Roles of sex-determining region Y-box 2 in cell pluripotency and tumor-related signaling pathways (Review)

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Abstract. The sex-determining region Y-box 2 (SOX2) gene, a member of the Sry-like high-mobility group box (SOX) gene family, encodes the transcription factor Sox2, which significantly contributes to the regulation of cell pluripotency. Sox2 is closely associated with early embryonic development, neural differentiation and other biological processes. An inreasing number of recent studies suggest that Sox2 exerts a positive effect on malignant tumors. According to these results, Sox2 is expected to become a novel target for cancer therapy by unveiling the mechanism through which it affects the biological behavior of tumors. Therefore, it is crucial to elucidate the detailed association of Sox2 with malignant tumors. The aim of this study was to review the role of Sox2 in pluripotency maintenance, early embryonic development and neural differentiation, as well as investigate the detailed mechanism through which Sox2 regulates cancer stem cells and tumorigenesis.

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Abbreviations: Sox2, sex-determining region Y-box 2; SOX, Sry-like high-mobility group box; CSCs, cancer stem cells; PI3K, phosphatidylinositol 3-kinase

Key words: sex-determining region Y-box 2, pluripotency, cancer stem cells, malignant tumor, signaling pathway

1. Introduction

Characteristics of sex-determining region Y-box 2 (Sox2). The transcription factor Sox2 belongs to the high-mobility group superfamily and is encoded by the SOX2 gene, a member of the SOX gene family, the members of which are conserved across species and involved in a number of developmental processes (1). Since the discovery of the first SOX gene, ~20 genes of the SOX gene family have been identified over the last 20 years in mammals and classified into groups according to protein specificity (2). The SOX2 gene was discovered and characterized in humans in 1994; it belongs to the SOXB1 group, which also includes SOX1 and SOX3. The SOXB1 group locates on chromosome 3q26.3-q27 and encodes a protein consisting of 317 amino acids (3,4). Although there are >80% sequence similarities across the SOXB1 family, SOX2 plays a distinct role in complex biological progress in a context-dependent manner, and is an indispensable factor for embryonic development.

Function of Sox2. Sox2 is well known as one of the pluripotency factors, also including Oct4, Nanog and Klf4. Sox2 contributes significantly to the regulation of cell pluripotency and is closely associated with early embryonic development, neural differentiation and sexual differentiation.

Sox2 is crucial for the maintenance of pluripotency in cells. Human pluripotent stem cells (hPSCs) include human embryonic stem cells (hESCs) and induced pluripotent stem cells (iPSCs) (5). These cells possess two important properties, namely limitless self-renewal in culture and the ability to differentiate into different types of blastoderm in the early embryo (6). These unique properties make hPSCs a research focus in the field of personalized medicine, drug screening or cell therapy (7,8) and the study of early human development, serving as a cell model to elucidate the molecular mechanisms regulating embryonic cell proliferation and differentiation (7). hPSCs are regulated by several stem genes, including SOX2. It has been reported that iPSCs were first produced from fibroblasts through regulation of Sox2. Liu et al (8) reported that, in human iPSCs with human amniotic epithelial cells as feeder cells, overexpression of Sox2 maintained cell pluripotency via inhibition of endogenous microRNA-145 expression. Xu et al (9) also found that microRNA-145 regulates Sox2 and represses pluripotency in hESCs. These results indicate that Sox2 is indispensable in the maintenance of the pluripotency of hPSCs.

Sox2 plays an important role in embryogenesis. Sox2 is a factor indispensable to embryonic development. It was previously indicated that the main function of Sox2 is to help maintain the pluripotency of ESCs and repress trophectoderm and epiblast genes in later stages of development (10). During mouse embryogenesis, a zygote multiplies via mitosis, resulting in the formation of the multicellular morula (16-32 cells), which further develops to form the blastocyst (>32 cells). The blastoderm in the blastocyst is divided into the inner cell mass (ICM) and the trophectoderm. Cells in the ICM have the ability to differentiate into any type of cell found in the human body. Thus, ICM cells are classified as pluripotent. Sox2 was first identified in the morula, specifically in the ICM of the blastocyst and epiblast (11). Lack of Sox2 expression significantly affects the formation of the trophectoderm (11), indicating that Sox2 plays an important role in the formation of the embryo and the maintainance of cell pluripotence.

Sox2 plays an essential role in neural differentiation. Sox2 also appears to play an important role in the differentiation of the nervous system (10). Sox2 is expressed not only in ESCs, but also in several other types of stem cells, including neural stem cells, and it regulates the progress of primary neurogenesis, neural development and neural differentiation. Mizuseki et al (12) indicated in 1998 that Sox2 alone is not sufficient to induce neural tissue differentiation in ectodermal explants excised from blastulas. In 2009, Archer et al (13) demonstrated that there is an interaction among Sox1, Sox2, Sox3 and Oct4 during primary neurogenesis. Only Sox1 and Oct91 induce the formation of neurons; Sox2, however, is more effective compared with Sox1 in maintaining cells in a progenitor state. While the SoxB1s have overlapping functions, they are not strictly redundant, as they induce different sets of genes and are likely to partner with different proteins to maintain progenitor identity. Therefore, Sox2 is an important factor, although not the only factor, for primary neurogenesis, and it acts synergistically with other members of the SoxB1 family.

