Metabolic Markers of Breast Cancer: Enhanced Choline Metabolism and Reduced Choline-Ether-Phospholipid Synthesis

Rachel Katz-Brull, Dalia Seger, Dalia Rivenson-Segal, Edna Rushkin, and Hadassa Degani

Department of Biological Regulation, Weizmann Institute of Science, Rehovot 76100, Israel

ABSTRACT

Specific genetic alterations during malignant transformation may induce the synthesis and breakdown of choline phospholipids, mediating transduction of mitogenic signals. The high level of water-soluble choline metabolites in cancerous breast tumors, relative to benign lesions and normal breast tissue, has been used as a diagnostic marker of malignancy. To unravel the biochemical pathways underlying this phenomenon, we used tracer kinetics and $^{13}$C and $^{31}$P magnetic resonance spectroscopy to compare choline transport, routing, and metabolism to phospholipids in primary cultures of human mammary epithelial cells and in MCF7 human breast cancer cells. The rate of choline transport under physiological choline concentrations was 2-fold higher in the cancer cells. The phosphorylation of choline to phosphocholine and oxidation of choline to betaine yielded 10-fold higher levels of these metabolites in the cancer cells. However, additional incorporation of choline to phosphatidylcholine was similar in both cell types. Thus, enhanced choline transport and augmented synthesis of phosphocholine and betaine are dominant pathways responsible for the elevated presence of choline metabolites in cancerous breast tumors. Uniquely, reduced levels and synthesis of a choline-ether-phospholipid may also serve as a metabolic marker of breast cancer.

INTRODUCTION

Elevated concentrations of choline and choline metabolites (composite choline) were observed by MRS in a variety of malignancies (1, 2). This elevation has been particularly useful for differentiating between malignant and benign breast lesions, because the former contain a significantly higher level of composite choline than the latter (3–8). However, to fully exploit the diagnostic potential of breast MRS, it is necessary to elucidate the underlying biochemical mechanisms leading to this metabolic phenomenon.

Choline, a quaternary amine, is an essential nutrient supplied predominantly by the diet (9, 10). The capacity to take up and secrete high levels of choline and choline metabolites is a central function of mammary epithelial cells. During lactation, these cells are capable of concentrating choline from the plasma and, subsequently, secreting milk that is rich in choline-containing metabolites, primarily PCho and GPCho (11).

Active transport and diffusion are major mechanisms in the uptake of choline across cellular membranes. The diffusion capacity through membranes is related to the composition and special assembly of lipids, predominantly phospholipids and cholesterol, as well as of proteins. Routing of choline through its various metabolic pathways is cell and tissue specific (12). The intracellular metabolism of choline in the breast is partitioned among two major pathways: (a) synthesis of PtdCho; and (b) oxidation to produce the methyl donor betaine.

Choline metabolism and choline-derived metabolites can undergo extensive alterations as a result of malignant transformations. Progression of HMECs from a normal to a malignant phenotype was shown recently to be associated with a reversal in the ratio of PCho to GPCho, as well as an overall increase in the content of these two metabolites (13).

PCho is a precursor of choline-derived phospholipids, as well as a product of their hydrolysis. The synthesis and degradation of phospholipids may be induced by growth factors that play a major role in malignant transformations (14, 15). The level of PCho in human breast cancer cells was found to be ~10-fold higher than in HMECs (13, 16, 17). Moreover, high levels of PCho and other phosphoesters were detected in human breast cancer biopsies and in patients in vivo (1, 18). The high levels of PCho correlated with up-regulation and increased activity of choline kinase, and choline kinase inhibitors exhibited antitumor activity (19, 20). High choline transport was suggested as the cause for the elevated levels of PCho in breast cancer (21). However, the exact role of PCho in malignant transformation and the involvement of other choline metabolites in this transformation are not well understood.

