

Sox2 Promotes Malignancy in Glioblastoma by Regulating **Plasticity and Astrocytic** Differentiation^{1,2}

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Abstract

The high-mobility group-box transcription factor sex-determining region Y-box 2 (Sox2) is essential for the maintenance of stem cells from early development to adult tissues. Sox2 can reprogram differentiated cells into pluripotent cells in concert with other factors and is overexpressed in various cancers. In glioblastoma (GBM), Sox2 is a marker of cancer stemlike cells (CSCs) in neurosphere cultures and is associated with the proneural molecular subtype. Here, we report that Sox2 expression pattern in GBM tumors and patient-derived mouse xenografts is not restricted to a small percentage of cells and is coexpressed with various lineage markers, suggesting that its expression extends beyond CSCs to encompass more differentiated neoplastic cells across molecular subtypes. Employing a CSC derived from a patient with GBM and isogenic differentiated cell model, we show that Sox2 knockdown in the differentiated state abolished dedifferentiation and acquisition of CSC phenotype. Furthermore, Sox2 deficiency specifically impaired the astrocytic component of a biphasic gliosarcoma xenograft model while allowing the formation of tumors with sarcomatous phenotype. The expression of genes associated with stem cells and malignancy were commonly downregulated in both CSCs and serum-differentiated cells on Sox2 knockdown. Genes previously shown to be associated with pluripontency and CSCs were only affected in the CSC state, whereas embryonic stem cell self-renewal genes and cytokine signaling were downregulated, and the Wnt pathway activated in differentiated Sox2-deficient cells. Our results indicate that Sox2 regulates the expression of key genes and pathways involved in GBM malignancy, in both cancer stemlike and differentiated cells, and maintains plasticity for bidirectional conversion between the two states, with significant clinical implications.

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Abbreviations: α-SMA, actin alpha 2 smooth muscle (ACTA2); CNS, central nervous system; CSC, cancer stem-like cell; DEG, differentially expressed gene; ESC, embryonic stem cell; GFAP, glial fibrillary acidic protein; GBM, glioblastoma; GS, gliosarcoma; IF, intermediate filament protein; IHC, immunohistochemistry; NM, neurosphere media; NSC, neural stem cell; Sox2, SRY (sex determining region Y)-box 2; SVZ, subventricular zone; SDC, serum differentiated cell; TCGA, the Cancer Genome Atlas; TMA, tissue microarray; WHO, World Health Organization

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Introduction

In a variety of tumor types, a worse prognosis is associated with increased plasticity, loss of differentiation makers, and induction of stem cell phenotypes [1]. One key regulator of stemness is the highmobility group (HMG)-box transcription factor sex-determining region Y (SRY)-box 2 (Sox2), essential in early embryonic development [2]. Sox2 activity is critical for the maintenance of mouse embryonic stem cells (ESCs) [3] and is associated with the promoter region of 7% of protein-coding genes in human ESCs [4]. Sox2 activates and represses the expression of different gene sets in various tissues during development through HMG-box domainmediated DNA binding. Cooperative binding of other transcription factors contributes to target specificity [5]. Sox2 is highly expressed in neural stem cells (NSC), neural progenitors, and immature astrocytes in the subventricular zone (SVZ) of developing rodent brains and is required for maintenance and identity of NSC in the adult mouse brain [6]. Sox2 is re-expressed in astrocytes undergoing cell division in the mouse neocortex in response to brain injury [7]. In the adult human brain, Sox2 expression is restricted to regions with neurogenic potential, such as the SVZ [8]. Outside of normal development, Sox2 is an essential component of the transcription factor cocktail for reprogramming differentiated cells into iPS cells [9]. SOX2 gene is amplified or otherwise overexpressed in several tumor types [5]. Transcriptional regulation by Sox2 is context dependent, and its impact on malignancy is cancer-type specific. For example, Sox2 functions as an oncogene in lung and esophagus squamous cell carcinomas [10], whereas in ovarian cancer, its expression is restricted to cancer stemlike cells (CSCs) [11].

Glioblastoma (GBM), a WHO grade IV astrocytoma, is the most aggressive primary central nervous system tumor. Sox2 transcript and protein are upregulated in GBM tumors in relation to nontumor brain tissues [12,13], gene amplification is observed in 4% of GBM tumors profiled by The Cancer Genome Atlas (TCGA) [14], and promoter hypomethylation occurs with high frequency [12]. Sox2 is a component of the GBM proneural molecular subclass signature, along with other regulators of neural stem/progenitor cell fate [15]. Notable phenotypic plasticity of GBMs contributes to intratumor heterogeneity, encompassing cells in a range of developmental states, from CSCs to more differentiated neoplastic cells. These various developmental states can be partially captured ex vivo. Culturing GBM cells in serum-free medium supplemented with epidermal growth factor (EGF) and fibroblast growth factor (FGF) developed for NSCs allows the selection and propagation of neurospheres, a population enriched in CSC, defined by the ability to self-renew and differentiate into cells forming the bulk of the tumor, phenocopying the parental tumor in mouse orthotopic xenografts [16,17]. Sox2, ubiquitously expressed in GBM neurosphere cell cultures [16-21], has been shown to function in the proliferation of GBM CSC population and tumorigenicity [20]. Neurosphere cultures derived from adult mouse SVZ also express Sox2, and differentiation induced by FBS is accompanied by down-regulation of Sox2 mRNA and protein [22]. Similarly, culturing dissociated GBM tumors in FBScontaining media typically results in loss of the CSC phenotype and impaired tumorigenicity associated with down-regulation of NSC markers, including Sox2, and gain of more mature astrocytic markers [16,17,19].

Here, we verify that the endogenous expression of Sox2 in GBM biopsies and xenograft tumors is more widespread than expected for a marker of CSCs, raising the question as to whether in GBM tumors Sox2

performs distinct functions in differentiated and CSCs. Although few long-term serum-cultured GBM cells can regain expression of Sox2 [23], these cells accumulate genomic abnormalities leading to considerably divergence from the original tumor [16]. We have previously described a patient-derived low -passage serum-differentiated cell (SDC) line, which, contrary to typical GBM samples, retains Sox2 expression, along with the ability to dedifferentiate into neurospheres in vitro and to phenocopy the parental tumor in orthotopic xenografts [19]. These SDCs and CSCs originating from the same tumor specimen [19] constitute an adequate model to study Sox2 function in the two developmental states under an isogenic background. Gliosarcoma (GS), a WHO histologic variant [24] comprising approximately 1.8% to 2.8% of all GBMs, is characterized by biphasic malignant glial and sarcomatous components, resulting from mesenchymal metaplasia [25]. GSs are not distinguished from GBMs regarding clinical management and large-scale molecular profiling efforts, such as the TCGA, and are assigned to multiple molecular subclasses [26]. The availability of a GS model [19] presented an opportunity to contrast Sox2 functions in the glial and mesenchymal compartments, further highlighting its role in astrocytic tumorigenicity.

Materials and Methods

Tumor Samples and Cell Culture

Resected brain tumors were collected at Henry Ford Hospital (Detroit, MI) with written consent from patients in accordance with institutional guidelines and graded pathologically according to the WHO criteria. GBM tumors were dissociated, as previously described [27]. Dissociated cells were grown in Dulbecco's modified Eagle's medium (DMEM) (Life Technologies) supplemented with 10% FBS (HyClone) for traditional monolayer cultures or in neurosphere medium (NM), composed of DMEM/F-12 supplemented with N2 (Gibco), 0.5 mg/ml BSA (Sigma), 25 $\mu g/ml$ gentamicin (Gibco), 0.5% antibiotic/antimycotic (Invitrogen), 20 ng/ml basic fibroblast growth factor, and 20 ng/ml EGF (Peprotech). Cells were maintained in culture for up to passage 10 (low passage).

Immunohistochemistry

Immunohistochemistry was performed as described previously [19]. The following primary antibodies were used: anti-Sox2 (Cell Signaling Technology), anti–glial fibrillary acidic protein (anti-GFAP) (Abcam), anti-nestin (Millipore), anti-vimentin (Santa Cruz Biotechnology), anti– α -smooth muscle actin (anti– α -SMA) (Abcam), and anti–major histocompatibility complex I (anti-MHC I) (Abcam). Anti-Sox2 antibody specificity was verified by the absence of signal on Sox2 knockdown, as shown in Figure 4. Reticulin staining was performed with a kit (DakoCytomation). Images were captured using Nikon E800M microscope and DXM1200C digital camera. For immunofluorescence, secondary antibodies conjugated to DyLight 488 and DyLight 649 (Jackson ImmunoResearch Laboratories) were employed, nuclei were stained with DAPI (Invitrogen), and images captured using Nikon Eclipse 80i microscope equipped with epifluorescence.

Sox2 Knockdown and Ectopic Expression

Sox2 knockdown in low-passage 10% FBS cells was achieved by oligonucleotides targeting human SOX2-coding or nonsilencing control sequences cloned into BLOCK-iT Pol II miR RNAi expression vectors (Life Technologies) before cell transfection with Lipofectamine 2000 (Life Technologies). Sox2 knockdown in neurospheres was achieved using GIPZ Lentiviral shRNAmir (clones V3LHS_404430 and V3LHS_404432) and nonsilencing control

(RHS4346) (Thermo Scientific Open Biosystems). Sox2 ectopic expression was achieved by subcloning Sox2 cDNA from pCMV6-XL5-NM_002106.2 (OriGene) into pcDNA 3.1 mammalian expression vector (Invitrogen), under control of constitutive CMV promoter; the empty vector was used as control. Plasmid DNA constructs were verified by DNA sequencing and stably transfected into GBM monolayer cells using Lipofectamine 2000 (Invitrogen).

Neurosphere Formation Assay

Monolayer cells were harvested by trypsin treatment, washed three times in serum-free DMEM/F-12 medium, and plated in NM in regular tissue culture–treated flasks at a density of 10 cells per well. Neurosphere formation was monitored for 2 weeks, and neurospheres containing more than 32 cells were counted.

Cell Proliferation Assay

Cells were harvested and plated onto 96-well plates (500 cells per well for 10% FBS cultures and 1000 cells per well for neurosphere cultures). Cell viability was quantified for 7 days by intracellular ATP level measurements using CellTiter-Glo Luminescent Cell Viability Assay (Promega).

Xenograft Tumors

Following IACUC guidelines in an institutionally approved animal use protocol, dissociated GBM cells were implanted into 8-week-old female nude mice (NCRNU; Taconic Farms). Animals were anesthetized with a mixture of ketamine and xylazine. The injection site was manually drilled 2.5 mm to the right of the bregma and 1 mm posterior to the coronal suture. Dissociated cells (3×10^5 per mouse) were injected at a depth of 3 mm using a Hamilton syringe. Animals were monitored daily and killed on the first signs of neurologic deficit or weight loss greater than 20%. Xenograft tissue microarray (TMA) slides were obtained through participation in the Ivy GBM consortium.

Proteasome Inhibition

Exponentially growing cells were plated at a density of 1×10^5 cells/ml in growth medium. After 24 hours, medium was replaced with growth medium containing 10 μ M MG132 (Sigma). Cells were incubated for 4 to 24 hours at 37°C (5% CO₂) before whole-cell lysates were obtained.

Western Blot Analysis

Fifteen micrograms of total protein from whole-cell lysates were denatured in sodium dodecyl sulfate—gel loading buffer and separated by NuPAGE SDS-PAGE gel system (Life Technologies). Proteins were electrophoretically transferred to PVDF membranes (Life Technologies). Primary antibodies used were Sox2 (Chemicon) and β -actin (Sigma-Aldrich). Immobilon Western Chemiluminescent HRP substrate (Millipore) was used for detection.

Quantitative Real-Time Polymerase Chain Reaction Analysis

Total RNA was extracted with RNeasy Mini Kit (Qiagen), eluted in RNase-free water, and stored at -80°C . cDNA was prepared from 1 μg of DNAse I–treated RNA using SuperScript III and oligo(dT) (Invitrogen). Relative quantification of gene expression was performed by real-time polymerase chain reaction (PCR) with SYBR Green and ViiA 7 detection system (Applied Biosystems). β -Actin was used as internal reference, and relative mRNA levels were quantified by the $2^{(-\Delta\Delta Ct)}$ method. DNA oligo sequences are available on request.

Global Gene Expression Analysis

RNA samples were analyzed by Illumina Human HT12v4.0 Expression BeadChip, at the Wayne State University Applied Genomics Technology Center. The gene expression image files obtained from the Illumina iScan were uploaded to GenomeStudio (version 2010.3) using the Gene Expression module. Illumina array data were normalized using the rank invariant method, and the background was subtracted in BeadStudio. Normalization artifacts of negative expression values were set to a minimum null value. A single probe with the largest expression range across the samples was selected for genes with multiple probes. Fold change was calculated on the basis of the mean values for Sox2expressing cells (HF2303 parental and miRNAneg) and Sox2-deficient cells (Sox2miR1 and Sox2miR2). Genes with mean Illumina intensity levels lower than 20 for both groups were excluded. A stringent foldchange cutoff of 3.5 was used to generate the differentially expressed gene (DEG) lists. Global gene expression data files are available at GEO. MetaCore (Thomson Reuters) and IPA (Ingenuity Systems) software applications were used to further analyze the data sets. Microarray data are available through GEO (GSE51441).

TCGA Data Analysis

Preprocessed level 2 Agilent Human Gene Expression Microarray data were obtained from the TCGA Data Portal (https://tcga-data.nci.nih.gov/tcga/, July 2, 2012) for 517 GBM cases with no prior cancer diagnoses. Analysis was conducted at probe level. Pearson correlation coefficient between each of two Sox2 open reading frame probes (A_23_P159606, A_24_0379969, r = 0.96) and all other probes was conducted. A gene was considered to have consistent correlation if it showed correlation of ρ > 0.4 (positive) or ρ < -0.4 (negative) for at least 50% of its probes with either of the Sox2 probes. For visualization, consistently correlated probes were z score transformed and averaged to give a single value per gene. Hierarchical clustering of the genes was performed using the Pearson correlation and complete linkage. Molecular class was predicted from the metagene signatures for the four classes. Analysis was performed using R Software.

Statistical Analysis

Results are represented as means \pm SEM. Differences were analyzed by unpaired two-tailed Student's t test or one-way analysis of variance using Prism 5 (GraphPad Software). For Figures 4C and W1C, Dunn multiple-comparison test was used. ANCOVA was used for age-adjusted tests of means, and Cox regression was used for survival analysis (R Software). P < .05 was considered significant.

Results

Sox2 is Widely Expressed in GBM Tumors

Sox2 protein expression was assessed in newly diagnosed (n = 13) and recurrent (n = 12) GBMs and in grade III anaplastic oligodendroglioma (n = 9) and anaplastic astrocytomas (n = 5). The percentage of Sox2-positive cells in GBM tumors ranged from 9% to 85% (Figure 1, A and B). Sox2 was similarly highly expressed in the grade III gliomas (Figure 1B), in agreement with previous reports [28,29]. Representative images are shown in Figure 1C. The subclass assignment [15] was available for seven GBM samples (Figure 1A). In the proneural samples, 64% and 85% of cells were positive for Sox2 (Figure 1A), consistent with Sox2 being a component of the proneural subclass signature [15]. Sox2-positive cells ranged from 11.2% to 55.5% in the

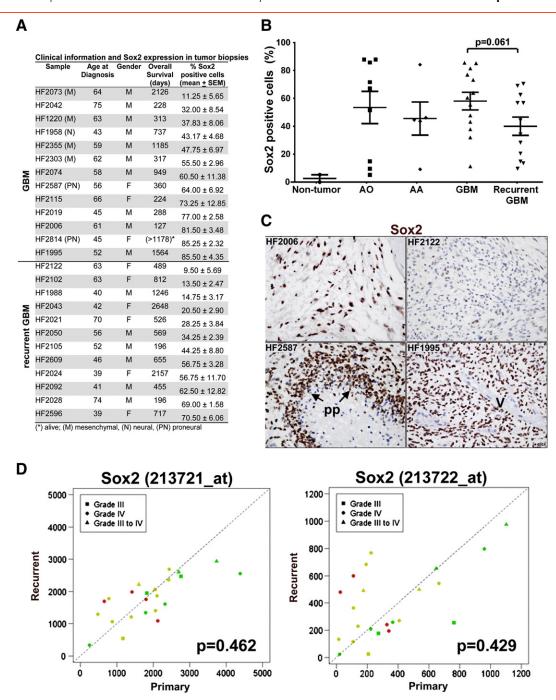


Figure 1. Sox2 expression in GBM tumors. Sox2 expression in 25 GBM biopsy specimens was assessed by immunohistochemistry. Sox2-positive nuclei were quantified in four randomly selected \times 400 optical fields per specimen. (A) Clinical information. (B) Scatterplot represents the percentage of Sox2-positive cells for each specimen (mean \pm SEM). Anaplastic oligodendroglioma (AO; n=9), anaplastic astrocytoma (AA; n=5), and nontumor brain (n=2) samples are included for reference. (C) Representative images show nuclear Sox2 expression in GBM biopsies; pseudopalisating necrosis (pp) and vessels (V) are indicated. (D) Graphical representation of Sox2 mRNA expression in 23 paired primary (horizontal axis) and recurrent (vertical axis) gliomas. Square, WHO grade III primary and recurrent gliomas (n=3); circle, primary and recurrent (grade IV) GBMs (n=16); triangle, primary grade III and recurrent GBM (n=4). Patients grouped by age (years): Green, <40; yellow, 40 to 60; red, >60.

mesenchymal tumors. An 18% decrease in the mean percentage of Sox2-positive cells in the recurrent GBM group relative to newly diagnosed was observed (P = .0609; 95% confidence interval = -0.9%-36.9%) (Figure 1B). When we adjusted for patient age at diagnosis (Figure 1A), a significant decrease in Sox2 expression was observed in the recurrent tumor group (21.7%, ANCOVA P = .0266; 95%

confidence interval = 2.8%- 40.6%). To further investigate how Sox2 expression changes on recurrence, transcriptome data for 23 paired primary and recurrent grade III and grade IV gliomas [30] were analyzed. Sox2 expression obtained from two Affymetrix U133 probes were not significantly different between primary and recurrent GBMs (213721_at, P = .4618; 213722_at, P = .4293) (Figure 1D). Sox2

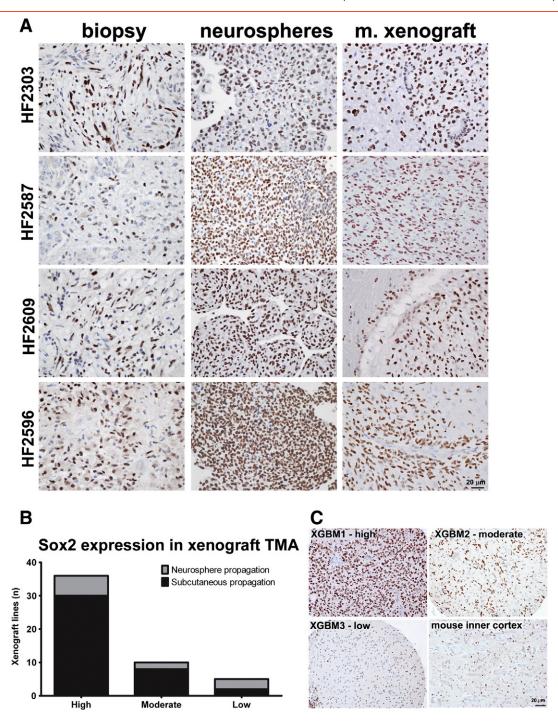


Figure 2. Sox2 expression in GBM neurospheres and orthotopic xenografts. (A) Sox2 expression in four sets of matched GBM biopsy, cultured neurosphere cells, mouse xenograft, two newly diagnosed (HF2303 and HF2587), and two recurrent tumors (HF2609 and HF2596). (B) Sox2-positive cells were quantified in TMA containing samples of xenograft tumors obtained from 51 GBM specimens propagated *in vitro* as neurosphere cultures or subcutaneously in nude mouse flanks, before intracranial implant in nude mice. Xenograft lines from subcutaneous and neurosphere propagation were ranked for Sox2 expression as high (>65% positive cells), moderate (between 20% and 65% positive cells), and low (<20% cells). (C) Representative images for each xenograft category in B and normal mouse brain (inner cortex). Scale, 20 μ m.

expression tended to decrease on recurrence for the younger patients and for the grade III tumors (Figure 1*D*).

Together, these observations indicate that Sox2 expression is not restricted to a small subpopulation of undifferentiated neoplastic cells, suggesting that Sox2 could regulate different programs in CSCs and more differentiated GBM cells.