2. Association between Sox2 and cancer stem cells (CSCs)

CSCs, also referred to as tumor-initiating cells or cancer stem-like cells, have been experimentally defined by their ability to initiate tumor formation upon implantation in immunocompromised mice and are considered to be an important factor in tumorigenesis, invasion and metastasis. Sox2 acts in CSCs via a number of signaling pathways. In 2012, Basu-Roy *et al* (14) reported that Sox2 is highly expressed in human and murine osteosarcoma cell lines, as well as in tumor samples, and is essential for the self-renewal of osteoblast progenitor cells. However, Sox2 antagonizes the Wnt pathway, which is a pro-differentiation pathway that may, in turn, reduce Sox2 expression. Rybak *et al* (15) reported in 2013 that the activation of epidermal growth factor receptor (EGFR) signaling pathways in prostate cancer stem cells (PCSCs) contributes to a significant upregulation of SOX2 at both the mRNA and the

protein levels, promoting the self-renewal of DU145 PCSCs. In addition, other studies indicated that the SOX2 expression status of CSCs is the basis of different pathological types of tumors (16,17). Therefore, Sox2 is an important factor in tumorigenesis via CSC amplification.

3. Correlation between Sox2 and malignant tumors in clinical statistics

An increasing number of studies suggest that Sox2 exerts a positive effect on pluripotency regulation, tumorigenesis, tumor invasion and metastasis in breast cancer (18), melanoma (19), laryngeal carcinoma (20), hepatocellular carcinoma (HCC) (21), ovarian cancer (22) and non-small-cell lung cancer (23).

Huang *et al* (18) evaluated 57 ductal carcinomas *in situ*, 552 invasive breast carcinomas and 107 corresponding metastatic lymph nodes and found that Sox2 expression was closely associated with clinicopathological parameters, including high histological grade, large tumor size, molecular subtypes with adverse outcome, negative hormone receptor status, high proliferation index and neuroendocrine marker expression.

Sun *et al* (21) evaluated 75 HCCs and discovered that high levels of Sox2 expression were correlated with metastasis and low survival rate in HCC. They also investigated the molecular role of Sox2 in human HCC cell lines with stepwise metastatic potential (MHCC97L, MHCC97H and MHCCLM3) and found that Sox2 activates epithelial-to-mesenchymal transition (EMT) in HCC cells (21).

Zhang *et al* (24) investigated the clinical role of Sox2 expression in ovarian carcinoma. Immunohistochemical staining of 540 human ovarian carcinoma samples for Sox2 was performed using tissue microarray and the results suggested that Sox2 expression was associated with high-grade carcinomas, particularly high-grade serous carcinoma, International Federation of Gynecology and Obstetrics stage and malignant mixed Müllerian tumors (P=0.048).

Several other malignant tumor clinical analyses have been performed in recent years on lung cancer, melanoma and brain tumors, providing evidence that Sox2 is closely associated with tumor invasion and metastasis and predicts poor prognosis in malignant tumor patients. However, the underlying mechanism remains to be elucidated.

4. Role of Sox2 in tumor-related signaling pathways

Sox2 regulates tumor growth, invasion and metastasis via a plurality of signaling pathways, including the phosphatidylinositol 3-kinase (PI3K)/protein kinase B (Akt), Wnt/ β -catenin, Notch, bone morphogenetic protein (BMP), estrogen receptor α (ER α) and mitogen-activated protein kinase kinase kinase kinase 4 (MAP4K4)-survivin signaling pathway. These signaling pathways form a complex regulatory network, indicating that Sox2 plays a decisive role in tumor bioinformatics (Table I).

PI3K/Akt signaling pathway. The PI3K/Akt and related signaling pathways are among the most well-known tumor-related signaling pathways; they are important in internalizing the effects of external growth factors and

Table I. Signaling pathways associated with Sox2 in different malignant tumors.

