## MicroRNA miR-100 Decreases Glioblastoma Growth by Targeting SMARCA5 and ErbB3 in Tumor-Initiating Cells

Technology in Cancer Research & Treatment Volume 19: 1-13 © The Author(s) 2020 Article reuse guidelines: sagepub.com/journals-permissions DOI: 10.1177/1533033820960748 journals.sagepub.com/home/tct



Bahauddeen M. Alrfaei, PhD<sup>1,2,3</sup>, Paul Clark, PhD<sup>1,4</sup>, Raghu Vemuganti, PhD<sup>1,2,5,6</sup>, and John S. Kuo, MD PhD<sup>1,2,4,5,7,8</sup>

#### Abstract

Glioblastoma multiforme (GBM) is the most aggressive and most frequently diagnosed malignant human glioma. Despite the best available standard of care (surgery, radiation, and chemotherapy), the median survival of GBM patients is less than 2 years. Many recent studies have indicated that microRNAs (miRNAs) are important for promoting or reducing/limiting GBM growth. In particular, we previously showed that GBMs express decreased levels of miR-100 relative to control tissue and that restoring miR-100 expression reduced GBM tumorigenicity by modulating SMRT/NCOR2 (Nuclear Receptor Corepressor 2). Here, we demonstrate that miR-100 overexpression decreases expression of the stem cell markers, nestin and L1CAM, and decreases proliferation of GBM tumor-initiating cells (cancer stem cells). We further show that miR-100-mediated anti-tumorigenic activity limits the activity of SMARCA5 and its downstream target STAT3 (known as mTOR-STAT3-Notch pathway). In addition, we report ErbB3 (Her3) as a putative miR-100 target, including inhibition of its downstream AKT and ERK signaling pathways.

#### Keywords

SMARCA5, microRNA, miR, ErbB3, GBM stem-like cells, tumor initiating cells

#### Abbreviation

AKT, Protein kinase B, PKB; ATM, ataxia telangiectasia mutated; Bfgf, Fibroblast Growth Factor; CNS, Central nervous system; DMEM, Dulbecco's Modified Eagle's Medium; EGF, Epidermal Growth Factor; EGFR, Epidermal growth factor receptor; ErbB2, Receptor tyrosine-protein kinase erbB-2; ErbB3, Erb-B2 Receptor Tyrosine Kinase 3; ERK, Extracellular Signal-Regulated Kinase; FGFR3, fibroblast growth factor receptor 3; GBM, Glioblastoma; HDAC, Histone deacetylase; LICAM, LI Cell Adhesion Molecule; MiR, microRNA; mRNA, Messenger Ribonucleic Acid; mTOR, Mammalian target of rapamycin; Notch, Notch homolog, translocation-associated (Drosophila); NSC, Neural stem cell; SMARCA5 / Sm5, (SWI/SNF Related, Matrix Associated, Actin Dependent Regulator of Chromatin, Subfamily A, Member 5); SMRT / NCOR2, Nuclear Receptor Co-Repressor 2; STAT3, Signal transducer and activator of transcription 3; TICs / CSCs, Tumor Initiating Cells / Cancer Stem Cells; UTR, Untranslated region

Received: May 15, 2020; Revised: August 12, 2020; Accepted: August 28, 2020.

- <sup>1</sup> Department of Neurological Surgery, University of Wisconsin School of Medicine and Public Health, Madison, WI, USA
- <sup>2</sup> Cellular and Molecular Pathology Training Program, University of Wisconsin School of Medicine and Public Health, Madison, WI, USA
- <sup>3</sup> Department of Cellular Therapy and Cancer Res, King Abdullah Int'l Medical Research Center (KAIMRC), King Saud bin Abdulaziz University for Health Sciences (KSAU-HS), Riyadh, Saudi Arabia
- <sup>4</sup> Carbone Cancer Center, University of Wisconsin School of Medicine and Public Health, Madison, WI, USA
- <sup>5</sup> Neuroscience Training Program, University of Wisconsin School of Medicine and Public Health, Madison, WI, USA
- <sup>6</sup> William S. Middleton Memorial Veterans Hospital, Madison, WI, USA
- <sup>7</sup> Department of Neurosurgery, Dell Medical School, The University of Texas at Austin, Austin, TX, USA

<sup>8</sup> Mulva Clinic for the Neurosciences, The University of Texas at Austin, Austin, TX, USA

#### **Corresponding Authors:**

Bahauddeen M. Alrfaei, King Abdullah International Medical Research Center (KAIMRC), King Saud bin Abdulaziz University for Health Sciences (KSAU-HS), MNGHA, Riyadh 11426, Saudi Arabia. Email: alrfaeiba@ngha.med.sa

John S. Kuo, Department of Neurosurgery, Dell Medical School, The University of Texas at Austin, 1601-B Trinity St, Stop Z1500, Austin, TX 78712, USA.

Email: john.kuo@austin.utexas.edu



Creative Commons Non Commercial CC BY-NC: This article is distributed under the terms of the Creative Commons Attribution-NonCommercial 4.0 License (https://creativecommons.org/licenses/by-nc/4.0/) which permits non-commercial use, reproduction and distribution of the work without further permission provided the original work is attributed as specified on the SAGE and Open Access pages (https://us.sagepub.com/en-us/nam/open-access-at-sage).

## Introduction

Glioblastoma multiforme (GBM), the most aggressive primary brain tumor, accounts for more than 50% of all detected malignant brain cancers and approximately 20% of all primary intracranial tumors.<sup>1,2</sup> Approximately 15,000 new cases of GBM and CNS malignancies are diagnosed annually in the USA.<sup>3</sup> Median patient survival is under 2 years even with the best standard of care.<sup>4,5</sup> The molecular mechanisms responsible for GBM growth and invasion are poorly understood.

