# **Energy transfer in "parasitic" cancer metabolism** Mitochondria are the powerhouse and Achilles' heel of tumor cells

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It is now widely recognized that the tumor microenvironment promotes **tumor microenvironment promotes cancer cell growth and metastasis via changes in cytokine secretion and extracellular matrix remodeling. However, the role of tumor stromal cells in providing energy for epithelial cancer cell growth is a newly emerging paradigm. For example, we and others have recently proposed that tumor growth and metastasis is related to an energy imbalance. Host cells produce energy-rich nutrients via catabolism (through autophagy, mitophagy and aerobic glycolysis), which are then transferred to cancer cells, to fuel anabolic tumor growth. Stromal cell derived L-lactate is taken up by cancer cells and is used for mitochondrial oxidative phosphorylation (OXPHOS), to produce ATP efficiently. However, "parasitic" energy transfer may be a more generalized mechanism in cancer biology than previously appreciated. Two recent papers in** *Science* **and** *Nature Medicine* **now show that lipolysis in host tissues also fuels tumor growth. These studies demonstrate that free fatty acids produced by host cell lipolysis are re-used via** β**-oxidation (**β**-OX) in cancer cell mitochondria. Thus, stromal catabolites (such as lactate, ketones, glutamine and free fatty acids) promote tumor growth by acting as high-energy onco-metabolites. As such, host catabolism via autophagy, mitophagy and lipolysis may explain** 

**the pathogenesis of cancer-associated cachexia and provides exciting new druggable targets for novel therapeutic interventions. Taken together, these findings also suggest that tumor cells promote their own growth and survival by behaving as a "parasitic organism." Hence, we propose the term "parasitic cancer metabolism" to describe this type of metabolic-coupling in tumors. Targeting tumor cell mitochondria (OXPHOS and** β**-OX) would effectively uncouple tumor cells from their hosts, leading to their acute starvation. In this context, we discuss new evidence that high-energy onco-metabolites (produced by the stroma) can confer drug resistance. Importantly, this metabolic chemo-resistance is reversed by blocking OXPHOS in cancer cell mitochondria, with drugs like Metformin, a mitochondrial "poison." In summary, parasitic cancer metabolism is achieved architecturally by dividing tumor tissue into at least two well-defined opposing "metabolic compartments:" catabolic and anabolic.**

### **Introduction**

The concept that tumor-host interactions are crucial in tumor progression is now well-accepted.<sup>1,2</sup> In fact, Stephen Paget<sup>3</sup> first proposed the "seed and soil" hypothesis in 1889, which states that cancer cells ("the seeds") metastasize systemically and

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**Figure 1.** Energy transfer in cancer metabolism. The tumor stroma generates catabolites that are transferred to cancer cells for anabolic growth. The stroma has high levels of autophagy, mitophagy, glycolysis and lipolysis, while epithelial cancer cells have high mitochondrial mass and activity (oxidative phosphorylation and β oxidation). Reproduced and modified with permission from reference 14.

grow best in the most tumor-promoting host organs or the most "fertile soil."4-6 In this context, tumor cells corrupt or transform their microenvironment in order to generate new blood vessels<sup>7,8</sup> to support their oxygen requirements.<sup>9,10</sup> Closely linked to this idea, we and others have begun to view cancer as a "parasitic disease" that "steals" energy-rich metabolites from the host microenvironment<sup>11-13</sup> (**Fig. 1**).

The tumor stroma, which is composed of fibroblasts, adipocytes, endothelial cells and macrophages, lies in extremely close proximity to cancer cells, and can directly promote tumor growth.15-20 Most research on the mechanism(s) by which the stroma promotes tumor growth has focused on changes in the extracellular matrix and the increased secretion of tumor promoting cytokines, such as IL-6, IL-8, SDF1, VEGF and TGFβ.<sup>20,21</sup> However, little is known regarding the metabolic properties of the tumor stroma. Several independent studies have recently highlighted the importance of metabolic coupling between cancer cells and other host cells to drive a form of "parasitic cancer metabolism." This metabolic-coupling promotes epithelial tumor cell growth and metastasis.

