


Emerging Roles of Long Noncoding RNA Regulator of Reprogramming in Cancer Treatment

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Abstract: Despite numerous advances in cancer treatment, the global prevalence and cancer-related mortality remain high. Understanding tumor initiation and progression mechanisms are critical as it will lead to the development of interventions for improving the prognosis of cancer patients. The roles of long noncoding RNAs (lncRNAs) in cancer have attracted immense research interest. Growing evidence indicates that lncRNA regulator of reprogramming (linc-ROR), a well-studied RNA, regulates the progression of various cancers, such as lung cancer (LC), hepatocellular carcinoma (HCC), breast cancer (BC), colorectal cancer (CRC), pancreatic cancer (PC), papillary thyroid carcinoma (PTC), or esophageal squamous cell carcinoma (ESCC). linc-ROR promotes the proliferation, invasion, migration and chemoresistance of cancer cells. Herein, we reviewed current literature on the modulatory functions and mechanisms of linc-ROR in cancer development. We highlight new linc-ROR-related therapeutic strategies in cancer treatment.

Keywords: cancer, long noncoding RNA, linc-ROR, mechanisms, clinical application

Introduction

Cancer remains a major public health challenge in the world.¹ Despite major investments into cancer research and management, its prognosis remains poor due to frequent relapse, metastasis, resistance to chemotherapy and lack of advanced diagnostic tools. Novel molecular targets that reverse chemoresistance or suppress metastasis, coupled with early diagnosis and treatment of cancer are urgently needed.

Protein-coding genes only account for less than 2% of the mammalian genome. Recently, research on the role of noncoding RNAs (ncRNAs) in cancer development has intensified.² ncRNAs such as long noncoding RNAs (lncRNAs), which contain >200 nucleotides, regulate multiple physiological and pathological processes. Although they lack protein-coding ability, lncRNAs regulate cis or trans transcription, organize nuclear domains and modulate protein expression.³ Recent studies have reported differentially expressed lncRNAs in cancer tissues compared to adjacent normal tissues. These differentially expressed lncRNAs have been shown to be associated with cancer initiation and development.⁴⁻⁶

The lncRNA regulator of reprogramming (linc-ROR) is upregulated in multiple human cancers regulating cancer initiation and development (Table 1). linc-ROR is located in chromosome 18 and possess 2.6 kb and four exons, with numerous binding sites between linc-ROR and miRNAs, and proteins.⁷ Typical targets of linc-ROR contain miR-145,⁸ miR-205,⁹ p53¹⁰ and so on. Linc-ROR was first reported in a study

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Table 1 Current Research on linc-ROR Associated with Malignant Tumor

Cancer Type	Binding miRNAs	Regulated Proteins	Function Role	Reference
Lung adenocarcinoma	miR-145	FSCN1	Chemoresistance↑, EMT↑	12
Non-small-cell lung cancer	–	–	–	23
Non-small-cell lung cancer	–	PI3K, Akt, mTOR, bcl-2	Chemoresistance↑	24
Breast cancer	–	–	EMT↑, chemoresistance↑	13
Breast cancer	–	–	EMT↑	14
Breast cancer	miR-205	ZEB1, ZEB2	EMT↑, chemoresistance↑	9
Breast cancer	–	DUSP7	Chemoresistance↑	28
Breast cancer	–	–	Autophagy↑, chemoresistance↑	29
Breast cancer	miR-34a	–	Chemoresistance↑	30
Breast cancer	–	TGF-β	Proliferation↑, invasion↑	31
Breast cancer	miR-145	MUC1	Invasion↑, migration↑	35
Breast cancer	miR-145	ARF6	Invasion↑, EMT↑	36
Breast cancer	–	–	–	37
Breast cancer	–	–	–	38
Hepatocellular carcinoma	miR-876-5P	FOXM1	Chemoresistance↑	40
Hepatocellular carcinoma	miR-145	RAD18	Radioresistance↑	44
Hepatocellular carcinoma	miR-145	ZEB2	Invasion↑, migration↑, EMT↑	45
Hepatocellular carcinoma	–	–	Resistance to hypoxia↑	46
Hepatocellular carcinoma	–	TGF-β	Chemoresistance↑	47
Colorectal cancer	miR-145	p53	Apoptosis↓, proliferation↑, radio-resistance↑	8
Colon cancer	miR-145	Sox2, Oct4 and Nanog	Proliferation↑, chemoresistance↑	49
Colon cancer	miR-145	–	Proliferation↑, invasion↑, migration↑	50
Pancreatic cancer	miR-124	PTBPI	Chemoresistance↑	53
Pancreatic cancer	–	ZEB1	Invasion↑, migration↑	56
Thyroid carcinoma	miR-145	TGF-β	Proliferation↑, invasion↑, EMT↑	59
Esophageal squamous cell carcinoma	miR-15b, miR-33a, miR-129, miR-145, and miR-206	Sox9	Proliferation↑, invasion↑, Migration↑, chemoresistance↑	61
Gastric cancer	–	–	Proliferation↑, invasion↑, stemness↑	15
Gastric cancer	–	–	Chemoresistance↑	63
Renal cancer	–	p53, c-Myc	Proliferation↑	65
Renal cancer	miR-206	VEGF	Proliferation↑, invasion↑, migration↑	66
Prostate cancer	miR-145	Oct4	Proliferation↑, invasion↑, stemness↑	69
Bladder cancer	–	ZEB1	Proliferation↑, invasion↑, migration↑	57
Ovarian cancer	–	–	Proliferation↑, invasion↑, migration↑, EMT↑	70
Ovarian cancer	–	–	–	71
Gallbladder cancer	–	TGF-β1	Proliferation↑, invasion↑, migration↑, EMT↑	72
Nasopharyngeal carcinoma	–	p53	Proliferation↑, invasion↑, chemoresistance↑	73
Glioma	–	SOX11, KLF4	Proliferation↓, stemness↓	74

