Energy disruptors: rising stars in anticancer therapy?

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The metabolic features of tumor cells diverge from those of normal cells. Otto Warburg was the first to observe that cancer cells dramatically increase their glucose consumption to generate ATP. He also claimed that cancer cells do not have functional mitochondria or oxidative phosphorylation (OXPHOS) but simply rely on glycolysis to provide ATP to the cell, even in the presence of oxygen (aerobic glycolysis). Several studies have revisited this observation and demonstrated that most cancer cells contain metabolically efficient mitochondria. Indeed, to sustain high proliferation rates, cancer cells require functional mitochondria to provide ATP and intermediate metabolites, such as citrate and cofactors, for anabolic reactions. This difference in metabolism between normal and tumors cells causes the latter to be more sensitive to agents that can disrupt energy homeostasis. In this review, we focus on energy disruptors, such as biguanides, 2-deoxyglucose and 5-aminoimidazole-4-carboxamide ribonucleotide, that interfere with the main metabolic pathways of the cells, OXPHOS, glycolysis and glutamine metabolism. We discuss the preclinical data and the mechanisms of action of these disruptors at the cellular and molecular levels. Finally, we consider whether these drugs can reasonably contribute to the antitumoral therapeutic arsenal in the future.

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INTRODUCTION

Cancer cells are characterized by uncontrolled and rapid proliferation. Deregulation of the cell division machinery requires metabolic adjustments to provide macromolecules and energy to fuel cell growth and division. In the presence of oxygen, glucose is converted via glycolysis into pyruvate, which is then transported to the mitochondria to be transformed into acetyl-CoA by pyruvate dehydrogenase for integration into the tricarboxylic acid cycle (TCA). The TCA provides intermediates for biosynthetic reactions, citrate, aspartate and two essential cofactors for the electron transport chain: NADH and FADH₂. Otto Warburg¹ was the first to demonstrate that the metabolism of cancer cells differs from normal cells. Even in the presence of oxygen, cancer cells reprogram their utilization of glucose and favor the production of lactic acid instead of transporting pyruvate into the mitochondria. Although Warburg named this process ‘fermentation’, the process is currently better known as ‘aerobic glycolysis’. This metabolic switch seems counterevative for rapidly dividing cells, which require large amounts of energy. Indeed, glycolysis is 18 times less efficient than mitochondrial oxidative phosphorylation for the production of ATP, and cells must adapt to compensate. To do so, the cells upregulate glucose uptake mainly via upregulating the expression of the glucose transporter Glut1. This avidity for glucose has proven useful for tumor detection and serves as a basis for detecting tumor cells by [¹⁸F] fluorodeoxyglucose positron emission tomography imaging.

The reliance on glycolysis is associated with the activation of oncogenic pathways. One of the most commonly altered signaling pathway in human cancer is the phosphoinositide 3-kinase (PI3K) pathway. This pathway is activated in response to growth factors and by mutations in the tumor suppressor gene Phosphatase and TENDsin homolog (PTEN). Once activated, the phosphoinositide 3-kinase pathway strongly promotes cancer cell proliferation and survival but also affects cell metabolism. The main effector of the phosphoinositide 3-kinase pathway is Akt. Akt is a regulator of glycolysis and plays a major role in the regulation of the bioenergetic balance. It stimulates glycolysis by increasing the expression and translocation of glucose transporters.² In addition, Akt indirectly activates the rate-limiting enzyme of glycolysis, phosphofructokinase-1, by phosphorylating phosphofructokinase-2, which produces fructose-2,6-bisphosphate, the most potent activator of phosphofructokinase-1.³ Finally, Akt is a strong activator of the mechanistic target of Rapamycin (mTOR) by phosphorylating and inhibiting tuberous sclerosis 2, the negative regulator of mTOR. mTOR is an indispensable catalytic subunit of two distinct protein complexes, mTOR complex 1 (mTORC1) and mTOR complex 2. Both complexes are key metabolic and bioenergetic checkpoints that integrate growth signaling and nutrient availability.⁴ Activated Akt strongly stimulates mTORC1, which positively regulates protein, lipid and nucleotide synthesis in response to sufficient nutrient and energy conditions⁵ (Figure 1). mTORC1 activation is a strong antiapoptotic and prosurvival signal.