MicroRNAs (miRNAs) are small, non-coding RNAs (16-22 nucleotides long) that repress protein translation by binding to the 3'UTRs of mRNAs.<sup>6,7</sup> Many miRNAs are known to be differentially expressed in various types of cancer<sup>8</sup>; it is therefore essential to understand their role in tumorigenesis.<sup>9</sup> At least 19 miRNAs have been identified as being linked to the pathogenesis of GBM.<sup>10,11</sup> Of these, miR-100 has been linked to several targets that are known to modulate GBM growth and/ or survival, such as fibroblast growth factor receptor 3 (FGFR3), silencing mediator of retinoic acid and thyroid hormone receptor (SMRT), and ATM Serine/Threonine Kinase (ATM; ataxia telangiectasia mutated).<sup>12-14</sup> In addition, miR-100 was reported to stimulate a positive therapeutic response in breast cancer stem-like cells undergoing hormonal therapy.<sup>15</sup> Moreover, it was demonstrated that gliomas' proliferation, apoptosis and angiogenesis were suppressed by inhibition of AKT and STAT3 signaling pathway.<sup>16</sup> The same pathways were effective against limiting glioma stem cells propagation.<sup>17</sup> Also, it has been reported that Akt-mTOR pathway regulates neurogenesis of neural stem cells.<sup>18</sup> The same pathway (Akt-mTOR) is involved in ErbB family activation and contributes to cancer stem cells resistance which make it eligible for EGFR-Targeted therapy.<sup>19</sup>

We previously demonstrated that downregulation of miR-100 promotes GBM growth and invasion, and that restoring miR-100 expression reduces GBM growth and survival.<sup>13</sup> We also showed that miR-100 limits GBM tumor proliferation and extends the survival of mice bearing orthotopic GBM xenografts by inhibiting the miR-100 target, SMRT/NCOR2.13 These findings were subsequently supported by Luan et al. (2015), who also confirmed the tumor suppressor activity of miR-100 and examined 2 GBM cell lines (U251 and T98G) in addition to 13 patient GBM specimens. They confirmed the decrease in endogenous levels of miR-100 in all tested tumor samples and in both cell lines, and compared GBM tumors to adjacent normal tissue. When miR-100 was overexpressed in GBM cell lines, reduced proliferation, migration, and chemosensitivity were observed. The authors concluded that miR-100 has anti-tumor activity against GBMs.

Since a single miRNA can regulate multiple target mRNAs, the beneficial effects of miR-100 might also be mediated by other target proteins in addition to SMRT/NCOR2. In this study, we have uncovered and evaluated additional pathways that might also be regulated by miR-100 and potentially contribute to its anti-tumorigenic activity. In particular, we evaluated the role of ErbB family members that we found are Technology in Cancer Research & Treatment

targeted by miR-100. Members of this protein family are known to be resistant to EGFR inhibitor therapy especially in GBM stem-like cells.<sup>19</sup> Two members of the ErbB family Her2 (ErbB2) and Her3 (ErbB3) are being targeted by immunotherapy in clinical trials designed to control GBM growth.<sup>20,21</sup>

In this study, we restored miR-100 levels in primary GBM cells by transfecting GBM tumor-initiating cells (TICs) with pre-miR-100, which generates 2 forms of miR-100: miR-100-5p and miR-100-3p. We found that the dominant isoform, miR-100-5p,<sup>22</sup> is down-regulated in GBM-derived TICs (also called GBM stem-like cells) known to be responsible for tumor progression and recurrence<sup>22</sup> because of radiation and chemo-resistance.<sup>23</sup> Thus, in this study, we tested the therapeutic utility of miR-100-5p overexpression for controlling GBM growth and also evaluated the downstream mechanism of miR-100-5p activity.

### **Materials and Methods**

### Isolation and Validation of GBM TICs

All human tumor specimens were collected after patient informed consent and with approval of University of Wisconsin-Madison Institutional Review Board No. (IRB 2012-0024-Certified, exempt). Patient-derived TICs were isolated from GBMs and validated as previously described.<sup>19,24-26</sup> Tumor tissue collected from the operating room was minced and chopped twice using a tissue chopper (Sorvall TC-2 Smith-Farquhar) and plated in medium (70%) Dulbecco's modified Eagle medium-high glucose, 30% Ham's F12, 1X B27 supplement, 5 µg/ml heparin, penicillinstreptomycin-amphotericin, and 20 ng/ml each of epidermal growth factor (EGF) and basic fibroblast growth factor (bFGF)). We used 3 primary (22, 33, and 44) GBM samples and 1 recurrent GBM sample (12.1). All of the GBM TICs were negative for EGFRvIII mutations.<sup>19</sup> A tumor-free neural stem cell (NSC) line prepared from fetal human cortical tissue and used as a source of normal control cells was a kind gift of Dr. Clive Svendsen.<sup>26</sup> Validation of TICs was done through neurosphere isolation and differentiation procedures beside staining for stem cells markers was done, Supp. Figure 1. More details on isolation, validation and neutrosphere's formation of TICs could be found in our previous publications.<sup>19,27,28</sup>

## Real-Time (Quantitative) PCR

Total RNA was isolated using an RNA isolation kit (Life Technologies). All probes and primers were purchased from Life Technologies, USA, and absolute quantitative PCR was conducted using TaqMan assays according to manufacturer's instructions (Life Technologies, USA). 18 s rRNA was used as a house-keeping control as described previously.<sup>19</sup> Relative expression was measured by subtraction targets ct from 18 s ct and the sum was further subtracted from control samples ct. The final results were applied on 2<sup>°</sup>(–delta delta CT). Calculation method was reported previously.<sup>29</sup> The primers has-miR-



**Figure 1.** Overexpression of pre-miR-100 reduces proliferation and targets SMARCA5. (A) qPCR shows that the expression of miR100-5p is lower in GBM TICs (12.1, 22, 33, and 44) compared to human neural stem cells lines (hnsc and hnsc2). (B) Percentage of normalized cell growth showing that the overexpression of pre-miR-100 reduces the number of GBM TICs compared to controls as determined by the MTS assay. (C) The targets of miR-100-5p as predicted by 3 algorithms: PICTAR, TARGET SCAN, and microRNA.org. The intersection of the circles represents the number of shared targets. (D) The overexpression of pre-miR-100 decreases proliferation in GBM TICs compared to the control. The results were normalized for pre-miR-100 expression (E) Immunoblots showing the reduction of SMARCA5-encoded HSNF2 H protein levels following the transient expression of pre-miR-100. The reduction in HSNF2 H levels does not occur when control miR is overexpressed. The boxes are included to make the blot easier to interpret. (F) The quantification of the immunoblot in panel (E) shows that HSNF2 H levels are reduced by 40-70% when pre-miR-100 is overexpressed. (G) Pre-miR-100 inhibits the luciferase signal from the SMARCA5 3'UTR reporter compared to controls that utilize a different miRNA or a reporter with a mutated target site. The host cells used were HEK293 T cells. Asterisk denotes statistical significance of p < 0.05.

100-5p, has-miR-100-3p, and has-pre-miR-100 were used for qPCRs assay and were commercially bought from Invitrogen, USA.

