# Mitochondrial biogenesis in epithelial cancer cells promotes breast cancer tumor growth and confers autophagy resistance

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Here, we set out to test the novel hypothesis that increased mitochondrial biogenesis in epithelial cancer cells would "fuel" enhanced tumor growth. For this purpose, we generated MDA-MB-231 cells (a triple-negative human breast cancer cell line) overexpressing PGC-1 $\alpha$  and MitoNEET, which are established molecules that drive mitochondrial biogenesis and increased mitochondrial oxidative phosphorylation (OXPHOS). Interestingly, both PGC-1 $\alpha$  and MitoNEET increased the abundance of OXPHOS protein complexes, conferred autophagy resistance under conditions of starvation and increased tumor growth by up to ~3-fold. However, this increase in tumor growth was independent of neo-angiogenesis, as assessed by immunostaining and quantitation of vessel density using CD31 antibodies. Quantitatively similar increases in tumor growth were also observed by overexpression of PGC-1 $\beta$  and POLRMT in MDA-MB-231 cells, which are also responsible for mediating increased mitochondrial biogenesis. Thus, we propose that increased mitochondrial "power" in epithelial cancer cells oncogenically promotes tumor growth by conferring autophagy resistance. As such, PGC-1 $\alpha$ , PGC-1 $\beta$ , mitoNEET and POLRMT should all be considered as tumor promoters or "metabolic oncogenes." Our results are consistent with numerous previous clinical studies showing that metformin (a weak mitochondrial "poison") prevents the onset of nearly all types of human cancers in diabetic patients. Therefore, metformin (a complex I inhibitor) and other mitochondrial inhibitors should be developed as novel anticancer therapies, targeting mitochondrial metabolism in cancer cells.

# Introduction

Mitochondria are not only the powerhouse of the cell, but they also play a critical role in the metabolism of proliferating cells.<sup>1</sup> Despite these known function(s) in cell growth and division, the role of mitochondria in carcinogenesis has been largely ignored for many years because of the "Warburg hypothesis."

Warburg's hypothesis, also known as "aerobic glycolysis," postulates that tumor cells shut down their oxidative mitochondrial metabolism and depend only on glycolysis, even under oxygenrich conditions.<sup>2</sup> This hypothesis discounts the fact that glycolysis produces a very low-yield of ATP, while cancer cells have high-energy demands for ATP. This implies that cancer cells cannot really depend solely on the glycolytic pathway to produce their energy. Utilizing the end products of glycolysis via oxidative mitochondrial metabolism [the TCA/citric acid cycle and oxidative phosphorylation (OXPHOS)] would be a much more efficient method of providing energy to cancer cells, which are in desperate need of ATP molecules.

Recently, we have directly shown that that oxidative mitochondrial activity is fact induced in cancer cells when they are co-cultured with fibroblasts.<sup>3-5</sup> As a consequence, we have proposed a new hypothesis, termed "two-compartment tumor metabolism," to resolve the Warburg paradox.<sup>3-5</sup> Our previous studies also indicated that cancer cells are able to recruit stromal cells by releasing reactive oxygen species (ROS) into their local microenvironment. As a result, cancer-associated fibroblasts (CAFs) activate autophagic signaling pathways in order to recycle their proteins and organelles, such as mitochondria (mitophagy). Without functional mitochondria, CAFs become glycolytic and secrete L-lactate and ketone bodies, which are then taken-up by nearby cancer cells.<sup>6</sup> Subsequently, these energy-rich nutrients are "burned" via oxidative mitochondrial metabolism in the cancer cells to produce energy and anabolic products.

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REPORT

Beclin-1

**Cathepsin B** 



EV

PGC-1α MitoNEET

Another important molecule for mitochondrial function and biogenesis is mitochondrial RNA polymerase, POLMRT.<sup>12</sup> POLRMT is responsible for the transcription of the 13 subunits of the OXPHOS complexes and acts as a RNA primase for mitochondrial DNA replication.<sup>13,14</sup> POLRMT requires transcription factor B2M (TFB2M) and transcription factor AM (TFAM) for mitochondrial RNA transcription.<sup>14</sup>

Here, we examined the role of mitochondrial biogenesis and function in breast cancer tumor growth. Using a preclinical animal model, we determined how the overexpression of mitoNEET, PGC-1 $\alpha$ , PGC-1 $\beta$  and POLMRT in epithelial breast cancer cells (MDA-MB-231) affects tumor growth. Remarkably, overexpression of these key molecules that drive mitochondrial biogenesis promoted tumor growth in nude mice. However, this increased growth was clearly independent of tumor vascularization. Our results demonstrate that mitochondrial biogenesis in epithelial cancer cells promotes breast cancer tumor growth, likely via increased energy production and nutrient reserves, which confers autophagy resistance.