We present herein comparative studies of the metabolic steps that determine the nature and distribution of choline metabolites in normal and cancerous mammary epithelial cells. Detailed radioactive tracer measurements and model-based analyses yielded the mechanisms and kinetic parameters of choline transport. Additional metabolic studies using $^{13}$C- and $^{31}$P-MRS and $^{13}$C-labeled choline enabled us to monitor the synthesis of PCho and betaine and characterize the composition and turnover of choline-derived phospholipids. Our results demonstrated enhanced transport of choline, augmented synthesis of PCho and betaine, and suppression of the synthesis of choline-derived ether lipids in breast cancer cells.

MATERIALS AND METHODS

Tissue Culture. MCF7 human breast cancer cells were cultured routinely in DMEM supplemented with 6% FCS and antibiotics, as described previously (21). T47D-clone 11 human breast cancer cells were routinely cultured in RPMI 1640 supplemented with 10% FCS, as described previously (17). Both media contained 28 µM choline, with an additional ~2 µM choline from the serum.

Primary cultures of HMECs were obtained from two sources: Clonetics Corp., San Diego, CA, which provided HMECs isolated from epithelial organoids of human breast tissue; and our laboratory. The cells from Clonetics were cultured in serum-free mammary epithelial cell growth medium supplemented with bovine pituitary extract, insulin, human epidermal growth factor, hydrocortisone, and antibiotics. For the second source of HMECs (HMEC’s), we isolated, separated, and cultivated the cells in our laboratory by processing breast tissue obtained from mammoplasty reduction surgery, as described previously (17).

Choline Transport. MCF7 and HMEC cells were preincubated for 3 h in choline-free DMEM supplemented with 2% FCS. Transport experiments were initiated by adding various concentrations of choline (1–400 µM) to the above medium and trace amounts (1–2 µCi) of [methyl-14C]choline chloride (Sigma}
Chemical Co., St. Louis, MO). For each choline concentration, samples were incubated with the tracer at 36 °C for 4, 8, and 12 min. After we cooled the samples to 0°C, they were washed three times with ice-cold PBS, and the cells were scraped with ice-cold methanol and mixed with scintillation fluid for additional β-counting. The initial transport rates were obtained by a linear fitting of the amount of radioactive choline at 4, 8, and 12 min (per cell number). The initial transport rates were further normalized to mg protein/cell, independently determined by the Bradford method (22), yielding a value of 220 ± 8 picogram/cell for MCF7 cells and 410 ± 50 picogram/cell for HMECs. The −2-fold higher protein content of HMECs, as well as the 2-fold higher NTP (predominantly ATP) correlated with the larger volume of the latter cells (17).

The kinetic parameters were calculated using Stein’s method for zero-trans transport (23). The mechanism(s) of transport was revealed through specific patterns of the plots of the initial rate $v$ versus $S$, and $1/k$ versus $S$. The kinetic parameters were calculated by a nonlinear fitting of each mode of plotting and are presented as the mean ± SD of the three plots.

Choline Metabolism. MCF7 cells and HMECs (1.5–2 × 10^6 and 2–4 × 10^7 cells, respectively) were incubated at 36 °C for 24 h in choline-free culture medium supplemented with 100 μM [1,2,3-13]choline, 99% 13C enriched (Cambridge Isotopes, Ltd., Andover, MA). The medium was then collected, and the cells were extracted using the dual-phase extraction method (24).

The methanol-water phase was treated with Chelex-100 (Sigma), lyophilized to dryness and kept at −20°C. Before the NMR measurement, the dried residue was redissolved in 0.5 ml of D_2O (99.9% enriched; Cambridge Isotopes, Ltd.) containing 5 μl of methanol as a standard for 13C measurements, at a pH of 8–8.3. After recording the 13C spectra and before the 31P measurements, 10 mM EDTA was added to the samples. The chloroform of the lipid phase was evaporated under N_2, and the dried residue was redissolved in a mixture of 0.4 ml of chloroform and 0.2 ml of methanolic EDTA solution (25).