Signaling pathways	Related malignant tumors	References
PI3k/Akt	Prostate cancer	(26), (27)
	Hepatocellular carcinoma	(28)
	Breast cancer	(29)
	Esophageal squamous cell carcinoma	(30)
Wnt/β-cantenin	Prostate cancer	(32)
	Glioblastoma	(33)
	Breast cancer	(32)
	Laryngeal cancer	(34)
Notch	Nasopharyngeal carcinoma	(38)
	Lung cancer	(17), (39)
BMP	Colorectal cancer	(40)
	Lung squamous cell carcinoma	(41)
$ER\alpha$	Breast cancer	(42)
MAP4K4-survivin	Lung cancer	(43)

Sox2, sex-determining region Y-box 2; PI3k, phosphatidylinositol 3-kinase; BMP, bone morphogenetic protein; $ER\alpha$, estrogen receptor α ; MAP4K4, mitogen-activated protein kinase kinase kinase kinase 4.

membrane tyrosine kinases. The activation of membrane kinases, including EGFR, by external growth factors initiates receptor dimerization and subsequent events to activate these intracellular pathways. PI3K participates in the regulation of proliferation, differentiation, apoptosis, glucose transport and other cellular functions. The most extensively investigated member of the PI3K family is type I PI3K, which may be activated by cell surface receptors. In mammalian cells, type I PI3K includes two subtypes, namely IA and IB. Recent publications indicated that signaling pathways including IA and its downstream molecules, such as Akt, are closely associated with the occurrence and development of malignant tumors. This pathway regulates tumor cell proliferation and survival; its abnormal activity not only causes cell malignant transformation, but is also associated with tumor cell migration, adhesion, angiogenesis and extracellular matrix degradation. Activated Akt regulates the expression of downstream molecules, including mammalian target of rapamycin (mTOR), extracellular signal-regulated kinase, Forkhead, guanosine diphosphate, insulin receptor substrate, glycogen synthase kinase 3 (GSK3), mitogen-activated protein kinase, nuclear factor-B, protein kinase C and signal transducer and activator of transcription, and then regulates cell proliferation, differentiation, apoptosis and migration. Sox2 was recently identified as a novel regulator of the PI3K/Akt pathway. Yu et al (25) reported in 2012 that Sox2 promotes the expression of CD49f, which regulates the ability of hMSCs to form spheres via activating the PI3K/Akt signaling pathway.

The role of the PI3K/Akt signaling pathway in malignant tumors is attracting increasing attention in the field of clinical and preclinical medicine. Morgan *et al* (26) reported in 2009 the prominent role of the PI3K/Akt/mTOR pathway in prostate cancer. Molecular changes in the PI3K/Akt/mTOR signaling pathway have been demonstrated to distinguish benign from malignant prostatic epithelium and are associated

with increasing tumor stage, grade and risk of biochemical recurrence. In addition, the PI3K/Akt/mTOR signaling pathway is associated with drug resistance in prostate cancer. Li *et al* (27) investigated the PC-3 prostate cancer cell line in 2013 and found that overexpression of Sox2 enhanced the resistance of PC-3 cells to paclitaxel through promoting cell proliferation and exhibiting an anti-apoptotic effect via activation of the PI3K/Akt pathway. Another study on HCC demonstrated that cyclin G1 expands liver tumor-initiating cells through Sox2 induction via Akt/mTOR signaling (28). Similar results were reported in breast cancer (29) and esophageal squamous cell carcinoma (SCC) (30), suggesting that Sox2 is correlated with CSCs via the PI3K/Akt pathway, although the detailed mechanism remains to be elucidated.

Wnt/β-catenin signaling pathway. The Wnt signaling pathway is among the most well-known tumor-related signaling pathways and it is a complex protein interaction network, most commonly involved in embryonic development and tumorigenesis. Three pathways, including the Wnt/β-catenin, planar cell polarity and Wnt/Ca2+ pathways, act synergistically but independently in physiological processes. Among these, the Wnt/β-catenin pathway is the one most closely associated with tumorigenesis. Activation of this pathway stabilizes β -catenin and, therefore, contributes to accumulation of this protein, which then enters the nucleus to activate the expression of target genes. In the absence of secreted glycolipoprotein Wnt, cytoplasmic β-catenin protein is constantly degraded by the action of the Axin complex, which is composed of the scaffolding protein Axin, the tumor suppressor adenomatous polyposis coli gene product, casein kinase 1 and GSK3 (31). Sox2 is a negative regulator of the Wnt pathway via binding to β-catenin and inducing the Wnt inhibitor Dickkopf-related protein (Dkk) 1. Li et al (32) reported that Sox2 promotes metastasis of breast and prostate cancer cells by promoting

EMT through WNT/ β -catenin. In addition, Sox2 affects the protein expression levels of Dkk3, dishevelled segment polarity protein (DVL) 1 and DVL3, which are regulators or downstream molecules of Wnt signaling (32). A previous study by Berezovsky *et al* (33) on glioblastoma multiforme (GBM) indicates that Sox2 reprograms differentiated cells into pluripotent cells and regulates the expression of key genes and Wnt pathways involved in GBM in cancer stem-like and differentiated cells. Another study by Yang *et al* (34) on HEp-2 laryngeal cancer cells revealed that the overexpression of Sox2 promotes cancer cell migration, invasion and EMT through the Wnt/ β -catenin pathway. However, the detailed association between the Sox2 and Wnt pathways has not been elucidated.