Thus, we have identified oxidative mitochondrial metabolism as a new "druggable" therapeutic target for the development of novel anticancer agents.

In addition, our findings directly support clinical studies demonstrating that metformin (a mitochondrial "poison" and complex I inhibitor) prevents the onset of nearly all types of human cancers in diabetic patients (reviewed within ref. 15). As a consequence, metformin, and other mitochondrial inhibitors, should be used to target mitochondrial metabolism in epithelial cancer cells.

## Results

Mitochondrial OXPHOS is increased in breast cancer cells during overexpression of PGC-1 $\alpha$  and mitoNEET. Recently,



**Figure 1.** Overexpression of PGC-1 $\alpha$  and MitoNEET induces the expression of mitochondrial OXPHOS complexes. (**A**) Using a lentiviral vector, we established MDA-MB-231 cell lines that stably overexpress MitoNEET, as seen by immunoblotting.  $\beta$ -actin is shown as a control for equal protein loading. (**B**) PGC-1 $\alpha$  and MitoNEET induce expression of mitochondrial OXPHOS complexes. Both PGC-1 $\alpha$  and mitoNEET-over-expressing MDA-MD-231 cells displayed elevated PGC-1 $\alpha$  expression. Most importantly, they also showed an elevation of mitochondrial complexes (OXPHOS).  $\beta$ -actin is shown as a control for equal protein loading. EV, empty vector (Lv105).

MitoNEET is an integral membrane protein component of the outer mitochondrial membrane that was found to be colocalized with pioglitazone.<sup>7</sup> It was named based on its mitochondrial localization and the presence of a unique amino acid sequence Asn-Glu-Glu-Thr (NEET) at its C terminus.<sup>7</sup> Recently, mitoNEET has been demonstrated to be a regulator of mitochondrial respiration.<sup>8</sup> Wiley et al. showed that mitochondria isolated from MitoNEET-null murine hearts displayed lower oxidative capacity.<sup>8</sup>

Furthermore, members of the peroxisome proliferator-activated receptor- $\gamma$  coactivators-1 (PGC-1) family play a central role

Cell Cycle



**Figure 3.** PGC-1 $\alpha$  and MitoNEET overexpression increases tumor growth. PGC-1 $\alpha$  and MitoNEET overexpressing MDA-MB-231 cells were subcutaneously injected into the flanks of nude mice. (**A**) Tumor volumes were measured on days 14, 19 and 21. Note that PGC-1 $\alpha$  and MitoNEET tumors demonstrate a trend of significantly larger volumes, as compared with control tumors. (**B**) Tumors were excised, weighed, and their volumes were measured. PGC-1 $\alpha$  tumor volumes and weights were 3- and 2.7-fold larger, as compared with control tumors, respectively. Also, MitoNEET tumor volumes and weights were 2.9- and 2.25-fold larger, as compared with control tumors, respectively. Also, MitoNEET tumor volumes and weights were 2.9- and 2.25-fold larger, as compared with control tumors, respectively. (**C**) Tumor angiogenesis and vessel quantification (number of vessels per field). Frozen sections were cut and immunostained with anti-CD31 antibodies. Quantification of CD31-positive vessels shows a non-significant reduction in angiogenesis in PGC-1 $\alpha$  tumors, while MitoNEET tumors displayed a significant reduction in CD31-positive vessels. These results indicate that PGC-1 $\alpha$  and MitoNEET stimulate tumor growth, but independently of neo-angiogenesis. EV, empty vector (Lv105).

we have suggested that increased mitochondrial function in cancer cells may directly promote tumor growth.<sup>3</sup> Here, to test this hypothesis more directly, we stably overexpressed PGC-1 $\alpha$  and MitoNEET in MDA-MB-231 cells, a human triple-negative breast cancer cell line. Then, immunoblot analysis was used to confirm the successful overexpression of both proteins (Fig. 1A and B).

PGC-1 $\alpha$  is a master transcription factor that stimulates mitochondrial biogenesis and respiratory function.<sup>8</sup> Moreover, mitoNEET, a mitochondrial outer membrane protein, regulates mitochondrial oxidative phosphorylation.<sup>8</sup> Interestingly, mitoNEET overexpressing MDA-MB-231 cells displayed elevated PGC-1 $\alpha$  expression. As expected, immunoblotting against various OXPHOS components showed that these mitochondrial complexes (I, III, IV and V) were increased in both PGC-1 $\alpha$  and MitoNEET-overexpressing MDA-MB-231 cells (Fig. 1B).