## **Cancer Cells Behave as Metabolic Parasites: Glycolysis in Fibroblasts Promotes Mitochondrial Metabolism in Cancer Cells, Fueling Tumor Growth**

Malaria infection is due to an intracellular parasite (Plasmodium) that induces oxidative stress in host cells,<sup>22</sup> resulting in the onset of host cell autophagy,<sup>23</sup> which supplies the parasites with highenergy metabolites and chemical building blocks. *T. cruzi* (Chagas disease), another intracellular parasite, also uses the same mechanism(s) involving oxidative stress<sup>24,25</sup> and host cell autophagy.<sup>26-28</sup> In addition, *T. cruzi* induces lipolysis of triglycerides in adipocytes, to generate free fatty acids that it can use as a fuel supply.<sup>29</sup>

Similarly, our group's recent results show that epithelial cancer cells are extracellular parasites that induce oxidative stress in adjacent stromal fibroblasts, by secreting hydrogen peroxide.<sup>11,30</sup> This oxidative stress causes stromal fibroblast activation, with the upregulation of HIF1- $\alpha$ activity driving autophagy, mitophagy and aerobic glycolysis in the tumor stroma.31-33 In parallel, oxidative stress also activates NFκB, further accelerating autophagy

and the local secretion of inflammatory cytokines from activated fibroblasts. Thus, oxidative stress, autophagy, aerobic glycolysis and inflammation are inextricably linked in the tumor stroma.<sup>30,31,34-37</sup> As such, the stroma provides catabolized nutrients to "fuel" the anabolic growth of tumor cells by enhancing their mitochondrial activity.30 L-lactate derived from glycolytic fibroblasts is transferred to cancer cells and is used to generate energy via oxidative mitochondrial metabolism (OXPHOS). Similarly, ketone bodies and glutamine derived from host cell catabolism can also fuel the mitochondrial activity of adjacent epithelial cancer cells<sup>11,38,39</sup> (**Fig. 1**). We have termed this new form of parasitic cancer metabolism the "reverse Warburg effect" (since increased glycolysis occurs in fibroblasts rather than tumor cells) or the "auotphagic tumor stroma model of cancer" (since tumor cells induce autophagy and mitophagy in adjacent fibroblasts).40-43 (For recent reviews on the conventional "Warburg effect," please see refs. 44–49).

Thus, stromal metabolites such as L-lactate, ketones and glutamine promote tumor growth by acting as high-energy onco-metabolites (Fig. 2).<sup>50</sup> As such, intracellular parasites and cancer cells use similar metabolic mechanism(s) for survival. These mechanistic insights have important implications for the design of novel therapeutic interventions for cancer treatment and prevention. For example, chloroquine is an effective anti-malarial drug (and also has anticancer activity), because it inhibits autophagy and cuts off "the fuel supply," by preventing energy transfer from host to parasite.<sup>11,42</sup>

Although this model of "parasitic metabolism" by tumors has only recently been proposed, energy transfer between cells to fuel growth is in fact not a new invention, but instead reflects the co-optation of a normal physiological process by tumor cells. Metabolic-coupling already exists physiologically in many different organ systems, such as in skeletal muscle, the brain and the ovary (**Fig. 3**). In skeletal muscle, fast-twitch fibers are glycolytic, and slow-twitch fibers are oxidative. Fasttwitch fibers produce L-lactate, which is transferred to slow-twitch fibers to fuel oxidative mitochondrial metabolism.<sup>51</sup>

This process is known as the "lactate shuttle." In the brain, a similar lactate shuttle exists. In this context, astrocytes are glycolytic and neurons are oxidative; this energy-sharing symbiosis has been termed "neuron-glia metabolic coupling."52,53 Finally, in the ovary, the granulosa cells are glycolytic and the oocyte is oxidative, and this metabolic coupling has been exploited for in vitro fertilization (IVF), to maintain viable oocytes.<sup>54,55</sup> Metaboliccoupling also exist between organ systems, such as in the muscle-kidney or muscleliver glutamine shuttles.<sup>56</sup> Therefore, metabolic coupling is a widely used normal physiologic process that has been adopted by epithelial tumor cells, to promote their own growth and survival of the tumor as a "parasitic organism."