Sox2 Expression is Conserved in Patient-Derived GBM Orthotopic Xenografts

To enrich for cells with CSC phenotype, neurosphere cultures from two newly diagnosed GBMs (HF2303 and HF2587) and two recurrent GBMs (HF2609 and HF2596) were implanted intracranially in immunocompromised mice. Sox2 was expressed in 55% to

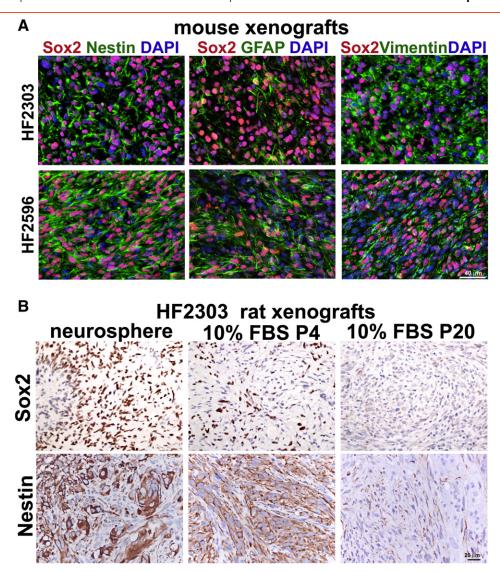


Figure 3. Sox2 is coexpressed with GFAP, nestin, and vimentin in neurosphere-derived GBM xenografts. (A) Coexpression of Sox2 (red) with the IF proteins (green) nestin, GFAP, and vimentin in HF2303 and HF2596 GBM mouse xenografts. Nuclei counterstained with DAPI. Scale, $40 \, \mu \text{m}$. (B) Sox2 and nestin are expressed in xenografts formed from HF2303 neurospheres or low-passage 10% FBS cultures (P4) and downregulated in higher passage (P20) 10% FBS serum cultures. Scale, $20 \, \mu \text{m}$.

70% of the nuclei for these biopsies (Figures 1 A and 2A). As expected, matched neurosphere cells were highly enriched in Sox2-positive cells, and biopsy expression pattern was replicated in the xenograft tumors (Figure 2A).

Sox2 expression was then evaluated in a TMA composed of samples from 51 mouse orthotopic GBM xenografts, including 40 originating from patient tumors propagated subcutaneously and 11 from neurosphere cultures. As shown in Figure 2B, 36 (71%) of the xenograft lines presented high percentage (>65%) of Sox2-positive nuclei, whereas only 5 (9.8%) of the xenografts displayed low (<20%) Sox2-positive nuclei. Representative images are shown (Figure 2C). These results indicate that robust Sox2 protein expression observed for the majority of the patient-derived orthotopic xenograft tumors is independent of the use of neurosphere culture or subcutaneous tumors as the preimplantation propagation method.

The intermediate filament (IF) protein nestin is a direct target of Sox2 regulation in neural stem and progenitor cells [31] and also

commonly expressed in a large percentage of cells in GBM tumors [19]. As expected, in neurosphere-derived GBM xenografts, Sox2 coexpressed with nestin (Figure 3A). Vimentin, the main IF protein of mesenchymal cells, also expressed in neural stem and progenitor cells, was coexpressed with Sox2 in the GBM xenografts (Figure 3A). GFAP, a major IF of mature astrocytes, also expressed in astrocytic progenitors, coexpressed with Sox2 (Figure 3A), as previously reported [32]. Sox2 coexpression with IF proteins markers of diverse lineages is in agreement with Sox2 not being restricted to undifferentiated tumor cells, underlying the plasticity of high-grade astrocytomas.

Whereas Sox2 is downregulated in the majority of GBM cells exposed to 10% FBS, we have previously characterized a GBM specimen, HF2303 (GS1), which retains Sox2 expression in low-passage SDCs [19]. Here, we show that Sox2 expression in xenograft tumors are consistent with the *in vitro* expression: Sox2-positive cells are observed in xenografts derived from HF2303 neurosphere/CSCs and low-passage SDCs (10% FBS P4) and not in higher passage

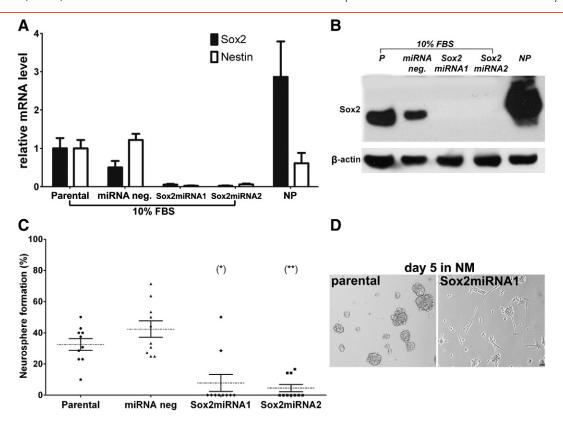


Figure 4. Sox2 knockdown prevents dedifferentiation of HF2303 GBM cells. (A) Sox2 and nestin mRNA levels in HF2303 cells (parental) and cells transfected with miRNA constructs targeting Sox2 (Sox2miRNA1 and Sox2miRNA2) or nonsilencing miRNA (miRNAneg) was evaluated by quantitative RT-PCR. HF2303 neurosphere cells (NP) were included for comparison. Results represent means \pm SEM; n=6. (B) Sox2 protein expression in whole-cell lysates determined by Western blot analysis confirms Sox2 knockdown in Sox2miRNA1 and Sox2miRNA2 lines. β-Actin was used as loading control. (C) The effect of Sox2 knockdown on dedifferentiation of HF2303 monolayers into neurospheres. Cells were seeded at low density in 96-well plates and cultured in NM. Neurosphere formation was monitored and quantified at day 12. Results represent means \pm SEM, for n=10 (*P < .05 and **P < .01), in relation to parental cells after Dunn multiple comparison test. (D) Representative images contrast Sox2-expressing and Sox2-deficient cells in NM culture. Bar, 20 µm.

SDCs (10% FBS P20, Figure 3B). Nestin expression in the tumors follows the Sox2 pattern (Figure 3B).

Sox2 Function in Plasticity of Differentiated GBM Cells

To study Sox2 function in differentiated patient-derived GBM cells, low-passage HF2303 SDCs were transfected with miRNA constructs targeting the Sox2-coding region (Sox2miRNA1 and Sox2miRNA2) or with a nonsilencing control (miRNAneg). Sox2 knockdown in HF2303_Sox2miRNA1 and HF2303_Sox2miRNA2 was verified by quantitative PCR (Figure 4A) and Western blot analysis (Figure 4B). Knocking down Sox2 expression in HF2303 cells significantly decreased the level of nestin transcript, a direct target of Sox2 (Figure 4A). Expression of Sox2 in isogenic HF2303 neurospheres (NP) are shown for comparison (Figure 4, A and B). We have previously demonstrated that HF2303 SDCs not only retain Sox2 expression but also the ability to dedifferentiate, forming neurospheres in a clonogenic assay [19]. Sox2 knockdown significantly impaired neurosphere formation in NM (Figure 4, C and D), whereas no effect in proliferation was observed (Figure W1).

Sox2 Expression is not Sufficient to Restore Plasticity and Tumorigenicity to Serum-Cultured Primary GBM Cells

We have previously described neurosphere/CSCs and SDCs from another GBM specimen (HF2414/GBM2) [19], demonstrating that

the CSCs express Sox2 and are tumorigenic, whereas the low-passage SDCs lose Sox2 expression, along with the ability to dedifferentiate in vitro, and to form tumors [19], typical observations for GBM SDCs [16]. To determine whether Sox2 ectopic expression was sufficient to restore plasticity and tumorigenicity to HF2414 SDCs, cells were stably transfected with Sox2 cDNA construct. Detectable levels of Sox2 protein expression were only attained in clones expressing very high levels of Sox2 transcript (HF2414-hSox2) in relation to clones expressing the empty vector (HF2414 V) (Figure 5, A and B). Endogenous Sox2 expression was negligible in HF2414 V cells and was not upregulated when cells were cultured in serum-free NM for 48 hours (Figure 5, A and B). Nestin mRNA was upregulated on ectopic Sox2 expression and further upregulated when the cells were transferred to NM (Figure 5A). The discrepancy between mRNA and protein levels (Figure 5, A and B) suggests that the posttranslational regulation of Sox2 expression is affected by culture conditions, because increased protein levels were observed when HF2414-hSox2 was transferred from 10% FBS to NM for 48 hours (Figure 5B), which was not mediated by an increase in mRNA expression (Figure 5A). Treatment with proteasome inhibitor MG132 (10 µM) increased Sox2 protein levels in HF2414-hSox2 cells (Figure 5B). Although predicted ubiquitin-binding sites exist in Sox2 (http://ubpred.org), we found no evidence of direct ubiquitination of Sox2 (not shown).

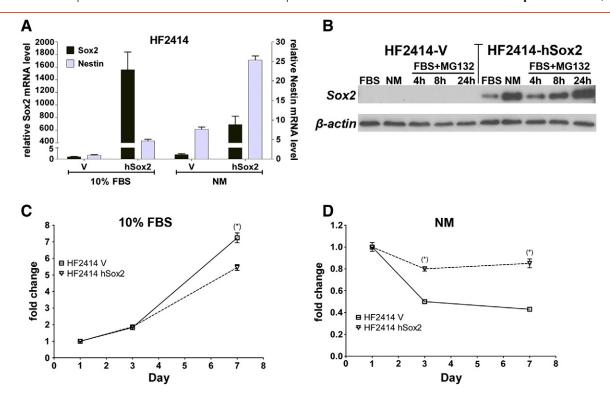


Figure 5. Sox2 ectopic expression in low-passage primary GBM monolayer line HF2414. HF2414 cells constitutively expressing Sox2 (HF2414-hSox2) or the empty vector (HF2414-V) were maintained in 10% FBS culture or transferred to NM for 48 hours. (A) Sox2 and nestin mRNA levels were determined by quantitative RT-PCR. Results represent means \pm SEM, for n=6. (B) Sox2 protein expression in whole-cell lysates was determined by Western blot analysis. Sox2 protein was detected in the HF2414-hSox2 grown in serum, and protein expression was increased when cells were transferred to NM or when proteasome inhibitor MG132 was added to the 10% FBS serum containing medium in a time-dependent manner. No Sox2 protein was observed in the HF2414-V cell line under the same culture conditions. (C) Cell proliferation in 10% FBS medium measured by CellTiterGlo was decreased for HF2414-hSox2 cells relative to HF2414-V cells. Mean \pm SE (n=10) is shown in the graph (*P<0.05). (D) When the cells were plated in low density in NM, no neurospheres formed, and there was a reduction in cell viability for 7 days, significantly more accentuated for HF2414-V. Mean \pm SE (n=10) is shown in the graph (*P<0.05).

Sox2 protein was not detected in HF2414-V cells, cultured under the same conditions (Figure 5*B*), in agreement with irreversible loss of endogenous Sox2 commonly observed for GBM SDCs. Ectopic expression of Sox2 had a negative effect in HF2414 cell proliferation in 10% FBS (Figure 5*C*) and was not sufficient to induce CSC phenotype or restore tumorigenicity to HF2414 SDCs (not shown) but led to significant increase in cell viability in NM (Figure 5*D*).

Sox2 Knockdown Alters Tumor Formation and Abolishes the Astrocytic Phenotype in Biphasic GS Tumor

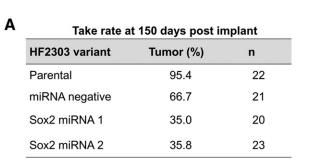
We have previously demonstrated that CSCs and low-passage SDCs from HF2303 GS tumor retain the intrinsic ability to generate the biphasic glial and sarcomatous tumor in orthotopic xenografts [19]. Sox2 deficiency in HF2303 SDCs reduced tumorigenicity, as observed by a 50% decrease intake rate relative to the control group (miRNAneg) (Figure 6A). GBM xenograft tumors were labeled with a human-specific marker (MHC I, Figure 6B). Distinct GFAP and reticulin-positive regions were observed in the Sox2-positive tumors (Figure 6C), as previously demonstrated [19]. Sox2 was expressed in both the glial and sarcomatous compartments of the xenograft tumors, although fewer positive cells were observed in the sarcomatous compartment (Figure 6C). No Sox2-positive tumor cells or GFAP expression was observed in the Sox2miRNA xenografts,

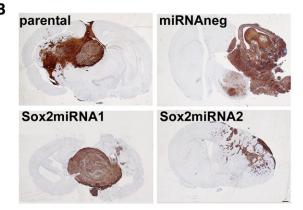
and the entire tumor consisted of the sarcomatous component, as evidenced by the reticulin staining (Figure 6C). Murine and human Sox2 proteins share 99.4% amino acid identity, leading to cross-species antibody labeling. Because Sox2-positive cells exist in the adult mouse brain (e.g., Figure 2C), colabeling of the human marker and Sox2 was used to distinguish between Sox2 expression in the human tumor (Figure 6, C and D, green arrowheads) and in mouse host brain tissue (Figure 6, C and D, black arrows), demonstrating that Sox2 protein was not detected in neoplastic cells of xenograft tumors from Sox2-deficient HF2303 cells. GS sarcomatous compartment can display diverse mesenchymal phenotypes. HF2303 mesenchymal component in the biopsy and xenografts is positive for α -SMA, a marker of myogenic lineage [19]. Sox2-deficient xenografts expressed α -SMA throughout the tumor (Figure 6E), further verifying the sarcomatous nature of the tumors.

Sox2 Regulates the Expression of Distinct Sets of Genes in Differentiated and CSC States

To better understand transcriptional regulation by Sox2 in differentiated GBM cells, genes differentially expressed between Sox2-positive HF2303 low-passage SDCs (parental and HF2303miRneg) and cells where Sox2 has been knocked down (HF2303_Sox2miRNA1 and HF2303_Sox2miRNA2) were derived from genome-wide transcriptome data (Illumina HT12v4.0). To

HF2303 mouse orthotopic xenografts





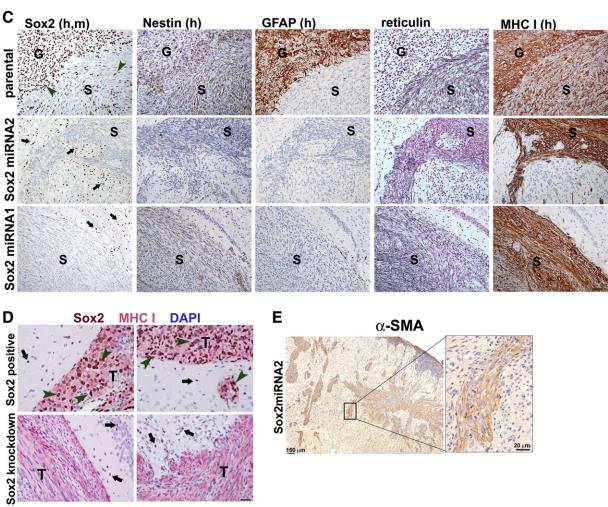
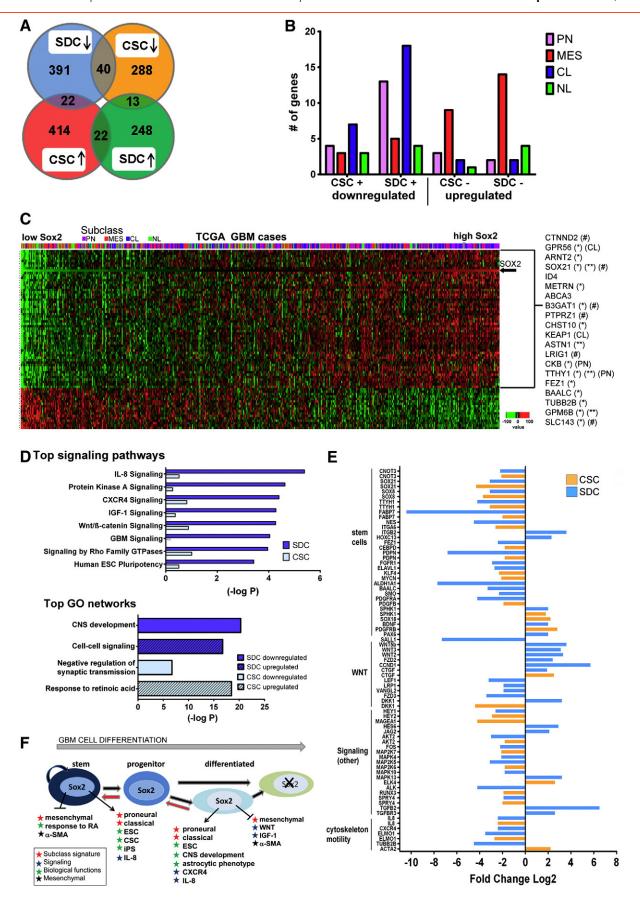


Figure 6. Sox2 loss affects the astrocytic compartment of GS. HF2303-parental, HF2303-mirRNAneg, HF2303-Sox2miRNA1, and HF2303-Sox2miRNA2 cells were implanted intracranially in immunocompromised mice (n=20-23).(A) Percentage of animals in each group that succumbed to tumor burden within 150 days post-implant. (B) Representative images of the xenograft tumors labeled with the human cell marker MHC I. Scale, 300 μ m. (C) Sequential coronal sections of FFPE xenograft brains were labeled with anti-Sox2 antibody recognizing both human and mouse (h,m) proteins, with human-specific antibodies against nestin, GFAP, and MHC I, and stained with reticulin. GFAP and reticulin are markers of the glial (G) and sarcomatous (S) phases of the biphasic GS tumors. Representative images for Sox2-positive and Sox2-negative tumors are shown. Scale, 40 μ m. (D) Colabeling of the human marker MHC I (red) and Sox2 (brown). Human cells expressing Sox2 in the tumor (T) (green arrowhead) and mouse brain cells expressing Sox2 (black arrow) are shown for representative images of Sox2-expressing and Sox2-deficient xenograft tumors. Scale, 20 μ m. (E) α-SMA is expressed throughout the tumor from Sox2-deficient HF2303 cells.



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compare these results with Sox2 activity in isogenic GBM CSCs, lentivirus-shRNA-mediated Sox2 knockdown in HF2303 neurospheres was attained (Figure W1), and differential gene expression between Sox2-positive (parental and NSV-shRNA) and Sox2deficient (Sox2shRNA1 and Sox2shRNA2) was similarly analyzed. Applying a 3.5-fold change cutoff (1.8 log₂), Sox2 knockdown affected the expression of 736 genes in SDCs (453 downregulated and 283 upregulated) and 799 genes in CSCs (341 downregulated and 458 upregulated) (Figure 7A). Table W1 contains the complete DEG lists. Only 62 transcripts were commonly regulated in SDCs and CSCs, whereas 35 presented opposite change in expression between the two groups (Figure 7A and Table W1). Overlap with Sox2 binding to cis-regulatory sequences was observed for 25.1% and 15.1% of DEGs in SDC and CSC, respectively (Table W1), suggesting a direct regulation by Sox2 for this subset. A total of 94 DEGs overlapped with GBM molecular subclass signature genes [15]. In Sox2-deficient cells, proneural and classic signature genes were notably downregulated in SDCs and to a lesser extent in CSCs, whereas mesenchymal signature genes were upregulated in both groups (Figure 7B and Table W1).

To compare the above results with genes correlated with Sox2 expression in tumor tissues, gene expression data for 517 GBMs in TCGA were analyzed to identify 56 and 17 genes presenting positive and negative correlation with Sox2 expression, respectively (Table W2 and Figure 7C). Significantly, 28.6% of the positively correlated genes overlapped with the DEGs in HF2303: 3 genes in both CSC and SDC states, 1 in CSCs, and the remaining 12 in the SDC state (Table W2 and Figure 7C), suggesting that genes correlated with Sox2 expression in the tumors are better represented in SDCs than in CSCs.

Top canonical signaling pathways and top GO network pathways representing the DEGs on Sox2 knockdown in CSCs and SDCs were identified using IPA and GeneGO/MetaCore, respectively (Figure 7D). Genes altered in SDCs were more connected in the pathway analyses than those in CSCs. Chemokine signaling was significantly affected by Sox2 knockdown in HF2303 cells. IL-8 expression was downregulated in both CSC and SDCs (Figure 7E), and DEG genes were enriched in IL-8 signaling components, especially in SDCs (Figure 7D). Chemokine (C-X-C motif) receptor 4 (CXCR4), a chemokine receptor involved in glioma cell invasion, was downregulated in SDC (Figure 7E), affecting CXCR4 signaling (Figure 7D).

Wnt receptors Frizzled homolog 2, as well as ligands Wnt2, Wnt3, and Wnt5B, were upregulated, whereas low-density lipoproteinrelated protein 1, a negative regulator, and lymphoid enhancerbinding factor 1, a transcription mediator of Wnt signaling, were

downregulated (Figure 7E). Up-regulation of Wnt target genes connective tissue growth factor, cyclin D1, and Dickkopf homolog 1 (Figure 7E) provide additional evidence that Wnt signaling is activated on Sox2 knockdown in SDCs (Figure 7D). Genes involved in GBM, insulin-like growth factor 1 (IGF-1), protein kinase A, and ρ GTPase signaling were also significantly enriched in the SDC data set (Figure 7D). The top GO networks indicate that Sox2 loss in SDCs results in down-regulation of genes involved in central nervous system development and up-regulation of genes involved in cell-cell signaling (Figure 7D). Sox2 knockdown in CSCs led to up-regulation of genes associated with response to retinoic acid (Figure 7D). Conversely, retinoic acid decreased Sox2 expression in GBM neurosphere cells [33].