When the nutrient supply is low, cells slow their metabolism to inhibit anabolic reactions and avoid energy shortage and death through the inhibition of mTORC1. The AMP-activated protein kinase (AMPK) is the main sensor of energy in cells. An increase in the AMP/ATP ratio induces the activation of AMPK⁷ AMPK phosphorylates and activates tuberous sclerosis 2, the negative regulator of mTOR. This signal opposes the effects of Akt and acts as a potent inhibitor of mTORC1. AMPK is a major adaptive kinase and a heterotrimeric enzyme complex; it comprises one catalytic α-subunit and two regulatory β and γ subunits. Upon an energy stress, AMP and ADP bind to the γ-subunit and induce a
conformational change that enhances its activity and the phosphorylation on Thr172 by the liver kinase B1 (LKB1). LKB1 is the upstream kinase required for the activation of AMPK in response to an energy stress. LKB1 is somatically mutated in some non-small-cell lung cancer and cervical carcinoma. Furthermore, germline mutations of LKB1 are responsible for the Peutz–Jeghers syndrome,8 which is characterized by the development of gastrointestinal and oral lesions. Importantly, AMPK can also be phosphorylated in response to calcium flux independently of LKB1 by the calcium/calmodulin-dependent protein kinase kinase 2. In addition, MAPKKK TAK1/MAP3K7 has been reported to phosphorylate Thr172 of AMPK.9

Considering the importance of the energy supply for cancer cells, the considerable interest in molecules interfering with the energy levels of cells is not surprising. Indeed, the induction of a strong energy stress in cancer cells leads to a major depletion of ATP, and consequently, to cell death. In this review, we will focus on three different promising molecules that interfere with energy cell metabolism: (i) biguanides (metformin and phenformin), used for the treatment of type II diabetes, (ii) 2-deoxyglucose, a potent inhibitor of glycolysis and (iii) Acadesine, also known as AICAR (5-aminoimidazole-4-carboxamide ribonucleotide), an energy restriction mimetic agent.

THE BIGUANIDES: MITOCHONDRIAL DISRUPTORS
The most popular drugs among this family of anti-diabetic molecules are metformin and phenformin. Phenformin was withdrawn from the market due to its toxic effects, mainly lactic acidosis. Phenformin was replaced by metformin, which is now the first-line treatment for type II diabetes, with more than 120 million prescriptions worldwide. Metformin, which has been used for decades in the treatment of diabetes, offers an excellent side effect profile. It does not induce hypoglycemia but does provoke some gastrointestinal disorders (diarrhea, bloating and nausea), which can be partially prevented by a progressive increase in the dosage. The rare cases of lactic acidosis reported (< 1/10 000) have occurred predominantly in patients with poor renal function. The mechanism of metformin action has been well studied in liver, adipose tissue, skeletal and heart muscles. The glucose-lowering effect is primarily a consequence of reduced hepatic glucose production, increased insulin sensitivity and glucose utilization by muscles and adipocytes, resulting in
decreased insulinemia. Most of these effects are the result of the activation of AMPK by metformin.

Biguanides regulate the AMPK/mTORC1 axis and reduce cancer incidence

The pioneering work of Zhou et al. in 2001 was the first demonstration that metformin induces the phosphorylation of AMPK on Thr172 (Figure 1). Then, the tumor suppressor LKB1 is required to phosphorylate AMPK in response to biguanides. The following question could then be asked: if metformin action requires LKB1 to activate AMPK, and AMPK inhibits mTORC1, what are the consequences in terms of cancer incidence in patients treated with metformin for decades? Evans et al. were the first to answer this question by demonstrating that metformin reduces the risk of cancer in a retrospective epidemiological study performed on 11,870 diabetic patients. Numerous retrospective studies followed, most of which examined specific cancers (breast, colon, pancreas and prostate) and drew the same conclusion. Of note, biases in some of the epidemiological studies have been reported, and the results should be considered with caution.

Nevertheless, the promising results of the experimental studies warrant the numerous ongoing clinical trials of metformin in cancer therapy.