Notch signaling pathway. The NOTCH gene was firstly discovered in *Drosophila*, in which partial loss of function of this gene caused a notch of the wing edge. The Notch signaling pathway is a highly conserved transduction pathway in the evolution and function of cell proliferation, differentiation and apoptosis regulation, and is involved in almost all tissues and organs. Notch is a transmembrane receptor, widely present in almost all known animal cells. The Notch pathway mediates cell-cell signal transfer and the corresponding signaling cascade reaction. The Notch signaling pathway is composed of Notch, Notch ligand (DSL protein) and CSL (a DNA-binding protein). Notch and its ligands are all single transmembrane proteins. Notch is cut by proteasome when the ligand binds to Notch of the adjacent cell, releasing intracellular Notch (ICN), which carries the nuclear localization signal in the cytoplasmic region. ICN then enters the nucleus and binds to CLS to regulate target gene expression. To date, the function of the Notch signaling pathway in the regulation of cell proliferation, differentiation and apoptosis has been extensively investigated, particularly in the inner ear (35-37). However, the number of studies on the interaction between Sox2 and the Notch pathway in tumor invasion and metastasis is limited. Zhang et al (38) investigated Notch1 signaling in nasopharyngeal carcinoma (NPC) and found that the Notch1 signaling-activated form was predominantly found in Sox2-negative cells. This finding indicated that the overexpression of Sox2 inhibits Notch1 signaling and contributes to the pathological self-renewal characteristics of CSCs in human NPC. By contrast, Chen et al (17) found that NOTCH3, c-Myc and WNT7B were significantly reduced after silencing SOX2 in A549 human lung cancer cells, suggesting that Notch signaling is the downstream pathway of Sox2. Another research on lung cancer revealed that the overexpression of Sox2 was able to inhibit Notch signaling in CC10+ cells, subsequently inhibiting K-Ras-induced tumor formation; inhibition of Notch signaling strongly inhibits adenocarcinoma formation, but promotes squamous hyperplasia in the alveoli (39). The abovementioned research results demonstrated that Sox2 may regulate the activity of the Notch signaling pathway and carcinogenesis. Of note, Sox2 may not only activate but also inhibit the Notch pathway, indicating that Sox2 exerts a two-way regulatory effect on Notch. The detailed mechanism, however, remains to be elucidated.

Other signaling pathways. The PI3K/Akt, Wnt/β-catenin and Notch signaling pathways are the most well-known pathways involved in tumorigenesis. In addition, there are

more signaling pathways correlated with Sox2 in regulating carcinogenesis. Fang et al (40) used ChIP-seq and functional analysis in SW620 colorectal cancer cells and found that Sox2 is associated with the BMP signaling pathway and a number of receptor-mediated signaling pathways, such as insulin-like growth factor 1 receptor and inositol 1,4,5-triphosphate receptor, type 2. A study conducted in 2014 by Fang et al (41) suggested that BMP4 is a downstream target of Sox2 in lung SCC. Sox2 is a negative regulator of BMP4 and overexpression of Sox2 may reduce the expression of BMP, promoting the occurrence and development of SCC (41). In addition, in breast cancer, Sox2 was found to be a new target of miRNA140, regulating breast tumor-initiating cells via the ERα signaling pathway (42). Chen et al (43) found that Sox2 regulates apoptosis through the MAP4K4-survivin signaling pathway in human lung cancer cells. In conclusion, Sox2 plays different roles in the regulation of tumor development, invasion and metastasis and drug resistance. The heterogeneity of research conclusions indicates the complexity of the network associating Sox2 with tumor-related signaling pathways. Therefore, the specific role of Sox2 in tumor-associated signaling pathways requires further investigation.

5. Conclusion and perspectives

Sox2 significantly contributes to the regulation of cell pluripotency and is closely associated with early embryonic development, particularly neural differentiation. Due to the similarities of the characteristics and functions between Sox2 and CSCs, the correlation between Sox2 and malignant tumors has been attracting increasing attention. An increasing number of studies suggest that Sox2 exerts a positive effect on pluripotency regulation, tumorigenesis, tumor invasion and metastasis in several types of malignant tumors. Sox2 regulates tumor growth, invasion and metastasis via a plurality of signaling pathways. These signaling pathways form a complex regulatory network, indicating that Sox2 plays a decisive role in tumor bioinformatics. Therefore, the roles of Sox2 in tumors require further elucidation. Sox2 and its upstream/downstream molecules in the transduction pathway network and the proteins interacting with Sox2 are expected to be novel targets for cancer therapy; the elucidation of the mechanism underlying the effect of Sox2 on the biological behavior of tumors is expected to lead to significant advances in cancer treatment.

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