PGC-1 $\alpha$  and mitoNEET confer autophagy resistance in breast cancer cells. Next, we wanted to test if PGC-1 $\alpha$  and mitoNEET will reduce autophagy in cancer cells, as they do in

skeletal muscles.<sup>16</sup> Interestingly, PGC-1 $\alpha$  and mitoNEET both protected cancer cells from autophagy, as Beclin-1 and Cathepsin B (markers of autophagy) were not upregulated under conditions of starvation (Fig. 2).

PGC-1  $\alpha$  and mitoNEET increase tumor growth in vivo, independently of angiogenesis. Traditionally, altered mitochondrial metabolism was considered as a secondary effect of proliferation. Here, we wanted to test if increased mitochondrial biogenesis and OXPHOS could directly promote tumor growth. PGC-1 $\alpha$  and mitoNEET-overexpressing MDA cells were injected in nude mice. Tumor volumes were then measured at various time points post-injection.

Tumors in which PGC-1 $\alpha$  and mitoNEET were overexpressed were significantly larger, as compared with control tumors (Fig. 3A). At 3 wk post-injection, PGC-1 $\alpha$  tumor volumes and weights were on average approximately 3-fold larger in than control tumors, which harbored the expression vector alone. Moreover, MitoNEET tumor volumes and weights were 2.9- and 2.25-fold larger than control tumors, respectively (Fig. 3B).



**Figure 4.** Overexpression of PGC-1 $\beta$  and POLRMT. Immunoblot analysis shows successful overexpression of PGC-1 $\beta$  and POLRMT in transduced MDA-MB-231 cells.  $\beta$ -actin is shown as a control for equal protein loading. EV, empty vector (Lv105).

To examine the possibility that the observed increase in tumor growth was due to increased angiogenesis, we immunostained tumor sections with CD31, a well-established marker used to quantitate vessel density. Interestingly, PGC-1 $\alpha$  tumors did not show any significant differences in vessel number, while mitoNEET tumors displayed decreased tumor vascularization (Fig. 3C). These results imply that the increased tumor growth caused by PGC-1 $\alpha$  and/or mitoNEET overexpression is independent of tumor vascularization.

PGC-1 $\beta$  and POLRMT also promote breast cancer tumor growth. Given the growth-promoting effects of PGC-1 $\alpha$  and mitoNEET, we decided to also study the functional effects of PGC-1 $\beta$  and POLRMT, a mitochondrial RNA polymerase,<sup>17</sup> on breast cancer tumor growth.

Using lentiviral vectors, MDA-MB-231 cells were stably transduced to overexpress either PGC-1 $\beta$  or POLRMT. Successful overexpression was evident by immunoblotting for total PGC-1 $\beta$ or POLRMT (Fig. 4). To test if overexpression of PGC-1 $\beta$  or POLRMT increases tumor growth, we subcutaneously injected PGC-1 $\beta$  or POLRMT-overexpressing MDA-MB-231 cells into the flanks of nude mice. The tumor growth curves for PGC-1 $\beta$ and POLRMT show a significant increase in tumor size (Fig. 5A). At 3 wk post-injection, PGC-1 $\beta$  and POLRMT tumor volumes were 2-fold larger than control tumors, which harbored the vector alone (Fig. 5B).

Finally, we examined if this observed increase in tumor growth is related to increased tumor angiogenesis. In fact, these tissue sections stained for CD31 displayed a significant decrease in vascularization in PGC-1 $\beta$  and POLRMT tumors (Fig. 5C). Thus, these data show that enhanced tumor growth induced by PGC-1 $\beta$  and POLRMT is not due to increases in neo-vascularization.

## Discussion

Traditionally, the "Warburg effect" has been used to describe cancer metabolism, whereby cancer cells become more dependent on glycolysis and less dependent on OXPHOS to produce their ATP.<sup>2</sup> However, this theory does not resolve the issue that cancer cells could not possibly depend solely on glycolysis, which has an extremely low yield of ATP. Recently, we and other researchers have highlighted the crucial role of mitochondrial metabolism in tumorgenesis.<sup>18-21</sup> In this study, we demonstrate the effect of elevated mitochondrial biogenesis and metabolism on tumor growth, using pre-clinical animal models. This was achieved by stably transducing MDA-MB-231 breast cancer cells to overexpress several positive regulators of mitochondrial biogenesis, including PGC-1 $\alpha$ , PGC-1 $\beta$ , mitoNEET and POLRMT. Interestingly, PGC-1 $\alpha$ , PGC-1 $\beta$ , mitoNEET and POLRMT overexpression all significantly increased tumor growth in nude mouse xenografts (Fig. 6).

