Paclitaxel causes *de novo* induction of invasive breast cancer cells by repolarizing

tumor-associated macrophages

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Abstract

Metastasis is the leading cause of breast cancer death, and tumor cells must migrate

and invade to metastasize. Breast cancer cells that express the pro-metastatic actin regulatory

protein MenaINV have an enhanced ability to migrate and intravasate within the primary tumor

and extravasate at secondary sites. Though chemotherapy improves patient survival, treatment

with paclitaxel leads to upregulation of MenaINV and an increase in metastasis in mice.

MenaINV expression can be induced in breast cancer cells through cooperative NF-κB/ Notch1

signaling with macrophages, which are often increased in tumors in response to chemotherapy.

MenaINV-expressing cells are also resistant to paclitaxel, raising the question of whether

paclitaxel increases MenaINV by *de novo* induction or by selectively killing non-MenaINV-

expressing cells. We hypothesized that paclitaxel causes *de novo* MenaINV induction by

increasing macrophage-tumor cell NF-κB/ Notch1 signaling. Understanding this pro-metastatic

effect of chemotherapy is crucial to refining treatment strategies.

In this study, we demonstrate that paclitaxel increases MenaINV expression by *de novo*

induction. Mechanistically, paclitaxel induces MenaINV by repolarizing tumor-associated

macrophages towards a pro-inflammatory phenotype. These pro-inflammatory macrophages

then participate in enhanced NF-κB/ Notch1 signaling with tumor cells, which leads to MenaINV

induction in the tumor cells. These results lay the groundwork for novel microenvironment-based

therapies to alleviate the pro-metastatic effects of chemotherapy in breast cancer.

Keywords: Breast cancer, macrophages, MenaINV, NF-κB, Notch1, paclitaxel, chemotherapy

Main text

Introduction

Breast cancer (BC) is the second leading cause of cancer death among women in the

United States, with most deaths resulting from metastasis (1). To metastasize, tumor cells must

migrate and intravasate within the primary tumor and extravasate at secondary sites. Tumor

cells accomplish these tasks using invadopodia – actin-rich protrusions with proteolytic activity

that degrade extracellular matrix. In BC cells, the actin regulatory protein MenaINV supports

invadopodium stability and degradative activity (2), enhancing cell migration and metastasis.

Furthermore, tumor cells that express MenaINV are more sensitive to chemotactic gradients in

the tumor microenvironment, which facilitate invadopodium formation and directional migration

towards blood vessels where they intravasate (3, 4). Indeed, BC cells that express MenaINV

have an enhanced ability to migrate, intravasate, and extravasate (4-6), and high MenaINV

expression is associated with poor outcomes in both mouse (4) and human (7, 8) BC.

Though chemotherapy improves patient survival, neoadjuvant paclitaxel increases the

percent MenaINV+ area in both mouse and human residual breast tumors, as well as lung

metastasis in mice (9). MenaINV-expressing cells are resistant to paclitaxel (10), so the

observed increase in MenaINV after paclitaxel treatment may be due to the selective killing of

non-MenaINV-expressing cells. However, we have shown that MenaINV is induced in tumor

cells via cooperative NF-κB/Notch1 signaling with macrophages (11). Chemotherapy-induced

tissue damage has been shown to cause a macrophage influx into tumors, potentially increasing

macrophage-tumor cell NF-κB/Notch1 signaling and MenaINV induction. Thus, in addition to the

selective killing of non-MenaINV-expressing cells, paclitaxel may also cause *de novo* induction

of MenaINV. We hypothesized that paclitaxel induces MenaINV expression by increasing

macrophage infiltration and macrophage-tumor cell NF-κB/Notch1 signaling. Here, we find that

paclitaxel causes macrophages to adopt a pro-inflammatory phenotype that promotes NF-

κB/Notch1 signaling and leads to *de novo* MenaINV induction in tumor cells.