Biguanides inhibit tumor growth and target cancer cell metabolism

Metformin and phenformin have a direct effect on tumor cells and inhibit the proliferation of numerous cancer cell lines. They both inhibit tumor growth and metastasis formation in multiple animal models that develop spontaneous or engrafted tumors. At the cellular level, metformin induces apoptosis, cell cycle arrest via the downregulation of cyclin D1 or autophagy, depending on the cancer cell line. However, the most striking discovery was certainly the effect of metformin on the energy metabolism of cancer cells. Two independent teams initially demonstrated that metformin inhibits complex 1 of the mitochondrial respiratory chain and targets oxidative phosphorylation (OXPHOS). Both studies showed that biguanides decrease oxygen consumption and impact the mitochondrial membrane potential. This effect was attributed to a mild but significant inhibition of complex 1. Whereas one study showed that the effect of metformin requires intact cells, the other study demonstrated that the inhibitory effect occurs in isolated mitochondria. This discrepancy was attributed to the fact that the two studies were performed in two different mitochondrial respiratory states. The mode of action of biguanides on complex 1 remained unsolved until the recent work of Bridges et al. In this study, metformin and other biguanides were found to inhibit ubiquinone reduction, one of the multiple steps of the catalytic reaction of complex 1. The discovery that metformin inhibits complex 1 activity in cancer cells even more strongly than Rotenone, a classical inhibitor of complex 1, opened new horizons for biguanides in the field of cancer metabolism. The inhibition seems specific to cancer cells, but more importantly, induces a strong decrease in the intracellular ATP concentration. Are the antiproliferative effects of metformin due to the energy stress it generates? To answer this question, Wheaton et al. have overexpressed the Saccharomyces cerevisiae protein ND11 in cancer cells. ND11 is a single-subunit NADH dehydrogenase and is resistant to metformin. It oxidizes NADH in a process similar to that of the multi-subunit mammalian complex 1. The authors showed both in vitro and in vivo that overexpression of ND11 reverses the antiproliferative and antitumoral action of metformin. As an energy disruptor, metformin has also been found to inhibit glucose production in hepatocytes and lipogenesis in prostate cancer. In response to the inhibition of complex 1 activity, cancer cells increase glucose consumption and glycolysis with an elevation of lactate production, decrease glucose oxidation and promote glutamine metabolism. Consequently, the inhibition of glutamine anaplerosis, a mechanism by which glutamine provides carbon to the TCA cycle via glutamate and α-ketoglutarate, by metformin further attenuates proliferation. Conversely, increasing glutamine metabolism rescues the antiproliferative effects of metformin. Together, these studies demonstrate that cancer cells activate compensatory pathways to counteract the metabolic chaos induced by metformin. Targeting these adaptations will improve the effect of biguanides and avoid resistance.

Biguanides target cancer stem cells

Cancer stem cells (CSCs) are localized in tumors, resistant to chemotherapy, and capable of self-renewal and differentiation. Importantly, CSCs are the cause of disease relapse. Biguanides appear to target this cancer cell population. The combination of metformin with chemotherapy has been shown to be more efficient than either drug alone in xenograft models using several cancer cell lines, and this treatment specifically targets CSCs. Furthermore, treatment with both drugs significantly prolongs the remission following xenograft implantation. This specific effect was confirmed in several other cancer models, including pancreas, breast and ovary. Interestingly, Sancho et al. have shown that CSCs rely mainly on OXPHOS and are unable to effectively induce glycolysis to compensate for reduced ATP production upon mitochondrial inhibition. The level of MYC expression controls this metabolic characteristic of CSCs; low MYC expression allows high PGCl-α expression, which results in enhanced mitochondrial biogenesis. Consequently, the observation that metformin specifically affects the viability of CSCs to a greater extent than non-CSCs is not surprising.

Biguanides and AMPK

Metformin and phenformin activate AMPK in most, if not all, cancer cells and consequently inhibit mTORC1 via tuberous sclerosis 2 protein. Several studies have shown that inhibition of AMPK with siRNA or compound C reverses the antiproliferative effects of metformin. However, we and others have found that metformin can mediate its effects independently of AMPK. We have shown that REDD1 (regulated in development and DNA damage responses 1) is upregulated in response to metformin and mediates its effects on mTORC1 and cell cycle arrest. Additional evidence for an AMPK-independent mechanism came from a study showing that metformin can still inhibit hepatic gluconeogenesis in mice lacking AMPK in the liver. According to Foretz et al., this inhibition is also independent of LKB1. Again, these reports demonstrate the diversity of the cellular actions of metformin, and elucidating the metabolic actions of metformin will certainly be beneficial for the treatment of several human pathologies.

