

Stemness is derived from thyroid cancer cells

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Risheng Ma, Thyroid Research Unit, Department of Medicine, Icahn School of Medicine at Mount Sinai and the James J Peters VA Medical Center, Room 2F-28, 130 West Kingsbridge Road, New York, NY 10468, USA e-mail: risheng.ma@mssm.edu **Background:** One hypothesis for thyroid cancer development is its derivation from thyroid cancer stem cells (CSCs). Such cells could arise via different paths including from mutated resident stem cells within the thyroid gland or via epithelial to mesenchymal transition (EMT) from malignant cells since EMT is known to confer stem-like characteristics. Furthermore, EMT is a critical process for epithelial tumor progression, local invasion, and metastasis formation. In addition, stemness provides cells with therapeutic resistance and is the likely cause of tumor recurrence. However, the relevance of EMT and stemness in thyroid cancer progression has not been extensively studied.

Methods: To examine the status of stemness in thyroid papillary cancer, we employed a murine model of thyroid papillary carcinoma and examined the expression of stemness and EMT using qPCR and histochemistry in mice with a thyroid-specific knock-in of oncogenic Braf (LSL-Braf(^{V600E})/TPO-Cre). This construct is only activated at the time of thyroid peroxidase (TPO) expression in differentiating thyroid cells and cannot be activated by undifferentiated stem cells, which do not express TPO.

Results: There was decreased expression of thyroid-specific genes such as Tg and NIS and increased expression of stemness markers, such as Oct4, Rex1, CD15, and Sox2 in the thyroid carcinoma tissue from 6-week-old BRAF^{V600E} mice indicating the dedifferentiated status of the cells and the fact that stemness was derived in this model from differentiated thyroid cells. The decreased expression of the epithelial marker E-cadherin and increased EMT regulators including Snail, Slug, and TGF- β 1 and TGF- β 3, and the mesenchymal marker vimentin demonstrated the simultaneous progression of EMT and the CSC-like phenotype. Stemness was also found in a cancer thyroid cell line (named Marca cells) derived from one of the murine tumors. In this cell line, we also found that overexpression of Snail caused up-regulation of vimentin expression and up-regulation of stemness markers Oct4, Rex1, and CD15, with enhanced migration ability of the cells. We also showed that TGF- β 1 was able to induce Snail and vimentin expression in the Marca cell thyroid cancer line, indicating the induction of EMT in these cells, and this induction of EMT and stemness was significantly inhibited by celastro a natural inhibitor of neoplastic cells.

Conclusion: Our findings support our earlier hypothesis that stemness in thyroid cancer is derived via EMT rather than from resident thyroid stem cells. In mice with a thyroid-specific knock-in of oncogenic Braf (LSL-Braf(V^{600E})/TPO-Cre), the neoplastic changes were dependent on thyroid cell differentiation and the onset of stemness must have been derived from differentiated thyroid epithelial cells. Furthermore, celastrol suppressed TGF- β 1 induced EMT in thyroid cancer cells and may have therapeutic potential.

Keywords: epithelial-mesenchymal transition, cancer stem cells, thyroid papillary carcinoma, celastrol, thyroid peroxidase, thyroglobulin, Braf

INTRODUCTION

The role of stem cells in thyroid cancer development remains unclear (1). There is no doubt that the thyroid gland retains a significant number of resident stem cells as shown in mice and human thyroid tissue (2–5) but the role of thyroid cancer stem cells (CSCs) in tumor formation and progression is less certain (1, 6). There are two major possibilities. The first is that neoplastic transformation of a normal resident thyroid stem cell takes place

and leads to tumor formation during which the stem cells undergo asymmetric divisions to secure their own survival as we have previously demonstrated (7). The second possibility is that stemness in thyroid cancer is derived directly from malignant thyroid epithelial cells via epithelial to mesenchymal transition (EMT). This is a well-known biologic process defined by the loss of epithelialspecific characteristics, the acquisition of a fibroblast-like morphology, reduced cellular adhesion, and increased motility during embryonic development (8). Cells undergoing EMT develop stem cell-like features such as the ability to self-renew (9). From a molecular point of view, EMT is characterized by the loss of epithelial markers, including E-cadherin and cytokeratins and the gain of mesenchymal-like gene expression program, such as fibronectin, vimentin, and N-cadherin (8). The use of a mouse model in which the malignant transformation can only occur in differentiated thyroid cells expressing thyroid peroxidase (TPO) is a simple approach to answering the question of where stemness is derived from in thyroid cancer. Our data point clearly to the important role of EMT in providing stem-like characteristics in papillary thyroid cancer.