Results

Paclitaxel causes *de novo* MenaINV induction

To test whether paclitaxel causes *de novo* MenaINV induction, we treated FVB-PyMT

mice bearing spontaneous breast tumors with paclitaxel and co-administered either DAPT – to

prevent MenaINV induction (11, 12) – or vehicle control (**Fig. 1A, Additional file 1: Materials**

and Methods). DAPT is a γ-secretase inhibitor used to block Notch signaling. However, given

the crosstalk between the Notch and NF-κB pathways, we (11) and others (13) have found that

DAPT also inhibits NF-κB activation. If paclitaxel increases the percent MenaINV+ area only by

selection, then dual DAPT/paclitaxel treatment would result in the same MenaINV increase as

paclitaxel alone through the selective killing of non-MenaINV-expressing cells.

Following treatment, mice were sacrificed, and their tumors were analyzed for MenaINV

expression, NF-κB activation (measured by p65 nuclear localization (**Fig. 1B**)), Notch1

activation (measured by cleaved Notch1 intracellular domain (N1ICD)), and macrophage

infiltration. As previously observed, paclitaxel-treated tumors expressed significantly more

MenaINV than controls (**Fig. 1C, D,** first two bars/panels), and this was accompanied by a

significant increase in NF-κB and Notch1 activation (**Fig. 1E-H**). When MenaINV induction was

inhibited with DAPT, paclitaxel no longer increased MenaINV (**Fig. 1C, D,** third bar/panel). This

indicates that paclitaxel causes *de novo* MenaINV induction.

We originally hypothesized that this induction resulted from increased macrophage

infiltration following paclitaxel treatment, but no difference was observed (**Fig. 1I**). This led us to

question the necessity of macrophages in paclitaxel-mediated MenaINV induction.

Macrophages are required for paclitaxel-mediated MenaINV induction

To test the role of macrophages in paclitaxel-mediated MenaINV induction, we used two

mouse models of BC: FVB-PyMT mice as above, and SCID mice bearing the triple-negative

HT17 patient-derived xenograft (14). Mice received paclitaxel and either clodronate liposomes –

to deplete macrophages – or PBS control liposomes (**Fig. 2A; Additional file 2: Fig. S1A**).

After treatment, tumors were analyzed for MenaINV expression, NF-κB and Notch1 activation,

and macrophage infiltration. Paclitaxel again caused a significant induction of MenaINV, and this

induction was prevented when macrophages were depleted (**Fig. 2B-D; Additional file 2: Fig.**

S1B, C). This indicates that macrophages *are* required for paclitaxel to induce MenaINV. NF-κB

and Notch1 signaling were also significantly decreased when macrophages were depleted

during paclitaxel treatment (**Fig. 2E-H; Additional file 2: Fig. S1D, E**), indicating that

macrophages are the primary source of additional NF-κB and Notch1 stimulation in the tumor

microenvironment following paclitaxel treatment. Our data indicate that for paclitaxel to induce

MenaINV, macrophages must be present, but not necessarily in higher proportions than in

untreated conditions. This led us to ask whether paclitaxel changes the phenotype of

macrophages to promote NF-κB/Notch1 signaling.

<u>Paclitaxel induces a pro-inflammatory phenotype in macrophages, which promotes NF-</u>

<u>κB/Notch1 signaling</u>

To investigate how paclitaxel affects the ability of macrophages to participate in NF-

κB/Notch1 signaling, we cultured RAW264.7 murine macrophages or THP-1 human

macrophages with 10 μM paclitaxel for 24 h and measured mRNA expression of Notch ligand

Jagged1 and secretion of TNFα, IL-1β, and TGF-β1, which activate NF-κB. TGF-β1 can also

enhance Notch signaling in several contexts (15). Paclitaxel significantly increased Jagged1

expression and TNFα/IL-1β secretion in both cell lines (**Fig. 3A-C; Additional file 2: Fig. S2A-**

C), and TGF-β1 secretion in RAW264.7 cells (**Fig. 3D**), but not THP-1 cells (**Additional file 2:**

Fig. S2D). These results indicate that paclitaxel causes macrophages to adopt a pro-

inflammatory phenotype that promotes their participation in NF-κB and Notch1 signaling, though

the exact phenotypic changes may differ between species.