2-DEOXYGLUCOSE: THE GLYCOLYTIC DISRUPTOR

2-Deoxyglucose (2-DG), an inhibitor of hexokinase (HK), blocks the first and rate-limiting reaction of glycolysis and competitively inhibits glucose uptake. 2-DG is phosphorylated by HK to form 2-deoxy-α-glucose-6-phosphate (2-DG-6-P), which cannot be metabolized by glycolysis but accumulates and inhibits HK. Targeting glycolysis in cancer therapy is very pertinent due to several metabolic and biological effects. First, the inhibition of glycolysis decreases the production of glycolytic intermediates, such as glucose-6-P, glyceraldehyde-3-P and 3-phosphoglycerate, which are the precursors of nucleic acids, phospholipids and
serine, respectively. Second, it decreases the antioxidant defenses of cancer cells. Third, it interferes with the N-glycosylation of proteins and induces endoplasmic reticulum (ER) stress. Finally, it induces an energy stress due to depletion of ATP.

2-DG interferes with anabolic reactions

The 2-DG-6-P produced by the phosphorylation of 2-DG by HK undergoes only the first enzymatic reaction in the pentose phosphate pathway to generate 2-DG-6-phosphogluconolactone, which cannot be further metabolized. The pentose phosphate pathway is a major metabolic pathway and is the principal source of NADPH and the source of ribose, the precursor of nucleotides. Glyceroldehyde-3-P is downstream of fructose 1,6-bisphosphate in glycolysis and generates glyceral-3-P, a precursor of phospholipids, which are the main components of the cellular membrane. Another important intermediate of glycolysis is 3-phosphoglycerate. 3-Phosphoglycerate is metabolized by 3-phosphoglycerate dehydrogenase, the first enzyme in the three-step serine biosynthetic pathway. 3-Phosphoglycerate dehydrogenase uses NAD as a cofactor to oxidize 3-phosphoglycerate into phosphohydroxypyruvate, which subsequent enzymes in the pathway convert into serine. Serine is essential for the synthesis of proteins and other biomolecules needed for cell proliferation, including nucleotides, phosphatidylserine and sphingosine. Overexpression of 3-phosphoglycerate dehydrogenase and increased biosynthesis of serine have been shown to be essential for the development of certain breast cancers.43 In conclusion, blocking glycolysis has a major impact on the principal anabolic pathways.

2-DG interferes with the antioxidant defenses

Reactive oxygen species (ROS) are primarily generated as by-products during mitochondrial electron transport. ROS cause numerous deleterious events, including inducing lipid peroxidation, DNA damage and protein oxidation. ROS thereby contribute to genomic mutations and an inability to differentiate and indirectly promote cell immortalization (which is a characteristic of cancer cells). Glycolysis, via the pentose phosphate pathway (which generates NADPH), attenuates the oxidative stress. NADPH provides reducing equivalents for glutathione- and thioredoxin-dependent peroxidase pathways,44 which are major antioxidants responsible for the detoxification of ROS and therefore protect against oxidative damage. Blocking glycolysis with 2-DG results in NADPH deficiency and induces the formation of ROS.45,46 Indeed, only one of the two potential molecules of NADPH is produced by the pentose phosphate pathway when 2-DG is added to the cells. As reported earlier, 2-DG-6-phosphogluconolactone (the product of the conversion of 2-DG-6-P) cannot be further metabolized and therefore produces only one molecule of NADPH from NADP+. Treatment with 2-DG reduces the antioxidant potential of cancer cells and causes them to become more sensitive to oxidative stress induced by radiotherapy and chemotherapy.