Epithelial to mesenchymal transition is also a commonly accepted key pathologic mechanism in epithelial tumor progression and has been well shown to allow acquisition of stem-like properties by cancer cells (8, 10, 11). The connection between the loss of E-cadherin and the gain of vimentin expression by cancer cells has been established by many studies (12) and has been associated with the development of invasive cancer, metastatic dissemination, and poor clinical prognosis in various human tumors (8) including thyroid carcinomas (13, 14). Several distinct mechanisms of E-cadherin down-regulation and vimentin up-regulation have been described. In particular, transforming growth factor-β (TGF-β) (including TGF-β1, TGF-β2, and TGF- β 3 isoforms) has been shown to regulate the expression of genes that play key roles in a large variety of biological phenomena, ranging from tissue remodeling to tumor initiation and progression (15). TGF-β1 induced overexpression of all stemness genes tested including Oct4, Nanog, Sox2, and CD133 and caused loss of epithelial morphology, assumption of a fibroblast-like appearance, and up-regulation of vimentin, Slug, Twist and down-regulation of cytokeratin and E-cadherin (16). Several zinc-finger transcriptional factors, such as Snail (17, 18), Slug (19), or Twist (20), serve as downstream effectors of the TGFB pathway and have been shown to induce the EMT through direct or indirect suppression of E-cadherin and up-regulation of vimentin transcription in cancer and thus contribute to tumor cell growth, migration, and invasion (8, 21).

In this report, we studied thyroid papillary cancer in mice with a thyroid-specific knock-in of oncogenic Braf [LSL-Braf(^{V600E})/TPO-Cre] (22) in order to determine if stemness was present in such tumors and whether EMT was the source of such characteristics.

MATERIALS AND METHODS

MICE

BrafV600E mice with a thyroid-specific knock-in of BrafV600E were established by crossing LSL-BrafV600E mice, in which a latent Braf mutant knock-in allele can be activated by Cre recombinase through excision of a floxed STOP cassette (23), with TPO-Cre mice, which express Cre under the control of the human TPO promoter (24), which is active only in thyroid follicular cells beginning at E14.5. Mice were in mixed genetic backgrounds. Both LSL-BrafV600E mice and TPO-Cre mice were kindly provided by Dr. James A. Fagin (Memorial Sloan-Kettering Cancer Center). All procedures were approved by the Institutional Animal Care Committee of Icahn School of Medicine at Mount Sinai.

MARCA CELLS – A BRAF^{V600E} CANCER CELL LINE

Primary tumors of Braf^{V600E} mice were minced and resuspended in RPMI 1640 with 10% fetal bovine serum (FBS) containing 100 U/ml type I collagenase (Sigma) and 1 U/ml dispase (Roche). Enzymatic digestion was carried out for 45 min at 37°C. Then, cells were seeded in RPMI 1640 supplied with 10% FBS, 100 IU penicillin/ml, and 100 μ g/ml streptomycin. After growing to confluence, cells were passaged. The doubling time of this line was 35.75 h and the cells retained the Braf mutation as shown by PCR genotyping of DNA isolated from WT and heterozygous LSL-BrafV600E (without Cre) thyroids and the Marca cells (data not shown).

PLASMIDS AND TRANSDUCTION

The pCMV6-Snail vector was electroporated into thyroid cancer cells (Neon transfection system; Invitrogen Life Technologies), and after 2 days, the cells were cultured in G418 (0.8 mg/mL) for 4 weeks until resistant clones were established. High-expressing clones were chosen for expansion into stable lines.

CELL PROLIFERATION AND VIABILITY ASSAYS

Cells were plated in 96-well plates, incubated for 24 h at 37°C, and treated with TGF- β 1 (R&D), 0.5 μ M Celastrol (Sigma) for 48 h. Proliferation of cells was measured using a Cell Counting Kit-8 (Dojindo) according to the manufacturer's protocol.

DIFFERENTIATION OF THYROID CSCs INTO ADIPOCYTES

Marca and MarcaSnail cells were cultured in medium alone or medium supplemented with $10 \,\mu$ g/mL transferrin, 0.5 mg/ mLinsulin, 0.2 nM T3, 1 μ M dexamethasone, and 1 mU/mL TSH for up to 15 days. Cells were then collected for analysis.