On Sox2 knockdown, α -SMA (actin, α 2) was upregulated in CSCs (Figure 7E) and in SDCs (by 3.2-fold, not shown), in agreement with the widespread expression of α -SMA observed for the Sox2-deficient GS xenografts (Figure 6E) and with the lower expression of Sox2 in the sarcomatous phase observed for Sox2positive tumors (Figure 6C). Genes commonly altered on Sox2 knockdown in both SDC and CSC states are enriched in ESC genes, including Sox21 and Tweety 1, both positively correlated with Sox2 expression in the TCGA data set (Figure 7C), CCR4-NOT transcription complex subunit 3 (CNOT3), fatty acid binding protein 7 (FABP7), and podoplanin (PDPN) (Figure 7E). Several genes associated with stem/progenitor cells were exclusively downregulated in SDCs: Brain and acute leukemia, cytoplasmic (BAALC), smoothened homolog (SMO), Sal-like 1 (SALL1), nestin (NES), and platelet-derived growth factor receptor α (PDGFR α) (Figure 7E), in agreement with enrichment of ESC self-renewal signaling genes in SDCs (Figure 7D). Genes downregulated in Sox2-deficient CSCs include n-Myc (MYCN), Krüppel-like factor 4 (KLF4), integrin-α6 (ITGA6), and aldehyde dehydrogenase 1 family, member A1 (ALDH1A1) (Figure 7E). A schematic model summarizing the main findings reported here is shown in Figure 7F.

Discussion

Sox2 is considered a CSC marker for GBMs due in part to its ubiquitous expression in tumorigenic neurosphere cultures [16,18-21] and analogous role in the determination of NSC identity [6]. However, Sox2 expression in 9% to 85% of GBM tumor cells reported here, and comparable to previous reports [13,28,29], indicates that Sox2 expression encompasses more differentiated neoplastic cells as well. To investigate Sox2 function in GBM differentiated cells, we used a rare low-passage SDC line derived from

Figure 7. Sox2 regulates distinct gene sets in GBM stem and differentiated cells. (A) Venn diagram shows the overlap of genes differentially expressed on Sox2 knockdown in HF2303 SDC and CSC. Down arrow, downregulated, positive correlation with Sox2 expression; up arrow, upregulated, negative correlation with Sox2 expression. (B) Genes differentially regulated in HF2303 CSC and SDC that overlap with the subclass signature. On Sox2 knockdown, proneural (PN) and classic (CL) signature genes are downregulated in SDC and modestly in CSC [positive (+) correlation], whereas mesenchymal (MES) signature genes are upregulated [negative (-) correlation] in both groups. The pattern of expression of neural (NL) signature genes did not correlate with Sox2 expression. (C) Sox2 expression in the TCGA GBM data set was significantly correlated with 73 genes. The 517 GBM cases are ordered according to the level of Sox2 expression and colored by their molecular class (red, mesenchymal; blue, classic; green, neural; purple, proneural). The mRNA expression data were converted to z scores, averaged across probes for each gene, and then ordered by hierarchical clustering. The heat map displays the standardized intensity with red high and green low. Select genes positively correlated with Sox2 expression are noted. Overlap with genes downregulated on Sox2 knockdown in HF2303 SDC and CSC are indicated by (*) and (**), respectively. Genes that are direct targets of Sox2 (#) or part of subclass signatures are indicated. (D) Signaling pathway analysis of the DEGs in CSC and SDC shows the top signaling pathways (IPA) and top GO networks (GeneGo/MetaCore). (E) Graph with selected DEGs between Sox2-expressing and Sox2-deficient HF2303 cells, represented as log₂ fold change. (F) Schematic representation of Sox2 role in GBM cell plasticity and transcriptional regulation in stemlike and differentiated GBM cells.

a patient with GBM that retains Sox2 expression, along with dedifferentitation and tumorigenic potential [19]. Tissue culture manipulation, such as the use of FBS to induce astrocytic differentiation, has provided insights into the differentiation potential of both NSCs [34] and GBM CSCs [35]. High-passage 10% FBS cell cultures incur accumulation of genomic abnormalities leading to a deviation of the GBM phenotype, even when these cells are tumorigenic [16,36]. Low-passage 10% FBS cultures commonly present down-regulation of NSC markers, including Sox2, and loss of tumorigenic potential [16,19], thus only partially representing the molecular profile of differentiated cells in the tumor.

Despite the caveats for the use of FBS as differentiation stimulus for GBM cells, we report a significant overlap of genes affected by Sox2 knockdown in SDCs and genes highly correlated with Sox2 expression in GBM tumor tissue profiled by TCGA. Several genes associated with ESCs and cancer were commonly downregulated in both CSCs and SDCs on Sox2 knockdown: Sox21, a known Sox2 target in GBMs [32] and mediator of Sox2 activity in several cell types, including colon cancer [37]; Tweety 1, a proneural subclass signature member and ER protein also required in embryonic and brain development [38]; CNOT3, involved in ESC self-renewal [39]; FABP7, expressed in radial glia cells and associated with GBM malignancy and cell migration [40]; and PDPN, involved in tumor invasion [41]. Several other genes associated with stem and progenitor cells were exclusively downregulated in Sox2-deficient SDCs: BAALC, restricted to neuroectoderm tissues and overexpressed in acute leukemia and GBM [42]; Shh receptor SMO; SALL1, a zinc finger transcription factor activator of NANOG in ESCs [43]; and the known Sox2 targets nestin and PDGFR\alpha [32]. Wnt pathway activation on Sox2 knockdown, reported here for GBM SDCs, has been observed in osteosarcomas [44]. Interestingly, Wnt activation has been shown to induce differentiation of GBM CSCs [45]. Activation of CXCR4 expression by Sox2 is in agreement with previous reports [21]. Paracrine IL-8 in perivascular niches has been recently shown to regulate GBM CSCs [46], and here, we report for the first time a strong correlation of Sox2 expression in GBM SDCs and CSCs with IL-8 expression and signaling. Activation of IGF-1 signaling on Sox2 knockdown in SDCs may explain the reported sensitivity of Sox2deficient GBM cells to the IGF-1R inhibition [21]. Loss of Sox2 led to down-regulation of several genes exclusively in CSCs, in particular transcription factors that act in concert with Sox2 to reprogram somatic cells into iPS cells (MYCN and KLF4) [47] and other genes previously associated with CSCs (ALDH1A1 and ITGA6) [48].

Sox2 is part of the gene signature for the GBM proneural molecular subclass [15]. Here, we show that Sox2 was correlated with the expression of both proneural and classic signature genes in the TCGA data set and in vitro functional studies. Furthermore, loss of Sox2 resulted in up-regulation of mesenchymal signature genes in vitro. No significant difference in Sox2 expression between recurrent and newly diagnosed GBMs was observed at protein level in this study or at transcript level in paired samples [30]. Mechanisms leading to the upregulation of Sox2 in GBM tumors include gene amplification [12,14,18], promoter hypomethylation [12], translational regulation [49], and activated signaling pathways [50]. Posttranslational regulation of Sox2 level by proteolysis has been previously observed in committed neuronal precursor cells [7]. Sox2 protein stabilization by MG132 treatment reported here has also been observed in mouse ESCs [51], and although the mechanism remains to be determined, it has possible implications for the clinical use of proteasome inhibitors.

Sox2 expression is retained in patient-derived orthotopic xenografts generated from neurosphere cells, in agreement with previous studies [52], and also from subcutaneously propagated tumors. It has been previously reported that Sox2 loss in GBM neurospheres impairs tumorigenicity [20,50]. Here, we show that Sox2 knockdown in GBM SDCs abolished dedifferentiation and acquisition of CSC phenotype in vitro and decreased tumorigenicity. Ectopic Sox2 expression was not sufficient to reprogram typical Sox2-deficient low-passage GBM SDCs to regain plasticity and tumorigenicity, possibly due to insufficient levels of required cofactors, similar to what has been observed in a high-passage GBM cell line [12]. Furthermore, we show that in the context of biphasic GSs, Sox2 specifically regulates the glial component, which displays histologic features typical of GBMs, whereas the loss of Sox2 did not affect the sarcomatous compartment, identified by reticulin staining of the dense extracellular matrix and composed of GFAP-negative spindle cells [24]. These findings have potential clinical implications in that any intervention aimed at decreasing Sox2 activity or downstream pathways in GS could favor metaplastic sarcomatous tumor growth.

We propose that Sox2 function in GBMs resemble not only its role in ESC and NSC identity, maintenance, and fate but also its function in committed progenitors and proliferating astrocytes [7]. As evidences continue to point to a "plastic CSC model" [53] for GBMs, our results indicate that Sox2 may be central to the maintenance of developmental plasticity during glial tumor progression, regulating dedifferentiation and acquisition of CSC properties. Furthermore, Sox2 regulates gene expression and signaling critical to malignancy in more differentiated cells.

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Supplementary materials

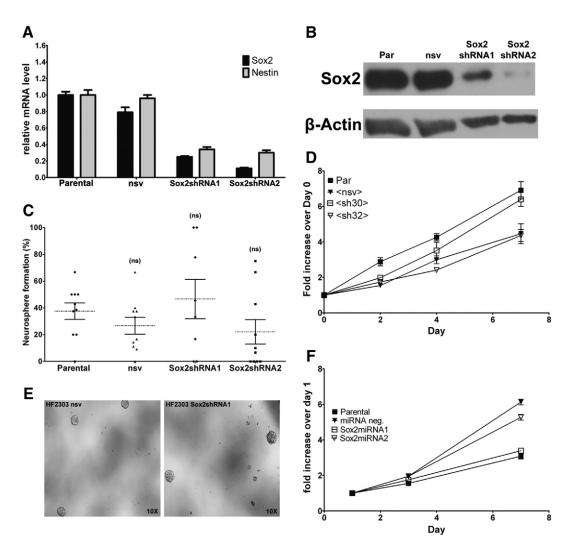


Figure W1. Sox2 knockdown in HF2303 neurospheres cells. (A) Sox2 and nestin mRNA levels in HF2303 (parental) cells and cells transfected with shRNA constructs targeting Sox2 (Sox2shRNA1 and Sox2shRNA2) or nonsilencing miRNA (NSV) were evaluated by quantitative reverse transcription (RT)–PCR. Results represent means \pm SEM, n=3. (B) Sox2 protein expression in whole-cell lysates determined by Western blot analysis confirms Sox2 knockdown in Sox2shRNA1 and Sox2shRNA2 lines; α-actin was used as loading control. (C) The effect of Sox2 knockdown on clonogenicity. Cells were seeded at low density in 96-well plates and cultured in NM. Neurosphere formation was monitored and quantified at day 11 (mean \pm SEM). ns, nonsignificant; P > .05 in relation to parental cells. (D) The effect of Sox2 knockdown on proliferation of HF2303 neurospheres. Cells were seeded at low density in 96-well players and cultured in NM. Cell viability was quantified by intracellular ATP level measured using CellTiter-Glo Luminescent Cell Viability Assay (Promega). (E) Representative images contrast Sox2-expressing and Sox2-deficient cells in NM. (F) The effect of Sox2 knockdown on proliferation of HF2303 monolayers. Cells were seeded at low density in 96-well players and cultured in FBS. Viability was measured on days 1, 3, and 7.

Table W1. DEGs in HF2303 GBM Cells on Sox2 Knockdown.

Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
ATP-binding cassette, subfamily A (ABC1), member 8	ABCA8	-4.3		
ATP-binding cassette, subfamily A (ABC1), member 9	ABCA9	-1.9		
ATP-binding cassette, subfamily B (MDR/TAP), member 9	ABCB9	-1.8		
ATP-binding cassette, subfamily D (ALD), member 1	ABCD1	-1.9		
Abhydrolase domain containing 11	ABHD11	-2.9		
Abhydrolase domain containing 3	ABHD3	-2.0		
ACN9 homolog (Saccharomyces cerevisiae)	ACN9	-2.3		CI
Acyl-CoA synthetase bubblegum family member 1	ACSEG	-5.8		CL
Acyl-CoA synthetase family member 2 ADAM metallopeptidase with thrombospondin type 1 motif, 1	ACSF2 ADAMTS1	-1.8 -2.2		
Administration Al receptor	ADAM131 ADORA1	-2.2 -2.1		
AE binding protein 1	AEBP1	-2.3		
Angiotensinogen (serpin peptidase inhibitor, clade A, member 8)	AGT	-5.6		
Adenosylhomocysteinase-like 2	AHCYL2	-3.0		
Allograft inflammatory factor 1-like	AIF1L	-3.9		
AKNA domain containing 1	AKNAD1	-2.2		
Aldo-keto reductase family 1 member 15 (ALD2)	AKR1B15	-3.1		
V-akt murine thymoma viral oncogene homolog 2	AKT2	-3.0		CL
Aldehyde dehydrogenase 1 family, member A1	ALDH1A1	-7.7	(1)	-
Anaplastic lymphoma kinase (Ki-1)	ALK	-4.2	(1)	
Ankyrin 2, neuronal	ANK2	-4.1		
Ankyrin repeat domain 20 family, member A4	ANKRD20A4	-3.8		
Ankyrin repeat domain 9	ANKRD9	-1.9		
Ankyrin repeat and sterile α motif domain containing 1B	ANKS1B	-2.3	(1)	NL
Adaptor-related protein complex 2, α 1 subunit	AP2A1	-2.1	(1)	
Adenomatosis polyposis coli down-regulated 1	APCDD1	-2.6	(-)	
Amyloid β (A4) precursor-like protein 1	APLP1	-2.2		
Arf-GAP with ρ-GAP domain, ANK repeat and PH domain-containing protein 3	ARAP3	-1.9		
ADP-ribosylation factor-like 4A	ARL4A	-2.3	(1)	
Aryl-hydrocarbon receptor nuclear translocator 2	ARNT2	-3.2		
Acetylserotonin O-methyltransferase-like	ASMTL	-1.8		
ATPase, class V, type 10B	ATP10B	-2.5		
B-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)	B3GAT1	-2.8	(1)	
Brain and acute leukemia, cytoplasmic	BAALC	-3.3		
Brain-specific angiogenesis inhibitor 3	BAI3	-3.3	(1)	PN
Bardet-Biedl syndrome 2	BBS2	-1.9		
Butyrylcholinesterase	BCHE	-4.0	(1)	
Brain expressed, X-linked 1	BEX1	-3.1		PN
Bloom syndrome	BLM	-2.2	(1)	CL
Bone marrow stromal cell antigen 2	BST2	-2.5		
BTB (POZ) domain containing 11	BTBD11	-3.7	(1)	
BTG family, member 2	BTG2	-3.0		
Chromosome 15 open reading frame 59	C15orf59	-2.6		
Chromosome 18 open reading frame 56	C18orf56	-3.8		
Chromosome 1 open reading frame 54	Clorf54	-3.2		MES
Chromosome 1 open reading frame 61	C1orf61	-3.5	(1)	
Chromosome 2 open reading frame 40	C2orf40	-2.1		
Chromosome 2 open reading frame 76	C2orf76	-1.9		
Chromosome 3 open reading frame 70	C3orf70	-4.9		
Chromosome 7 open reading frame 43	C7orf43	-2.1		
Chromosome 8 open reading frame 4	C8orf4	-3.4	(1)	
Carbonic anhydrase II	CA2	-5.3		
Calcium channel, voltage-dependent, P/Q type, α 1A subunit	CACNA1A	-2.1	(1)	
Calcium channel, voltage-dependent, β 2 subunit	CACNB2	-2.1	(1)	
Cell adhesion molecule 4	CADM4	-3.7		
Ca2+-dependent activator protein for secretion 2	CADPS2	-2.5	(1)	
Calcitonin receptor-like	CALCRL	-4.9	(1)	CT.
Calcium/calmodulin-dependent protein kinase (CaM kinase) II β	CAMK2B	-2.0	(1)	CL
Calcium/calmodulin-dependent protein kinase II inhibitor 1	CANDO	-2.7	(1)	
Caspase recruitment domain family, member 8	CARD8	-2.7		MEC
Caspase 1, apoptosis-related cysteine peptidase (interleukin 1, β, convertase)	CASP1	-2.1		MES
Coiled-coil domain containing 74B	CCDC74B	-2.0		
CD276 molecule	CD276	-1.8	(1)	
CD83 molecule	CD83	-2.4	(1)	
Cat eye syndrome chromosome region, candidate 1	CECR1	-4.2 3.6		
Cell cycle exit and neuronal differentiation 1	CEND1	-3.4		
Centrosomal protein 112kDa (CCDC46) Complement factor I	CEP112	-2.0		
	CFI	-2.9		
•	CHCHD10	2.2		
Coiled-coil-helix-coiled-coil-helix domain containing protein 10, mitochondrial Cell adhesion molecule with homology to L1CAM (close homolog of L1)	CHCHD10 CHL1	-2.2 -7.9	(1)	

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Cholinergic receptor, nicotinic, α 9	CHRNA9	-2.8		
Carbohydrate (keratan sulfate Gal-6) sulfotransferase 1	CHST1	-3.8		
Carbohydrate sulfotransferase 10	CHST10	-2.5		
Carbohydrate (N-acetylglucosamine-6-O) sulfotransferase 2	CHST2	-2.8		
Creatine kinase, brain Chloride channel 2	CKB CLCN2	-1.9 -2.7		
Claudin 23	CLCN2 CLDN23	-2.6		
CAP-GLY domain containing linker protein 2	CLIP2	-2.8		CL
Calsyntenin 2	CLSTN2	-2.5	(1)	
CCR4-NOT transcription complex, subunit 3	CNOT3	-2.2		
Canopy 4 homolog (zebrafish)	CNPY4	-2.5		
Coagulation factor C homolog, cochlin (Limulus polyphemus)	COCH	-2.1		
Collagen, type XXII, α 1	COL22A1	-2.2	(1)	
Collagen, type IV, α 5 (Alport syndrome)	COL4A5	-2.3	(1)	
Collagen, type IV, \alpha 6	COL4A6	-4.8	(1)	
Collagen, type IX, \alpha 2	COL9A2 CORO2B	-2.2 -2.0	(1)	
Coronin, actin binding protein, 2B DNA-directed RNA polymerase III subunint RPC9	CRCP	-2.0 -1.9	(1)	
Cysteine-rich protein 1 (intestinal)	CRIP1	-3.0		
Crystallin, γ S	CRYGS	-4.5		
Cysteine and glycine-rich protein 2	CSRP2	-2.9		
Chemokine (C-X3-C motif) ligand 1	CX3CL1	-2.4	(1)	
Coxsackie virus and adenovirus receptor	CXADR	-2.3	(1)	
Chemokine (C-X-C motif) ligand 14	CXCL14	-3.3		
Chemokine (C-X-C motif) receptor 4	CXCR4	-2.4	(1) (3)	
Cytochrome b5 reductase 2	CYB5R2	-3.5		
Cytochrome c, somatic	CYCS	-1.9	(1)	
Cytochrome P450, family 51, subfamily A, polypeptide 1	CYP51A1	-2.2		
Dysbindin (dystrobrevin binding protein 1) domain containing 1	DBNDD1	-2.2		
Dehydrogenase/reductase (SDR family) member 3	DHRS3	-2.2		
DIRAS family, GTP-binding RAS-like 3	DIRAS3 DLG3	-2.1 -2.6		
Discs, large homolog 3 (neuroendocrine-dlg, <i>Drosophila</i>) DLGAP1 antisense RNA 2	DLG3 DLGAP1-AS2	-2.3		
Distal-less homeobox 1	DLX1	-1.8		
DNA-directed RNA polymerase III subunit RPC9	DNAJC30	-1.8		
Dynamin 3	DNM3	-1.8	(1)	PN
Dipeptidyl-peptidase 4 (CD26, adenosine deaminase complexing protein 2)	DPP4	-2.1		
Dorsal inhibitory axon guidance protein	DRAXIN	-4.0		
Dystrobrevin, α	DTNA	-2.8	(1)	
Dual specificity phosphatase 4	DUSP4	-2.8		
Dual specificity phosphatase 6	DUSP6	-2.9		
Endothelin receptor type B	EDNRB	-5.0		
Endonuclease/exonuclease/phosphatase family domain containing 1	EEPD1 EIF1AY	-1.8 -4.2		
Eukaryotic translation initiation factor 1A, Y-linked ELAV (embryonic lethal, abnormal vision, <i>Drosophila</i>)-like 1 (Hu antigen R)	ELAVL1	-4.2 -2.7		
Engulfment and cell motility 1	ELMO1	-3.5	(1)	
Elastin (supravalvular aortic stenosis, Williams-Beuren syndrome)	ELN	-2.8	(1)	
Elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2	ELOVL2	-8.3	(1)	CL
ER membrane protein complex subunit 10	EMC10	-1.9		
Energy homeostasis associated	ENHO	-2.1		
Enolase superfamily member 1	ENOSF1	-1.8		
Epoxide hydrolase 4	EPHX4	-3.1		
Estrogen-related receptor γ	ESRRG	-2.3	(1)	
Fatty acid binding protein 5 (psoriasis-associated)	FABP5	-1.9	(1)	
Fatty acid binding protein 7, brain	FABP7	-10.4		
Family with sequence similarity 196, member A	<i>FAM196A</i> FAM212B	-3.7 -2.4		
Family with sequence similarity 212, member B Family with sequence similarity 43, member A	FAM43A	-2.4 -4.7	(1)	
Family with sequence similarity 4.5, member C	FAM69C	-5.8	(1)	
Family with sequence similarity 84, member B	FAM84B	-2.6	(1)	
Fasciculation and elongation protein ζ 1 (zygin I)	FEZ1	-2.4		
Fin bud initiation factor	FIBIN	-4.9		
Hypothetical gene supported by AK094963	FLJ37644	-2.5		
Fibronectin leucine rich transmembrane protein 3	FLRT3	-3.3		
V-fos FBJ murine osteosarcoma viral oncogene homolog	FOS	-2.2		
Forkhead box S1	FOXS1	-2.1		
FERM and PDZ domain containing 3	FRMPD3	-3.2		C*
Frizzled homolog 3 (Drosophila)	FZD3	-3.4		CL
Γ-aminobutyric acid (GABA) A receptor, β 1	GABRB1	-2.9		
Galactose-3-O-sulfotransferase 4 UDP-N-acetyl-α-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1)	GAL3ST4 GALNT1	-2.0 -1.8	(1)	
ODI-14-acctyr-a-1-garactosamme-porypeptide 14-acctyrgaractosammyntansierase 1 (Ganvac-11)	GALIVII	-1.0	(1)	