2-DG interferes with the metabolism of glycoproteins

The glycosylation of proteins is an important biological process that promotes protein stability, proper protein folding and cell adhesion. Glycosylation consists of the addition of N-glycans to the nitrogen of asparagine or arginine of the proteins and occurs in the lumen of the ER. During the N-glycosylation reaction, mannose is converted to mannose-glucosamine diphosphate (GDP), an important intermediate in glycansynthesis.47 Glucose participates in this process because glucose-6-P can be converted to mannose-6-phosphate and then to mannose-GDP. 2-DG is structurally similar to mannose and leads to the formation of 2-DG-GDP, which competes with mannose-GDP in the formation of oligosaccharide chains. Thus, the abnormal glycoproteins formed from 2-DG disrupt the N-glycosylation process and prevent normal protein folding. An accumulation of unfolded/misfolded proteins results in the activation of the unfolded-protein response, which acts to prevent ER stress. On the one hand, it inhibits protein synthesis, while on the other hand, it decreases the degradation of abnormal proteins. Treatment with 2-DG induces ER stress and the expression of ER-stress specific markers, including the C/EBP homologous (CHOP) protein, which plays an important role in ER-stress-induced cell death. Many studies suggest that the toxicity of 2-DG can be attributed to the inhibition of glycosylation. Indeed, addition of exogenous mannose can rescue cells from 2-DG-induced cell death but does not reverse the decrease in ATP.48,49

2-DG induces an energetic stress

The inhibition of glycolysis annihilates the two ATP-producing reactions in glycolysis: (1) the conversion of 1,3-bisphosphoglycerate into 3-phosphoglycerate, which generates two molecules of ATP from two molecules of ADP; and (2) the transformation of phosphoenolpyruvate into pyruvate by pyruvate kinase, which leads to the formation of two molecules of ATP. Ultimately, the net production of ATP by glycolysis is two molecules of ATP because HK and phosphofructokinase-1 both require one molecule of ATP. Importantly, blocking glycolysis decreases the concentration of pyruvate, the precursor of acetyl-CoA and citrate, which integrate into the TCA to ultimately produce ATP. Therefore, the inhibition of glycolysis affects OXPHOS via a reduction in TCA flux.

These actions lead to a decrease in intracellular ATP concentration, cell cycle blockade, inhibition of cell growth and cell death.50,51

The mimetic agent: AICAR

AICAR or Acadesine is a drug that directly activates AMPK. Inside the cells, AICAR is phosphorylated by adenosine kinase into 5-amino-4-imidazolecarboxamide ribotide (ZMP). ZMP is an analog of AMP and thus mimics several of its effects.52 Unlike biguanides and 2-DG, AICAR induces the phosphorylation and activation of AMPK but does not disturb the energy status of the cells. AICAR has been used for decades as an activator of AMPK and was recently used as a doping agent that increases running capacity without any training.53,54

AICAR interferes with biosynthetic pathways

AICAR has been shown to regulate several aspects of metabolism. Most of the pioneering studies on the activation of AMPK were performed using this drug in normal cells and insulin-sensitive organs. AICAR inhibits anabolic processes and promotes catabolic, ATP-generating reactions. For instance, AICAR inhibits lipid and cholesterol synthesis in the liver through the phosphorylation and inactivation of acetyl-CoA carboxylase and 3-hydroxy-3-methylglutaryl-CoA reductase (HMG-CoA reductase), respectively.55 On the other hand, a key biological effect of AICAR is the stimulation of fatty acid oxidation. AMPK phosphorylates and inactivates acetyl-CoA carboxylase, which catalyzes the formation of malonyl-CoA, an inhibitor of carnitine palmitoyl-transferase-1.56 Carnitine palmitoyl-transferase-1 is an enzyme associated with the outer membrane of the mitochondria that allows the entry of long-chain fatty acids into the mitochondria to be oxidized. AICAR is also a strong inhibitor of protein synthesis, again through AMPK, which phosphorylates and inactivates eukaryotic elongation factor-2. Numerous studies have also shown that AICAR inhibits the translation of proteins though its negative action on mTORC1. AICAR modulates the expression of transcription factors, such as SREBP1c and ChREBP, leading to the inhibition of lipogenesis. In skeletal muscle and heart, AICAR promotes glucose uptake and lipid oxidation. The stimulation of glucose uptake is
attributed to an increase in glucose transporter expression and translocation to the membrane.\textsuperscript{57} Given the antihyperlpidemic and antihyperglycemic effects of AICAR and the anti-diabetic properties of the compound in mice,\textsuperscript{58} clinicians have started to use AICAR for the treatment of type II diabetes.\textsuperscript{59–61}