The large volumes of the culture medium were concentrated by sequential lyophilizations. The resulting dried residue was dissolved in 0.5 ml of D_2O (99.9% enriched; Cambridge Isotopes, Ltd.) containing 5 μl of methanol as a standard for 13C measurements, at a pH of 8–8.3. After recording the 13C spectra and before the 31P measurements, 10 mM EDTA was added to the samples. The chloroform of the lipid phase was evaporated under N_2, and the dried residue was redissolved in a mixture of 0.4 ml of chloroform and 0.2 ml of methanolic EDTA solution (25). The large volumes of the culture medium were concentrated by sequential lyophilizations. The resulting dried residue was dissolved in 0.5 ml of D_2O (99.9% enriched; Cambridge Isotopes, Ltd.) containing 5 μl of methanol as a standard for 13C measurements, at a pH of 8–8.3. After recording the 13C spectra and before the 31P measurements, 10 mM EDTA was added to the samples. The chloroform of the lipid phase was evaporated under N_2, and the dried residue was redissolved in a mixture of 0.4 ml of chloroform and 0.2 ml of methanolic EDTA solution (25).

Lipid extraction of T47D and the second source of HMEC's was performed using a modification of Folch's method, which extracted phospholipids with an efficiency similar to that of the dual-phase extraction method described above (24).

NMR Spectroscopy. High-resolution NMR spectra were recorded on a DMX-500 spectrometer operating at 11.7 Tesla (Bruker Analytic GMBH, Karlsruhe, Germany). 13C spectra of the water-soluble metabolites and the lipids were recorded at 125.7 MHz, using a 1H/13C dual probe and applying 60° pulses, 2.4-s repetition time, and continuous composite pulse proton decoupling. 31P NMR spectra of the water-soluble metabolites and the lipids were recorded at 202.4 MHz by applying 45° pulses, a repetition time of 2 s, and continuous composite pulse proton decoupling. The signals of α-NTP (at −10.03 ppm) and the signal of added phenylphosphonic acid served as references for chemical shift and concentration, respectively. Signal intensity and area in NMR spectra were measured with XWIN-NMR (Bruker Biospin MRS GmbH). The amount of NTP was determined from the combined area of γ-NTP + β-nucleotide disphosphate. The nonphosphorylated metabolites were referenced to NTP through the use of the PCho signal, which appeared in both the 13C and the 31P spectra. The areas of the signals in both the 31P and the 13C spectra were converted to concentration units, percent enrichment, or metabolite ratio, taking into account 13C enrichment as well as differences attributable to relaxation and nuclear Overhauser enhancement. The final results are presented as the mean ± SE.

RESULTS
Choline Transport
The mechanisms contributing to choline transport into MCF7 human breast cancer cells and HMECs, as well as the rate constants associated with each mechanism, were determined by applying a tracer kinetic method. The initial rate of choline uptake ($v$) was measured as a function of choline concentration in the medium ($S$). Plots of $v$ versus $S$ (Fig. 1, A and B) as well as $S/v$ and $Stv$ versus $S$ revealed the presence of a saturable, Michaelis-Menten-like mechanism as well as a diffusion mechanism, according to:

$$v = \frac{V_{\text{max}}S}{K_m + S} + K_D S,$$

where $K_m$ and $V_{\text{max}}$ define the rate constant and maximum rate of the saturable transport, and $K_D$ the diffusion rate constant. Fitting the data according to the above equation (Fig. 1, A and B) yielded similar $K_m$s in both MCF7 cells and HMECs (20 ± 5 μM and 14 ± 9 μM, respectively); however, $V_{\text{max}}$ was 2-fold faster in the cancer cells (Table 1). Thus, MCF7 breast cancer cells exhibited increased choline transport activity and stimulation of choline uptake in the physiological range (10–30 μM) of choline (Fig. 1, C and D).