In further support of the idea that a "lactate shuttle" also exists in human tumors, we have shown that the distribution of lactate transporters is highly compartmentalized in human breast cancers (**Fig. 4**). MCT4, which functions in the extrusion of L-lactate and is a marker of oxidative stress and aerobic glycolysis, is largely confined to cancer-associated fibroblasts in the tumor stroma.<sup>14,57</sup> In contrast, MCT1, which drives L-lactate uptake is specifically localized to epithelial cancer cells.57 Importantly, MCT transporters have a broad specificity and can also function in the shuttling of ketone bodies from the tumor stroma to epithelial cancer cells.

Two of our most recent papers published in the journal *Cell Cycle* directly demonstrate that human breast cancer cells use OXPHOS, mitochondria and parasitic metabolic-coupling.14,58

Since we believe that both skeletal muscle and human tumors are constructed on the same metabolic principles (with adjacent glycolytic and oxidative compartments), we subjected tumors to a specific histochemical stain that has been used for over 50 years to visualize these two compartments in muscle.<sup>14</sup> This histochemical stain allows detection of the functional activity of mitochondrial complex IV [cytochrome *C* oxidase (COX)], a key component of the respiratory chain and OXPHOS.14 In skeletal muscle, fasttwitch fibers are glycolytic and are COXnegative. In contrast, slow-twitch fibers are



**Figure 2.** Onco-metabolites derived from the tumor stroma promote anabolic cancer cell growth via the TCA cycle and oxidative mitochondrial metabolism. Note that various stromally derived onco-metabolites (L-lactate, ketones, free fatty acids and glutamine; shown in red) all feed into the TCA/Krebs cycle via either Acetyl-CoA or Alpha-Keto-Glutarate, promoting oxidative mitochondrial metabolism (OXPHOS) in epithelial cancer cells. The end result is highly efficient ATP production in aggressive cancer cells.





oxidative and are COX-positive. **Figure 5** (left part) shows the distribution of these two metabolic fiber types.

Similar results were obtained in human breast cancers.14 The tumor stroma is largely COX-negative and hence more glycolytic. Conversely, breast cancer epithelial cell "nests" are COX-positive and are therefore more oxidative.<sup>14</sup> A representative image is shown in **Figure 5** (right part). These findings provide striking functional data to support the idea



**Figure 4.** Shuttling the onco-metabolite L-lactate from the tumor stroma to epithelial cancer cells. Previously, we proposed that a "lactate shuttle" also exists in human breast cancers. More specifically, the distribution of lactate transporters is highly compartmentalized in human breast cancers. Note that MCT4 (a marker of aerobic glycolysis and L-lactate secretion) is largely confined to cancer-associated fibroblasts in the tumor stroma. Conversely, MCT1 (a marker of L-lactate uptake) is localized to epithelial cancer cells. Because of their broad specificity, the same MCT transporters can also function in the shuttling of ketone bodies from the tumor stroma to epithelial cancer cells. Arrows indicate the direction of energy transfer. Reproduced and modified with permission from reference 57.



**Figure 5.** Metabolic compartmentalization of mitochondrial activity in skeletal muscle and human breast cancer. Frozen sections from skeletal muscle tissue or human breast cancers were subjected to a routine histochemical stain that detects the functional activity of mitochondrial complex IV [cytochrome *C* oxidase (COX)]. This allows the visualization of oxidative mitochondrial metabolism in tissue sections. COX-positive cells are positively stained brown (see red arrows). In skeletal muscle, note that fast-twitch fibers are glycolytic (Gly) and are COX-negative, while slow-twitch fibers are oxidative (Ox) and are COX-positive. In breast cancers, the tumor stroma is glycolytic (Gly) and is COX-negative, while epithelial tumor cell nests are oxidative (Ox) and are COX-positive. These results support the idea that tumors show metabolic compartmentalization and specialization, as occurs in skeletal muscle tissue. Reproduced and modified with permission from reference 14.

that tumors use metabolic-coupling and shuttle L-lactate, resulting in a net energy transfer from the tumor stroma to epithelial cancer cells.