PGC-1 $\alpha$  and PGC-1 $\beta$  are transcriptional co-activators that have a central role in maintaining metabolic pathways, such as oxidative metabolism, energy homeostasis and glucose and lipid metabolism in normal cells.<sup>22</sup> Recently, PGC-1 $\beta$  knockdown in tamoxifen-resistant MCF-7 cells resulted in a cell line that was sensitive to tamoxifen-treatment.<sup>23</sup> Similarly, Shiota et al. showed that silencing of PGC-1 $\alpha$  inhibited LNCaP prostate cancer cell proliferation.<sup>24</sup>

It is worth pointing out that PGC-1 family members induce superoxide dismutase (SOD), catalase (CAT) and glutathioneperoxidase (GPX), all critical protein antioxidants.<sup>25,26</sup> This action protects cancer cells against elevated mitochondrial ROS. Alternatively, PGC-1 $\alpha$  has been shown to inhibit ROS by inducing uncoupling proteins, which leak protons across the inner membrane of mitochondria to lower their membrane potential, eventually decreasing ROS produced by the mitochondria. Interestingly, this antioxidant role can protect cancer cells from self-released ROS, allowing them to activate neighboring CAFs via oxidative stress.

Interestingly, type 2 diabetic patients suffer from oxidative stress due to reduced mitochondrial mass and oxidative capacity in skeletal muscle.<sup>27-29</sup> Pioglitazone, a thiazolidinedione member that is used as an insulin-sensitizing drug for type 2 diabetes, improves mitochondrial respiration in type 2 diabetic patients.<sup>29</sup> Classically, thiazolidinedione family members are thought to be agonists for peroxisomal proliferator-activated receptor (PPAR)- $\gamma$  expression, which, in turn, induces the expression of PGC-1 $\alpha$ .<sup>30</sup>

Recently, pioglitazone was found to co-localize with an ironcontaining outer mitochondrial membrane protein, mitoNEET.<sup>7</sup> It is possible that pioglitazone induces mitochondrial respiration through mitoNEET binding, as mitoNEET-null mice demonstrated lower mitochondrial capacity.<sup>8,29</sup> Accordingly, our results demonstrate that mitoNEET-overexpressing cancer cells exhibited an increase in mitochondrial complexes I, III, IV, V and a decrease in autophagy. Strikingly, mitoNEET tumors were ~3-fold larger than control tumors. Thus, we established a novel role for mitoNEET in promoting tumor growth. These results provide evidence that mitochondrial metabolism and OXPHOS are driving factors for cancer growth and are not a side effect of increased cellular proliferation.

Moreover, POLMRT is a mitochondrial bacteriophage-related RNA polymerase that is needed for the transcription and replication of mtDNA.<sup>31,32</sup> POLRMT is responsible for transcription

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OXPHOS subunits. of several Interestingly, nuclear factor NRF-1, which is regulated by PGC-1 family members, stimulates POLRMT to increase mitochondrial replication.14 Moreover, POLRMT depletion affects mtDNA replication.32 Our results demonstrate that POLRMToverexpressing cancer cells drive a 2-fold increase in tumor growth in nude mice. This growth was independent of tumor angiogenesis. Thus, our current studies are consistent with a new role for POLRMT as a tumor promoter.

Finally, we observed that overexpression of PGC-1 $\alpha$  and mitoNEET, which induce mitochondrial biogenesis, conferred autophagy resistance under conditions of starvation. In addition, overexpression of PGC-1a and mitoNEET increased tumor growth by ~3-fold. Similarly, loss of Beclin-1, a required autophagy gene and tumor suppressor, is known to confer autophagy resistance in cancer cells and increase tumor growth.33-35 Thus, autophagy resistance in cancer cells—mediated by either (1) increased mitochondrial power or (2) genetic loss of required autophagy genes-has the same net effect, to increase tumor growth rates.

In conclusion, we studied several molecules that induce mitochondrial biogenesis in cancer cells. We have shown that all four of these molecules

promote tumor growth. Thus, we propose that increased mitochondrial biogenesis functions as an "oncogene" or tumor promoter in cancer cells. Induced oxidative mitochondrial metabolism would allow cancer cells to more efficiently utilize imported L-lactate, ketone bodies, glutamine and fatty acids, which are all produced by neighboring activated CAFs or related stromal cells. Thus, targeting cancer cell mitochondria with mitochondrial "poisons" could halt this metabolic coupling, effectively shutting-down the "engine" or "powerhouse" driving tumor growth.