Neoadjuvant chemotherapy alters macrophage subtype composition in patients

<u>Neoadjuvant chemotherapy alters macrophage subtype composition in patients</u>

We next investigated whether chemotherapy also repolarizes macrophages in BC

patients. We identified a previously published dataset of single-cell RNA sequencing reads from

human BC patient biopsies taken either before treatment (treatment-naïve) or after neoadjuvant

chemotherapy (NAC), as described in the original publication (16). We annotated the

macrophage compartment of this dataset based on ontogeny and functional gene signatures, as

previously described (17), allowing our analysis to account for a spectrum of macrophage

polarizations (**Fig. 3E**). UMAP visualization showed clear clustering between treatment groups,

indicating distinct transcriptional profiles (**Fig. 3F**).

Both treatment groups exhibited macrophage heterogeneity. However, the macrophage

population in NAC tumors was dominated by interferon-primed monocyte-derived (Mac_IFN)

and tissue-resident (RTM_IFN) macrophages, which are characterized by high expression of

interferon-primed genes, including *CCL8, IDO1*, and *CXCL9* (17) (**Fig. 3G**). The Mac_IFN

subtype increased from 13.7% of macrophages in the treatment-naïve group to 34.5% of

macrophages in the NAC group, while the RTM_IFN subtype rose from 26.8% to 30.9%.

Together, these interferon-primed macrophages account for 65.4% of all macrophages in NAC

tumors, compared to 40.5% in treatment naïve tumors.

Gene set variation analysis (GSVA) revealed that these two subtypes were highly

upregulated in pathways related to inflammatory cytokine secretion (**Fig. 3H**). The Mac_IFN

subtype was also upregulated in Notch signaling genes, which include the Notch ligands (**Fig.**

3H). These results confirm that NAC promotes an inflammatory phenotype in macrophages in

human BC, which increases their ability to participate in NF-κB/Notch1 signaling, and

complement our previous finding that NAC increases MenaINV expression in human BC (9).

Taken together, our results reveal that chemotherapy skews macrophages towards a

pro-inflammatory phenotype, increasing their ability to participate in NF-κB/Notch1 signaling

with tumor cells, which subsequently leads to MenaINV induction (**Fig. 4**).

Discussion

In this study, we discovered that neoadjuvant paclitaxel induces the expression of the

pro-metastatic protein MenaINV in BC cells by promoting NF-κB/Notch1 signaling with

macrophages. We previously observed a paclitaxel-driven increase in MenaINV (9, 11).

However, because MenaINV expression confers resistance to paclitaxel (10), it was unknown

whether this increase resulted from *de novo* induction or selective killing of non-MenaINV-

expressing cells. Here, we show that the paclitaxel-driven increase in MenaINV is the result of

de novo induction; when the pathways involved in MenaINV induction were blocked during

paclitaxel treatment, no increase in MenaINV was observed (**Fig. 1C**). These results indicate

that paclitaxel *induces* MenaINV expression.

We have previously shown that Notch1 signaling between macrophages and tumor cells

is crucial for robust MenaINV induction (11, 12). The Notch1 receptor and its ligands are

membrane-bound, so Notch1 signaling only occurs between adjacent cells. Given this and our

previous finding that macrophages are necessary for the paclitaxel-mediated MenaINV increase

(11), we hypothesized that paclitaxel causes a macrophage influx into the tumor, increasing

opportunities for macrophage-tumor cell Notch1 signaling. Interestingly, we found that, while

macrophages are necessary for paclitaxel to induce MenaINV (**Fig. 2B-D**), an increase in

macrophage proportion is not (**Fig. 1C, I**). We conclude that, rather than increasing

macrophage-tumor cell NF-κB/Notch1 signaling by increasing macrophage infiltration, paclitaxel

repolarizes macrophages, allowing them to participate in such signaling more readily (**Fig. 3**).