## MiRNA and siRNA Transfection

Cells were transfected with miRNA precursors, siRNAs, miRNA mimics and the miR-100 isoforms miR-100-5p and miR-100-3p (all from Life Technologies) using 15 pmol Pep-Mute reagent (SignaGen Labs, Rockville MD) per million cells as described previously.<sup>19,20</sup>

#### Cell Viability Assay

GBM TICs were grown as spheres, disseminated into single cells, then inoculated in 96-well plates at a density of 20,000 cells/well and temperature of 37 oC. After 1 day of growth, the cells were transfected with pre-miR-100 or with a control miR. Following 2 days of growth in culture medium containing 4.5 g glucose DMEM/F12, 20 pg EGF and 20 pg bFGF, cell numbers were quantified using an MTS assay (CellTiter 96 Aqueous, Promega, USA) as per the manufacturer's instructions.

### Cell Proliferation Assay

The Click-iT EdU assay (which is similar to the BrdU assay) was performed according to the manufacturer's (Invitrogen) instructions. Twenty thousand cells were plated and then transfected with pre-miR-100 or control miRs in combination with the SMRT/NCOR2 expression vector (pSMRT, Fisher Scientific) after 1 day of growth. The Click-iT EdU assay was performed following an additional 2 days of growth in culture medium containing DMEM/F12, EGF and bFGF.

#### Luciferase Reporter Assay

293 T cells were co-transfected with pre-miR (15 pmol) and a luciferase reporter plasmid (1 ug) containing the 3'UTR of SMARCA5 or ErbB3 mRNA in 96-well culture plates. Twentyfour to 48 hours after transfection, the light switch luciferase assay (Switch Gear Genomics, Menlo Park, CA) was performed according to the manufacturer's instructions. The signal was detected using a microplate luminometer (Turner Biosystems, Inc., CA) running Veritas software version 1.9.2. The reporter plasmids contained the 3'UTRs of SMARCA5 or ErbB3 mRNA, which include the miR-100 seed sequence. The control plasmids (negative controls) contained a mutated seed sequence.

### Western Blotting

Western blotting was performed as previously described.<sup>19,30</sup> Cell lysates were collected, and the protein concentration was determined by the Bradford assay (Bio-Rad, CA). The protein samples were electrophoresed on SDS-PAGE gels, transferred to PVDF membranes, and analyzed using antibodies against  $\alpha$ -tubulin, phospho-AKT (S473), total-AKT, phospho-ERK (Thr202/Tyr204), total-ERK, phospho-STAT3 (Tyr705),

STAT3, p21, SMARCA5 (Santa Cruz Biotechnology, USA), SMRT/NCOR2 (Santa Cruz Biotechnology, USA), ErbB3 (Santa Cruz Biotechnology, USA), ErbB2 (Santa Cruz Biotechnology, USA), human nestin (Santa Cruz Biotechnology, USA), and L1CAM (Fisher, USA). All antibodies were purchased from Cell Signaling, USA, unless otherwise indicated.

## Tumor Xenograft Assay

UW-Madison institution-approved animal protocol was followed for all experimental procedures. Tumor xenografts were generated via stereotactic implantation of tumor cells as described previously.<sup>13</sup> Briefly, GBM cells were enzymatically dissociated into single cells. One million cells were suspended in 5  $\mu$ l of PBS and stereotactically implanted at 0.33  $\mu$ l/min into the right striatum of anesthetized immunodeficient 6-8 weeks old NOD-SCID mice (Jackson lab) at the following coordinates referenced from bregma: 0 mm anteroposterior, +2.5 mm mediolateral, and -3.5 mm dorsoventral.<sup>31</sup> Xenograft growth was detected and verified by MRI, and brains containing the xenografts were obtained from the animals after death.

## Inducible and Stable Expression of Pre-miR-100 and Immunohistochemical Analysis

Tumor xenograft model has been previously described.<sup>13</sup> Briefly, The 22 T and U87 GBM cell lines were orthotopically implanted into immunodeficient NOD-SCID mice. Control and miR-100 doxycycline inducible overexpression vectors (Gene-Copoeia, MD) were created and then integrated into the genomes of tumor cells with lentiviruses. Later, at least 1 million cells were implanted into mouse brains. The cells holding vectors resulted in a 3-fold increase in expression above baseline miR-100 levels were isolated. Those vectors were validated as described previously.<sup>13</sup> When severe clinical symptoms were detected, or the mice were moribund, they were sacrificed. Immunohistochemistry was carried out on mouse brain 16 days after transplantation. The brains of the mice were formalinfixed, paraffin-embedded, sectioned (into 5 µm-thick sections), and stained with hematoxylin, as described previously.<sup>32</sup> The animal implantation procedure, validation and Ki-67 index were previously reported.<sup>13</sup>

#### Statistics

The statistical analyses were performed using Student's t-test and 1-way ANOVA/Tukey's multiple comparison post-tests. All error bars represent the standard error of the mean (S.E.M.), and the significance level (\*) was P < 0.05.

#### Results

## Expression of miR-100-5p Is Down-Regulated in GBM Cells

Quantitative PCR revealed that expression of endogenous miR-100-5p was reduced by approximately 50–80% in the 4 GBM

Rank	MicroRNA.org	PICTAR	TARGETSCAN 5.2
1	TMPRSS13 (NM_001077263)	SMARCA5 (NM_003601)	THAP2 (NM_031435)
2	SMARCA5 (NM_003601)	BAZ2A (NM_013449)	KBTBD8 (NM_032505)
3	ANKAR (NM_144708)	HS3ST3B1 (NM_006041)	HS3ST3B1 (NM_006041)
4	ICK* (NM 014920)	HS3ST2 (NM 006043)	HS3ST2 (NM 006043)
5	AP1AR* (NM 018569)	FRAP1 (NM 004958)	CTDSPL (NM 005808)
6	NCOR2* (NM_001077261)	EIF2C2 (NM_012154)	SMARCA5 (NM_003601)

Table 1. Predicted Targets of MIR-100.

A list of the top 6 predicted targets of microrna-100-5p (miR-100-5p) and their transcript ID numbers as they rank according to 3 different algorithms: microRNA. org, PICTAR, and TARGETSCAN. The asterisk (\*) denotes targets that have the same score and thus can be given the same rank as the highest ranked candidate. Shading represents repeated appearance of SMARCA5.

tumor cell populations (12.1, 22, 33, and 44) evaluated relative to the levels of endogenous miR-100-5p in 2 different control normal human neural stem cell (hnsc) lines (n = 3/group; P < 0.05; Figure 1A). The results were normalized for pre-miR-100 expression.