RNA ISOLATION AND RT-PCR

Total RNA was extracted from BRAF^{V600E} mice thyroid tumor tissues and normal thyroid gland, and cultured human thyroid cancer cells using the RNeasy system (Qiagen Ltd). cDNA synthesis was performed using the SuperScript III system (Invitrogen Corp.). Quantitative (q)RT-PCR was performed by using the SYBER Green PCR system on a StepOnePlus instrument (Applied Biosystems). Relative expression levels of each RT-PCR product was analyzed using the $2^{-\Delta\Delta CT}$ method and normalized to the expression of the housekeeping gene GAPDH. Data presented (mean) are from three independent experiments in which all sample sets were analyzed in triplicate.

IMMUNOHISTOCHEMISTRY

Paraffin-embedded, formalin-fixed tumor tissues from Braf^{V600E} mice and normal thyroid gland from same strain mice were sectioned into 6 μ m thick slices. Sections were deparaffinized in xylene and rehydrated in a series of graded ethanol, and their antigens were retrieved by heating the slides in a citrate buffer (pH 6.0). Endogenous peroxidase activity was quenched with 0.3% H₂O₂ in methanol for 10 min at room temperature. The sections were then incubated for 20 min with 2.5% normal house blocking serum. This procedure was followed by incubation with the primary antibodies at proper dilution overnight at 4°C. Negative controls were obtained by replacing the primary antibodies with control isotype IgG. Staining was visualized using the ImmPRESSTM reagent

(Vector Laboratories, Burlingame, CA, USA) detection system with peroxidase substrate solution. Slides were counterstained in hematoxylin and coverslipped in synthetic media.

WOUND-HEALING ASSAY

Cells were plated in six-well culture plates in complete culture medium and grown to 90% confluence. A wound was made by scrapping with a sterilized 100 μ l pipette tip in the middle of the cell monolayer. Cells were then cultured with fresh complete culture medium containing 5 ng/ml TGF β with or without 0.5 μ M celastrol treatment for 24 h. After that, the ability of cells to migrate into the cleared section was observed and photographed.

RESULTS

EXPRESSION OF THYROID-SPECIFIC GENES AND STEM CELL MARKERS

qRT-PCR analysis were performed for thyroid-specific markers in total mRNA from 6-week old BRAF^{V600E} murine thyroid tumors and wild type normal thyroid. Expression of Tg and NIS was markedly decreased in the tumor tissue (**Figure 1A**), while the stem cell markers (Oct4, Rex1, CD15, and Sox2), although detectable in normal thyroid, were greatly increased in the neoplastic tissues (**Figure 1B**). These data indicated the undifferentiated state of the BRAF^{V600E} tumor specimens and indicated the presence of cells with CSC-like features. These results were confirmed by immuno-histochemical analysis of Tg, which was barely detectable in the tumor tissue while Oct4 was overexpressed (**Figures 1C–F**).

EXPRESSION OF EMT MARKERS IN THYROID CANCER

The hallmark of EMT is the down-regulation of E-cadherin and the up-regulation of vimentin expression. A significant loss of E-cadherin gene expression and an increase in vimentin gene expression was seen in the BRAF^{V600E} thyroid tumors compared to normal thyroid (**Figure 2A**) and similar data were obtained by immunohistochemistry where vimentin staining was highly positive only in BRAF^{V600E} tumor specimens (**Figures 2B–E**). Morphologically, thyroid cancer cells differed from the polarized, epithelial shape of the wild type thyroid cells, and appeared as a spindle-shaped phenotype (**Figures 2C,E**). The data demonstrated that while E-cadherin was well expressed in normal thyroid tissue, its expression was markedly decreased and vimentin expression up-regulated in the thyroid carcinomas from the transgenic mice over-expressing BRAF^{V600E} (25). Such information supported the transition of thyroid cells into undifferentiated CSC-like cells by passing through EMT (9, 26).