Interestingly, the diffusion rate constant $K_D$ was higher in HMECs, indicating changes in cell membrane properties during malignant transformation. However, this mechanism contributed significantly to choline transport only at high concentrations of choline, much above the physiological range (Fig. 1, C and D).

Choline Metabolism
Choline Routing. Differential routing and incorporation of choline into intracellular water-soluble metabolites were investigated in MCF7 cells and HMECs. 13C labeling of choline metabolites was monitored after incubation with 100 μM [1,2,3-13C]choline (high above the $K_m$ of choline transport). The two 13C methylene signals of intracellular choline, PCho and betaine, were clearly observed in the 13C spectra of the water phase extracts of MCF7 cells (Fig. 2A).

Table 1 Kinetic parameters of choline transport in HMECs and MCF7 cells

<table>
<thead>
<tr>
<th></th>
<th>HMEC</th>
<th>MCF7</th>
</tr>
</thead>
<tbody>
<tr>
<td>$K_m$ (μM)</td>
<td>12 ± 9</td>
<td>20 ± 5</td>
</tr>
<tr>
<td>$V_{\text{max}}$ (nmol/mg protein/h)</td>
<td>8.8 ± 3.4</td>
<td>20 ± 4</td>
</tr>
<tr>
<td>$V_{\text{max}}$ (nmol/mg protein/h/μM)</td>
<td>83 ± 15</td>
<td>44 ± 19</td>
</tr>
<tr>
<td>$R^2$</td>
<td>0.94* 0.86* 0.84*</td>
<td>0.99* 0.99*</td>
</tr>
</tbody>
</table>

*The saturable kinetic parameters, $K_m$ and $V_{\text{max}}$, as well as the nonsaturable rate constant $K_D$, were determined by a nonlinear least squares curve fitting of: $v$ initial rate, $v$ versus choline concentration; $S$; $S/v$ versus $S$ and $Stv$ versus $S$. Results are presented as the mean ± SD of the values obtained in the three curve fittings. The corresponding values in units per cell are: $V_{\text{max}}$ of 3.6 ± 1.4 and 4.4 ± 0.8 fmol/cell/h in HMEC and MCF7, respectively; and $K_D$ of 34 ± 6 and 9.7 ± 4.1 fmol/cell/h in HMEC and MCF7, respectively.

Fig. 1. Variation in the initial rate of choline transport with choline concentration in MCF7 cells and HMECs. The solid lines in A (MCF7 cells) and B (HMECs) demonstrate the best curve fitting to the kinetic equation: $v = \frac{V_{\text{max}}S}{K_m + S} + K_D S$. C and D, calculation of choline transport in MCF7 and HMECs, respectively, at 0–100 μM choline, using the transport rate constants described in Table 1. I, initial rate of choline uptake (equation above); II, saturable choline uptake; III, passive diffusion of choline.
13C-enriched betaine was also detected in the spectra of the concentrated incubation medium of these cells (Fig. 2B), indicating partial transfer of this metabolite to the medium. The 13C spectra of HMECs demonstrated the presence of intracellular 13C-enriched choline and PCho; however, betaine was not detected in the cell extracts or the medium.

In the spectra of both the normal and the cancer cells, we did not observe incorporation of the 13C-enriched choline into GPCho, a breakdown product of PdCho, indicating a slow turnover and breakdown of PdCho. Both cell types were shown to contain GPCho, with the level in the cancer cells ~10-fold higher than in HMECs (17).

Quantitative analysis of the 13C spectra showed that the pools of choline, PCho, and betaine were fully 13C enriched in both cell types. However, the content of these metabolites was markedly different (Fig. 3). In MCF7 cells, most of the choline was metabolized to PCho, which accumulated in the cells to a very high level. The remaining choline was oxidized to betaine, but a large fraction of this metabolite (60%) was found in the medium. On the other hand, in HMECs most of the choline remained in its free form, and only a small fraction was metabolized to PCho, whereas oxidation to betaine was too low to be detected.