## Restoring MiR-100 Levels Decreased Cell Viability and Proliferation

When the TIC cell lines were transiently transfected with premiR-100, we observed a 20–50% decrease in cell viability (n = 3/group; p < 0.05; Figure 1B) and a 20–40% decrease in cell proliferation (n = 3/group; p < 0.05; Figure 1D) compared to the respective controls transfected with a control miRNA. It worth noting that transient transfection using pre-miR-100 generates 200-300% more of pre-miR-100 expression than Hnsc baseline.

## SMARCA5 mRNA Is a Target of miR-100-5p

Using 3 target identification algorithms (PICTAR, TARGETS-CAN, and microRNA.org), we found 18 common putative targets of miR-100-5p (Figure 1C). SMARCA5 was the only common target that was among the top 6 targets identified by all 3 algorithms (Table 1). When the TICs (12.1, 22, 33, and 44) were transfected with pre-miR-100, the protein levels of HSNF2 H (encoded by SMARCA5) were approximately 40-70% lower than observed in the control transfected cell lines (p < 0.05; n = 3; Figure 1E, F). We further confirmed the miRtarget relationship by co-expressing a SMARCA5 3'UTR luciferase reporter vector with pre-miR-100 or with miR-100-5p. When 293 T cells were co-transfected with the SMARCA5 3'UTR and with either pre-miR-100 or miR-100-5p, SMARCA5 3'UTR luciferase activity was inhibited by 45% (p < 0.05; n = 3) compared to cells transfected with the control miRNA (Figure 1G).

## Reduced Activation of Signaling Proteins AKT, ERK, and ErbB3 Are Involved in miR-100-5p Expression

In the cell lines treated with pre-miR-100, there was a significant reduction in the phosphorylation of AKT and ERK relative to the cell lines transfected with the control premiRNA (Figure 2A-D; p < 0.05; n = 3/group). Treatment with siRNAs that are specific to SMRT or SMARCA5 also significantly inhibited AKT and ERK phosphorylation (Figure 2A-D; p < 0.05; n = 3). Furthermore, the knockdown of ErbB3 (it functions upstream of AKT and ERK) significantly suppressed the phosphorylation of AKT and ERK in TICs (Figure 2A-D).

Pre-miR-100 releases 2 mature miRNAs, miR-100-5p and miR-100-3p. Overexpression of Pre-miR-100 reduces ErbB3 expression in all TICs lines. Interestingly, ErbB3 expression decreases when SMRT, and SMARCA5 get inhibited which are targets of miR-100 (Figure 3A and B; p < 0.05; n = 3). Since ErbB3 is upstream of AKT and ERK, it is possible that loss of ErbB3 leads to a reduction in downstream signaling. Knockdown of SMRT and SMARCA5 (Sm5) genes by inhibitory siRNA, reduce ErbB3 levels. Our bioinformatics analysis showed that the miR-100-3p targets ErbB3 (Figure 3C). We found that expression of miR-100-3p was 40–80% lower in GBM TICs than in control NSCs (Figure 3D; p < 0.05; n = 3). A luciferase reporter assay showed that miR-100-3p significantly inhibited the expression of the ErbB3 3'UTR vector (Figure 3E; P < 0.002; n = 3).

## SMRT and SMARCA5 Controls STAT3

When the GBM TIC lines were transfected with pre-miR-100, SMRT siRNA, or SMARCA5 siRNA, phosphorylation of STAT3 (a marker of STAT3 activation) was completely inhibited in all 4 of the tested cell lines (Figure 4A, B; p < 0.05; n = 3).

## Overexpression of miR-100-5p Upregulates p21

Western blot analysis showed that expression of the cell cycle inhibitor p21 (WAF1 or cyclin-dependent kinase inhibitor 1) was upregulated by 50–300% when the various GBM TICs were treated with pre-miR-100, miR-100-5p, or SMRT siRNA (Figure 4C, D; P < 0.05; n = 3).

### Pre-miR-100 Treatment Decreases Stem Cell Markers

Nestin and L1CAM are both stem cell markers that are normally expressed by GBM TICs. When the GBM TICs were



**Figure 2.** The activity of the AKT and ERK pathways is reduced following the overexpression of pre-miR-100. (A) Immunoblot showing the reduction in phospho-AKT protein levels following the transfection of GBM TICs with pre-miR-100 or with the following siRNAs: siErbB3, siSMRT, or siSm5 (siSMARCA5). This pattern is not observed in GBM TICs transfected with control miRNA. The boxes are included to make the blot easier to interpret. (B) Quantification of protein levels in panel (A) reveals that their reduction ranges from 20% to 60%. (C) Immunoblot showing the inhibition of phospho-ERK following the transfection of GBM TICs with pre-miR-100 or siErbB3. No inhibition of phospho-ERK is observed in the control. The boxes are included to make the blot easier to interpret. Total ERK (T-ERK) was measured to ensure that effect is only at phosphorylation level and not total protein. (D) Quantification of the protein levels in panel (C) shows that the reduction of phospho-ERK ranges from 10% to 70%. Asterisk denotes statistical significance of p < 0.05.

transiently transfected with pre-miR-100 or with miR-100-5p, there was a 40–75% decrease in nestin levels (Figure 5A, B; P < 0.05; n = 3) and a 60% decrease in L1CAM levels (Figure 5C, D; p < 0.05; n = 3) compared to the control miRNA-transfected cell lines.

# Stable Overexpression of Pre-miR-100 Prevents ErbB2 and ErbB3 Expression

As mentioned above tumor xenograft model has been previously described.<sup>13</sup> The results of inducing pre-miR-100 around



**Figure 3.** Overexpression of pre-miR-100 targets ErbB3 mRNA and decreases ErbB3 protein levels. (A) Verification of ErbB3 inhibition following transfection with pre-miR-100, siErbB3, siSMRT, and Sm5 (siSMARCA5). The boxes are included to make the blot easier to interpret. (B) Quantification of panel (A) shows a 30%-80% reduction in ErbB3 (Her3) protein levels following transfection with pre-miR-100, siErbB3, siSMRT, or Sm5 (SMARCA5 siRNA). The control does not reduce ErbB3 expression. (C) Diagram showing the predicted binding of miR-100-3p to ErbB3 mRNA. (D) Internal expression of miR-100-3p in GBM TICs (12.1, 22, 33, and 44) compared to human neural stem cells (hnsc) as detected by qPCR. (E) miR-100-3p lessens the signal from the ErbB3 3'UTR luciferase reporter compared to controls that utilize a different miRNA or a reporter with a mutated target site. The host cells used were HEK293 T cells. Asterisk denotes statistical significance of p < 0.05. Backgrounds of some images were modified for clarity.