Furthermore, using a bioinformatics approach, we demonstrated that epithelial tumor cells laser captured from primary human breast cancers had upregulated mitochondrial genes not seen in tumor fibroblasts. This cancer cell mitochondrial gene signature consisted largely of OXPHOS-related gene transcripts that were upregulated (> 4-fold) in epithelial cancer cells, compared with adjacent stromal tissue. Using this OXPHOS-signature, we showed that this type of oxidative mitochondrial metabolism is significantly upregulated in most human breast cancers ( $> 2,000$  patients examined;  $p < 10^{-20}$ ), in both  $ER(+)$  and  $ER(-)$  tumor subtypes.<sup>14</sup> In ER(-) patients, this MITO/OXPHOS signature was specifically associated with tumor metastasis.<sup>14</sup>

In addition, we have used an shRNA approach to genetically create glycolytic fibroblasts, by knocking-down the major mitochondrial transcription factor, namely, TFAM.58 As predicted, these TFAM-deficient fibroblasts underwent metabolic re-programming toward a more glycolytic state, with loss of respiratory chain components as well as increased hydrogen peroxide and L-lactate production.58 These glycolytic fibroblasts also significantly promoted tumor growth using a human tumor xenograft model, employing MDA-MB-231 breast cancer cells.58

## **Adipocytes Provide Fatty Acids to Ovarian Cancer Cells to Fuel Mitochondrial** b**-Oxidation and Promote Tumor Growth**

A new study by Nieman et al. published in *Nature Medicine* has now evaluated the growth-promoting properties of the omentum in metastatic ovarian cancer from a metabolic perspective.<sup>59</sup> They show that the proximity of adipocytes to ovarian cancer cells leads to lipolysis in adipocytes, with the release of free fatty acids.<sup>59</sup> Ovarian cancer cells adjacent to these transformed or activated adipocytes show increased fatty acid uptake and utilization via β-oxidation in

mitochondria,<sup>59</sup> demonstrating that metabolic-coupling also occurs between these two cellular compartments (summarized in **Figs. 1 and 2**).

The lipolytic enzyme, hormonesensitive lipase (HSL) and the fatty acid transport protein, FABP4, were found to be upregulated in adipocytes in omental metastases as compared with primary ovarian cancers.<sup>59</sup> FABP4(-/-) mice injected intraperitoneally with ovarian cancer cells, had a significant reduction in tumor burden as compared with wild-type mice.<sup>59</sup> This highlights, for the first time, the critical pro-tumorigenic importance of metabolic-coupling between adipocytes and ovarian cancer cells.

The investigators also demonstrated in vivo and in vitro that homing, migration and invasion of ovarian cancer cells is induced by omental adipocytes via IL-8, IL-6, MCP-1 and TIMP-1, since neutralizing antibodies led to decreased homing.<sup>59</sup> This was confirmed in vivo for IL-8 by using an IL-8 receptor inhibitor. Inflammatory cytokines such as IL-6 and IL-8 are involved in promoting catabolic processes, including lipolysis and glycolysis.60-63 In addition, tumor cells can induce the production of inflammatory cytokines, including IL-6 and IL-8, and transform the stroma to a highly catabolic state with increased autophagy.<sup>35</sup> These studies once again stress the link between inflammation and autophagybased catabolism in the stroma and tumor progression.

In summary, metabolic coupling between adipocytes and cancer cells favors tumor growth, and this can be reversed by inhibiting lipid transport from adipocytes to carcinoma cells.<sup>59</sup>

# **Adipocyte Lipolysis Promotes Cancer-Associated Cachexia and Leads to a Generalized Catabolic State, Driving Tumor Growth**

Cancer-associated cachexia (CAC) is a well-known complication of cancer that frequently is the cause of death in cancer patients. CAC leads to abnormal whole body catabolic metabolism, similar to starvation.<sup>64</sup> The mechanism(s) underlying tumor-associated cachexia are poorly



**Figure 6.** Uncoupling parasitic cancer metabolism. Drugs such as chloroquine (which inhibits autophagy) and metformin (which inhibits lipolysis), will prevent energy transfer to cancer cells and tumor growth. In this scenario, mitochondrial poisons (such as metformin, arsenic and others) could also be used to uncouple tumor cells from the energy-producing host stroma.

understood, and no widely effective strategies, including increased calorie intake, exist to manage the condition.<sup>65</sup>