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Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
UDP-N-acetyl-α-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 (GalNAc-T4)	GALNT4	-1.8		CL
$UDP\textit{-N-}acetyl-\alpha-D-galactosamine: polypeptide \textit{N-}acetylgalactosaminyltransferase-like 1}$	GALNTL1	-4.5		
Growth associated protein 43	GAP43	-5.2	(1)	
Growth arrest-specific 8	GAS8	-3.0 -2.5		
Gap junction protein, β 2, $26kDa$ Glycosyltransferase 25 domain containing 2	<i>GJB2</i> GLT25D2	-2.5 -2.0		
Guanine nucleotide binding protein (G protein), α inhibiting activity polypeptide 1	GNAI1	-2.1	(1)	NL
Guanine nucleotide binding protein (G protein), γ 11	GNG11	-3.2	(1)	
Guanine nucleotide binding protein (G protein), γ 2	GNG2	-5.2	(1)	
Guanine nucleotide binding protein (G protein), γ 7	GNG7	-2.1		CL
Glycoprotein M6A	GPM6A	-4.8		PN
Glycoprotein M6B	GPM6B	-3.7		
G protein–coupled receptor 162 G protein–coupled receptor 56	GPR162 GPR56	-2.0 -3.2		
Growth factor receptor-bound protein 10	GRB10	-1.9		
Hyaluronan synthase 3	HAS3	-4.9		
Histone deacetylase 4	HDAC4	-3.0	(1)	
Hepatocyte cell adhesion molecule	HEPACAM	-3.9		
Hairy/enhancer-of-split related with YRPW motif 1	HEY1	-2.6	(1)	
HHIP antisense RNA 1	HHIP-AS1	-4.8		
Huntingtin interacting protein 1	HIP1	-3.0		
High-mobility group AT-hook 1	HMGA1	-1.8		
Homer homolog 1 (Drosophila)	HOMER1	-6.5	(1)	
Hippocalcin-like 1	HPCAL1 HS3ST1	-2.2	(1)	
Heparan sulfate (glucosamine) 3-O-sulfotransferase 1 Heat shock 70kDa protein 6 (HSP70B')	HSPA6	-2.7 -2.1	(1)	
Heat shock γοκορ protein σ (1191766) Heat shock protein, α-crystallin-related, B6	HSPB6	-4.1		
Indoleamine 2,3-dioxygenase 1	IDO1	-2.8		
Iduronate 2-sulfatase (Hunter syndrome)	IDS	-1.9		
Intraflagellar transport 57 homolog (Chlamydomonas)	IFT57	-2.4		
Intraflagellar transport 81 homolog (Chlamydomonas)	IFT81	-1.8		
Insulin-like growth factor 2 mRNA binding protein 3	IGF2BP3	-5.7		
Insulin-like growth factor binding protein 2, 36kDa	IGFBP2	-6.0		
Insulin-like growth factor binding protein 3	IGFBP3	-1.9	(1)	
Insulin-like growth factor binding protein 5	IGFBP5 IGSF1	-2.0		
Immunoglobulin superfamily, member 1 Immunoglobulin superfamily, member 5	IGSF1 IGSF5	-2.0 -2.1		
Interleukin 17 receptor B	IL17RB	-2.0		
Interleukin-1 receptor accessory protein-like 1	IL1RAPL1	-8.5		PN
Interleukin-27 receptor, α	IL27RA	-2.2	(1)	
Interleukin-33	IL33	-5.0		
Interleukin-8	IL8	-2.4	(1)	
Inositol 1,4,5-triphosphate receptor, type 2	ITPR2	-2.4	(1)	
Potassium voltage-gated channel, subfamily F, member 1	KCNF1	-2.0	(1)	CL
Kv channel interacting protein 4	KCNIP4	-2.5	(1)	
Potassium inwardly-rectifying channel, subfamily J, member 2 Potassium channel, subfamily K, member 12	KCNJ2 KCNK12	-2.0 -2.8	(1)	
Potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4	KCNN4	-2.5		
Potassium channel tetramerisation domain containing 12	KCTD12	-2.4		
Potassium channel tetramerisation domain containing 13	KCTD13	-2.6		
Potassium channel tetramerisation domain containing 17	KCTD17	-2.2		
Lysine (K)-specific demethylase 5D	KDM5D	-2.8		
KIAA0408	KIAA0408	-2.3		
KIAA1147	KIAA1147	-1.8		
KIAA1211	KIAA1211	-2.6	(4)	
Kinesin family member 13B	KIF13B	-2.0	(1)	
Kinesin family member 26B Kinesin family member 5C	KIF26B KIF5C	-2.5 -5.0		
Kelch domain containing 9	KLHDC9	-2.6		
Kelch-like 29 (Drosophila)	KLHL29	-2.5		
Kynureninase (L-kynurenine hydrolase)	KYNU	-2.6	(1)	MES
Laminin, α 1	LAMA1	-3.1		
Lymphoid enhancer-binding factor 1	LEF1	-3.2		
LEM domain containing 1	LEMD1	-5.2		
Lectin, galactoside-binding, soluble, 9B	LGALS9B	-1.9		
Leucine-rich repeat LGI family, member 2	LGI2	-3.9		
Leukemia inhibitory factor receptor α	LIFR	-2.3		
LIM domain kinase 1 Long intergenic popprotein coding RNA 341	LIMK1 LINC00341	-2.9 -2.1		
Long intergenic nonprotein coding RNA 341 Long intergenic nonprotein coding RNA 623	LINC00341 LINC00623	-2.1 -3.2		
Lipase, endothelial	LIPG	-3.1		
•	LOC100128288	-1.8		

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SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
LON peptidase N-terminal domain and ring finger 1	LONRF1	-2.7		
LON peptidase N-terminal domain and ring finger 2	LONRF2	-2.9	(1)	
Lysophosphatidic acid receptor 1	LPAR1	-1.9		
Latrophilin 3	LPHN3	-2.0	(1)	PN
Lipoprotein lipase	LPL	-4.2		
Lipid phosphate phosphatase-related protein type 5	LPPR5	-4.4		
Low-density lipoprotein–related protein 1 (α-2-macroglobulin receptor)	LRP1	-1.9	(4)	
Low-density lipoprotein–related protein 1B (deleted in tumors)	LRP1B	-2.5	(1)	
Low-density lipoprotein receptor–related protein 4 Leucine rich repeat containing 4C	LRP4 LRRC4C	-3.0 -3.4	(1)	
Leucine rich repeat containing 4C Leucine rich repeat neuronal 2	LRRN2	-4.5	(1)	
Leucine rich repeat neuronal 3	LRRN3	-2.5		
LY6/PLAUR domain containing 1	LYPD1	-2.7		
Leucine zipper, putative tumor suppressor 1	LZTS1	-4.8		
Mal, T cell differentiation protein	MAL	-4.8		
MAM domain containing 2	MAMDC2	-3.2	(1)	
Mannosidase, α, class 1C, member 1	MAN1C1	-3.9		
Mitogen-activated protein kinase kinase 5	MAP2K5	-3.1		
Mitogen-activated protein kinase 10	MAPK10	-1.8	(1)	
Mitogen-activated protein kinase 4	MAPK4	-2.1		
Mitogen-activated protein kinase 8 interacting protein 1	MAPK8IP1	-2.8		
MARCKS-like 1	MARCKSL1	-2.1		PN
Midkine (neurite growth-promoting factor 2)	MDK	-2.4	(4)	O.
Mesenchyme homeobox 2	MEOX2	-2.0	(1)	CL
Meteorin, glial cell differentiation regulator	METRN MFSD6	-3.1 -1.9		
Major facilitator superfamily domain containing 6 Matrix Gla protein	MGP	-1.9 -1.9		
MKL/myocardin-like 2	MKL2	-2.9		
Megalencephalic leukoencephalopathy with subcortical cysts 1	MLC1	-2.4		
Musashi homolog 1 (Drosophila)	MSI1	-2.1		
Microseminoprotein, prostate associated	MSMP	-4.6		
Metallothionein 1X	MT1X	-2.2		
Myosin binding protein C, slow type	MYBPC1	-5.0		NL
Myosin VB	MYO5B	-1.9		
Neurocan	NCAN	-2.9		
Noncompact myelin associated protein	NCMAP	-2.5		
NDRG family member 4	NDRG4	-3.2		
NEL-like 2 (chicken)	NELL2	-2.5		
Nestin	NES	-4.5	(5)	CL
Nuclear factor I/B	NFIB	-3.0	(1)	
Nuclear factor of κ light polypeptide gene enhancer in B cells inhibitor, β	NFKBIB	-2.7	(1)	
Neuroligin 4, X-linked	NLGN4X	-1.8	(1)	
Neuroligin 4, Y-linked	NLGN4Y	-7.3 -3.6		
Nicotinamide nucleotide adenylyltransferase 2	NMNAT2		(1)	
Nicotinamide nucleotide adenylyltransferase 3 Neuro-oncological ventral antigen 1	NMNAT3 NOVA1	-2.1 -2.7	(1) (1)	
Neuronal PAS domain protein 2	NPAS2	-3.9	(1)	
Neuronal PAS domain protein 3	NPAS3	-3.3	(1)	CL
Neuropeptide B	NPB	-2.6	(-)	
Neuropeptide Y receptor Y5	NPY5R	-6.6		
Notch-regulated ankyrin repeat protein	NRARP	-3.2		
Neuronal cell adhesion molecule	NRCAM	-4.3		
Neurogranin (protein kinase C substrate, RC3)	NRGN	-2.8	(1)	
Neuropilin 2	NRP2	-1.8	(1)	
Netrin 1	NTN1	-2.2	(1)	
OAF homolog (Drosophila)	OAF	-2.4		
Olfactomedin-like 3	OLFML3	-2.6		
Oxysterol binding protein-like 6	OSBPL6	-2.4	(1)	
Orthopedia homeobox	OTP	-2.2		
Prostate androgen-regulated mucin-like protein 1	PARM1	-4.2		
Protocadherin 20	PCDH20	-3.4	(1)	
Protocalherin 9	PCDH9	-6.4	(1)	
Protocadherin γ subfamily A, 5	PCDHGA5	-2.5		
Protocadherin γ subfamily C, 3 Phosphodiesterase 1B, calmodulin-dependent	PCDHGC3 PDE1B	-2.0 -5.2		
Phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, <i>Drosophila</i>)	PDE4B	-3.2 -2.1	(1)	
Phosphodiesterase 8B	PDE8B	-3.3	(1)	
Platelet-derived growth factor receptor, α polypeptide	PDGFRA	-4.2	(2)	
PDZ and LIM domain 4	PDLIM4	-2.4	\=/	
Podoplanin	PDPN	-6.8		MES
Paternally expressed 10	PEG10	-2.6		

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Pellino homolog 1 (<i>Drosophila</i>)	PELI1	-2.9		PN
GPI deacylase	PGAP1	-2.8	(1)	
PiggyBac transposable element derived 5	PGBD5	-2.9		NL
Pleckstrin homology-like domain, family A, member 1	PHLDA1	-2.0		
Phosphatidylinositol glycan anchor biosynthesis, class F Protein (peptidylprolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	PIGF PIN4	-3.8 -2.1		
Pigeon homolog (<i>Drosophila</i>)	PION	-2.2		
Phosphatidylinositol transfer protein, cytoplasmic 1	PITPNC1	-3.2		
Protein kinase (cAMP-dependent, catalytic) inhibitor α	PKIA	-2.6	(1)	
Protein kinase (cAMP-dependent, catalytic) inhibitor β	PKIB	-3.3		
Phospholipase C, γ 1	PLCG1	-3.1		
Phospholipase D family, member 3	PLD3	-3.0	(1)	
Pleckstrin homology domain containing, family B (evectins) member 1 Pleckstrin homology domain containing, family G (with pGef domain) member 1	PLEKHB1 PLEKHG1	-3.1 -2.4	(1)	
Prostate transmembrane protein, androgen induced 1	PMEPA1	-4.5		
Paraneoplastic antigen MA2	PNMA2	-3.4	(1)	
Podocalyxin-like 2	PODXL2	-1.9		PN
POM (POM121 homolog, rat) and ZP3 fusion	POMZP3	-4.4		
Periostin, osteoblast specific factor	POSTN	-5.7	(1)	
Phosphatidic acid phosphatase type 2C	PPAP2C	-2.5		
Protein phosphatase 1, catalytic subunit, β isoform	PPP1CB	-1.8		
Protein kinase, cAMP-dependent, catalytic, β Protein kinase C, θ	PRKACB PRKCQ	-1.8 -2.2		
Proline rich 7 (synaptic)	PRR7	-2.2		
Prostaglandin F2 receptor negative regulator	PTGFRN	-2.3	(1)	
Prothymosin, α (gene sequence 28)	PTMA	-1.9		
Pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1)	PTN	-2.4		
Protein tyrosine phosphatase, nonreceptor type 12	PTPN12	-1.9		
Protein tyrosine phosphatase, nonreceptor type 13 (APO-1/CD95 (Fas)-associated phosphatase)	PTPN13	-2.4	(1)	
Protein tyrosine phosphatase, receptor type, D	PTPRD PAR26	-6.4	(1)	
RAB36, member RAS oncogene family RAN binding protein 2	RAB36 Ranbp2	-3.5 -2.3		
Rap guanine nucleotide exchange factor (GEF) 5	RAPGEF5	-5.4		
RAS, dexamethasone-induced 1	RASD1	-1.9		
Ras interacting protein 1	RASIP1	-4.1		
Ras association (RalGDS/AF-6) domain family 2	RASSF2	-3.0		
Retinol binding protein 7, cellular	RBP7	-2.5		
Regulator of calcineurin 2 Receptor accessory protein 1	RCAN2 REEP1	-2.2 -2.1	(1)	PN
Replication factor C (activator 1) 2, 40kDa	RFC2	-2.1 -1.9	(1)	riv
Raftlin family member 2	RFTN2	-4.9		
Regulator of G-protein signaling 2, 24kDa	RGS2	-3.2	(1)	
Rhomboid domain containing 3	RHBDD3	-2.2		
Ras homolog gene family, member U	RHOU	-3.0	(1)	
Regulating synaptic membrane exocytosis 4	RIMS4	-2.6		
P family GTPase 2	RND2	-1.9		
Ring finger protein 112 Reticulon 1	RNF112 RTN1	-2.0 -4.2		
Sal-like 1 (<i>Drosophila</i>)	SALL1	-7.3	(1)	
Sterile α motif domain containing 13	SAMD13	-2.7		
Sterile α motif domain containing 5	SAMD5	-3.4		
Secretogranin II (chromogranin C)	SCG2	-3.0		
Sodium channel, voltage-gated, type IV, β	SCN4B	-2.4		
Sodium channel, voltage-gated, type IX, α subunit	SCN9A	-1.9	(1)	
Scrapic responsive protein 1	SCRG1	-5.0 2.5	(1)	
Sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A Sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E	SEMA3A SEMA3E	-2.5 -5.1	(1)	
Sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A	SEMA6A	-2.5	(1)	CL
Selenoprotein N, 1	SEPN1	-2.2	(1)	OL.
Septin 3	SEPT3	-2.2		
Septin 5	SEPT5	-2.0		
SET binding protein 1	SETBP1	-2.0	(1)	
Sideroflexin 5	SFXN5	-3.7	(1)	
Sarcoglycan, E	SGCE SCV222	-6.4		
Homolog of rat pragma of Rnd2 SH3 and PX domains 2A	SGK223 SH3PXD2A	-2.4 -2.1	(1)	
SHC (Src homology 2 domain containing) family, member 4	SHC4	-2.1 -6.8	(1) (1)	
SLAIN motif family, member 1	SLAIN1	-3.9	(1)	
Solute carrier family 10 (sodium/bile acid cotransporter family), member 4	SLC10A4	-3.3		
Solute carrier family 1 (glial high affinity glutamate transporter), member 3	SLC1A3	-1.9	(1)	
Solute carrier family 25, member 13 (citrin)	SLC25A13	-2.0	(1)	
Solute carrier family 25 (mitochondrial carrier), member 18	SLC25A18	-3.6		

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Solute carrier family 26, member 7	SLC26A7	-2.4		
Solute carrier family 27 (fatty acid transporter), member 1	SLC27A1	-2.0		
Solute carrier family 35, member B4	SLC35B4	-2.1		
Solute carrier family 35, member F1	SLC35F1	-3.1		
Solute carrier family 39 (zinc transporter), member 12	SLC39A12	-3.0		
Solute carrier family 45, member 3	SLC45A3	-5.6	4.3	
Solute carrier family 4, anion exchanger, member 3	SLC4A3	-3.8	(1)	
Solute carrier organic anion transporter family, member 2A1	SLCO2A1	-3.7		DAT
Solute carrier organic anion transporter family, member 5A1	SLCO5A1	-2.5		PN
SLIT and NTRK-like family, member 3 Sarcolipin	SLITRK3 SLN	-4.1 -6.4		
SWI/SNF-related, matrix-associated, actin-dependent regulator of chromatin, subfamily d, member 3	SMARCD3	-1.9	(1)	
Structural maintenance of chromosomes 6	SMC6	-2.6	(1)	
Smoothened homolog (<i>Drosophila</i>)	SMO	-2.3	(1)	CL
Sine oculis binding protein homolog (Drosophila)	SOBP	-1.9	· /	
Suppressor of cytokine signaling 2	SOCS2	-2.5	(1)	CL
Sortilin-related receptor, L(DLR class) A repeats-containing	SORL1	-3.1		
SRY (sex-determining region Y)-box 15	SOX15	-3.5		
SRY (sex-determining region Y)-box 2	SOX2	-4.0		PN
SRY (sex-determining region Y)-box 21	SOX21	-3.1	(1) (2)	
SRY (sex-determining region Y)-box 3	SOX3	-4.4		
SRY (sex-determining region Y)-box 8	SOX8	-3.1		
SPARC-like 1 (mast9, hevin)	SPARCL1	-2.4		
Sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) 2	SPOCK2	-4.0		
Spondin 1, extracellular matrix protein	SPON1	-4.9		
Secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1)	SPP1	-5.2	(4)	
Sprouty homolog 1, antagonist of FGF signaling (<i>Drosophila</i>)	SPRY1	-3.1	(1)	CT.
Sprouty homolog 2 (Drosophila)	SPRY2	-2.2	(1)	CL
Sprouty homolog 4 (<i>Drosophila</i>)	SPRY4	-1.9 -2.0	(1)	PN
SLIT-ROBO P GTPase activating protein 3 Sorcin	SRGAP3 SRI	-2.8	(1)	PIN
SFRS protein kinase 2	SRPK2	-2.8 -4.0	(1) (1)	
Sushi-repeat-containing protein, X-linked	SRPX	-2.6	(1)	
Single stranded DNA binding protein 4	SSBP4	-2.5	(1)	
ST6 β -galactosamide α -2,6-sialyltranferase 1	ST6GAL1	-3.9	(1)	
Stanniocalcin 1	STC1	-2.3		
Serine/threonine kinase 32A	STK32A	-2.2		
Storkhead box 2	STOX2	-3.2		
Syntaxin 1A (brain)	STX1A	-2.1		
Syntabulin (syntaxin-interacting)	SYBU	-2.1		
Synemin, intermediate filament protein	SYNM	-2.2		
TAF6 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 80kDa	TAF6	-3.4		
Transgelin 3	TAGLN3	-3.4		
Transcobalamin II; macrocytic anemia	TCN2	-2.8		
Tyrosyl-DNA phosphodiesterase 1	TDP1	-2.4		
Transforming growth factor, α	TGFA	-2.5	(1)	
Thrombospondin 2	THBS2	-2.6		
TIMP metallopeptidase inhibitor 4	TIMP4	-4.5	(4)	
Tousled-like kinase 1	TLK1	-2.0	(1)	
Transmembrane 7 superfamily member 2	TM7SF2	-2.1	(1)	
Transmembrane protein 108	TMEM108	-2.3	(1)	
Transmembrane protein 132A	TMEM132A TMEM163	-2.2 -3.2	(1)	
Transmembrane protein 163 Transmembrane protein 255A	TMEM165 TMEM255A	-2.6	(1)	
Transmembrane protein 26	TMEM253A	-2.3		
Transmembrane protein 55A	TMEM55A	-1.9		
Transmembrane protein 86A	TMEM86A	-4.0		
Transmembrane and tetratricopeptide repeat containing 2	TMTC2	-5.2	(1)	
Tumor necrosis factor, α-induced protein 6	TNFAIP6	-4.6	(-)	
Tumor necrosis factor (ligand) superfamily, member 15	TNFSF15	-2.6		
Tumor necrosis factor (ligand) superfamily, member 4 (tax-transcriptionally activated glycoprotein 1, 34kDa)	TNFSF4	-6.9		
TP53 target 1 (nonprotein coding)	TP53TG1	-1.8		
Tumor protein D52	TPD52	-1.8	(1)	
Tubulin polyglutamylase complex subunit 2	TPGS2	-2.7		
Thiamin pyrophosphokinase 1	TPK1	-6.3	(1)	
Tribbles homolog 2 (Drosophila)	TRIB2	-2.0	(1)	CL
TLR4 interactor with leucine-rich repeats	TRIL	-2.9		
Tripartite motif-containing 36	TRIM36	-1.9	(1)	
Tripartite motif-containing 47	TRIM47	-3.1		
Tripartite motif-containing 9	TRIM9	-3.3		
Trichorhinophalangeal syndrome I	TRPS1	-2.2	(1)	