AICAR and cancer

Most of the metabolic effects described above provide evidence for a beneficial effect of AICAR in cancer cells. Indeed, numerous studies have shown that AICAR promotes cell cycle arrest and/or apoptosis in many cancer cell lines.\textsuperscript{15,62–66} Interestingly, in some cases, the antitumoral effects of AICAR are independent of AMPK. Santidrian et al. have demonstrated that AICAR induces apoptosis in chronic lymphocytic leukemia, even when the catalytic unit of AMPK is deleted. Furthermore, the authors have shown that this proapoptotic effect is present in p53 mutated cells and is preceded by an increase in the proapoptotic proteins NOXA, BIM and PUMA.\textsuperscript{67} Recently, much attention has been paid to the effects of AICAR in hematological cancers, specifically in lymphoma, acute lymphoblastic leukemia and T-cell lymphoblastic lymphoma.\textsuperscript{68–70} In this model (as well as in other cancers, such as prostate cancer), AICAR is a much stronger activator of AMPK and inducer of apoptosis than biguanides or 2-DG.\textsuperscript{15} Notably, the concentration of AICAR required for the induction of apoptosis in cancer cells is within the millimolar range, which implies obvious problems for clinical use.

THE OTHER SIDE OF AMPK ACTIVATION

Although many studies support the tumor-suppressive role of AMPK, emerging evidence suggests that under certain circumstances, activation of AMPK protects against cell death. Indeed, activated AMPK confers metabolic adaptation to tumor cells which reduce anabolic reactions. Under nutrient starvation activated AMPK promotes cell survival and inhibits apoptosis.\textsuperscript{71} Not only AMPK activation maintains ATP levels but it also maintains NADPH levels which is crucial since it participates in redox regulation and cell survival. When activated, AMPK inhibits acetyl-CoA carboxylase and consequently the consumption of NADPH for fatty acid synthesis. In addition, itactivates fatty acid oxidation which generates NADPH.\textsuperscript{72} Constitutive adaptation to energy stress has been shown in pancreatic cancer cells, and this tolerance to low glucose was attributable to the high expression level of AMPK. Knockdown of the catalytic unit of AMPK diminished the ability of cells to grow in glucose-free media.\textsuperscript{73} More recently, Park et al.\textsuperscript{74} showed that the downregulation of the AMPK a1 and a2 subunits inhibits prostate cancer cell growth and induced apoptosis. This other aspect of AMPK activation is important and requires attention when considering using AMPK activators for therapeutic use.

WHAT IS THE FUTURE FOR METABOLIC DISRUPTORS IN THE CLINIC?

Metformin

The remarkable efficiency of biguanides to overcome cell and tumor proliferation in experimental studies and the evidence from epidemiological studies have prompted many clinicians to start clinical trials. Metformin can be prescribed by any physician and is safe and freely available. As a result of this excitement and easy access to the drug, more than 100 clinical studies have been launched (see clinicaltrial.gov). Under this unique scenario, some studies have not been as rigorous as other clinical trials using novel compounds. Therefore, when the results of the trials are available, we will need to carefully examine their design. Encouraging results concerning a small number of patients have recently been published. Hosono et al.\textsuperscript{75} have demonstrated that metformin (at the low 250 mg/day dosage for 1 month) decreases the number and size of colorectal aberrant crypt foci (an endoscopic surrogate marker of colorectal cancer) in non-diabetic patients. More recently, in patients with prostate cancer treated with metformin between the day of prognosis and the radical prostatectomy, Joshua et al.\textsuperscript{76} have observed a reduction in Ki67 staining and a significant decrease in the phosphorylation of P-4EBP1, a target of mTORC1. Conversely, the recent results of the first clinical trial performed on more than 100 patients with pancreatic cancer were disappointing. The addition of metformin at a conventional anti-diabetic dose (500 mg twice a day) to classical chemotherapy (erlotinib and gemcitabine) had no advantage for the survival of patients with advanced pancreatic cancers.\textsuperscript{77} Of note, pancreatic tumor cells mainly rely on glycolysis; therefore, metformin may not be a good candidate for this cancer. The coming year should be crucial to conclude whether metformin has a true beneficial effect in cancer.