# Materials and Methods

Materials. Antibodies were all purchased from commercial sources: anti-PGC-1 $\alpha$  (NBP-1–04676, Novus), anti-mitoNEET (ab118027, Abcam), anti-PGC-1 total (ab72230, Abcam), anti-POLRMT (HPA006366, Sigma), anti-OXPHOS (MS601, Mitoscience), anti- $\beta$  actin (A5441, Sigma), anti-Beclin-1 (NBP1–00085, Novus) and anti-Cathepsin B (FL-339, Santa Cruz Biotechnology).





Cell culture and transfection. MDA-MB-231 cells were cultured in Dulbecco's modified Eagle's medium (DMEM), supplemented with 10% fetal bovine serum in a 37°C humidified atmosphere containing 5% CO<sub>2</sub>, unless otherwise noted. Lentiviral particles were produced after 48 h of transfecting the GeneCopoeia 293Ta lentiviral packaging cell line with lentiviral plasmids (EX-NEG-Lv105) (empty vector), PGC-1 $\alpha$  (EX-U0564-Lv105), MitoNEET (EX-V0831-Lv105), PGC-1 $\beta$  (EX-Mm13777-Lv105) and POLRMT (EX-Z7565-Lv105) (all obtained from GeneCopoeia, Inc.) using the Lenti-Pac HIV Expression Packing Kit (GeneCopoeia, Inc.) according to manufacturer's recommendation. MDA-MB-231 breast cancer cells were transduced with lentivirual particles in the presence of 5  $\mu$ g/ml of Polybrene (Santa Cruz Biotech). Successfully transduced cells were selected with 2  $\mu$ g/ml of puromycin.

Immunoblotting. Cells were lysed by scraping into cell lysis buffer (10 mM Tris, pH 7.5, 150 mM NaCl, 1% Triton X-100 and 60 mM n-octyl-glucoside), containing protease inhibitors (Boehringer Mannheim). To induce autophagy,



**Figure 6.** Mitochondrial biogenesis increases tumor growth. Schematic diagram highlighting that MitoNEET, PGC-1 ( $\alpha$ - and  $\beta$ -isoforms) and POLRMT all drive mitochondrial biogenesis, and effectively promote tumor growth, when expressed in cancer cells. Thus, the therapeutic use of mitochondrial inhibitors or "poisons" (such as metformin) will inhibit tumor growth. Importantly, numerous clinical studies have now shown that metformin (a mitochondrial compex l inhibitor) prevents the onset of nearly all types of human cancers in diabetic patients.

cells were starved for 5 h using Hank's Balanced Salt Solution (HBSS) supplemented with 40 mM HEPES. Samples were incubated on a rotating platform at 4°C and were then centrifuged at 12,000  $\times$  g for 10 min (at 4°C) to remove insoluble debris. Protein concentrations were analyzed using the BCA reagent (Pierce). Samples were then separated by SDS-PAGE (12% acrylamide) and transferred to nitrocellulose. All subsequent wash buffers contained 10 mM Tris, pH 8.0, 150 mM NaCl, 0.1% Tween 20, which was supplemented with 5% nonfat dry milk (Carnation) for the blocking solution and 1% bovine serum albumin (Sigma) for the antibody diluent. Horseradish peroxidase-conjugated secondary antibodies were used to

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visualize bound primary antibodies with an ECL detection kit (Pierce).

Animal studies. All animals studied were maintained in a pathogen-free environment/barrier facility at the Kimmel Cancer Center at Thomas Jefferson University under National Institutes of Health (NIH) guidelines. All animal protocols were approved by the Institutional Animal Care and Use Committee (IACUC). Tumor cells (MDA-MB-231;  $1 \times 10^6$  cells) in 100 µl of sterile PBS were injected into the flanks of athymic NCr nude mice (NCRNU; Taconic Farms; 6–8 wk of age). Mice were then sacrificed at 21 d post-injection; tumors were excised to determine their weights and volumes.

Angiogenesis quantification. Frozen sections were processed, as we described previously.<sup>36</sup> Then the anti-CD31 antibody was incubated overnight at 4°C. Finally, the sections were counterstained with hematoxylin for 5–10 sec, air-dried and mounted with cover-slips. To quantify tumor angiogenesis, CD31-positive vessels were enumerated in 8–10 fields within the central area of each tumor using a 20X objective lens and an ocular grid (0.25 mm<sup>2</sup> per field). The total numbers of vessel per unit area was calculated using ImageJ, and the data was represented graphically.

Statistical analysis. Statistical significance was examined by the Student's t-test. Values of  $p \le 0.05$  were considered significant. All p values were two-tailed (2T) unless otherwise noted.

## Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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