EXPRESSION OF EMT INDUCERS IN THYROID CANCER

TGF- β is a multifunctional cytokine that plays a dual role in cancer; in early stages it inhibits tumor growth, whereas later it promotes invasion and metastasis formation (27–29). TGF- β is also a known major inducer of EMT and its pro-invasive action is thought to be via EMT induction of a series of transcriptional repressors including Snail, Slug, and Twist (28, 30) (**Figure 3A**). We found that TGF- β , Snail, and Slug were highly expressed in the murine thyroid cancer tissue (**Figures 3B,C**). TGF- β binds to TGF- β RII that subsequently phosphorylates the TGF- β RI activin-receptor like kinase 5, which in turn phosphorylates Smad2 and Smad3 (28). Then the complex of Smad proteins translocates into the nucleus where it is able to interact with the Snail and Slug transcription factors and jointly regulate target genes, inducing down-regulation of the epithelial marker E-cadherin and up-regulation of the





FIGURE 2 | Expression of EMT markers in normal mouse thyroid and thyroid tumor. (A) Expression of E-cadherin and vimentin in normal mouse thyroid and BRAF^{veove} mice thyroid tumor by qRT-PCR. The data were normalized using GAPDH as an endogenous control. The results



mesenchymal marker vimentin (31). We found that the expression of TGF- β 1, TGF- β 2R, and TGF- β 3 were all increased in BRAF^{V600E} mice thyroid tumors compared to wild type normal thyroid tissue (**Figure 3B**).

Expression of Snail has been detected in many different types of primary human cancers, including breast, colon, and stomach cancer (32). Here, we observed that expression of both Snail (sixfold) and Slug (twofold) was increased in BRAF^{V600E} murine thyroid

E-cadherin (B,C) and vimentin (D,E) in normal mouse thyroid (B,D) and thyroid tumor (C,E).

three parallels. (B-E) Representative images of immunostaining of

tumor tissue compared to normal thyroid (**Figure 3C**). The results demonstrated an inverse relationship between E-cadherin and TGF- β 1 or Snail expression (33) consistent with TGF- β 1 contributing to the induction of EMT in thyroid cancer via the Snail family of transcription factors.

ECTOPIC EXPRESSION OF SNAIL ENHANCES EMT IN THYROID CANCER CELLS

Increased expression of Snail, Slug, Twist, ZEB1, and ZEB2 expression in thyroid cancers has been well demonstrated (34, 35). Snail appears to play a fundamental role in EMT and its expression is associated with invasiveness, metastases, tumor recurrence, and poor prognosis (36). Snail induction of EMT and CSC-like properties has been described in other cancers including breast and squamous cell carcinoma (37, 38). To study the function of Snail in thyroid cancer cells, we developed a thyroid cancer cell line (the Marca cell line) from the murine tumors initiated by oncogenic Braf (LSL-Braf(V600E)/TPO-Cre) and transfect the cells with pCMV6-SNAIL. Over-expression of Snail after transfection was demonstrated by qRT-PCR (Figure 4A). To test whether the expressed Snail was functional, we performed qRT-PCR for the EMT markers. Expression of the mesenchymal marker vimentin was markedly increased by over-expression of Snail compared to untransfected cells, as detected by qRT-PCR, but did not affect E-cadherin expression (Figure 4B). The stem cell markers Oct4, Rex1, and CD15 were also increased in cells over-expressing Snail compared to the control cells (Figure 4C). Furthermore, after being cultured in adipocyte differentiation medium, early adipocyte differentiation markers such as the fatty acid-binding protein 4 (FABP4) and CCAAT-enhancer-binding protein α (CEBP α) were markedly increased in cells overexpressing Snail compared to the control cells (Figure 4D). The results indicated the multipotential of cells over-expressing Snail. The same morphological changes of elongated and spindle-like



shapes appeared to be further induced by such over-expression. In addition, the migration of cells across an artificial wound was increased with over-expression of Snail (**Figure 4E**). The results indicated that Snail over-expression induced EMT and the CSC-like phenotype in thyroid cancer cells and this enhanced the potential for cancer cell migration.

CELASTROL SUPPRESS TGF-_{β1} INDUCED EMT

As described, there is evidence that TGF- β can induce EMT, which participates in the early stage of metastasis formation in a variety of epithelial tumors and this action is mediated by the Snail signaling pathway (15, 16). Celastrol, a compound used in traditional Chinese medicine, has been reported to suppress cancer cell migration by inhibiting TGF- β 1 induced EMT (39). We found that celastrol $(0.5\,\mu M)$ did not reduce the cancer cell line viability (Figure 5A) but attenuated the effects of TGF- β 1 on stem cell gene expression (Figures 5B,C). While treatment with TGF-B1 induced prominent morphological changes in the thyroid cancer cells, including elongated and spindle-like shapes, they were noticeable suppressed by co-treatment with celastrol (Figure 5D). Co-incubation with celastrol also reversed TGF-β1mediated induction of cell migration. The results were scored using a double-blind method by analyzing several areas of the wound (Figure 5E). These findings showed that celastrol was able to inhibit the effects of TGF- β 1 induced EMT in BRAF^{V600E} cancer cells.