3 folds above base line showed lower expression of ErbB2 and ErbB3 in mice bearing GBM cells transfected with the premiR100-expression vector than in mice transplanted with GBM cells transfected with the control vector (Figure 6 A, B; P < 0.05; n = 8).

#### Discussion

We previously reported that the down-regulation of miR-100, and the subsequent reduction in levels of its target SMRT/ NCOR2, can contribute to GBM tumorigenicity.<sup>13</sup> We also



**Figure 4.** Overexpression of pre-miR-100 reduces phospho-STAT3 and increases p21 (WAF1). (A) Immunoblot showing a reduction in phospho-STAT3 when GBM TICs are transfected with pre-miR-100, siErbB3, siSMRT, or Sm5 (SMARCA5). This pattern is not observed following transfection with the controls. The boxes are included to make the blot easier to interpret. Total STAT3 (T-STAT3) shows no change in total protein level. (B) Quantification of the protein levels in panel (A) shows a reduction 60%–90% of phospho-STAT3 except for the control and siErbB3. (C) p21 protein levels are elevated in response to miR-100-5p overexpression or SMRT silencing. (D) Quantification of the protein levels in panel (C) shows elevated p21 levels compared to the control by 20%–300%. Asterisk denotes statistical significance of p < 0.05. Backgrounds of some images were modified for clarity.

showed previously that restoring miR-100 levels reduces GBM tumorigenicity and extends the survival of animals bearing orthotopic GBM xenografts.<sup>13</sup> It worth noting that normal distribution of miR-100-5p expression in normal tissue is the

highest in CNS compared to other organs based on TissueAtlas database (See supp. Figure 1).<sup>33</sup> Our results here demonstrate that miR-100 overexpression, by using pre-miR-100 to replenish miR-100 levels in GBM cells, activates the cell cycle



**Figure 5.** Overexpression of miR-100-5p reduces stem cell markers. (A) Immunoblot showing a reduction in the stem cell marker nestin, following the overexpression of pre-miR-100 or of miR-100-5p but not of the control miRNA. The boxes are included to make the blot easier to interpret. (B) Quantification of protein levels in panel (A) represent reduction of 40%–80% as compared to control. (C) Immunoblot showing the reduction in protein levels of the stem cell marker L1CAM when GBM TICs were transfected with pre-miR-100 (miR-100-5p), siSMRT, or Sm5 (siSMARCA5). This reduction was not observed after transfection with the control miRNA. The boxes are included to make the blot easier to interpret. (D) Quantification of the protein levels in panel (C) showing L1CAM reduction of 30%–70% compared to the control. Asterisk denotes statistical significance of p < 0.05. Backgrounds of some images were modified for clarity.

inhibitor p21 and inhibits pathways that are important for cell survival and proliferation (e.g., AKT, ERK, and STAT3), and targets SMARCA5 and ErbB3 (Her3).

We observed lower abundance of miR-100-5p in patientderived GBM TICs compared to human neural stem cells, and found that transfection with pre-miR-100 decreased the proliferation and viability of GBM TICs. Further, in silico analysis to TCGA database we found that median expression of miR-100-5p for GBM is 10.75 units while for low grade glioma is 7806.94 units. Low grade glioma is around 780 times higher. These data suggest that miR-100 probably plays an important role in limiting GBM growth by modulating 1 or



**Figure 6.** Overexpression of pre-miR-100 decreases the expression of ErbB2 and ErbB3 *in vivo*. (A) Mouse NOD-SCID GBM xenografted brain tissue stably overexpressing pre-miR-100 exhibits less ErbB2 (Her2) expression than the control tissue overexpressing the vector alone. The darker the brown staining, the higher expression of Her2 antigen. The more positive stain is scored with more pluses. The pre-miR-100 group in picture 2, is scored with 1 plus (+), and the control tissue is scored with 3 pluses (+ + +) in picture 1. (B) GBM xenograft tissue stably overexpressing pre-miR-100 exhibits less ErbB3 (Her3) expression than control tissue overexpressing the vector alone. The pre-miR-100 group is scored with questionable positivity (+/-) in picture 4, and the control group is scored with 2 pluses (+ +) in picture 3. Scale bar: 50  $\mu$ m.

more of its targets, in addition to previously identified SMRT/ NCOR2.<sup>13</sup>

By further refining the bioinformatics analysis of miR-100-5p targets, we identified SMARCA5 (SWI/SNF-related matrixassociated actin-dependent regulator of chromatin subfamily A member 5) as a target of miR-100-5p. When we combined the target predictions provided by the microRNA.org, PICTAR, and TARGETSCAN algorithms, only 18 targets were identified in common (Figure 1C). We have calculated the prediction score from the 3 different systems and reported the top 6 candidates. Of these, SMARCA5 was the only 1 with a highest score collectively (Table 1). SMARCA5 is reported to be highly expressed in proliferating stem and progenitor cells, and its activity disappears or is reduced considerably in terminally differentiated cells.<sup>34,35</sup> In addition, in silico analysis of circSMARCA5 by Barbagallo et al. revealed SMARCA5 as a potential malignancy inhibitor to GBM expansion.<sup>36</sup> Thus, we investigated whether SMARCA5 also plays a role in the down-regulation of miR-100-5p in GBM tumors and in the subsequent effect on GBM growth. Luciferase reporter assays confirmed that SMARCA5 is a target of miR-100 (Figure 1G). In addition, we have demonstrated that miR-100 expression reduces the expression of SNF2 H (the *SMARCA5*-encoded protein) in GBM TICs by approximately 50% (Figure 1G). This finding suggests that miR-100-5p silences the *SMARCA5*-encoded transcript and that the SNF2 H protein might play a role in GBM tumorigenesis.

It was reported that SMARCA5 interacts with HDAC3<sup>37,38</sup> which modulates AKT. It is also known that AKT controls cell survival and proliferation.<sup>39,40</sup> In our study, we observed that the knockdown of SMRT or of SMARCA5, or overexpression of pre-miR-100, reduced the phosphorylation of AKT and ERK in GBM TICs. ErbB (Her) family proteins are known to act upstream of both the AKT and ERK pathways.<sup>41,42</sup> The

knockdown of SMRT or SMARCA5, or transfection with premiR-100, significantly decreased ErbB3 (Her3) protein levels in GBM TICs. However, the precise mechanism by which ErbB3 protein levels are reduced is not clear because ErbB3 lies upstream of the AKT and ERK pathways. Interestingly, our bioinformatics analysis revealed that ErbB3 is a target of miR-100-3p, the less predominant form of miR-100 that is released from pre-miR-100. In this work, ErbB3 3'UTR luciferase vector experiments confirmed this miR-target relationship (Figure 3E). The data related to miR-100-3p was an interesting observation that was not explored previously. TCGA has no miR-100-3p expression data listed.<sup>43</sup> Thus, we can not correlate it with patients survival. However, assessing our data showed better outcome with miR-100 precursor (miR-100-5p + miR-100-3p) than miR-100-5p alone. The precursor in this case is more effective because it targets 3 or more genes rather than 2 oncogenic genes. The miR-100 precursor targets SMRT, SMARCA5, and EebB3 While miR-100-5p targets ErbB3 in our assays.

