate or maintain their viability in the absence of glutamine. Subsequent experiments showed that glutamine is involved in oxidative mitochondrial metabolism.⁵² In order to be oxidized, glutamine first loses its amide group to form glutamate by glutaminase (GLS), which then loses its amine group to form α -ketoglutarate by glutamate dehydrogenase, in the TCA cycle. Some cancer cells generate more than 50% ATP by oxidizing α -ketoglutarate in mitochondria. For example, in lymphoma cells, even in the absence of glucose or under hypoxia, glutamine is able to fully sustain the mitochondrial TCA cycle for ATP production.⁵³ During glutamine oxidation in the TCA cycle, glutamine metabolism contributes to mitochondrial ROS production. To combat ROS oxidative stress, glutamine can be converted directly into GSH by the enzyme glutathione cysteine ligase (GCL). The reduced form of GSH has an antioxidant role and plays a key role in controlling cellular redox homeostasis (Fig. 3).⁵⁴

In addition to ATP production, real-time ¹³C NMR studies showed that glutamine provides the major source of oxaloacetate (**Fig. 3**). Oxaloacetate is the substrate that condenses with acetyl-coA to form citrate, which serves as a precursor for fatty acids synthesis.⁵⁵ The fatty acids are used for energy production through β -oxidation or lipids synthesis for membrane production in highly proliferating cancer cells. Fatty-acid synthesis involves one key enzyme, fatty-acid synthase (FASN), which is overexpressed in cancer. Several FASN inhibitors such as C93 and FAS31 have been tested in pre-clinical studies (**Table 1**). Treatment of tumor cells with these FASN inhibitors effectively suppresses the growth in small lung cancer and mouse melanoma models.^{56,57} By contributing to citrate production, the role of glutamine as a carbon source provides the mitochondria with precursors for maintaining mitochondrial membrane potential, which is critical for glutamine-dependent cancer cells. Glutamine carbon is also converted into lactate acid and secreted from cells. Conversion of glutamine into lactate acid requires malic enzyme, which decarboxylates malic acid and produces NADPH to meet the need in the proliferating cells (**Fig. 3**).⁵⁸

Glutamine also contributes to macromolecular synthesis by means other than production of oxaloacetate, NADPH and supply of carbon to mitochondria. Real-time ¹³C NMR studies showed that glutamine also serves as a nitrogen source.58 Theamido and amino groups of glutamine contribute to metabolic intermediates in the synthesis of non-essential amino acids, nucleotide biosynthesis, hexosamines and ammonia (Fig. 3). Yuneva and colleagues reported that supplementation of glutamine-starved Myc-driven cells with TCA intermediates could rescue cell viability but not proliferation.⁵⁹ Similarly, Meng and colleagues reported that supplementation of glutamine-starved Hep3B cells with alternate nitrogen sources such as alanine or asparagine could rescue cell proliferation.⁶⁰ In nucleotide biosynthesis, the role of glutamine as nitrogen donor for both purine and pyrimidine has been implicated in support of proliferation. Gaglio and colleagues showed that the K-Ras transformed fibroblasts cultured in the glutamine-deprived media exhibit reduced cellular proliferation and decreased progression into S phase. The growth potential of the transformed cells was restored by adding the four deoxyribonucleotides.⁶¹ These results suggest that the role of glutamine as a nitrogen donor might explain some of the proliferation-inhibitory effects of glutamine deprivation.

Many types of cancer cells are sensitive to glutamine deprivation, including pancreatic cancer cells, lung cancer cells andglioma cells. Pre-clinical tests of L-asparaginase can significantly deplete blood glutamine levels and has some success in the treatment of pediatric acute lymphoblastic leukemia (ALL) (Table 1). However, L-asparaginase treatment has shown significant toxicity and side effects, such as increasing mental confusion and early signs of coagulopathy.⁶² Novel glutamine metabolism-based therapies for cancer are needed which do not impair normal tissues. Glutamine-dependent cancer cells always make use of mitochondria to produce precursors from glutamine for the synthesis of lipids, proteins and nucleotides through activities of the mitochondrial electron transport chain. Thus, inhibition of mitochondrial respiration chain by mitoformin can also inhibit glutamine metabolism and slow the growth of glutamine-dependent cancer cells.63 The oncogene Myc regulates several steps in the glutamine metabolism and the expression level of Myc has been demonstrated to be linked to the increased glutaminolysis through transcription program. L-y-glutamyl*p*-nitroanilide (GPNA), an inhibitor of SLC1A5 which is a direct target of Myc, can inhibit glutamine uptake and suppress glutamine-dependent mTOR activation⁶⁴ (Table 1). Glutamine metabolism is mediated by three types of enzymes: glutaminase (GLS), glutamate dehydrogenase and aminotransferase. Active Rho GTPases can increase GLS for transformation.⁶⁵ One compound 968, a small molecular inhibitor of Rho GTPase, can



Figure 3. Glutamine metabolism in cancer cells. High throughput glutamine uptake feeds cancer cell growth and proliferation with a large pool of carbon and nitrogen for the biosynthesis of the nonessential amino acids and fatty acids for membrane production. Carbon precursors derived from glutaminolysis (oxaloacetate and glutamate) serve as the carbon substrate for amino acid biosynthesis and lactate. Glutamate donates its amine group to these carbon substrates to produce non-essential amino acids (NEAA), nucleotide and α -ketoglutarate. Glutamine can be converted directly into GSH which has antioxidant role and plays a key role in controlling cellular redox homeostasis.

inhibit growth, migration and invasive activity of transformed fibroblasts and human cancer cells and shrink tumors in mouse xenograft models without obvious adverse effects⁶⁶ (**Table 1**). Another recent study indicates that glutamine dehydrogenase may not be the rate-limiting step in glutamine metabolism. The major route of glutamine-derived carbon that enters the TCA cycle is through transamination.⁶⁷ Thus, aminotransferase appears to be a promising target for cancer therapy.