DISCUSSION

In this study, using mice with a thyroid-specific knock-in of oncogenic Braf (LSL-Braf(V600E)/TPO-Cre), the thyroid epithelial cells became transformed and progressed to invasive carcinomas with a very short latency as shown by neoplastic transformation present at birth. These mice also become profoundly hypothyroid due to dysregulation of genes involved in thyroid hormone biosynthesis and consequently develop high TSH levels (22). We demonstrated the down-regulation of thyroid-specific markers (Tg and NIS) in their cancer tissue and this was reflected in isolated thyroid cancer cells derived from the tumors. We also found that expression of the epithelial marker E-cadherin was diminished. In contrast, there was up-regulation of stem cell markers (Oct4, Rex1, CD15, and Sox2) and the mesenchymal marker vimentin as evidenced by immunohistochemistry and RT-PCR. These data confirmed that EMT was induced in the thyroid carcinoma model of $Braf^{V600E}$ mice (25) and, in particular, also demonstrated that thyroid cells acquire differentiated CSC-like properties by passing through EMT rather than deriving from resident stem cells which would not be activated by the TPO-dependent construct initiating the neoplastic response (9, 22, 26).

Epithelial to mesenchymal transition is a normal morphological event during embryonic development, tissue remodeling, and wound-healing but has also been shown to occur in neoplastic cells, especially in metastases (8). By the EMT processes, epithelial cells acquire fibroblast-like properties and exhibit reduced



intercellular adhesion and increased motility providing cells undergoing EMT with cancer stem cell properties including enhanced invasive abilities and chemoresistance (9, 11). Such cells exhibit down-regulated E-cadherin and up-regulated vimentin expression as part of the acquisition of these changes, which have been shown to be induced by TGF- β 1 (40) or increased expression of the transcription factor Snail (30, 37, 41).

Hence, EMT can be provoked by signals that cells receive from their microenvironment, such as TGF-B (15, 42) and several transcription factors have been implicated in the control of EMT in addition to Snail and including Slug, Twist, ZEB1/2, SIP1, and E12/E47 (29, 36, 43) all of which can be induced by TGF- β (18). In agreement, we found enhanced expression of the Snail family of EMT inducers (including Snail and Slug) and TGF-\u03b31, TGF-\u03b33, and TGF-\u03b3RII in the murine papillary thyroid carcinoma samples as shown in previous studies (13, 35, 44). Recent work has also demonstrated that thyroid cancer, which developed in BRAF^{V600E} mice were susceptible to TGF- β 1 induced EMT (25) and BRAF^{V600E} promoted invasiveness of thyroid cancer cells by decreasing E-cadherin expression through a Snail-dependent mechanism (45). We also demonstrated that TGF-B1 or over-expression of Snail dramatically increased the mesenchymal marker vimentin and stem cell markers in thyroid cancer cells derived from BRAF^{V600E} mice. Furthermore, celastrol, a plant triterpene derived from the root of "Thunder of God Vine"

(Tripterygium wilfordii Hook F.), exhibited anticancer potential and inhibited TGF- β 1 induced EMT in the neoplastic cells (39, 46). Our findings demonstrated that celastrol attenuated TGF- β 1 induced EMT and stemness related gene changes and cellular migration. However, the mechanism of this inhibition requires further clarification.

In conclusion, our results suggested that the thyroid papillary cancer of BRAF^{V600E} mice undergo EMT and dedifferentiate to acquire stem cell-like features. TGF- β 1 treatment and overexpression of Snail in thyroid cancer cells derived from BRAF^{V600E} mice further reinforced EMT, while celastrol attenuated the TGF- β 1 effects. Our findings provide strong evidence that thyroid cancer stemness in murine papillary cancer is derived via the EMT process and that thyroid CSCs are not derived from mutated thyroid stem cells. Hence, these data support the stochastic model where every malignant thyroid cancer cell has the potential to act as a CSC rather than CSCs being a biologically distinct subset within the malignant cell population.

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