The study by Das et al. recently published in the journal *Science* presents important new data regarding CAC. By studying the effects of adipose tissue metabolism on muscle and fat mass,<sup>66</sup> the investigators discovered that inhibition of lipolysis halts muscle and fat-related cachexia, as observed in lipase-knockout mice. Triglycerides are metabolized to fatty acids and glycerol by adipose triglyceride lipase (ATGL) and hormone-sensitive lipase (HSL). Inhibition of ATGL and, to a lesser degree, of HSL, prevented cachexia and weight loss.<sup>66</sup> In fact, ATGLknockout mice had similar weights as control mice that were not injected with cancer cells.<sup>66</sup> Also, by studying white adipose tissue (WAT) in cancer patients, they found that ATGL activity was highest and WAT mass was the lowest in patients who died with CAC, as compared with other cancer patients.<sup>66</sup> These data directly show that metabolic-coupling exists between different organ systems in cancer patients. It is well-known that chemokines can alter the metabolism of different organs, but this study goes on to show that the metabolites themselves act as chemokines, inducing changes in the metabolism of other organs.<sup>66</sup>

# **Clinical Implications of Metabolic Coupling for Cancer Treatment**

The discovery of metabolic-coupling between tumor cells and their hosts opens many new avenues for drug discovery involving novel therapeutic targets. Thus, like diabetes, cancer should be viewed as a systemic metabolic disease of energy imbalance. In fact, similar metabolic profiles (with insulin resistance and increased lipolysis) are found in patients with diabetes mellitus and those with advanced cancers and cachexia.67-69 Anti-diabetic drugs such as metformin, which inhibits both lipolysis and mitochondrial oxidative phosphorylation,70,71 need to be further investigated for the effective management of cancer patients with "parasitic metabolism" or "metabolic coupling," between cancer cells and the host (**Fig. 6**).

Metformin is a known "mitochondrial poison" that mechanistically functions as a complex I inhibitor that blocks mitochondrial oxidative phosphorylation. As a consequence, Metformin also activates AMP kinase<sup>72</sup> and increases insulinsensitivity, thus decreasing glucose blood levels,73 which would be conceptually beneficial in treating cancers. Epidemiologic studies support the use of metformin in cancer treatment and suggest a protective



**Figure 7.** Visualizing the anti-mitochondrial effects of Metformin in human breast cancer tumor tissue. Frozen sections from human breast cancers were subjected to mitochondrial Complex I (NADH) activity staining. NADH-positive cells are positively stained blue (see red arrows). In breast cancers, note that the tumor stroma is glycolytic and is NADH-negative, while epithelial tumor cell nests are oxidative and are NADH-positive. Note that treatment with Metformin, a known complex I inhibitor, prevented the NADH-staining of epithelial cancer cell nests. Positive controls with skeletal muscle were performed in parallel to ensure the specificity of the staining procedure and allowed us to detect complex I-positive muscle fibers (blue), which represent oxidative slowtwitch fibers. Reproduced and modified with permission from reference 14.



**Figure 8.** Stromal onco-metabolites confer drug resistance by promoting mitochondrial health or well-being, providing an escape from stress-induced apoptosis. Simple stromally derived metabolites (such as L-lactate, ketones and glutamine) promote mitochondrial "health" in cancer cells, effectively shutting off their apoptotic machinery, resulting in protection against cell death, even when challenged with anticancer drugs. In contrast, we can overcome metabolite-induced drug resistance in cancer cells by using mitochondrial poisons (such as Metformin).

effect against cancer development<sup>74</sup> and cancer progression in diabetics receiving metformin.75 Thus, the anticancer activity of metformin may stem from its antimitochondrial activity, thereby preventing cancer cells from using the energy-rich onco-metabolites (L-lacate, ketones,

glutamine and fatty acids) derived from the tumor stroma.

Recently, we used a mitochondrial complex I activity stain to functionally visualize the anti-mitochondrial activity of Metformin in human breast cancer tumor samples.14 As with complex IV

staining, complex I mitochondrial activity was mainly localized to the epithelial cancer cell compartment.<sup>14</sup> Strikingly, tumor tissue frozen sections treated with Metformin revealed a near complete absence of Complex I activity, $14$  consistent with the idea that Metformin does indeed function as a Complex I inhibitor in vivo (**Fig. 7**).