Choline Incorporation to Lipids. 31P and 13C NMR of lipid cell extracts monitored the incorporation of the 13C-labeled choline into PtdCho and other choline-derived lipids. In the 31P spectra of both the normal and the cancer cells, we observed signals of [1,2-13C]-labeled PtdCho and signals at 13C natural abundance of PtdCho, phosphatidylinositol, sphingomyelin, phosphatidylserine, peak 5, sphingomyelin, peak 6, phosphatidylcholine, peak 7, cholesterol-phospholipid, peak 8 and 10, [1,2-13C]PtdCho; peak 9, PtdCho. Inset in A, expansion of phosphatidylethanolamine-plasmalogen and phosphatidylethanolamine region. The spectra were acquired with 48,100 scans (A) and 22,500 scans (B).
either MCF7 or T47D breast cancer cells (Table 2). On the basis of previous studies, this signal was assigned to a choline-ether-phospholipid (26, 27), presumably 1-alkyl-2-acetyl-sn-glycero-3-PCho (28).

In the $^{13}$C spectra of the lipid phase extracts of MCF7 cells, we observed incorporation of the $^{13}$C-enriched methylenes of choline into PtdCho (Fig. 5A), with no signs of their incorporation into other lipids. However, in the $^{13}$C spectra of HMECs, unique $^{13}$C methylene signals of nonenriched and [1,2-$^{13}$C]choline-enriched lipid were detected, in addition to the presence of $^{13}$C-enriched PtdCho (Fig. 5B). We have tentatively assigned these signals to the choline-ether-phospholipid, detected exclusively in the $^{31}$P spectra of HMECs. This assignment was based on: (a) a characteristic chemical shift; (b) $^{13}$C$_1$-$^{13}$C$_2$ splitting (40 Hz); (c) splitting attributable to the neighboring phosphate (5.1 Hz); and (d) similarity in the areas of the signals ($\sim$10% of PtdCho) in $^{31}$P and $^{13}$C spectra.

Quantitative analysis of the signals in the $^{31}$P and $^{13}$C spectra indicated similarities in the extent of $^{13}$C labeling of PtdCho in the cancer and normal cells (39 ± 2% and 33 ± 5%, respectively; mean ± SD). Thus, although transport and phosphorylation of choline were faster in the cancer cells, the turnover of PtdCho appeared to proceed at about the same rate in both the normal and cancer cells, indicating similar control of the rate-limiting, cytosine-diphosphocholine synthesis. The extent of $^{13}$C labeling of the unique HMEC choline-ether-phospholipid (11 ± 3%) was of the same order of magnitude as that of PtdCho, indicating continuous, significant incorporation of choline into both phospholipids. This finding suggests that this unique lipid plays an important role in the normal functioning of mammary cells.

A substantial difference was found in the levels of the ethanolamine-ether-phospholipid phosphatidylethanolamine plasmalogen in the normal cells and in the cancer cells (Table 2). It was thus concluded that the capacity of normal mammary cells to synthesize ether-phospholipids was markedly suppressed in their malignant counterpart.

**DISCUSSION**

In this study, we focused on investigating alterations in choline transport, oxidation, and phosphorylation induced by malignant transformation of HMECs. In addition, we searched for differences in membrane phospholipid composition and choline incorporation into those lipids in normal and cancerous mammary cells.

A sensitive radioactive labeling method and detailed measurements taken to determine the kinetic parameters of choline transport demonstrated the presence of both a saturable and a nonsaturable transport mechanism with significantly different rate parameters. The maximum rate of the saturable component of choline transport was $>2$-fold faster in the cancer cells, whereas the diffusion rate constant was higher in the normal cells (Table 1). The saturable mechanism predominated in the physiological range of choline in human plasma. It is therefore reasonable to propose that, in breast cancer patients, choline uptake into a malignant lesion is faster than its uptake into the surrounding normal glandular tissue. In addition, the increased choline kinase activity demonstrated in transformed cells (29) may further augment the level of PCho in these cells, thus increasing further the intensity of the composite choline signal of breast cancer, as recorded by MRS, to detectable levels.