Recent studies have shown that SMARCA5 needs to be highly expressed for stem cell self-renewal<sup>44</sup> and that HDAC3 is required for the function of the SMARCA5-encoded protein SNF2 H.<sup>38</sup> Furthermore, SMRT is essential for HDAC3 activity,<sup>45-47</sup> and the induction of a DNA-damage response by SMRT inhibition leads to increased p21 levels.48-50 Consistent with this finding, we observed that p21 protein levels were increased by 100% in GBM TIC lines transfected for miR-100-5p overexpression, or with transfected with an siRNA that silences SMRT relative to control cells (Figure 4C). This p21 induction resulting from miR-100 activity suggests that GBM TIC proliferation will be inhibited. There is an inverse relationship between p21 levels and STAT3 activation,<sup>51-53</sup> and our results showed that the overexpression of either pre-miR-100 or miR-100-5p in GBM TICs almost completely blocked STAT3 phosphorylation (Figure 4A, B). This suggests that miR100-mediated STAT3 inhibition can exert downstream effects on GBM growth. It worth noting that STAT3 phosphorylation at position Tyr705 works under interaction of mTOR-STAT3-Notch signaling pathway known to inhibit glioma growth and regulate stem cells.54,55

The Notch1 pathway regulates SMRT activity which is known to be involved in cancer stemness.<sup>56,57</sup> Further, SMARCA5 inhibition stimulates differentiation<sup>58,59</sup> in GBM TICs which were tested for the loss of stem cell markers. The protein levels of both nestin and L1CAM were significantly lower in GBM TIC cells overexpressing miR-100-5p than in control cells (Figure 5A and D). As expected, knockdown of SMARCA5 and SMRT also diminished L1CAM protein levels (Figure 5D). The inhibition of L1CAM (CD171) was previously reported to decrease DNA damage repair (ATM; a protein kinase that is recruited and activated by DNA doublestrand breaks) in GBM TICs through inhibition of NBS1 which activates ATM.<sup>60</sup> This suggests that the loss of miR-100-5p in GBMs plays a role in maintaining the 'stem cell-like' status of GBM TICs. Similarly, miR-100 was found to target the SMARCA5-encoded protein SNF2 H in adenocarcinoma

cells.<sup>61</sup> Thus, SMARCA5 may partially play an important role in cancer by maintaining the de-differentiated state of cancer stem-like cells.

Mice implanted with GBM TICs that were transfected with stable pre-miR-100 expression or inducible pre-miR-100 expression vectors showed tumor xenograft immunostaining that revealed decreased ErbB2 (Her2) and ErbB3 (Her3) protein levels in the pre-miR-100 group compared to the control group (Figure 6A, B).

This study provides evidence that miR-100 indirectly inhibits 3 major pathways (STAT3, AKT, and ERK) in GBM TICs. STAT3 is inhibited by the up-regulation of p21 that results from the miR-100-mediated down-regulation of SMRT and/ or SMARCA5. AKT and ERK are inhibited by the suppression of either HDAC or ErbB3. The proliferation of GBM TICs was markedly reduced, and differentiation was induced following transfection with pre-miR-100.

In conclusion, our studies show that miR-100 targets many pathways that might contribute to tumorigenicity. A recent randomized clinical trial showed that miR-100 is also responsible for vitamin D-induced tumor suppression in primary prostate cancer.<sup>62</sup> Furthermore, many studies have linked miR-100 with sensitizing tumors to radiotherapy.<sup>12,63,64</sup> Findings reported in this study suggest that altering miR-100 levels and its downstream effects (possibly with pre-miR-100 or analogs) might have potential clinical applications in GBM treatment strategies.

#### Acknowledgments

Appreciation to TCGA database and OncoLnc which have been used to analyze and describe data in this manuscript.<sup>41</sup> In addition, human miRNA tissue atlas has been used to analyze and describe data in this manuscript.

#### **Declaration of Conflicting Interests**

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

#### **Ethical Approval Statement**

All human tumor specimens were collected after patient informed consent and with approval of University of Wisconsin-Madison Institutional Review Board No. (IRB 2012-0024—Certified, exempt). Also, all animal experiments were approved by institutional animal care and use committee of University of Wisconsin-Madison.

#### Funding

The author(s) disclosed receipt of the following financial support for the research, authorship, and/or publication of this article: We appreciate support from Saudi Ministry of National Guard RC13/258/R (King Abdullah Int'l Medical Research Center, KAIMRC) to BMA, and NIH T32GM007507, UL1RR025011, RC4AA020476, NCI HHSN261201000130C, P30CA014520 grants, the Wisconsin Partnership Program core grant support from Center for Stem Cell and Regenerative Medicine, from the University of Wisconsin (Graduate School and Dept. of Neurological Surgery), and the HEADRUSH Brain Tumor Research Professorship and Roger Loff Memorial Fund for GBM Research to JSK. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

#### ORCID iD

Bahauddeen M. Alrfaei D https://orcid.org/0000-0002-5740-9187

#### Supplemental Material

Supplemental material for this article is available online.