#### **Reversing Drug Resistance: Targeting Cancer Cell Mitochondria for Destruction**

Importantly, stromal-epithelial metabolic-coupling also induces chemo-resistance.76,77 Thus, resistance to standard anticancer therapies may be both a metabolic and stromal phenomenon, related to mitochondrial "health" or "well-being" in cancer cells.76,77

Mitochondria fulfill a dual role in cell metabolism. During anabolic cell growth, they are the "powerhouse" of the cell and fuel proliferation. However, during periods of stress and cell death (apoptosis), they act as "sensors" to decide whether a given cell will undergo suicide or programmed cell death. In this context, the mitochondria are also the cell's executioners. Most chemotherapeutic agents act via the induction of apoptosis in cancer cells, placing mitochondria at the center stage of both cancer cell metabolism and drug resistance (**Fig. 8**).

In this context, we have shown that energy-rich metabolites derived from stromal cells maintain mitochondrial "wellbeing" in cancer cells, thereby conferring drug resistance.76,77 More specifically, simple metabolites (such as L-lactate, ketones and glutamine) promote mitochondrial "health" in cancer cells, effectively shutting off their apoptotic machinery, resulting in protection against cell death, even when challenged with toxic drugs.<sup>76,77</sup> Similarly, we can overcome metabolite-induced chemo-resistance in cancer cells by using mitochondrial poisons (such as Metformin or Arsenic) or by using drug combinations that function as mitochondrial poisons (Tamoxifen + Dastatinib).76,77 Conversely, we can genetically pheno-copy the protective effects of these onco-metabolites by overexpressing recombinant proteins that maintain mitochondrial "health," such as TIGAR,78,79 directly conferring chemoresistance in cancer cells.76,77

Thus, metabolic-coupling promotes tumor growth and makes epithelial cancer cells more resistant to conventional therapies. As such, drugs that can target mitochondrial function in cancer cells or drugs that halt glycolysis, lipolysis or catabolism in the surrounding tumor stroma may be beneficial in preventing tumor progression and metastasis.

Ultimately, this new concept of "parasitic cancer metabolism" (**Fig. 9**) could radically change how we treat cancer patients and stimulate new metabolic strategies for cancer prevention and therapy.13,43,50,81

#### **Note Added In Proof**

After this Perspective was accepted for publication in *Cell Cycle*, another paper appeared in the journal *Science*, showing that baseline mitochondrial dysfunction in cancer cells is a positive predictor of the therapeutic response to conventional chemotherapy. Thus, as we proposed based on our experiments, mitochondrial dysfunction in cancer cells sensitizes them to chemotherapy, resulting in a good clinical response and increased overall survival in cancer patients. As such, mitochondrial dysfunction (i.e., the conventional Warburg effect) in tumor cells predicts chemo-sensitivity in cancer patients, whereas mitochondrial "health and wellbeing" in cancer cells (i.e., indicative of "Parasitic Cancer Metabolism") predicts chemo-resistance. These clinical findings directly support our new paradigm, which is based on (1) mitochondrial function and (2) energy transfer in cancer metabolism.

For further details, please see: Chonghaile TN, Sarosiek KA, Vo TT, Ryan JA, Tammareddi A, Moore VD, et al. Pretreatment mitochondrial priming correlates with clinical response to cytotoxic chemotherapy. Science 2011; 334:1129-33; PMID:22033517.

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**Figure 9.** Metabolic compartments in parasitic cancer metabolism. In summary, we believe that cancer cells act as metabolic parasites and extract nutrients from host cells by inducing catabolic processes (autophagy, mitophagy, aerobic glycolysis and lipolysis). As a consequence, the tumor stroma shows a shift toward aerobic glycolysis, and epithelial cancer cells show functional hyperactivation of oxidative mitochondrial metabolism (OXPHOS). In support of this model, cancerassociated fibroblasts and the tumor stroma overexpress PKM2 (a rate-limiting glycolytic enzyme, left panel). Conversely, breast cancer epithelial cells upregulate MT-CO1 (a key component of mitochondrial complex IV, right panel). The metabolic compartmentalization of PKM2 and MT-CO1 were visualized by immunostaining with specific antibody probes (brown reaction product). Reproduced and modified with permission from references 14 and 80.

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