Choline uptake into mammary epithelial cells was investigated previously in cells isolated from lactating rats (30). In the rat lactating cells, as in HMECs, saturable and nonsaturable mechanisms were found to be operative, with the former predominating at physiological concentrations (30). The mean $K_m$ of the saturable component ($35 \pm 16 \mu M$) was within the range of that found here for HMECs and MCF7 cells (Table 1). However, the $V_{\text{max}}$ of transport ($1.24 \pm 0.19$ nmol/mg protein/h) was lower, presumably because of differences in species and isolation protocols.

The nonsaturable transport in HMECs and MCF7 cells contributed to the uptake at high external choline concentrations and presumably occurred via passive diffusion, as was also suggested for the lactating rat mammary cells (30).

The phospholipid distribution of both normal HMECs and MCF7 human breast cancer cells revealed similarities in composition of diacyl phospholipids and sphingomyelin but pointed up a substantial decrease in the amount of the ether-phospholipids in the cancer cells. Levels of both choline-ether-phospholipids and ethanolamine-ether-phospholipids were significantly lower in the cancer cells relative to HMECs and to amounts detected in mammary fibroblasts (Ref. 17;
data not shown). The role played by ether phospholipids in mammary cells is currently unclear. Furthermore, it is also not known how a lack of these phospholipids might affect breast cancer etiology, despite interest in the function of ether lipids in cancer therapy (31). However, it appears reasonable to propose that the enhanced rates of passive diffusion of the positively charged choline through the cell membranes of HMECs, relative to that of MCF7, could be associated with the unique alterations in the phospholipid subclasses making up the membrane. In addition, our novel finding that human breast cancer cells contain reduced amounts of ether phospholipids might be related to the differential anticancer activity of alkyl-lysophospholipids (31), which in turn is associated with disturbances in membrane phospholipid metabolism (32–36).

In this study, we confirmed that most of the choline in MCF7 cells was converted to PCho and then routed through the cytokine-diphosphocholine pathway to form choline phospholipids (21, 37). In addition, we showed that a significant amount of choline (~25%) was oxidized to betaine, and most of it was found in the growth medium of the cells. It therefore appears that the two non-intersecting pathways, phosphorylation and oxidation of choline, are augmented in the course of malignant transformation of mammary cells.

In summary, our results demonstrated enhanced transport of choline, an augmented synthesis of PCho and betaine, and a unique suppression of the synthesis of choline-derivative ether lipids in MCF7 and T47D human breast cancer cells. These biochemical changes support the diagnostic utilization of the composite choline magnetic resonance signal as a marker for breast cancer.

ACKNOWLEDGMENTS

We thank Barbara Morgenstern for editing the manuscript.

REFERENCES


Metabolic Markers of Breast Cancer: Enhanced Choline Metabolism and Reduced Choline-Ether-Phospholipid Synthesis

Rachel Katz-Brull, Dalia Seger, Dalia Rivenson-Segal, et al.


Updated version
Access the most recent version of this article at:
http://cancerres.aacrjournals.org/content/62/7/1966

Cited articles
This article cites 30 articles, 10 of which you can access for free at:
http://cancerres.aacrjournals.org/content/62/7/1966.full#ref-list-1

Citing articles
This article has been cited by 13 HighWire-hosted articles. Access the articles at:
http://cancerres.aacrjournals.org/content/62/7/1966.full#related-urls

E-mail alerts
Sign up to receive free email-alerts related to this article or journal.

Reprints and Subscriptions
To order reprints of this article or to subscribe to the journal, contact the AACR Publications Department at pubs@aacr.org.

Permissions
To request permission to re-use all or part of this article, use this link http://cancerres.aacrjournals.org/content/62/7/1966.
Click on "Request Permissions" which will take you to the Copyright Clearance Center’s (CCC) Rightslink site.