This aligns with previous reports that paclitaxel activates TLR4 receptors on macrophages,

increasing inflammatory signaling (18), and complements our previous finding that TNFα

increases MenaINV expression in tumor cells *in vitro* (11).

Chemotherapy has long been known to exert both pro- and anti-tumoral effects on

macrophages. Indeed, prior evidence links chemotherapy-induced pro-inflammatory

macrophages with elevated innate and adaptive anti-tumoral immune response (19). Here, we

show that chemotherapy-induced pro-inflammatory macrophages also have the pro-tumoral

effect of inducing the pro-metastatic protein MenaINV in tumor cells. As drug discovery

platforms and protein-targeting capabilities advance, it is crucial to investigate the exact

mechanisms behind both the pro- and anti-tumoral responses to chemotherapy to refine

treatment strategies.

List of abbreviations: *BC*, breast cancer; *N1ICD*, Notch1 intracellular domain; *NAC*,

neoadjuvant chemotherapy; *Mac_IFN*, interferon-primed monocyte-derived macrophages;

RTM_IFN, interferon-primed tissue-resident macrophages; *GSVA*, gene set variation analysis.

Figure Legends

Figure 1. Paclitaxel increases MenaINV through *de novo* induction: (**A**) Experimental

design of paclitaxel and DAPT treatments in PyMT mice. Figure created with BioRender.com.

(**B**) Representative image of PyMT breast tumor stained for p65. Yellow arrow points to cell with

active NF-κB signaling (nuclear p65). Red arrow points to cell with only inactive NF-κB signaling

(cytoplasmic p65). (**C, D**) Quantification and representative images of MenaINV (measured by

% MenaINV+ area) in tumors from mice treated with control (ctrl), paclitaxel (ptx), and paclitaxel

+ DAPT (ptx + DAPT). (**E, F**) Quantification and representative images of NF-κB activation

(measured by p65 nuclear localization) in tumors from mice treated with control and paclitaxel.

(**G, H**) Quantification and representative images of Notch1 activation (measured by expression

of cleaved Notch1 intracellular domain) in tumors from mice treated with control and paclitaxel.

(**I**) Macrophage infiltration (measured by % IBA1+ area) in tumors from mice treated with

control, paclitaxel, and paclitaxel + DAPT. Data are reported as mean +/- SEM. Data in **C** and **I**

were analyzed using a one-way ANOVA. Data in **E** and **G** were analyzed using a *t*-test. *$p <$

0.05. Scale bars = 20 µM. Each point on the graphs represents the average value from 10 fields

of view taken from an individual mouse.

Figure 2. Macrophages are required for paclitaxel-mediated MenaINV induction: (**A**)

Experimental design of paclitaxel and clodronate treatments in PyMT mice. Figure created with

BioRender.com. (**B, C**) Quantification and representative images of MenaINV in tumors from

mice treated with control (ctrl), paclitaxel (ptx), and paclitaxel + clodronate liposomes (ptx +

CLD). (**D**) Macrophage infiltration (# IBA1+ cells per high power field (HPF)) in tumors treated

with control, paclitaxel, and paclitaxel + clodronate. (**E, F**) Quantification and representative

images of NF-κB activation in tumors from mice treated with control, paclitaxel, and paclitaxel +

clodronate liposomes. (**G, H**) Quantification and representative images of Notch1 activation in

tumors from mice treated with control, paclitaxel, and paclitaxel + clodronate liposomes. Data

are reported as mean +/- SEM. Data were analyzed using a one-way ANOVA. *$p < 0.05$, **$p <$

0.01, ***$p < 0.001$, ****$p < 0.0001$. Scale bars = 20 µM. Each point on the graphs represents the

average value from 10 fields of view taken from an individual mouse.