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Thiosulfate sulfurtransferase (rhodanese)-like domain containing 1	TSTD1	-4.6		
Tubulin tyrosine ligase-like family, member 4	TTLL4	-2.8		
Tweety homolog 1 (Drosophila)	TTYH1	-4.2		
Tweety homolog 3 (Drosophila)	TTYH3	-2.4	(1)	
Tubulin, β 2B	TUBB2B	-4.5		
Ubiquitin specific peptidase 14 (tRNA-guanine transglycosylase)	USP14 VANGL2	-2.1		
Vang-like 2 (van gogh, <i>Drosophila</i>) VGF nerve growth factor inducible	VANGL2 VGF	-1.9 -2.0		
Visinin-like 1	VSNL1	-2.0 -1.9	(1)	
V-set and transmembrane domain containing 2 like	VSTM2L	-1.9	(1)	
Von Willebrand factor A domain containing 1	VWA1	-3.0		
Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	WAS	-2.5		
WAS/WASL interacting protein family, member 1	WIPF1	-4.0		MES
Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, γ polypeptide	YWHAG	-1.9		
Zinc finger and BTB domain containing 7C	ZBTB7C	-6.5	(1)	
Zic family member 2 (odd-paired homolog, <i>Drosophila</i>)	ZIC2	-3.3		
Zinc finger, matrin type 4	ZMAT4	-4.2	(1)	
Zinc finger protein 226	ZNF226	-1.9		
Zinc finger protein 3	ZNF3	-2.1		
Zinc finger protein 331 Zinc finger protein 573	ZNF331 ZNF573	-2.0 -1.8		
Zinc finger protein 975 Zinc finger protein 615	ZNF615	-2.0		
Zinc finger protein 679	ZNF679	-4.4		
Zinc finger protein 679 Zinc finger protein 680	ZNF680	-2.4	(1)	
Zinc finger protein 681	ZNF681	-2.0	(1)	
ABI gene family, member 3 (NESH) binding protein	ABI3BP	2.1	(1)	
Ankyrin repeat and BTB (POZ) domain containing 1	ABTB1	2.5		
Adenosine A2b receptor	ADORA2B	2.0		
Adrenergic, α-1B-, receptor	ADRA1B	2.0	(1)	
Adrenergic, β-2-, receptor, surface	ADRB2	4.0	(1)	
	AGSK1	1.8		
Ajuba LIM protein	AJUBA	2.4		
Adenylate kinase 4	AK4	1.9		
Aldo-keto reductase family 1, member C1	AKR1C1	6.3		
Aldo-keto reductase family 1, member C3 (3-α hydroxysteroid dehydrogenase, type II) Aldo-keto reductase family 1, member C4 (chlordecone reductase; 3-α hydroxysteroid dehydrogenase, type I; dihydrodiol dehydrogenase 4)	AKR1C3 AKR1C4	2.2 3.7		
Activated leukocyte cell adhesion molecule	ALCAM	2.2	(1)	
Angiopoietin-like 2	ANGPTL2	2.0		
Ankyrin repeat domain 1 (cardiac muscle)	ANKRD1	4.0		
Alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150)	ANPEP	2.3		
Annexin A11	ANXA11	2.7		NII
Annexin A3	ANXA3	4.9		NL
Annexin A6 Adenomatosis polyposis coli down-regulated 1-like	ANXA6 APCDD1L	2.0 2.3	(1)	
P GTPase activating protein 30	ARHGAP30	1.8	(1)	
P GDP dissociation inhibitor (GDI) β	ARHGDIB	2.8		
P guanine nucleotide exchange factor (GEF) 10	ARHGEF10	2.5		
Aspartate β-hydroxylase	ASPH	2.2	(1)	
Argininosuccinate synthetase 1	ASS1	2.1		
Arginine vasopressin-induced 1	AVPI1	2.1		
AXL receptor tyrosine kinase	AXL	2.3	(1)	
Brain abundant, membrane attached signal protein 1	BASP1	2.6	(1)	NL
Breast cancer anti-estrogen resistance 3	BCAR3	2.9		
Brain-derived neurotrophic factor	BDNF	2.0		
BEN domain containing 7	BEND7	2.1		
Chromosome 11 open reading frame 75	C11orf75	3.2		
Chromosome 11 open reading frame 87	C11orf87	1.9		
Calmodulin-like 4	CALML4 CASP4	3.0 2.3		MES
Caspase 4, apoptosis-related cysteine peptidase Castor zinc finger 1	CASF4 CASZ1	2.9		MES
Caveolin 1, caveolae protein, 22kDa	CASZ1 CAV1	1.9		
Coiled-coil domain containing 147	CCDC147	2.3		
Coiled-coil domain containing 80	CCDC80	2.8		
Cyclin D1	CCND1	5.7	(1)	
CD163 molecule-like 1	CD163L1	3.0		
CD24 molecule	CD24	4.9		
CDC14 cell division cycle 14 homolog B (Saccharomyces cerevisiae)	CDC14B	2.3		
Cadherin 11, type 2, OB-cadherin (osteoblast)	CDH11	2.0	(1)	
Cadherin 13, H-cadherin (heart)	CDH13	4.4	(1)	
Complement factor B	CFB	2.9		
Chorionic gonadotropin, β polypeptide	CGB	2.2		

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Chorionic gonadotropin, β polypeptide 1	CGB1	4.3		
Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2	CITED2	2.1	(1)	
Cytidine monophospho-N-acetylneuraminic acid hydroxylase, pseudogene	CMAHP	2.2		
CKLF-like MARVEL transmembrane domain containing 7	CMTM7	2.5		
Calponin 1, basic, smooth muscle	CNN1	3.8		
Calponin 2	CNN2	2.0		MES
Collagen, type XI, α 1	COL11A1	1.8	(1)	
Collagen, type XII, α 1	COL12A1	2.9	(1)	MEC
Collagen, type V , α 1	COL 643	2.9	(1)	MES
College, type VI, a 2	COL6A2 COL8A1	1.8 1.9		
Collagen, type VIII, α 1 Coronin, actin binding protein, 2A	CORO2A	2.0		
Carboxypeptidase A4	CPA4	3.9	(1)	
Cadherin-like and PC-esterase domain containing 1	CPED1	2.2	(1)	
Colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog	CSF1R	2.9		
Chondroitin sulfate proteoglycan 4	CSPG4	3.7		
C-terminal binding protein 2	CTBP2	2.6	(1)	
Connective tissue growth factor	CTGF	1.9	(-)	
Cathepsin H	CTSH	2.2		
Chemokine (C-X-C motif) receptor 7	CXCR7	2.8	(1)	
Cytochrome P450, family 26, subfamily B, polypeptide 1	CYP26B1	2.4	(1)	
Cysteine-rich, angiogenic inducer, 61	CYR61	2.7		
Cytohesin 3	CYTH3	1.8		
Disabled homolog 2, mitogen-responsive phosphoprotein (<i>Drosophila</i>)	DAB2	3.4		MES
DEP domain containing 7	DEPDC7	2.7		
Deiodinase, iodothyronine, type II	DIO2	2.0		
DIX domain containing 1	DIXDC1	1.9		
Dickkopf homolog 1 (Xenopus laevis)	DKK1	3.2	(4)	
DnaJ (Hsp40) homolog, subfamily B, member 12	DNAJB12	2.1		
Dedicator of cytokinesis 2	DOCK2	1.9	(1)	
Dermatan sulfate epimerase	DSE	2.2		MES
Deltex 3-like (Drosophila)	DTX3L	1.9		
Dual specificity phosphatase 1	DUSP1	2.0	(1)	
Endothelin converting enzyme 2	ECE2	3.7	(1)	
EH-domain containing 4	EHD4	2.2	(1)	
Elongation factor, RNA polymerase II, 2	ELL2	1.8	(1)	
ELM2 and Myb/SANT-like domain containing 1	ELMSAN1	1.9		
Ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)	ENPP2	2.8		NL
V-ets erythroblastosis virus E26 oncogene homolog 2 (avian)	ETS2	4.2	(1)	
Eyes absent homolog 1 (Drosophila)	EYA1	4.0		
Coagulation factor III (thromboplastin, tissue factor)	F3	2.6		
Family with sequence similarity 133, member A	FAM133A	2.3		
Family with sequence similarity 24, member B	FAM24B	3.3		
Fibrillin 1	FBN1	1.9	(4)	
Fibroblast growth factor 13	FGF13	2.4	(1)	
Filamin A interacting protein 1-like	FILIP1L	1.9		
Hypothetical LOC642477 Filmonography Invasion and American Processin 2	FLJ39632 FLRT2	3.4	(1)	
Fibronectin leucine rich transmembrane protein 2 Folate receptor 1 (adult)	FOLR1	3.0 3.5	(1)	
Forkhead box C1	FOXC1	2.9		
Fraser syndrome 1	FRAS1	3.4		
Follistatin	FST	3.5		
FXYD domain containing ion transport regulator 5	FXYD5	3.3		MES
Frizzled homolog 2 (Drosophila)	FZD2	2.4	(1)	IVILO
Growth arrest-specific 6	GAS6	4.1	(1)	
Guanylate binding protein 2, interferon-inducible	GBP2	2.7		
Growth differentiation factor 15	GDF15	1.8		
Glucose-fructose oxidoreductase domain containing 1	GFOD1	2.8		
GDNF family receptor α 1	GFRA1	2.6		
Gap junction protein, γ 12, 47kDa	GJC2	2.3		
G protein-coupled receptor 1	GPR1	2.4	(1)	
G protein-coupled receptor, family C, group 5, member C	GPRC5C	2.3	(1)	
Glutathione peroxidase 3 (plasma)	GPX3	2.3		
GRAM domain containing 3	GRAMD3	2.7		
Gasdermin A	GSDMA	2.1		
H2A histone family, member Y2	H2AFY2	2.2		
HLA complex P5	HCP5	2.3		
Hairy and enhancer of split 6 (Drosophila)	HES6	2.9		
Homeobox C13	HOXC13	2.3		
Homeobox C6	HOXC6	2.1		
Heat shock 27kDa protein 3	HSPB3	2.1		

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Heat shock 27kDa protein family, member 7 (cardiovascular)	HSPB7	6.0		
Immediate early response 3	IER3	2.2		
Interferon induced transmembrane protein 1	IFITM1	2.5		
Insulin-like growth factor binding protein 4	IGFBP4	3.5		
Interleukin 13 receptor, α 2	IL13RA2	3.4	(1)	
Interleukin 15	IL15	2.1	(1)	
Interleukin 7 receptor Integrin, α 11	IL7R ITGA11	2.4 2.2		
Integrin, β 2 (complement component 3 receptor 3 and 4 subunit)	ITGB2	3.6		MES
Jagged 2	JAG2	2.1		IVILO
Junctophilin 2	JPH2	2.0		
Kazrin, periplakin interacting protein	KAZN	3.0		
Potassium voltage-gated channel, Shab-related subfamily, member 1	KCNB1	3.4		
Potassium voltage-gated channel, delayed-rectifier, subfamily S, member 1	KCNS1	1.8		
KIAA1522	KIAA1522	2.0	(1)	
Kruppel-like factor 11	KLF11	1.9		
Killin, p53-regulated DNA replication inhibitor	KLLN	1.9		
Keratin 18 pseudogene 55	KRT18P55	3.0		
Keratin 80	KRT80	3.8		
Keratin 81	KRT81	6.8		
Lactamase, β	LACTB	2.0		
Layilin	LAYN	1.9		
Leprecan-like 1	LEPREL1	3.2		
LIM and senescent cell antigen-like domains 2	LIMS2	4.6		
Long intergenic nonprotein coding RNA 263	LINC00263	2.8		
	LOC400464	2.0		
	LOC400879	2.0		
Y 1 - 1	LOC645638	3.6) (FC
Lysyl oxidase	LOX	3.0	(1)	MES
Leucine rich repeat neuronal 1	LRRN1	1.9	(1)	
Melanoma antigen family C, 2	MAGEC2	2.4	(1)	MES
Mitogen-activated protein kinase 13	MAPK13 MARCH3	3.2 1.9		MES
Membrane-associated ring finger (C3HC4) 3 Mab-21 domain containing 2	MB21D2	3.4		
Meis homeobox 3 pseudogene 1	MEIS3P1	1.8		
Microfibrillar-associated protein 4	MFAP4	1.8		
Microtubule associated monoxygenase, calponin and LIM domain containing 2	MICAL2	3.0	(1)	
MHC class I polypeptide-related sequence B	MICB	2.1	(1)	
Mohawk homeobox	MKX	3.3		
Melanophilin	MLPH	2.9		
Matrix metallopeptidase 3 (stromelysin 1, progelatinase)	MMP3	3.2		
Mesothelin	MSLN	3.8		
Methionine sulfoxide reductase B3	MSRB3	3.7		
Mucin 1, cell surface associated	MUC1	1.9		
Matrix-remodelling associated 8	MXRA8	2.2		
Myosin, light chain 1, alkali; skeletal, fast	MYL1	3.7	(1)	
Myosin, light chain 9, regulatory	MYL9	3.4		
Myosin IC	MYO1C	2.9		
Nucleosome assembly protein 1-like 3	NAP1L3	1.8		
Neuroblastoma, suppression of tumorigenicity 1	NBL1	3.4		
Neutral cholesterol ester hydrolase 1	NCEH1	2.0		
Neutrophil cytosolic factor 2 (65kDa, chronic granulomatous disease, autosomal 2)	NCF2	3.6		MES
NIMA (never in mitosis gene a)-related kinase 7	NEK7	2.8	(4)	
Nexilin (F actin binding protein)	NEXN NEXN	2.5	(1)	DNI
NK2 homeobox 2 Neuromedin U	NKX2-2 NMU	1.9 2.1		PN
N-terminal Xaa-Pro-Lys <i>N</i> -methyltransferase 1 Netrin G1	NTMT1 NTNG1	1.9 3.8	(1)	
Nuclear casein kinase and cyclin-dependent kinase substrate 1	NUCKS1	4.7	(1)	
Opioid growth factor receptor-like 1	OGFRL1	2.4		
Oxytocin receptor	OXTR	1.9		
Procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), α polypeptide II	P4HA2	2.0		MES
P antigen family, member 5 (prostate associated)	PAGE5	4.2		
Palmdelphin	PALMD	3.1		
3'-phosphoadenosine 5'-phosphosulfate synthase 2	PAPSS2	3.5		
Paired box 6	PAX6	2.0		
Platelet-derived growth factor receptor-like	PDGFRL	2.3		
PDZ and LIM domain 1 (elfin)	PDLIM1	2.9		
Phosphoglucomutase 3	PGM3	1.8		
Pleckstrin homology-like domain, family B, member 2	PHLDB2	2.3		
Paired-like homeodomain 2	PITX2	4.7	(1)	
Turica mic nomeodomani 2				

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Pleiomorphic adenoma gene-like 1	PLAGL1	2.0		
Pleckstrin homology domain containing, family F (with FYVE domain) member 1	PLEKHF1	2.7		
Phospholipid scramblase 4	PLSCR4	2.6		
Plexin A2	PLXNA2	1.9	(1)	
Phosphatidic acid phosphatase type 2 domain containing 1A	PPAPDC1A	2.4		
Prickle homolog 1 (<i>Drosophila</i>)	PRICKLE1	2.9	(1)	
Proline rich 16	PRR16	2.6	(1)	
Protease, serine, 23	PRSS23 PSG4	2.6 5.0		
Pregnancy specific β-1-glycoprotein 4 Protein tyrosine phosphatase, receptor type, F	PTPRF	2.4		
Protein tyrosine phosphatase, receptor type, M	PTPRM	1.9	(1)	
RAB32, member RAS oncogene family	RAB32	2.1	(1)	MES
RAB3B, member RAS oncogene family	RAB3B	5.1		11120
Retinoic acid early transcript 1G	RAET1G	2.2		
Retinoic acid receptor, β	RARB	3.9		
Retinoic acid receptor responder (tazarotene induced) 2	RARRES2	2.6		
Ras association (RalGDS/AF-6) domain family 7	RASSF7	1.8		
RNA binding motif, single stranded interacting protein	RBMS3	1.9	(1)	
RNA binding protein with multiple splicing	RBPMS	2.5	(1)	
Reticulocalbin 3, EF-hand calcium binding domain	RCN3	1.9		
Reversion-inducing-cysteine-rich protein with kazal motifs	RECK	1.9		
RAS-like, estrogen-regulated, growth inhibitor	RERG	4.1		
Raftlin, lipid raft linker 1	RFTN1	3.7		
RGM domain family, member B	RGMB	1.9		
Regulator of G-protein signaling 10	RGS10 RHBDF2	1.8 2.4		
Rhomboid 5 homolog 2 (<i>Drosophila</i>) Ribonuclease, RNase A family, 4	RNASE4	2.4		
Ring finger protein 150	RNF150	2.7		
Arginyl aminopeptidase (aminopeptidase B)	RNPEP	1.8		
Ribosomal protein S6 kinase, 90kDa, polypeptide 2	RPS6KA2	2.2		
Related RAS viral (r-ras) oncogene homolog 2	RRAS2	2.1	(1)	
Sodium channel, nonvoltage-gated 1 α	SCNN1A	3.5	(-)	
Serologically defined colon cancer antigen 3	SDCCAG3	2.3		
Sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B	SEMA3B	3.1		
Serpin peptidase inhibitor, clade B (ovalbumin), member 7	SERPINB7	3.1	(1)	
SERTA domain containing 4	SERTAD4	3.2		
SERTAD4 antisense RNA 1	SERTAD4-AS1	3.3		
Surfactant associated 1, pseudogene	SFTA1P	2.3		
SH2 domain containing 4A	SH2D4A	2.4		
SH3-domain GRB2-like 3	SH3GL3	7.2	(1)	NL
SH3 domain containing ring finger 2	SH3RF2	2.8		
Solute carrier family 1 (neutral amino acid transporter), member 5	SLC1A5	2.0		
Solute carrier family 22 (organic cation transporter), member 4	SLC22A4	1.8		
Solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 24	SLC25A24 SLC25A43	1.9		
Solute carrier family 25, member 43 Solute carrier family 38, member 1		2.0 3.2	(1)	
Solute carrier family 6 (neurotransmitter transporter, glycine), member 9	SLC38A1 SLC6A9	2.0	(1)	CL
Solute carrier family 9 (neurotransmitter transporter, glycine), member 7	SLC9A7	2.3		CL
SMAD family member 6	SMAD6	3.1	(1)	
Smg-6 homolog, nonsense mediated mRNA decay factor (C. elegans)	SMG6	2.1	(1)	
Sosondowah ankyrin repeat domain family member A	SOWAHA	2.5		
SPANX family, member C	SPANXC	2.9	(1)	
SPANX family, member E	SPANXE	4.9		
SPATA31 subfamily C, member 2	SPATA31C2	2.2		
Sphingosine kinase 1	SPHK1	2.0		
Serine palmitoyltransferase, long chain base subunit 3	SPTLC3	3.0		
Serglycin	SRGN	3.1		
Slingshot homolog 3 (Drosophila)	SSH3	2.0		CL
Sjogren syndrome/scleroderma autoantigen 1	SSSCA1	2.2	(1)	
ST6 (α - N -acetyl-neuraminyl-2,3- β -galactosyl-1,3)- N -acetylgalactosaminide α -2,6-sialyltransferase 3	ST6GALNAC3	2.0	(1)	
StAR-related lipid transfer (START) domain containing 5	STARD5	1.9	(1)	
StAR-related lipid transfer (START) domain containing 8	STARD8	2.2	(1)	
Stanniocalcin 2	STC2	3.5	(1)	
Syntaxin binding protein 5 (tomosyn)	STXBP5	2.7	(1)	
Syncoilin, intermediate filament protein	SYNC	2.3		
Transgelin TFA domain family member 4	TAGLN TEAD4	3.3 3.1		
TEA domain family member 4 Teneurin transmembrane protein 4	TENM4	2.2		
Transforming growth factor, β 2	TGFB2	6.5		
Transforming growth factor, β receptor III	TGFBR3	2.6		
	THBS1	2.7	(1)	MES
Thrombospondin 1	1 HB21	2./	(1)	MES