Amino-oxyacetic acid (AOA) (Table 1), an inhibitor of transaminase, can selectively suppress the proliferation of MDA-MB-231 cells and inhibit the growth of MDA-MB-231 breast tumor xenografts in mouse model.⁶⁸ These studies suggest that selective inhibition of one step of glutamine metabolism (such as glutaminase or glutamine aminotransferase) might produce an anticancer effect without the toxicity associated with inhibition of the full glutamine metabolism chain. Based on the characteristics of tumors exhibiting increased glutamine metabolism, novel glutamine-based imaging techniques have recently emerged. Glutamine PET tracers ¹⁸F-(2S,4R)4-fluoroglutamine and L-[5-11C]-glutamine have been shown to be taken up by glutaminolytic cancer cell lines and tumors in the mouse model and can be used to image glutamine metabolism both in vitro and in vivo.⁶⁹ This relatively recent technology will facilitate the ongoing development of drugs that target glutamine metabolism pathway.

Conclusions and Perspectives

Modulating cancer metabolism is a novel therapeutic strategy for suppression of tumor growth. The increasing interest in cancer metabolism has already generated a series of new potentially useful therapeutic agents. Nevertheless, there are potential difficulties and concerns due to the low selectivity and specificity of the current generation of agents. It is known that in addition to cancer cells, some normal tissues (brain, retina or testis), stem cells and immune cells also display glycolysis, mitochondrial oxidative phosphorylation and glutaminolysis. Therefore, most current metabolism-based therapeutic agents show some toxic effects on normal cells. The question for the next decade of metabolic cancer research is: What is the best way to specifically target the metabolism of cancer cells? First, a better understanding is necessary of the key metabolic differences between cancer cells and non-malignant cells. A better understanding of how different cancer cells adapt these processes to fulfill their energy requirements may help improve the selectivity of cancer metabolism-based therapy. These efforts may lead to generation of highly specific cancer metabolismtargeted agents that uniquely induce cancer cell death. For example, Myc addicts cancer cells to glutamine by preventing them from supplying the TCA with other nutrients.⁵⁹ Thus, glutamine addiction is often exploited for metabolic therapy.

Second, we need a better understanding of how oncogenic activation controls cancer cell metabolism for proliferation. The metabolic type of a given tumor may vary widely from oxidative to glycolytic due to oncogenes activated or the microenvironmental change.³⁴ For example, under stressful environments, cancer cells can take advantage of oncogene-induced autophagy or mitophagy for energy demand. Some mutations in BRAF or KRAS can sustain cancer cell growth in low glucose conditions.⁷⁰ In this regard, key metabolic oncogene regulators could serve as therapeutic targets without impairing normal tissues.

Third, heterogeneity of metabolic profiles of different cancer types needs to be further clarified, such as brain cancer (glycolytic type), lung cancer (OXPHOS type) and others. If the metabolic profile primarily used for energy production is better elucidated, targeting the predominant bio-energetic pathway might lead to more effective utilization of the metabolismtargeted agents. Hence, re-activation of mitochondrial oxidative metabolism in glycolytic tumors or reduced mitochondrial content and OXPHOS capacity in "OXPHOS" type of tumors may be considered as potential therapeutic strategies.

Fourth, delivering the metabolism-targeted agents to the tumor sites by specific delivery systems will enhance the anticancer efficiency of those drugs and decrease their side effects. Although the success of nanoparticle interventions has currently been primary limited to laboratory tests in human cancer cell lines and mouse xenograft models, this new approach shows promise to provide an alternative strategy for clinical cancer diagnosis and therapy.

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Cancer cells display metabolic flexibility, which provides ability for a given cancer cell to alternate between glycolysis and OXPHOS in response to physiological needs. It is becoming increasingly understood that cancer cells can develop resistance to inhibition of one metabolic pathway through upregulation of other alternate metabolic pathways. Combinational approaches that target the increased metabolic flexibility of cancer cells are thus worth exploring. Bezielle (BZL101), a candidate oral drug that is distinguished from other similar agents due to its ability to inhibit both glycolysis and OXPHOS, has shown promising efficacy and excellent safety in the early phase clinical trials for advanced breast cancer.⁷¹ Similarly, the combined use of 2-DG and metformin in a recent study led to significant cell death and tumor growth inhibition, associated with decreases in cellular ATP.72 These studies indicate that deprivation of tumor bio-energetics by dual inhibition of energy pathways might be an effective novel therapeutic approach for a board spectrum of human tumors. In conclusion, altered energy metabolism in cancer cells provides a unique opportunity to develop novel and more effective anticancer therapies.

Disclosure of Potential Conflicts of Interest

The authors declare no conflicts of interest.

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