#### References

- Van Meir EG, Hadjipanayis CG, Norden AD, Shu HK, Wen PY, Olson JJ. Exciting new advances in neuro-oncology: the avenue to a cure for malignant glioma. *CA Cancer J Clin.* 2010;60(3): 166-193.
- Sauvageot CME, Weatherbee JL, Kesari S, et al. Efficacy of the HSP90 inhibitor 17-AAG in human glioma cell lines and tumorigenic glioma stem cells. *Neuro Oncol.* 2008;11(2):109-121.
- Kruchko C, Ostrom QT, Gittleman H, Barnholtz-Sloan JS. The CBTRUS Story: Providing Accurate Population-Based Statistics on Brain and Other Central Nervous System Tumors for Everyone. Oxford University Press; 2018.
- Stupp R, Mason WP, van den Bent MJ, et al. Radiotherapy plus concomitant and adjuvant temozolomide for glioblastoma. *N Engl J Med.* 2005;352(10):987-996.
- Hegi ME, Diserens AC, Gorlia T, et al. MGMT gene silencing and benefit from temozolomide in glioblastoma. *N Engl J Med.* 2005; 352(10):997-1003.
- Bartel DP.MicroRNAs: target recognition and regulatory functions. *Cell*. 2009;136(2):215-233.
- Filipowicz W, Bhattacharyya SN, Sonenberg N. Mechanisms of post-transcriptional regulation by microRNAs: are the answers in sight? *Nat Rev Genet*. 2008;9(2):102-114.
- Lu J, Getz G, Miska EA, et al. MicroRNA expression profiles classify human cancers. *Nature*. 2005;435(7043):834-838.
- Volinia S, Calin GA, Liu CG, et al. A microRNA expression signature of human solid tumors defines cancer gene targets. *Proc Natl Acad Sci U S A*. 2006;103(7):2257-2261.
- Visani M, de Biase D, Marucci G, et al. Expression of 19 micro-RNAs in glioblastoma and comparison with other brain neoplasia of grades I–III. *Mole Oncol.* 2014;8(2):417-430.
- Brower JV, Clark PA, Lyon W, Kuo JS. MicroRNAs in cancer: glioblastoma and glioblastoma cancer stem cells. *Neurochem Int.* 2014;77:68-77.
- Luan Y, Zhang S, Zuo L, Zhou L. Overexpression of miR-100 inhibits cell proliferation, migration, and chemosensitivity in human glioblastoma through FGFR3. *Onco Targets Ther.* 2015; 8:3391.
- Alrfaei BM, Vemuganti R, Kuo JS. MicroRNA-100 targets SMRT/NCOR2, reduces proliferation, and improves survival in glioblastoma animal models. *PLOS ONE*. 2013;8(11):e80865.
- Ng WL, Yan D, Zhang X, Mo YY, Wang Y. Over-expression of miR-100 is responsible for the low-expression of ATM in the human glioma cell line: M059 J. DNA Repair. 2010;9(11): 1170-1175.

- Petrelli A, Carollo R, Cargnelutti M, et al. By promoting cell differentiation, miR-100 sensitizes basal-like breast cancer stem cells to hormonal therapy. *Oncotarget*. 2015;6(4):2315.
- Tan Y, Huang N, Zhang X, et al. KIAA0247 suppresses the proliferation, angiogenesis and promote apoptosis of human glioma through inactivation of the AKT and Stat3 signaling pathway. *Oncotarget*. 2016;7(52):87100-87113.
- Moon SH, Kim DK, Cha Y, Jeon I, Song J, Park K-S. PI3K/Akt and Stat3 signaling regulated by PTEN control of the cancer stem cell population, proliferation and senescence in a glioblastoma cell line. *Int J Oncol.* 2013;42(3):921-928.
- Liang Q, Luo Z, Zeng J, et al. Zika virus NS4A and NS4B proteins deregulate Akt-mTOR signaling in human fetal neural stem cells to inhibit neurogenesis and induce autophagy. *Cell Stem Cell*. 2016;19(5):663-671.
- Clark PA, Iida M, Treisman DM, et al. Activation of multiple ERBB family receptors mediates glioblastoma cancer stem–like cell resistance to EGFR targeted inhibition. *Neoplasia*. 2012; 14(5):420-428.
- Ahmed N, Brawley V, Hegde M, et al. Autologous HER2 CMV bispecific CAR T cells are safe and demonstrate clinical benefit for glioblastoma in a Phase I trial. *J Immuno Cancer*. 2015; 3(Suppl 2):O11-O11.
- Li G, Mitra S, Wong AJ. The epidermal growth factor variant III peptide vaccine for treatment of malignant gliomas. *Neurosurg Clin N Am.* 2010;21(1):87-93.
- Guo L, Zhao Y, Yang S, Zhang H, Chen F. An integrated analysis of miRNA, lncRNA, and mRNA expression profiles. *BioMed Res Int*. 2014;2014:345605.
- Bao S, Wu Q, McLendon RE, et al. Glioma stem cells promote radioresistance by preferential activation of the DNA damage response. *Nature*. 2006;444(7120):756-760.
- Lee J, Kotliarova S, Kotliarov Y, et al. Tumor stem cells derived from glioblastomas cultured in bFGF and EGF more closely mirror the phenotype and genotype of primary tumors than do serumcultured cell lines. *Cancer Cell*. 2006;9(5):391-403.
- Ignatova TN, Kukekov VG, Laywell ED, Suslov ON, Vrionis FD, Steindler DA. Human cortical glial tumors contain neural stemlike cells expressing astroglial and neuronal markers in vitro. *Glia*. 2002;39(3):193-206.
- Svendsen CN, ter Borg MG, Armstrong RJE, et al. A new method for the rapid and long term growth of human neural precursor cells. *J Neurosci Methods*. 1998;85(2):141-152.
- Zorniak M, Clark PA, Leeper HE, et al. Differential expression of 2',3'-cyclic-nucleotide 3'-phosphodiesterase and neural lineage markers correlate with glioblastoma xenograft infiltration and patient survival. *Clin Cancer Res.* 2012;18(13):3628-3636.
- Pointer KB, Clark PA, Eliceiri KW, Salamat MS, Robertson GA, Kuo JS. Administration of Non-torsadogenic human Ether-à-gogo-related gene inhibitors is associated with better survival for high hERG–expressing glioblastoma patients. *Clinical Cancer Research*. 2017;23(1):73.
- Rao X, Huang X, Zhou Z, Lin X. An improvement of the 2O(delta delta CT) method for quantitative real-time polymerase chain reaction data analysis. *Biostat Bioinforma Biomath*. 2013; 3(3):71-85.