Figure 3. Paclitaxel alters macrophage phenotype, promoting NF-κB and Notch1

signaling: (**A**) Jagged1 mRNA expression in RAW264.7 macrophages treated with 10 µM

paclitaxel. Secretion of (**B**) TNFⓍ, (**C**) IL-1β, and (**D**) TGF-β1 from RAW264.7 macrophages

treated with paclitaxel. (**E-H**) single-cell RNA-sequencing analysis of macrophages in the tumors

of breast cancer patients who were treatment-naïve (naïve) or had undergone neoadjuvant

chemotherapy (NAC). (**E**) UMAP (uniform manifold approximation and projection) of tumor-

associated macrophages colored by macrophage subtype. Subtypes beginning with "Mac" are

monocyte-derived macrophages (based on expression of monocyte markers *CCR2, VCAN,*

S100A6, and *CD52*). Subtypes beginning with "RTM" are tissue-resident macrophages (based

on expression of tissue residency markers *TCF12, MS4A4A, GATM, LYVE1, FOLR2, and PLTP,*

as described in the original publication (17)). AgPres = antigen presenting, Alv-like = alveolar-

like, Angio = angiogenic, Hypo = hypoxia-related, IFN = interferon-primed, LA = lipid-associated,

Rec = recruited, IM = interstitial macrophage, MT = metallothioneins. (**F**) UMAP of tumor-

associated macrophages colored by treatment group. (**G**) Stacked bar graph showing each

macrophage subtype as a percentage of all macrophages in tumors of treatment-naïve and

NAC-treated BC patients. NAC results in increases in the Mac_IFN and RTM_IFN subtypes

(black outlines). (**H**) Gene set variation analysis (GSVA) shows that the macrophage subtypes

that dominate the NAC group (Mac_IFN and RTM_IFN, magenta outline) have increased

activation of pro-inflammatory pathways involved in NF-κB and Notch signaling (green outlines).

Data in **A-D** are reported as mean +/- SEM and were analyzed using the Mann-Whitney U test.

*p < 0.05. Each replicate was normalized to its control. All *in vitro* experiments were repeated 4

times.

Figure 4. Paclitaxel skews macrophages towards pro-inflammatory phenotype,

increasing macrophage-tumor cell NF-κB/Notch1 signaling, which induces MenaINV.

Paclitaxel causes macrophages to take on a pro-inflammatory phenotype. Pro-inflammatory

macrophages express higher levels of TNF and Jagged1, which activate NF-κB and Notch1

signaling in tumor cells, respectively. Increased levels of NF-κB and Notch1 activation in tumor

cells leads to increased MenaINV expression. Figure created with BioRender.com.

Supplementary Information

Additional file 1: Materials and Methods (.pdf)

Additional file 2: Supplemental figures 1-2 and supplemental figure legends (.pdf)

Declarations

Ethics approval and consent to participate

All procedures were conducted in accordance with National Institutes of Health regulations and

approved by the Albert Einstein College of Medicine Institutional Animal Care and Use

Committee.

Consent for publication

Not applicable

Availability of data and materials

The datasets used and/or analyzed during the current study are available from the

corresponding author on reasonable request.

Competing interests

The authors declare that they have no competing interests.

Funding

This study was supported by grants from the NIH (R01 CA255153, K99/R00 CA237851), SIG

OD019961, the Gruss-Lipper Biophotonics Center, the Integrated Imaging Program for Cancer

Research, The Evelyn Gruss-Lipper Charitable Foundation, and The Helen & Irving Spatz

Foundation.

Authors' contributions

MFD, MHO, and DE performed conceptualization. MFD, GSK, CLD, SS, JL, and RJE provided

methodology. RJE developed and validated the MenaINV antibody. MFD and JL carried out

formal analysis. SS & DE provided software and computational image analysis. MFD, GSK, and

CLD performed investigation. MFD wrote the manuscript. GSK, JSC, MHO, and DE performed

funding acquisition. MHO and DE supervised the study. All authors read and approved the final

manuscript.

<u>Acknowledgments</u>

We thank the Histopathology Core Facility for their assistance with sample processing and the

Analytical Imaging Facility for their imaging services. We thank Luis Rivera Sanchez for

assistance with macrophage quantification and Ved Sharma for contributing to sample

generation. We thank Junya Zhang for performing the scRNA-seq analysis and Dianne Cox for

advising on macrophage biology and methodology.

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