SDC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
TIMP metallopeptidase inhibitor 3 (Sorsby fundus dystrophy, pseudoinflammatory)	TIMP3	4.6		
Transmembrane protein 255B	TMEM255B	2.0		
Transmembrane protein 44	TMEM44	1.9		
Transmembrane protein 45A	TMEM45A	2.3	(1)	
Tumor protein p53 inducible protein 11	TP53I11	2.8	(1)	
Two pore segment channel 2	TPCN2	2.0		
Tropomyosin 1 (α)	TPM1	2.0		
Tropomyosin 2 (β)	TPM2	1.8		
Tetraspanin 10	TSPAN10	2.9		
Tetraspanin 4	TSPAN4	2.1		
Tuftelin 1	TUFT1	2.5		
Thioredoxin interacting protein	TXNIP	1.8		
Urothelial cancer associated 1	UCA1	2.4		
Vascular endothelial growth factor C	VEGFC	1.8		
WAS protein family, member 3	WASF3	1.9		
Wingless-type MMTV integration site family member 2	WNT2	3.3		
Wingless-type MMTV integration site family, member 3	WNT3	3.1		
Wingless-type MMTV integration site family, member 5B	WNT5B	3.6		
WW and C2 domain containing 1	WWC1	2.1		
Yip1 interacting factor homolog A (Saccharomyces cerevisiae)	YIF1A	1.9		
Zinc finger, CCHC domain containing 5	ZCCHC5	3.1	(1)	
Zinc finger protein 618	ZNF618	2.2		
Zinc finger protein 730	ZNF730	1.8		

CSC

Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
AP2 associated kinase 1	AAK1	-2.5	(1)	
ATP-binding cassette, subfamily A (ABC1), member 2	ABCA2	-2.5		
Acetylcholinesterase (Yt blood group)	ACHE	-2.2		
Acid phosphatase 6, lysophosphatidic	ACP6	-2.7		
Acyl-CoA synthetase short-chain family member 1	ACSS1	-2.0		
Activin A receptor, type IIA	ACVR2A	-3.5	(1)	
Adenosine deaminase	ADA	-2.1		
ADAM metallopeptidase domain 15	ADAM15	-2.1		
ADAM metallopeptidase with thrombospondin type 1 motif, 6	ADAMTS6	-2.2	(1)	
ADAM metallopeptidase with thrombospondin type 1 motif, 8	ADAMTS8	-3.9	(1)	
Angiotensinogen (serpin peptidase inhibitor, clade A, member 8)	AGT	-2.1	. ,	
Allograft inflammatory factor 1-like	AIF1L	-2.3		
Adenylate kinase 4	AK4	-2.5		
A kinase (PRKA) anchor protein 10	AKAP10	-1.8		
V-akt murine thymoma viral oncogene homolog 2	AKT2	-1.8		CL
Aldehyde dehydrogenase 4 family, member A1	ALDH4A1	-2.1		OL.
Aldehyde dehydrogenase 6 family, member A1	ALDH6A1	-2.7		
Antagonist of mitotic exit network 1 homolog (Saccharomyces cerevisiae)	AMN1	-2.1		
Angiopoietin 2	ANGPT2	-4.3		
P guanine nucleotide exchange factor (GEF) 39	ARHGEF39	-1.8		
ADP-ribosylation factor-like 4A	ARL4A	-2.9	(1)	
Arylsulfatase G	ARSG	-1.8	(1)	
Ankyrin repeat and SOCS box-containing 9	ASB9	-3.6	(1)	
Asparaginase like 1	ASRGL1	-2.1	(1)	
Astrotactin 1	ASTN1	-2.0		
Autophagy related 13	ATG13	-3.6		
ATPase type 13A4	ATP13A4	-3.6		
ATPase type 13/4 ATPase, Na+/K+ transporting, β 2 polypeptide	ATP18A4 ATP1B2	-2.3		
	ATP162 ATP9B	-2.5 -2.1	(1)	
ATPase, class II, type 9B	ATP9B ATPBD4		(1)	
ATP binding domain 4		-1.9	(1)	
UDP-GlcNAc:βGal β-1,3-N-acetylglucosaminyltransferase 2	B3GNT2	-2.0	(1)	
Bardet-Biedl syndrome 9	BBS9 BCAP29	-3.1		
B cell receptor-associated protein 29		-6.0		
B-box and SPRY domain containing	BSPRY BTC	-2.1	(1)	
Bcellulin		-2.2	(1)	
Chromosome 10 open reading frame 12	C10orf12	-2.4		
Chromosome 11 open reading frame 63	C11orf63	-2.4		
Chromosome 19 open reading frame 6	C19orf6	-1.8		
Chromosome 2 open reading frame 80	C2orf80	-5.8		
Chromosome 6 open reading frame 15	C6orf15	-2.0		
Chromosome 6 open reading frame 170	C6orf170	-1.9		
Carbonic anhydrase II	CA2	-4.3		
Carbonic anhydrase IV	CA4	-2.4	(1)	NL

FRAS1 related extracellular matrix protein 2

GRB2-associated binding protein 2 $UDP\text{-}N\text{-}acetyl\text{-}\alpha\text{-}D\text{-}galactosamine}; polypeptide \textit{N-}acetylgalactosaminyltransferase 13 (GalNAc-T13)}$

FRMD6 antisense RNA 1

GRB2-associated binding protein 1

FREM2

GAB1

GAB2

GALNT13

FRMD6-AS1

-3.8

-2.1

-4.6

-1.9

-2.9

TABLE W1. (continued)					
CSC					
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signatur	
Calcium channel, voltage-dependent, P/Q type, α 1A subunit	CACNA1A	-2.8	(1)		
Calcium channel, voltage-dependent, β 1 subunit	CACNB1	-2.6			
Cell adhesion molecule 1	CADM1	-4.5			
Ca2+-dependent activator protein for secretion 2	CADPS2	-2.0	(1)		
Calpain 9	CAPS	-2.8	(1)		
Calcyphosine	CAPS CCDC103	-2.3 -4.3			
Coiled-coil domain containing 103 Coiled-coil domain containing 94	CCDC94	-4.5 -2.2			
CD14 molecule	CD14	-2.2		MES	
CD276 molecule	CD276	-2.4		WILD	
CD79b molecule, immunoglobulin-associated β	CD79B	-2.1			
Cadherin 6, type 2, K-cadherin (fetal kidney)	CDH6	-3.1	(1)	CL	
Carcinoembryonic antigen-related cell adhesion molecule 21	CEACAM21	-2.2			
CCAAT/enhancer binding protein (C/EBP), δ	CEBPD	-1.8	(1)		
CUGBP, Elav-like family member 1	CELF1	-2.4			
Centrosomal protein 152kDa	CEP152	-1.8			
Centrosomal protein 170kDa	CEP170	-2.7	(1)		
Centrosomal protein 70kDa	CEP70	-2.1			
Complement factor I	CFI	-2.2			
Chorionic gonadotropin, β polypeptide	CGB	-2.1			
Chromogranin B (secretogranin 1)	CHGB	-2.3	(1)		
C-type lectin domain family 4, member A	CLEC4A	-1.9			
C-type lectin domain family 9, member A	CLEC9A	-2.0			
CCR4-NOT transcription complex, subunit 3	CNOT3	-2.1			
Cannabinoid receptor 1 (brain)	CNR1	-2.1	(1)		
Cordon-bleu WH2 repeat protein	COBL	-3.5	(1)		
Component of oligomeric golgi complex 5	COG5 COL11A2	-2.3 -2.0	(1)		
Collagen, type XI, α 2 Collagen, type V, α 1	COLTAZ COL5A1	-2.0 -1.8	(1) (1)	MES	
Carnitine palmitoyltransferase 1A (liver)	CPT1A	-1.9	(1)	WILS	
Crystallin, μ	CRYM	-4.5		NL	
Cold shock domain containing C2, RNA binding	CSDC2	-3.0	(1)	112	
CTAGE family, member 11, pseudogene	CTAGE11P	-2.2	(1)		
CTS telomere maintenance complex component 1	CTC1	-2.4			
Chemokine (C-X-C motif) ligand 3	CXCL3	-1.8			
Dachsous 1 (Drosophila)	DCHS1	-2.1			
Doublecortin-like kinase 2	DCLK2	-2.6			
Decorin	DCN	-3.0	(1)		
DEAD (Asp-Glu-Ala-Asp) box polypeptide 51	DDX51	-1.9			
DENN/MADD domain containing 1A	DENND1A	-2.5	(1)		
Der1-like domain family, member 3	DERL3	-8.2			
Dehydrogenase/reductase (SDR family) member 12	DHRS12	-3.7			
DEXH (Asp-Glu-X-His) box polypeptide 58	DHX58	-2.3			
Diaphanous homolog 3 (Drosophila)	DIAPH3	-2.1	(1)		
DIS3 mitotic control homolog (Saccharomyces cerevisiae)	DIS3	-1.8			
Dickkopf homolog 1 (Xenopus laevis)	DKK1	-4.4	(4)		
Dynein, axonemal, assembly factor 3	DNAAF3	-5.2			
DnaJ (Hsp40) homolog, subfamily C, member 12	DNAJC12	-2.5			
Estrogen receptor binding site associated, antigen, 9	EBAG9	-2.0			
Extracellular matrix protein 2, female organ and adipocyte specific	ECM2 EFEMP1	-3.0 -1.9	(1)		
EGF-containing fibulin-like extracellular matrix protein 1 Engulfment and cell motility 1	ELMO1	-2.7	(1) (1)		
Exoribonuclease 1	ERI1	-2.1	(1)		
ERI1 exoribonuclease family member 2	ERI2	-2.1			
Ets variant gene 3	ETV3	-2.9	(1)		
Fatty acid binding protein 7, brain	FABP7	-2.0	\-/		
Family with sequence similarity 153, member B	FAM153B	-3.7			
Family with sequence similarity 5, member B	FAM5B	-2.6	(1)		
Fanconi anemia, complementation group A	FANCA	-2.7			
Fatty acyl CoA reductase 2	FAR2	-2.0			
Fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)	FGFR1	-2.9			
Filamin A interacting protein 1-like	FILIP1L	-2.1			
Hypothetical protein FLJ22184	FLJ22184	-2.8			
Fibronectin leucine rich transmembrane protein 3	FLRT3	-1.9			
Fibromodulin	FMOD	-2.0			
Forkhead box A2	FOXA2	-2.0	(1)		
Forkhead box R1	FOXR1	-8.3			
CDAC1 I. I. III	EDEM (2	2.0			

(1) (1)

(1)

TABI	.E. W1	. (continue	ď

TABLE W1. (continued)				
CSC Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
UDP-N-acetyl-α-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	GALNT3	-2.8		
Growth associated protein 43	GAP43	-4.2	(1)	
Gap junction protein, β 6, 30kDa	GJB6	-3.0		
Guanine nucleotide binding protein (G protein), γ 7	GNG7	-2.5		CL
G patch domain containing 2	GPATCH2	-2.2		
Glycoprotein M6B	GPM6B	-3.2		
Glutathione peroxidase 3 (plasma)	<i>GPX3</i> GRIK1	-2.0 -1.8		CL
Glutamate receptor, ionotropic, kainate 1 G protein-coupled receptor kinase 5	GRK5	-1.8 -1.8	(1)	CL
Germ cell associated 1	GSG1	-2.0	(1)	
H1 histone family, member N, testis-specific	H1FNT	-2.8		
HemK methyltransferase family member 1	HEMK1	-2.6		
Hairy/enhancer-of-split related with YRPW motif 2	HEY2	-2.9	(1)	
Hypoxia inducible factor 3, α subunit	HIF3A	-2.2		
Major histocompatibility complex, class II, DM β	HLA-DMB	-5.2		
Heat shock 27kDa protein 3	HSPB3	-2.1		
HUS1 checkpoint homolog b (S. pombe)	HUS1B	-3.4		
Huntingtin interacting protein K	HYPK ICA1	-1.9 -3.0		
Islet cell autoantigen 1, 69kDa Interferon, α-inducible protein 27	IFI27	-3.0 -2.3		
Insulin-like growth factor binding protein 7	IGFBP7	-3.1	(1)	
IGF-like family member 3	IGFL3	-4.8	(1)	
Immunoglobulin superfamily, member 11	IGSF11	-3.2	(1)	
Interleukin 8	IL8	-2.4	(1)	
InaD-like (Drosophila)	INADL	-2.9		
Insulinoma-associated 1	INSM1	-2.2		
Inositol hexakisphosphate kinase 3	IP6K3	-1.8		
Interferon regulatory factor 4	IRF4	-1.9		
Integrin, \alpha 6	ITGA6	-2.6		
Kelch repeat and BTB (POZ) domain containing 3	KBTBD3	-2.0		
Potassium channel tetramerisation domain containing 21	KCTD21	-5.2	(1)	
KIAA0319-like Kruppel-like factor 4 (gut)	KIAA0319L KLF4	-3.4 -2.3	(1)	
Kruppel-like factor 7 (ubiquitous)	KLF4 KLF7	-2.5 -1.8		
Kelch domain containing 8A	KLHDC8A	-2.1		CL
Kelch-like 23 (Drosophila)	KLHL23	-2.0		O.L
Kelch-like 5 (Drosophila)	KLHL5	-3.8		
Killer cell lectin-like receptor subfamily F, member 1	KLRF1	-4.5		
Like-glycosyltransferase	LARGE	-5.1	(1)	
Lysocardiolipin acyltransferase 1	LCLAT1	-1.8		
Leptin (obesity homolog, mouse)	LEP	-1.9	(1)	
Leprecan-like 1	LEPREL1	-3.0		
Leucine-rich repeat LGI family, member 2	LGI2	-3.5		
Long intergenic nonprotein coding RNA 261 Long intergenic nonprotein coding RNA 282	LINC00261 LINC00282	-3.1 -2.1		
Long intergenic nonprotein coding RNA 324	LINC00282 LINC00324	-1.9		
Long intergenic nonprotein coding RNA 478	LINC00478	-2.2		
Long intergenic nonprotein coding RNA 612	LINC00612	-1.8		
Long intergenic nonprotein coding RNA 623	LINC00623	-2.9		
	LOC100129502	-1.9		
	LOC100130276	-2.8		
	LOC100131096	-2.3		
	LOC100132354	-2.2		
	LOC255167	-2.1		
	LOC283710	-2.9		
	LOC285696	-3.7		
	LOC389765	-2.5		
	LOC441155 LOC645355	-2.9 -2.1		
	LOC653562	-7.3		
	LOC728755	-2.2		
	LOC730441	-3.4		
Lysyl oxidase-like 3	LOXL3	-2.3		
Plasticity related gene 1	LPPR4	-2.7		
Leucine rich repeat containing 37, member A3	LRRC37A3	-3.1		
Leucine rich repeat neuronal 1	LRRN1	-3.9	(1)	
Leucine rich repeat neuronal 3	LRRN3	-3.0		
Lumican	LUM	-2.2		
Lymphocyte antigen 6 complex, locus G5C	LY6G5C	-2.2		
Lymphocyte antigen 6 complex, locus H Malanoma antigen family A 11	LY6H MACEALL	-3.8		
Melanoma antigen family A, 11 Mal, T cell differentiation protein	MAGEA11 Mal	-4.2 -2.1		
MANSC domain containing 1	MANSC1	-2.1		
THE TOO COMMENT CONCERNING 1	MUNIOCI	-2.0		

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TABLE W1. (continued)				
CSC			0 - 100	
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Microtubule-associated protein 2	MAP2	-2.9	(1)	PN
Mitogen-activated protein kinase kinase 6	MAP2K6	-1.8		
Mitogen-activated protein kinase kinase 7	MAP2K7	-2.1	(1)	
Microtubule-associated protein 6	MAP6	-2.6	(1)	DNI
Microtubule-associated protein τ	MAPCUA	-2.7	(1)	PN
Membrane-associated ring finger (C3HC4) 2 Microtubule associated serine/threonine kinase family member 4	MARCH2 MAST4	-1.8 -3.4		
MEF2B neighbor	MEF2BNB	-2.7		
Malignant fibrous histiocytoma amplified sequence 1	MFHAS1	-2.8	(1)	
Myocardial infarction associated transcript (nonprotein coding)	MIAT	-3.1	(-)	
MLLT4 antisense RNA 1 (head to head)	MLLT4-AS1	-1.9		
Metallophosphoesterase 1	MPPE1	-2.0		
Mucin 12, cell surface associated	MUC12	-1.9		
Myeloid-associated differentiation marker	MYADM	-2.1		
Myosin binding protein C, slow type	MYBPC1	-2.3		NL
V-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)	MYCN	-2.1	(1)	
Myosin IE	MYO1E	-1.8	(1)	CI
Myosin VC	MYO5C	-2.2	(1)	CL
N-myc downstream regulated gene 1 Nicotinamide nucleotide transhydrogenase	NDRG1 NNT	-1.8 -3.3	(1)	
NADPH oxidase activator 1	NOXA1	-3.3 -2.3	(1)	
Nuclear pore complex interacting protein family, member B15	NPIPL2	-2.6	(1)	
Neuropeptide Y receptor Y5	NPY5R	-2.5		
Neuronal cell adhesion molecule	NRCAM	-2.2		
Nuclear receptor binding SET domain protein 1	NSD1	-2.2		
Neurotensin	NTS	-7.2		
2'-5'-oligoadenylate synthetase 2, 69/71kDa	OAS2	-1.8		
Protein kinase C and casein kinase substrate in neurons 2	PACSIN2	-2.0		
Peptidyl arginine deiminase, type II	PADI2	-5.2		
Protocadherin γ subfamily A, 5	PCDHGA5	-2.2		
Protocadherin γ subfamily B, 6	PCDHGB6	-2.4		
Pecanex-like 4 (Drosophila)	PCNXL4	-5.6	(4)	
Phosphodiesterase 8B	PDE8B	-6.0	(1)	
Platelet-derived growth factor β polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	PDGFB	-1.9	(1)	
PDZ and LIM domain 5 Podoplanin	PDLIM5 PDPN	-2.3 -1.8	(1)	MES
PDZ domain containing 2	PDZD2	-1.9		IVILO
PDZ domain containing RING finger 3	PDZRN3	-3.0		
PHD finger protein 8	PHF8	-3.4		
Phosphatidylinositol-4-phosphate 5-kinase-like 1	PIP5KL1	-2.1		
Phospholipase A2, group IID	PLA2G2D	-2.3		
Polo-like kinase 2 (<i>Drosophila</i>)	PLK2	-1.8		
Phospholipid transfer protein	PLTP	-3.2		
Postmeiotic segregation increased 2 pseudogene 1	PMS2P1	-2.0		
POM121 transmembrane nucleoporin-like 8 pseudogene	POM121L8P	-4.3		
Protein phosphatase 2 (formerly 2A), regulatory subunit B", α	PPP2R3A	-2.2		
Proline rich 13	PRR13	-2.1		
Protein tyrosine phosphatase type IVA, member 2	PTP4A2	-2.5		
Poliovirus receptor-related 2 (herpesvirus entry mediator B)	PVRL2	-4.4		
Pyrroline-5-carboxylate reductase 1 Pyrin and HIN domain family, member 1	PYCR1 PYHIN1	-4.2 -3.8		
RAB24, member RAS oncogene family	RAB24	- 3.8 - 1.9		
RAB3A, member RAS oncogene family	RAB3A	-1.9		
RAB3A interacting protein (rabin3)	RAB3IP	-3.5	(1)	
Retinoic acid receptor responder (tazarotene induced) 1	RARRES1	-2.6	(1)	
RNA binding motif (RNP1, RRM) protein 3	RBM3	-3.2		
Retinol binding protein 7, cellular	RBP7	-2.6		
Regulator of calcineurin 2	RCAN2	-3.0		
Replication factor C (activator 1) 5, 36.5kDa	RFC5	-2.8		
Raftlin family member 2	RFTN2	-2.4		
Regulator of cell cycle	RGCC	-2.3		
Rhomboid, veinlet-like 2 (<i>Drosophila</i>)	RHBDL2	-2.3		
P family GTPase 2	RND2	-2.3		
Ring finger protein 14	RNF14	-2.0		
Ring finger protein 148	RNF148	-1.8		
RNA, Ro-associated Y4	RNY4	-2.3		
Ribosomal protein L37a	RPL37A	-2.8		
Ribosomal protein S6 kinase, 90kDa, polypeptide 2 Radical S adenosal methioning domain compaining 2	<i>RPS6KA2</i> RSAD2	-2.4 -3.4		
Radical S-adenosyl methionine domain containing 2 Runt-related transcription factor 3	RSAD2 RUNX3	- 3.4 - 1.9		
Scavenger receptor class A, member 3	SCARA3	-1.9 -1.8		
Secretogranin III	SCG3	-7.1		PN
	5545	,		**1