2-DG

Based on Warburg’s observation, 2-DG was investigated as a monotherapy in the 1950s, with no conclusive results.\textsuperscript{78} Because most cancer cells have normal capacities for using alternative sources of carbons through oxidative phosphorylation, this outcome is not surprising. More recent studies have tried to determine the tolerable and optimal dose of 2-DG for use in patients. Adverse side effects include fatigue, dizziness, restlessness and hypoglycemic symptoms. Depending on the study, the dose of 2-DG administered daily can vary from 45 to 300 mg/kg and is always combined with other anticancer treatments, such as radiotherapy or chemotherapy.\textsuperscript{79–81} Only one of these studies commented on the efficiency of 2-DG as an antitumoral agent. Indeed, Raez et al.\textsuperscript{80} found one patient with partial response after 1 week of 2-DG alone and 12 patients with stable disease among a total of 34 patients. Surprisingly, no other clinical trials using 2-DG are registered on the NIH clinical trial website.

AICAR

Before the discovery of the action of AICAR on AMPK, this drug was described as a very potent cardioprotective agent with a different mechanism of action compared with standard nucleoside analogs, such as fludarabine.\textsuperscript{82} So far, only one clinical trial using AICAR in cancer has been reported in the literature, and few are mentioned on the clinicaltrial.gov website. The reported phase I/II clinical trial was performed in patients with B-cell chronic lymphocytic leukemia, which is the most frequent type of leukemia in the elderly in western countries. The symptoms of the disease arise from a clonal excess of B lymphocytes. Ex vivo, B cells have been found to be very sensitive to the proapoptotic effects of AICAR.\textsuperscript{83} A total of 24 patients were enrolled in the clinical trial, and AICAR had a very safe profile for doses between 50 and 250 mg/kg. Encouraging results were obtained; a reduction in palpable lymph nodes was observed in most patients.\textsuperscript{83} Interestingly, clinical trials combining two of these metabolic disruptors have not been performed. The combination of metformin, an inhibitor of OXPHOS, and 2-DG, the inhibitor of glycolysis, sounds promising. This approach relies on the induction of a major energetic stress targeting the two main ATP-producing pathways. In vitro and in vivo experiments have established the relevance of such a combination\textsuperscript{84,85} with a strong reduction in tumor growth.

CONCLUSION

Although the use of the metabolic disruptors alone is very efficient in isolated cancer cells, the use of these agents as monotherapy is certainly not warranted. Firstly, proposing a treatment with solely
an anti-diabetic drug or a glucose analog or a doping agent to patients with advanced cancer is ethically questionable in a clinical trial. Secondly, the metabolic disruptors significantly improve the antitumoral effect of classical therapies, including chemotherapy and radiation, in the animal experiments. Thirdly, these metabolic drugs act as sensitizers and cause tumor cells to become more susceptible to a second hit.

In the future, better understanding the mode of action of these drugs not only in cancer cells but also on stromal cells will be important to overcome tumor heterogeneity and improve the efficiency of metabolic disruptors. Most laboratory in vitro studies use high concentrations of metformin, 2-DG and AICAR that are not compatible with the concentrations found in serum. Derivatives that improve the pharmacokinetics of these drugs are urgently needed. Metformin is highly polar and requires specific transporters to enter the cells, namely organic cation transporters. The responsiveness of tumor cells to metformin greatly depends on the expression of these organic cation transporters. In addition, Birsoy et al. have found that cell lines sensitive to low glucose are also defective in OXPHOS and that this trait confers sensitivity to biguanides. Thus, the expression of organic cation transporters and the metabolic characteristics of the cancer cells should be investigated and known prior to treatment with biguanides.

The tumor microenvironment plays a central role in the aggressiveness of tumors. The metabolic disruptors have been shown to interfere with stromal cell metabolism. For example, Beneteau et al. have elegantly demonstrated that 2-DG in combination with etoposide promotes an antitumoral immune response. In this study, only tumor cells treated with both agents ‘vaccinated’ mice against tumor growth. More recently, metformin has been shown to increase CD8(+) tumor-infiltrating lymphocytes, which promote tumor rejection. Naïve CD8(+) T lymphocytes treated with metformin migrate into the tumor and exert an antitumoral effect following adoptive transfer. A better understanding of the global effects of metabolic disruptors is needed, and the results from the ongoing clinical trials will certainly orientate the research in the field.

CONFLICT OF INTEREST
The authors declare no conflict of interest.

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