- Wheeler DL, Huang S, Kruser TJ, et al. Mechanisms of acquired resistance to cetuximab: role of HER (ErbB) family members. *Oncogene*. 2008;27(28):3944-3956.
- Vivanco I, Mellinghoff IK. Epidermal growth factor receptor inhibitors in oncology. *Curr Opin Oncol.* 2010;22(6):573-578.
- Fischer AH, Jacobson KA, Rose J, Zeller R. Hematoxylin and eosin staining of tissue and cell sections. *Cold Spring Harbor Protocols*. 2008;2008(5):pdb.prot4986.
- Ludwig N, Leidinger P, Becker K, et al. Distribution of miRNA expression across human tissues. *Nucleic Acids Res.* 2016;44(8): 3865-3877.
- Tarantino C, Paolella G, Cozzuto L, et al. MiRNA 34a, 100, and 137 modulate differentiation of mouse embryonic stem cells. *FASEB J.* 2010;24(9):3255-3263.
- Stopka T, Skoultchi AI. The ISWI ATPase Snf2 h is required for early mouse development. *Proceed National Acad Sci.* 2003; 100(24):14097-14102.
- Barbagallo D, Caponnetto A, Cirnigliaro M, et al. CircSMARCA5 inhibits migration of glioblastoma multiforme cells by regulating a molecular axis involving splicing factors SRSF1/ SRSF3/PTB. *Int J Mole Sci.* 2018;19(2):480.
- Hakimi M-A, Bochar DA, Schmiesing JA, et al. A chromatin remodelling complex that loads cohesin onto human chromosomes. *Nature*. 2002;418(6901):994-998.
- Alenghat T, Yu J, Lazar MA. The N-CoR complex enables chromatin remodeler SNF2 H to enhance repression by thyroid hormone receptor. *EMBO J.* 2006;25(17):3966-3974.
- Soriano FX, Hardingham GE. In cortical neurons HDAC3 activity suppresses RD4-dependent SMRT export. *PLoS ONE*. 2011;6(6): e21056.
- Bardai FH, Price V, Zaayman M, Wang L, D'Mello SR. Histone deacetylase-1 (HDAC1) is a molecular switch between neuronal survival and death. *J Biol Chem*. 2012;287(42):35444-35453.
- Lee-Hoeflich ST, Crocker L, Yao E, et al. A central role for HER3 in HER2-amplified breast cancer: implications for targeted therapy. *Cancer Res.* 2008;68(14):5878-5887.
- Serra V, Scaltriti M, Prudkin L, et al. PI3 K inhibition results in enhanced HER signaling and acquired ERK dependency in HER2-overexpressing breast cancer. *Oncogene*. 2011;30(22): 2547-2557.
- Anaya J. OncoLnc: linking TCGA survival data to mRNAs, miR-NAs, and lncRNAs. *Peer J Comp Sci.* 2016;2:e67.
- Mulder KW, Wang X, Escriu C, et al. Diverse epigenetic strategies interact to control epidermal differentiation. *Nat Cell Biol*. 2012;14(7):753-763.
- Bhaskara S, Knutson SK, Jiang G, et al. Hdac3 is essential for the maintenance of chromatin structure and genome stability. *Cancer Cell*. 2010;18(5):436-447.
- Watson PJ, Fairall L, Santos GM, Schwabe JWR. Structure of HDAC3 bound to co-repressor and inositol tetraphosphate. *Nature*. 2012;481:335-340.
- You SH, Lim HW, Sun Z, Broache M, Won KJ, Lazar MA. Nuclear receptor co-repressors are required for the histonedeacetylase activity of HDAC3 in vivo. *Nat Struct Mol Biol.* 2013;20(2):182-187.

- Jepsen K, Gleiberman AS, Shi C, Simon DI, Rosenfeld MG. Cooperative regulation in development by SMRT and FOXP1. *Genes & Development*. 2008;22(6):740-745.
- Neganova I, Vilella F, Atkinson SP, et al. An important role for CDK2 in G1 to S checkpoint activation and DNA damage response in human embryonic stem cells. *Stem Cells*. 2011; 29(4):651-659.
- 50. Armas P, Margarit E, Mouguelar VS, Allende ML, Calcaterra NB. Beyond the binding site: in vivo identification of tbx2, smarca5 and wnt5b as molecular targets of CNBP during embryonic development. *PLoS One.* 2013;8(5):e63234.
- Barré B, Avril S, Coqueret O. Opposite regulation of Myc and p21 waf1 transcription by STAT3 proteins. *J Biol Chem.* 2003;278(5): 2990-2996.
- Sinibaldi D, Wharton W, Turkson J, Bowman T, Pledger WJ, Jove R. Induction of p21WAF1/CIP1 and cyclin D1 expression by the Src oncoprotein in mouse fibroblasts: role of activated STAT3 signaling. *Oncogene*. 2000;19(48):5419-5427.
- Satriotomo I, Bowen KK, Vemuganti R. JAK2 and STAT3 activation contributes to neuronal damage following transient focal cerebral ischemia. *J Neuro Chem.* 2006;98(5):1353-1368.
- Zou M, Hu C, You Q, Zhang A, Wang X, Guo Q. Oroxylin A induces autophagy in human malignant glioma cells via the mTOR-STAT3-Notch signaling pathway. *Mole Carcin*. 2015; 54(11):1363-1375.
- Androutsellis-Theotokis A, Leker RR, Soldner F, et al. Notch signalling regulates stem cell numbers in vitro and in vivo. *Nature*. 2006;442(7104):823-826.
- Ulasov IV, Nandi S, Dey M, Sonabend AM, Lesniak MS. Inhibition of sonic hedgehog and notch pathways enhances sensitivity of CD133+ glioma stem cells to temozolomide therapy. *Mole Med.* 2011;17(1):103-112.
- Borggrefe T, Oswald F. The notch signaling pathway: transcriptional regulation at notch target genes. *Cell Mole Life Sci.* 2009; 66(10):1631-1646.
- Stopka T, Blafkova J, Zakova D, et al. Cloning and expression of murine hematopoietic specific chromatin remodeling gene SMARCA5. *Exp Hema*. 2000;28(7 Suppl 1):119.
- Gigek CO, Lisboa LCF, Leal MF, et al. SMARCA5 methylation and expression in gastric cancer. *Cancer Invest*. 2011;29(2):162-166.
- Cheng L, Wu Q, Huang Z, et al. L1CAM regulates DNA damage checkpoint response of glioblastoma stem cells through NBS1. *EMBO J.* 2011;30(5):800-813.
- Mueller AC, Sun D, Dutta A. The miR-99 family regulates the DNA damage response through its target SNF2 H. Oncogene. 2013;32(9):1164-1172.
- Giangreco AA, Vaishnav A, Wagner D, et al. Tumor suppressor microRNAs, miR-100 and -125b, are regulated by 1,25dihydroxyvitamin D in primary prostate cells and in patient tissue. *Cancer Prevent Res.* 2013;6(5):483-494.
- Feng B, Wang R, Chen LB. MiR-100 resensitizes docetaxelresistant human lung adenocarcinoma cells (SPC-A1) to docetaxel by targeting Plk1. *Can Lett.* 2012;317(2):184-191.
- Cui S, Boren T, Chan G, Indermauer1 M, Dressman HK, Lancaster JM. MicroRNAs that underlie ovarian cancer development and response to chemotherapy. *Can Res.* 2007;67(9 Suppl):4514.