CSC TABLE W1. (continued)				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Secretagogin, EF-hand calcium binding protein	SCGN	-2.1		
Sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F	SEMA4F	-2.6		
Serine incorporator 2	SERINC2	-3.4		
Surfactant associated 1, pseudogene	SFTA1P	-2.8		
SH2B adaptor protein 2	SH2B2	-2.7	(1)	
SH3-domain kinase binding protein 1	SH3KBP1	-2.2		
SH3 domain containing ring finger 2	SH3RF2	-3.1		
Signal-induced proliferation-associated gene 1	SIPA1	-2.8	(1)	
Signal-regulatory protein α	SIRPA	-4.8	(1)	
Solute carrier family 16, member 4 (monocarboxylic acid transporter 5)	SLC16A4	-1.8		
Solute carrier family 2 (facilitated glucose transporter), member 12	SLC2A12	-2.6	(1)	
Solute carrier family 30 (zinc transporter), member 1 Solute carrier family 35, member F1	SLC30A1 SLC35F1	-3.6 -4.9	(1)	
Solute carrier family 38, member 8	SLC38A8	-3.0		
Solute carrier family 39 (zinc transporter), member 4	SLC39A4	-2.6		
Solute carrier family 5 (iodide transporter), member 8	SLC5A8	-4.4		
Solute carrier family 6 (proline IMINO transporter), member 20	SLC6A20	-2.2		
Solute carrier family 6 (neurotransmitter transporter, glycine), member 9	SLC6A9	-5.3		CL
Solute carrier organic anion transporter family, member 1C1	SLCO1C1	-2.4		
Secretory leukocyte peptidase inhibitor	SLPI	-3.7		
Small integral membrane protein 1	SMIM1	-2.2		
Small nucleolar RNA, H/ACA box 28	SNORA28	-1.8		
Small nucleolar RNA, C/D box 114-3	SNORD114-3	-1.8		
Small nucleolar RNA, C/D box 14C	SNORD14C	-2.1		
Small nucleolar RNA, C/D box 6	SNORD6	-5.9		
Small nucleolar RNA, C/D box 95	SNORD95	-2.0		
Sorbin and SH3 domain containing 1	SORBS1	-1.8	(1)	DAY.
SRY (sex-determining region Y)-box 2	SOX2	-1.6	(1)(2)	PN
SRY (sex-determining region Y)-box 21	SOX21 SOX2-OT	-4.3	(1)(2)	
SOX2 overlapping transcript (nonprotein coding) SRY (sex-determining region Y)-box 8	SOX2-01	-2.0 -3.7		
Sperm associated antigen 1	SPAG1	-2.0		
SPARC-like 1 (mast9, hevin)	SPARCL1	-2.2		
Spermatogenesis and centriole associated 1	SPATC1	-2.2		
Speedy/RINGO cell cycle regulator family member E3	SPDYE3	-2.7		
Sprouty homolog 4 (Drosophila)	SPRY4	-2.0	(1)	
SLIT-ROBO P GTPase activating protein 1	SRGAP1	-2.1	(1)	
ST8SIA6 antisense RNA 1	ST8SIA6-AS1	-1.9		
STAU2 antisense RNA 1	STAU2-AS1	-3.2		
Synapse differentiation inducing 1	SYNDIG1	-2.2		
TBC1 domain family, member 17	TBC1D17	-2.5		
Transducin (β)-like 3	TBL3	-1.8		
Tyrosyl-DNA phosphodiesterase 1	TDP1	-1.9		
Trefoil factor 3 (intestinal)	TFF3	-3.0		
Transmembrane protein 100	TMEM100	-1.9	(4)	
Transmembrane protein 106A	TMEM106A	-1.9	(1)	
Transmembrane protein 156	TMEM156	-3.1		
Transmembrane protein 161B Transmembrane protein 164	TMEM161B TMEM164	-2.2 -3.0		
Transmembrane protein 178A	TMEM104 TMEM178A	-1.9		
Tumor protein p63	TP63	-5.4		
Tumor protein D52	TPD52	-3.8	(1)	
Tripartite motif-containing 24	TRIM24	-1.8	(1)	
tRNA methyltransferase 44 homolog (Saccharomyces cerevisiae)	TRMT44	-2.8		
Transient receptor potential cation channel, subfamily M, member 3	TRPM3	-2.8	(1)	
Tetraspanin 14	TSPAN14	-2.1		
Testis specific protein, Y-linked 1	TSPY1	-2.4		
Tweety homolog 1 (Drosophila)	TTYH1	-3.1		
Taxilin γ 2, pseudogene	TXLNG2P	-2.0		
UDP glucuronosyltransferase 2 family, polypeptide B11	UGT2B11	-4.6		
Urotensin 2	UTS2	-2.3		
Vaccinia related kinase 3	VRK3	-4.9		
WD repeat domain 49	WDR49	-2.0	(1)	
WD repeat domain 74	WDR74	-3.0		
WD repeat domain 86	WDR86	-2.4		
WD repeat domain 89	WDR89	-3.3		
X antigen family, member 1D Zing finger and BTB domain containing 40	XAGE1D ZRTR40	-6.3		
Zinc finger and BTB domain containing 40 Zinc finger with KRAB and SCAN domains 5	ZBTB40 ZKSCAN5	-7.8 -3.2		
Zinc finger with KKAB and SCAN domains 3 Zinc finger protein 182	ZNF182	-5.2 -5.5		
Zinc finger protein 102 Zinc finger protein 519	ZNF519	-2.3		
Zinc finger protein 575 Zinc finger protein 555	ZNF555	-1.9		
0 1		/		

TABLE W1. (continued)					
CSC Description	Symbol	Fold Change	Sox2 Target	Subclass	
	.,	(Log ₂)	(Ref)	Signature	
Zinc finger protein 608	ZNF608	-2.0	(1)		
Zinc finger protein 678	ZNF678	-2.3			
Zinc finger protein 717	ZNF717	-2.3			
Zinc finger protein 761	ZNF761	-1.8			
Zinc finger protein 880	ZNF880	-2.4			
Zinc finger protein 890, pseudogene	ZNF890P	-2.0			
Zinc finger and SCAN domain containing 12	ZSCAN12	-1.8			
Zinc finger and SCAN domain containing 16	ZSCAN16	-2.4			
Zinc finger, X-linked, duplicated B	ZXDB	-1.8			
Aminoadipate aminotransferase	AADAT	5.0	(1)		
Acyl-Coenzyme A dehydrogenase family, member 8	ACAD8	1.8			
Actin, α 2, smooth muscle, aorta	ACTA2	2.2	(4)		
ADAM metallopeptidase domain 28	ADAM28	2.3	(1)		
ADAM metallopeptidase with thrombospondin type 1 motif, 18	ADAMTS18	1.8			
Adenylate cyclase 9	ADCY9	3.3			
Alcohol dehydrogenase 6 (class V)	ADH6	2.3	(1)		
AF4/FMR2 family, member 3	AFF3	2.7	(1)		
1-acylglycerol-3-phosphate <i>O</i> -acyltransferase 3	AGPAT3	2.5			
Aldo-keto reductase family 1 member 15 (ALD2)	AKR1B15	1.9			
Aminolevulinate, δ-, synthase 2 (sideroblastic/hypochromic anemia)	ALAS2	2.9			
Angiopoietin-like 3	ANGPTL3	2.3			
Angiopoietin-like 4	ANGPTL4	1.9			
Ankyrin 1, erythrocytic	ANK1	2.4			
Anoctamin 2	ANO2	2.1			
Annexin A8-like 2	ANXA8L2	2.5			
Adaptor-related protein complex 1 associated regulatory protein	APPAR	3.3			
Amyloid β (A4) precursor protein-binding, family B, member 2 (Fe65-like)	APCDD11	8.0	(1)		
Adenomatosis polyposis coli down-regulated 1-like	APCDD1L APOE	1.9	(1)		
Apolipoprotein E		1.9			
P GDP dissociation inhibitor (GDI) β	ARHGDIB ARL9	2.2			
ADP-ribosylation factor-like 9		1.8			
Armadillo repeat containing 8	ARMC8	1.9			
Achaete-scute complex homolog 2 (<i>Drosophila</i>)	ASCL2 ASPRV1	2.8 2.1			
Aspartic peptidase, retroviral-like 1 Argininosuccinate synthetase 1	ASS1	2.7			
ATPase, class V, type 10B	ATP10B	2.4			
Attractin	ATT 10B	2.1			
BAI1-associated protein 2-like 1	BAIAP2L1	2.1	(1)		
Biglycan	BGN	2.5	(1)		
Bine-homocysteine methyltransferase 2	ВНМТ2	1.9			
BR serine/threonine kinase 2	BRSK2	2.1			
Bone marrow stromal cell antigen 1	BST1	2.0			
Biotinidase	BTD	2.4			
Blood vessel epicardial substance	BVES	2.1			
Chromosome 11 open reading frame 31	C11orf31	4.5			
Chromosome 17 open reading frame 99	C17orf99	3.2			
Chromosome 1 open reading frame 115	C1orf115	2.1			
Chromosome 1 open reading frame 141	Clorf141	3.4			
Chromosome 1 open reading frame 170	C1orf170	1.9			
Chromosome 20 open reading frame 201	C20orf201	2.3			
Chromosome 21 open reading frame 49	C21orf49	2.0			
Chromosome 21 open reading frame 67	C21orf67	2.2			
Chromosome 2 open reading frame 40	C2orf40	5.2			
Chromosome 2 open reading frame 81	C2orf81	2.7			
Chromosome 3 open reading frame 52	C3orf52	2.2			
Chromosome 8 open reading frame 46	C8orf46	4.7			
Chromosome 9 open reading frame 106	C9orf106	2.0			
Calcium binding protein 7	CABP7	3.9			
Calcium channel, voltage-dependent, L type, α 1C subunit	CACNA1C	2.3	(1)		
Cancer antigen 1	CAGE1	3.5			
Caspase 6, apoptosis-related cysteine peptidase	CASP6	2.8			
Coiled-coil domain containing 163, pseudogene	CCDC163P	2.3			
Coiled-coil domain containing 42B	CCDC42B	1.8			
Coiled-coil domain containing 57	CCDC57	3.2	(1)		
Chemokine (C-C motif) receptor 2	CCR2	2.1			
CD163 molecule-like 1	CD163L1	2.2			
CD302 molecule	CD302	2.4			
CDC14 cell division cycle 14 homolog A (Saccharomyces cerevisiae)	CDC14A	5.0			
CDC42 effector protein (P GTPase binding) 5	CDC42EP5	2.1			
Cadherin 15, M-cadherin (myotubule)	CDH15	6.0	(1)		
CDP-diacylglycerol-inositol 3-phosphatidyltransferase (phosphatidylinositol synthase)	CDIPT	1.9			
Cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)	CDKN2D	2.2			

CSC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Cdon homolog (mouse)	CDON	1.9		
Chromodomain protein, Y-linked, 1	CDY1	2.0		
Carcinoembryonic antigen-related cell adhesion molecule 2, pseudogene	CEACAM22P CEP112	3.5		
Centrosomal protein 112kDa (CCDC46) Centrosomal protein 85kDa-like	CEP112 CEP85L	2.2 6.7		
Cripto, FRL-1, cryptic family 1	CFC1	3.6		
Chorionic gonadotropin, β polypeptide 1	CGB1	1.9		
ChaC, cation transport regulator homolog 2 (E. coli)	CHAC2	2.7		
Carbohydrate (N-acetylgalactosamine 4-sulfate 6-O) sulfotransferase 15	CHST15	2.9		
C-type lectin domain family 4, member C	CLEC4C	4.7		
Clavesin 2	CLVS2 CMTM4	4.8 2.7		
CKLF-like MARVEL transmembrane domain containing 4 Cannabinoid receptor interacting protein 1	CNRIP1	3.0		
Collagen, type I, α 1	COL1A1	2.6		MES
Collagen, type I, α 2	COL1A2	2.9	(1)	MES
Collagen, type IV, α 6	COL4A6	2.5	(1)	
Collagen, type V, α 3	COL5A3	3.9		
Collagen, type VI, \alpha 2	COL6A2	2.8		
Collagen, type IX, \alpha 2	COL9A2	1.8		
COMM domain containing 8	COMMD8 CORO1A	2.1 2.0		
Coronin, actin binding protein, 1A Carboxypeptidase B1 (tissue)	CPB1	3.3		
Cytokine receptor-like factor 2	CRLF2	2.8		
Crystallin, y S	CRYGS	2.9		
Chondrosarcoma associated gene 1	CSAG1	2.9		
Connective tissue growth factor	CTGF	2.5		
Cathepsin H	CTSH	5.2		
Cytokine-like 1	CYTL1	2.5		
D-2-hydroxyglutarate dehydrogenase	D2HGDH	2.6		
DAB2 interacting protein	DAB2IP	3.0		
Death associated protein-like 1 Deleted in azoospermia 4	DAPL1 DAZ4	2.2 3.1		
DCN1, defective in cullin neddylation 1, domain containing 2 (Saccharomyces cerevisiae)	DCUN1D2	2.5		
Defensin, β 106A	DEFB106A	5.9		
Defensin, β 128	DEFB128	2.7		
DEP domain containing 7	DEPDC7	2.1		
DiGeorge syndrome critical region gene 5 (noncoding)	DGCR5	4.0		
Disrupted in renal carcinoma 3	DIRC3	2.0		
DKFZP434L187 protein	DKFZP434L187 DLK2	1.9 2.9		
Δ-like 2 homolog (<i>Drosophila</i>) Dmx-like 1	DMXL1	2.0		
DNA (cytosine-5-)-methyltransferase 3 β	DNMT3B	2.5		
Deoxynucleotidyltransferase, terminal, interacting protein 2	DNTTIP2	3.2		
Dipeptidyl-peptidase 8	DPP8	5.4		
Dermatopontin	DPT	2.0		
Dpy-19-like 2 pseudogene 2 (C. elegans)	DPY19L2P2	3.1		
Dorsal inhibitory axon guidance protein	DRAXIN	2.1		
Dopamine receptor D2	DRD2	3.5		
Dystonin E2F transcription factor 8	DST E2F8	1.9 1.9		
Epstein-Barr virus induced gene 3	EBI3	2.1		
Ephrin-B1	EFNB1	2.4		
ELK4, ETS-domain protein (SRF accessory protein 1)	ELK4	2.6		
EP400 N-terminal like	EP400NL	2.2		
Envoplakin	EVPL	2.3		
Coagulation factor VIII, procoagulant component (hemophilia A)	F8	6.1		
Fatty acid binding protein 4, adipocyte	FABP4	2.0		
Family with sequence similarity 110, member D Family with sequence similarity 159, member B	FAM110D Fam159B	2.5 2.7		
Family with sequence similarity 174, member A	FAM174A	2.7		
Family with sequence similarity 179, member A	FAM179A	2.5		
Family with sequence similarity 196, member A	FAM196A	1.8		
Family with sequence similarity 27-like	FAM27L	2.1		
Family with sequence similarity 65, member C	FAM65C	4.1		
Family with sequence similarity 71, member C	FAM71C	5.8		
Family with sequence similarity 72, member A	FAM72A	1.8	(1)	
Fibulin 1 FRYO22 antisense RNA 1	FBLN1 FRYO22-AS1	2.6	(1)	
FBXO22 antisense RNA 1 Fc receptor-like 3	FBXO22-AS1 FCRL3	3.2 2.0		
	FLJ31945	6.2		
	FLJ37786	2.6		
	FLJ3//00	2.0		
	FLJ45248	1.8		

TABLE	W1.	(continued)
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CSC TABLE W1. (contin	uea)			
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Fibronectin type III domain containing 5	FNDC5	2.6		
Folate receptor 1 (adult)	FOLR1	2.0		
Forkhead box K2	FOXK2	5.2	(1)	
Forkhead box protein O6	FOXO6	4.8		
Forkhead box S1	FOXS1	3.2		
Fraser syndrome 1	FRAS1	2.6		
FSHD region gene 2 family, member C	FRG2C	2.6		
Frizzled-related protein	FRZB	2.4		
FtsJ methyltransferase domain containing 2	FTSJD2	2.5		
FXYD domain containing ion transport regulator 5	FXYD5	1.9		
Glucose-6-phosphatase, catalytic subunit	G6PC	2.7		
Γ-aminobutyric acid (GABA) receptor, ρ 1	GABRR1	5.2		
G antigen 1	GAGE1	6.9		
GRINL1A combined protein Glutamine-fructose-6-phosphate transaminase 2	GCOM1 GFPT2	3.2 2.6		
Glycoprotein, α-galactosyltransferase 1 (GGTA1), noncoding RNA.	GGTA1P	2.6		
GTPase, IMAP family member 4	GIMAP4	2.1		
Gap junction protein, β 2, 26kDa	GJB2	2.0		
GLI pathogenesis-related 1 (glioma)	GLIPR1	3.0		
Guanine nucleotide binding protein (G protein), α 14	GNA14	1.8		
Golgin A6 family, member C	GOLGA6C	3.5		
Golgin A6 family-like 9	GOLGA6L9	2.1		
Golgin A8 family, member I	GOLGA8I	3.5		
Glycosylphosphatidylinositol anchored high density lipoprotein binding protein 1	GPIHBP1	2.0		
Glycoprotein M6A	GPM6A	2.2		PN
G protein-coupled receptor 128	GPR128	2.3		111
G protein-coupled receptor 18	GPR18	1.9		
Growth factor receptor-bound protein 14	GRB14	1.9		
GREB1 protein	GREB1	2.2		
Glutamate receptor, ionotropic, N-methyl D-aspartate 2B	GRIN2B	3.0		
General transcription factor IIH, polypeptide 2, 44kDa	GTF2H2	3.8		
Glycogenin 2	GYG2	2.0		
Hyaluronan synthase 3	HAS3	3.4		
HAUS augmin-like complex, subunit 5	HAUS5	2.2		
Hemoglobin, α 1	HBA1	1.9		
Hyperpolarization activated cyclic nucleotide-gated potassium channel 1	HCN1	2.3		
Histone deacetylase 7	HDAC7	1.9		
High density lipoprotein binding protein (vigilin)	HDLBP	6.8	(1)	
Hairy and enhancer of split 2 (Drosophila)	HES2	2.7		
Histone cluster 1, H2ag	HIST1H2AG	3.0		
Major histocompatibility complex, class I, A	HLA-A	3.4		
Major histocompatibility complex, class II, DR α	HLA-DRA	2.2		
H2.0-like homeobox	HLX	3.6		
Histamine N-methyltransferase	HNMT	2.0	(1)	
Homeobox B4	HOXB4	2.2		
Harakiri, BCL2 interacting protein (contains only BH3 domain)	HRK	4.2		
Heat shock transcription factor, Y-linked 1	HSFY1	5.1		
Heat shock protein 90kDa α (cytosolic), class B member 4 (pseudogene)	HSP90AB4P	1.8		
Histatin 3	HTN3	2.4		
Hydrocephalus inducing homolog (mouse)	HYDIN	3.5		
Insulin-like growth factor binding protein 3	IGFBP3	2.3	(1)	
Interleukin 15 receptor, α	IL15RA	2.1		MES
Interleukin 17 receptor B	IL17RB	4.4		
Interleukin 1 receptor, type II	IL1R2	2.1	(1)	
Interleukin 1 receptor accessory protein	IL1RAP	1.9	(1)	
Interleukin 2 receptor, β	IL2RB	2.4		
Interleukin 5 (colony-stimulating factor, eosinophil)	IL5	2.9		
Inhibitor of growth family, member 1	ING1	1.9		
Inversin	INVS	2.4		
Interferon regulatory factor 5	IRF5	2.3		
Interferon regulatory factor 6	IRF6	1.8		
KAT8 regulatory NSL complex subunit 1-like	KANSL1L	2.2		
Potassium inwardly-rectifying channel, subfamily J, member 4	KCNJ4	2.6		
Potassium large conductance calcium-activated channel, subfamily M β member 3	KCNMB3	4.6	(1)	
Potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3	KCNS3	1.9	(1)	
Potassium channel, subfamily V, member 2	KCNV2	2.5		
KIAA1033	KIAA1033	1.8		
KIAA1211-like	KIAA1211L	2.2		
Killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1	KIR3DL1	2.5		
Klotho	KL VI VI 4	2.6		
Kallikrein-related peptidase 14	KLK14	4.6	(1)	
Laminin, β 3	LAMB3	2.3	(1)	

CSC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Leukocyte cell derived chemotaxin 1	LECT1	1.8		
LIM homeobox 1	LHX1	8.0	(1)	
LIM homeobox 6	LHX6	2.2		
Long intergenic nonprotein coding RNA 265	LINC00265	2.2		
Long intergenic nonprotein coding RNA 426	LINC00426	1.9		
Long intergenic nonprotein coding RNA 427	LINC00427	2.4		
Long intergenic nonprotein coding RNA 545	LINC00545	1.8		
Lines homolog (Drosophila)	LINS	2.0		
LIM homeobox transcription factor 1, α	LMX1A	3.2	(1)	
	LOC100128239	3.2		
	LOC100128292 LOC100131551	2.1 1.9		
	LOC100131331 LOC100132418	1.8		
	LOC100506060	1.8		
	LOC148696	1.9		
	LOC286083	2.0		
	LOC402160	3.6		
	LOC440895	2.3		
	LOC644248	2.4		
	LOC645177	3.1		
	LOC728724	2.6		
	LOC93432	2.2		
Leucine-rich repeats and guanylate kinase domain containing	LRGUK	2.7		
Low density lipoprotein-related protein 12	LRP12	3.3	(1)	
Leucine rich repeat containing 32	LRRC32	4.5	(1)	
LY6/PLAUR domain containing 6B	LYPD6B	8.0		
Mab-21-like 1 (C. elegans)	MAB21L1	3.1		CL
MAFF interacting protein (pseudogene)	MAFIP	2.8		
Melanoma antigen family A, 1 (directs expression of antigen MZ2-E)	MAGEA1	2.7	(1)	
Melanoma antigen family A, 3	MAGEA3	2.4	(1)	
Melanoma antigen family B, 1	MAGEB1	2.1		
Melanoma antigen family C, 2	MAGEC2	1.9	(1)	
Mago-nashi homolog B (<i>Drosophila</i>)	MAGOHB	4.3	(4)	
Mannosidase, β A, lysosomal-like	MANBAL	2.2	(1)	
Microtubule-associated protein 1 light chain 3 γ	MAPILC3C	3.1		
Membrane-associated ring finger (C3HC4) 1	MARCH1	3.3		
MARVEL domain containing 1	MARVELD1 MB	2.4 3.7		
Myoglobin MCF 2 call line derived transforming sequence	MCF2	7.3		
MCF.2 cell line derived transforming sequence Minichromosome maintenance domain containing 2	MCMDC2	2.0		
Mucolipin 3	MCOLN3	2.2		
Methyltransferase like 2B	METTL2B	2.3		
Microfibrillar-associated protein 4	MFAP4	7.1		
The contract protein 1	MGC72080	2.5		CL
Matrix Gla protein	MGP	2.0		
MER1 repeat containing imprinted transcript 1	MIMT1	2.6		
MKL/myocardin-like 2	MKL2	2.0		
Myeloid/lymphoid or mixed-lineage leukemia 3	MLL3	2.1	(1)	
Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10	MLLT10	3.3	(1)	
Motilin	MLN	3.0		
Membrane metallo-endopeptidase	MME	3.8		
Matrix metallopeptidase 13 (collagenase 3)	MMP13	3.1	(1)	
Matrix metallopeptidase 23A (pseudogene)	MMP23A	2.2		
Membrane protein, palmitoylated 4 (MAGUK p55 subfamily member 4)	MPP4	2.8		
Mitochondrial ribosomal protein L3	MRPL3	1.9		
Mitochondrial ribosomal protein S12	MRPS12	3.0		
Membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I,	MS4A2	2.2		
receptor for; β polypeptide)				
Mucin 19, oligomeric	MUC19	5.0		
MAX interactor 1	MXI1	5.0	(1)	
V-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)	MYCL1	2.0	(1)	
Myosin, light chain 10, regulatory	MYL10	1.8		
Myosin IB Myosin IC	MYO1B MYO1C	4.6 2.3		
NLR family, apoptosis inhibitory protein	NAIP	2.4		
Neuroblastoma breakpoint family, member 15	NBPF15	2.8		
Neurocalcin δ	NCALD	1.8		PN
Neutrophil cytosolic factor 1, (chronic granulomatous disease, autosomal 1)	NCF1	2.6		114
Nonprotein coding RNA 185	NCRNA00185	1.8		
NEBL antisense RNA 1	NEBL-AS1	1.8		
NEL-like 1 (chicken)	NELL1	2.1	(1)	
Nucleolar complex associated 4 homolog (Saccharomyces cerevisiae)	NOC4L	2.1	(*)	
NADPH oxidase organizer 1	NOXO1	2.2	(1)	
o			V-7	

Sodium channel, voltage-gated, type I, β

Selectin E (endothelial adhesion molecule 1)

Sidekick homolog 2 (chicken)

TABLE W1. (continued) CSC Description Symbol Fold Change Sox2 Target Subclass (Ref) (Log₂) Signature Neuropeptide S NPS 2.2 (1) Neuronal pentraxin II NPTX2 3.1 (1) Neurexin 2 NRXN2 2.1 PN Neurotensin receptor 1 (high affinity) NTSR1 2.8 Optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia) OPA3 2.7 Olfactory receptor, family 11, subfamily A, member 1 OR11A1 2.1 Olfactory receptor, family 2, subfamily A, member 9 pseudogene OR2A9P 4.6 Olfactory receptor, family 51, subfamily E, member 2 OR51E2 3.6 OTU domain containing 1 OTUD1 2.1 Purinergic receptor P2X, ligand-gated ion channel, 3 P2RX3 P antigen family, member 5 (prostate associated) PAGE5 1.9 Paralemmin 3 PALM3 4.7 Par-3 partitioning defective 3 homolog (C. elegans) PARD3 3.0 PCDH11X Protocadherin 11 X-linked 3.5 Protocadherin α 12 PCDHA12 5.5 Protocadherin y subfamily B, 7 PCDHGB7 2.9 Phosducin-like 3 pseudogene 4 PDCL3P4 2.2 **PDGFRB** Platelet-derived growth factor receptor, β polypeptide 2.8 PDXDC2P Pyridoxal-dependent decarboxylase domain containing 2, pseudogene 19 PDZ domain containing 1 PDZK1 3.1 Platelet/endothelial cell adhesion molecule (CD31 antigen) PECAM1 1.9 Pepsinogen 3, group I (pepsinogen A) PGA3 3.5 Placental growth factor, vascular endothelial growth factor-related protein PGF 2.7 PHLDA3 Pleckstrin homology-like domain, family A, member 3 2.0 Phosphatidylinositol 4-kinase type 2 β PI4K2B 2.3 Phospholipase A2, group IIC PLA2G2C 2.5 MFS Plasminogen activator, urokinase PLAU 1.8 **PLAUR** MES Plasminogen activator, urokinase receptor 2.5 PLCL2 2.4 Phospholipase C-like 2 Pleckstrin homology domain containing, family G (with PGef domain) member 3 PLEKHG3 2.3 Premelanosome protein **PMEL** 2.6 POM121L1P POM121 transmembrane nucleoporin-like 1, pseudogene 2.6 Proopiomelanocortin (adrenocorticotropin/ β-lipotropin/ α-melanocyte stimulating hormone/ **POMC** 2.0 β-melanocyte stimulating hormone/ β-endorphin) PPIE Peptidylprolyl isomerase E (cyclophilin E) 19 PPP1R9B Protein phosphatase 1, regulatory (inhibitor) subunit 9B 2.0 PR domain containing 1, with ZNF domain PRDM1 44 Protein kinase C, ε PRKCE 2.2 (1) Proline rich 5 (renal) PRR5 4.2 (1) Prostaglandin D2 synthase 21kDa (brain) PTGDS 5.5 Prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase) PTGS1 5.2 PTPRE 1.8 Protein tyrosine phosphatase, receptor type, E Protein tyrosine phosphatase, receptor type, O PTPRO (1) 1.9 Pyridine nucleotide-disulphide oxidoreductase domain 2 PYROXD2 1.8 RAB20, member RAS oncogene family RAB20 2.2 (1) RAB30, member RAS oncogene family RAB30 1.8 RAET1G Retinoic acid early transcript 1G 6.2 RAN guanine nucleotide release factor RANGRF 1.9 Rap guanine nucleotide exchange factor (GEF) 5 RAPGEF5 2.2 Retinoic acid receptor, α RARA 2.2 RAS protein activator like 2 RASAL2 2.7 (1) RASD family, member 2 RASD2 2.6 Ras association (RalGDS/AF-6) domain family 6 RASSF6 4.1 Retinoblastoma binding protein 6 RBBP6 2.4 RCC1 domain containing 1 RCCD1 2.0 RELT tumor necrosis factor receptor RELT 3.0 RFX7 Regulatory factor X, 7 1.8 Rhomboid 5 homolog 2 (Drosophila) RHBDF2 2.1 RIPPLY1 2.2 Ripply1 homolog (zebrafish) Ring finger protein 186 RNF186 2.1 Retinitis pigmentosa 1 (autosomal dominant) RP1 2.4 (1) RPRMI. 2.4 Reprimo-like Ras-related associated with diabetes RRAD 2.2 Runt-related transcription factor 1; translocated to, 1 (cyclin D-related) RUNX1T1 2.1 (1) Sterile a motif domain containing 14 SAMD14 2.3 SET binding factor 1 pseudogene 1 SBF1P1 2.9 Small Cajal body-specific RNA 13 SCARNA13 2.3 Small Cajal body-specific RNA 9-like SCARNA9L 2.6 Secretogranin V (7B2 protein) SCG5 2.6

(1)

SCN1B

SDK2

SELE

2.9

2.2

1.9

CSC				
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature
Sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3D	SEMA3D	3.2	(1)	
Serpin peptidase inhibitor, clade A (α-1 antiproteinase, antitrypsin), member 3	SERPINA3	4.9		
Serpin peptidase inhibitor, clade I (neuroserpin), member 1	SERPINI1	4.0		NL
SH3 and PX domains 2A	SH3PXD2A	6.2	(1)	
Src homology 2 domain containing E	SHE	3.2		
Src homology 2 domain containing F	SHF	2.1		
SIX homeobox 4	SIX4	1.8		
Solute carrier family 16, member 10 (aromatic amino acid transporter)	SLC16A10	2.5	(1)	
Solute carrier family 16, member 14 (monocarboxylic acid transporter 14)	SLC16A14	1.9		
Solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7 Solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15	SLC17A7 SLC25A15	2.5 2.3		
Solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 24	SLC25A13	7.4		
Solute carrier family 25 (introduction at earlier, phosphate earlier), member 24 Solute carrier family 25, member 52	SLC25A52	2.5		
Solute carrier family 26, member 4	SLC26A4	2.5		
Solute carrier family 27 (fatty acid transporter), member 4	SLC27A4	3.5		
Solute carrier family 39 (zinc transporter), member 12	SLC39A12	3.0		
Solute carrier family 39 (zinc transporter), member 3	SLC39A3	1.8		
Solute carrier family 39 (zinc transporter), member 8	SLC39A8	2.0		
Solute carrier family 45, member 3	SLC45A3	2.2		
Solute carrier family 46, member 2	SLC46A2	2.3		
Solute carrier family 5 (sodium/glucose cotransporter), member 2	SLC5A2	5.7		
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2	SMARCC2	2.5		
Small integral membrane protein 8	SMIM8	2.6		
Snail homolog 2 (<i>Drosophila</i>)	SNAI2	2.0		
Small nuclear RNA activating complex, polypeptide 3, 50kDa	SNAPC3	3.9		
Stannin	SNN	2.0		
Small nucleolar RNA, H/ACA box 72	SNORA72	2.6		
Small nucleolar RNA, C/D box 101	SNORD101	1.9		
Small nucleolar RNA, C/D box 113-6	SNORD113-6	3.4		
Sorting nexin family member 21	SNX21	2.2		
Sosondowah ankyrin repeat domain family member A	SOWAHA	3.2		
SRY (sex-determining region Y)-box 18	SOX18	2.2		MEC
SP100 nuclear antigen	SP100 SPAG16	2.2 2.0		MES
Sperm associated antigen 16 SPATA31 subfamily D, member 5, pseudogene	SPATA31D5P	2.3		
Speedy homolog A (<i>Drosophila</i>)	SPDYA	2.5		
Speedy/RINGO cell cycle regulator family member E6	SPDYE6	3.3		
Sphingosine kinase 1	SPHK1	1.8		
Spondin 1, extracellular matrix protein	SPON1	2.0		
Steroidogenic acute regulatory protein	STAR	3.5		
Serine/threonine kinase 16	STK16	4.3		
Serine/threonine kinase 17b	STK17B	2.1	(1)	
Suppressor of G2 allele of SKP1 (Saccharomyces cerevisiae) pseudogene 3	SUGT1P3	2.4		
Sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3	SULT1A3	2.1		
Sad1 and UNC84 domain containing 3	SUN3	5.2		
Sushi domain containing 2	SUSD2	2.9		
Supervillin	SVIL	6.8	(1)	
Synaptonemal complex central element protein 3	SYCE3	2.0		
Synaptotagmin III	SYT3	2.0		
Transgelin	TAGLN	2.6		
T cell acute lymphocytic leukemia 2	TAL2	4.3		
TBC1 domain family, member 26	TBC1D26	2.7		
Transducin (β)-like 1Y-linked	TBL1Y	3.9		
T cell, immune regulator 1, ATPase, H+ transporting, lysosomal V0 subunit A3	TCIRG1	1.9		MES
Testis development related 1 (nonprotein coding)	TDRG1	3.9	(1)	
TEK tyrosine kinase, endothelial (venous malformations, multiple cutaneous and mucosal)	TEK	2.0	(1)	
Testis-specific kinase 2	TESK2	7.3		
Tet methylcytosine dioxygenase 3	TET3	5.4		
THAP domain containing 4	THAP4 TJP3	2.5 3.2		
Tight junction protein 3 Transmembrane 4 L six family member 19	TM4SF19	2.0		
Transmembrane channel-like 4	TMC4	4.9		
Transmembrane protein with EGF-like and two follistatin-like domains 2	TMEFF2	2.3	(1)	
Transmembrane protein 107	TMEM107	1.8	(1)	
Transmembrane protein 110	TMEM110	2.2		
Transmembrane protein 120B	TMEM120B	2.0		
Transmembrane protein 191B	TMEM191B	2.7		
Transmembrane protein 65	TMEM65	1.8		
Thioredoxin-related transmembrane protein 1	TMX1	5.5		
Tumor necrosis factor, α-induced protein 2	TNFAIP2	1.9		
Tumor necrosis factor, α-induced protein 8	TNFAIP8	1.9		MES
	TNFRSF18	2.5		
Tumor necrosis factor receptor superfamily, member 18	111110110	2.7		

THE REAL PROPERTY.	
TABLE W1.	(continued)

TABLE W1. (continued) CSC						
Description	Symbol	Fold Change (Log ₂)	Sox2 Target (Ref)	Subclass Signature		
TNNI3 interacting kinase	TNNI3K	2.7				
Topoisomerase (DNA) I pseudogene 1	TOP1P1	2.9				
TOX high-mobility group box family member 2	TOX2	2.6				
Tumor protein p53 regulated apoptosis inducing protein 1	TP53AIP1	2.1				
Trafficking protein particle complex 6B	TRAPPC6B	1.9				
TRNA aspartic acid methyltransferase 1	TRDMT1	2.4				
Tripartite motif-containing 51E, pseudogene	TRIM51EP	2.7				
Transient receptor potential cation channel, subfamily M, member 1	TRPM1	3.5				
Testis-specific transcript, Y-linked 5	TTTY5	2.4				
Tubulin, α 8	TUBA8	2.3				
Tubulin, β pseudogene 5	TUBBP5	4.4				
Urocortin 2	UCN2	3.1				
Vestigial like 2 (Drosophila)	VGLL2	1.9				
Vitelline membrane outer layer 1 homolog (chicken)	VMO1	2.0	(1)			
Vanin 2	VNN2	4.8				
WASH and IL9R antisense RNA 1	WASIR1	4.0				
WD repeat domain 63	WDR63	4.1				
WD repeat domain 87	WDR87	1.8				
WAP four-disulfide core domain 2	WFDC2	2.9				
YME1-like 1 (Saccharomyces cerevisiae)	YME1L1	2.2				
·	YPLR6490	2.4				
Zinc finger, CCHC domain containing 12	ZCCHC12	2.1				
Zinc finger, FYVE domain containing 27	ZFYVE27	2.4				
ZMIZ1 antisense RNA 1	ZMIZ1-AS1	2.0				
Zinc finger, MYND-type containing 15	ZMYND15	2.7				
Zinc finger protein 138	ZNF138	2.9				
Zinc finger protein 48	ZNF48	2.2				
Zinc finger protein 493	ZNF493	2.3				
Zinc finger protein 570	ZNF570	3.8				
Zinc finger protein 607	ZNF607	2.0				
Zinc finger protein 793	ZNF793	3.6				
Zinc finger protein 815, pseudogene	ZNF815P	2.9				
Zinc finger protein 847, pseudogene	ZNF847P	2.7				
Zona pellucida-like domain containing 1	ZPLD1	2.1				

Bold, genes commonly regulated in SCS and CSC; italic, genes regulated in oposite directions in SDC and CSC.

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 $\textbf{Table W2.} \ \ \text{Genes Highly Correlated with Sox2 mRNA Expression in the TCGA Data}.$

Gene ID	Correlation	In Vitro Sox2 Knockdown DEG	Sox2 Target (Ref)	Verhaak Subclass
ABCA3	Positive			_
ARNT2	Positive	SDC		
ASTN1	Positive	CSC		
B3GAT1	Positive	SDC	(1)	
BAALC	Positive	SDC		
BAI2	Positive			
BCAN	Positive			PN
C10rf61	Positive	SDC	(1)	PN
C3orf39	Positive			
CHST10	Positive	SDC		
СКВ	Positive	SDC		PN
CLIP3	Positive			
CTNND2	Positive			
DBI	Positive			
DCLK2	Positive			
DDR1	Positive			
DENND5A (RAB61P1)	Positive		(1)	
DNAJB5	Positive		(1)	
DTX1	Positive			
EBF4	Positive			
FAM181B (MGC33846)	Positive		(1)	
FEZ1	Positive	SDC	(1)	
FXYD6		SDC		PN
GDF1	Positive		(1)	rn
	Positive			
GNA11	Positive	222 202		
GPM6B	Positive	SDC, CSC		CY.
GPR56	Positive	SDC		CL
HEPN1	Positive			
ID4	Positive			
JAKMIP2	Positive			C.Y.
KEAP1	Positive		(1)	CL
LRIG1	Positive	00.0	(1)	
METRN	Positive	SDC		-
MLC1	Positive			CL
MT3	Positive			
NFIX	Positive			
NLRP1	Positive			
NR2F1	Positive		(1)	
OLIG1	Positive			
OSBPL6	Positive	SDC	(1)	
PCDHGA1	Positive			
POU3F2	Positive			
PTPRZ1	Positive		(1)	
RGMA	Positive			
SHC3	Positive			
SLC1A3	Positive	SDC	(1)	
SOX2	Positive	SDC,CSC	(1)	PN
SOX21	Positive	SDC,CSC	(1) (2)	
TBCB	Positive			
TCEA2	Positive			
TTYH1	Positive	SDC, CSC		PN
TUBA4A	Positive			
TUBB2A	Positive			
TUBB2B	Positive	SDC		
TUBB6	Positive			
TUBB8	Positive			
ZBTB47	Positive			
ANXA3	Negative	SDC		NL
ATF1	Negative			***
B3GNT2	Negative		(1)	
C4orf32	Negative		(1)	
C6orf211	Negative		(-/	
FEM1B	Negative			
ITGA4	Negative			MES
MOB1A				IVILES
	Negative			
MST4	Negative			
NUDT15	Negative			
PCYOX1	Negative			

TABLE W2. (continued)						
Gene ID	Correlation	In Vitro Sox2 Knockdown DEG	Sox2 Target (Ref)	Verhaak Subclass		
PTPN2	Negative					
RALB	Negative		(1)			
STT3B	Negative					
TBC1D8B	Negative					
THOC7	Negative					
TMF1	Negative					

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^{1.} Fang X, Yoon JG, Li L, Yu W, Shao J, Hua D, Zheng S, Hood L, Goodlett DR, Foltz G,et al. (2011). The SOX2 response program in glioblastoma multiforme: an integrated ChIP-seq, expression microarray, and microRNA analysis. BMC Genomics 12,11.

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