that used induced pluripotent stem cells (iPSCs). The study showed that linc-ROR exerts vital effects towards maintaining stem cell pluripotency.⁷ Besides, it has been shown that linc-ROR promotes the development of many cancers. Epithelial-to-mesenchymal transition (EMT) plays an important role in aggressive traits of cancer cells and has been shown to be upregulated in almost all cancers.¹¹ Increased expression of linc-ROR induces EMT process and the acquisition of docetaxel-resistance in lung cancer (LC).^{12–14} In addition, aberrant expression of linc-ROR was associated with oncogenesis and progression of breast

cancer (BC),^{13,14} gastric cancer (GC),¹⁵ or colorectal cancer (CRC),¹⁶ among others. Accumulating evidence indicates that linc-ROR may serve as a cancer diagnostic and prognostic biomarker.^{17,18} The onset, maintenance and modulation of cancer stem cells (CSCs) were also reported to be tightly regulated by linc-ROR.^{7,19}

Together, these findings suggest that linc-ROR is clinically important as they can be targeted to control cancer development. This review summarizes the latest findings on the roles of linc-ROR in cancer, with an emphasis on the underlying mechanisms and clinical applications.

linc-ROR-Related Cancers

Non-Small-Cell Lung Cancer

LC is one of the most commonly diagnosed malignancies worldwide and the leading cause of cancer-related deaths.¹ In 2017, LC along with stroke and ischemic heart disease were named the three leading causes of short life expectancy in China.²⁰ Although tremendous advancements have been made in the management of cancer, LC treatment outcome remains poor. It is therefore critical to explore new and effective clinical biomarkers as well as novel therapeutic targets for this cancer.

Non-Small-Cell LC (NSCLC), the main subtype of LC, accounts for more than 80% of diagnosed cases of LC.^{21,22} Compared to the normal lung tissues, NSCLC tissues have higher expression of linc-ROR, hence advanced TNM stage and poor prognosis.²³ This suggests that linc-ROR promotes the progression of cancer and may serve as a prognostic biomarker in LC. Resistance to chemotherapy often leads to poor outcomes. Accumulating evidence suggests that linc-ROR is associated with chemoresistance in LC.^{12,24} For instance, by acting as a molecular sponge of miR-145, linc-ROR increases the expression of Fascin1 (FSCN1), a target gene of miR-145, thereby contributing to the acquisition of docetaxel-resistance in lung adenocarcinoma cells.¹² Linc-ROR also induces EMT in docetaxel-resistant cells, and promotes the invasion and migration of NSCLC cells.¹² Shi et al showed that the knockdown of linc-ROR improved sensitivity of NSCLC to cisplatin by regulating the PI3K/Akt/mTOR signaling pathway.²⁴ These findings suggest that linc-ROR may reverse the chemoresistance of NSCLC.

Breast Cancer

BC is the most common malignant tumor and leading cause of cancer-related deaths in women worldwide.¹ Despite significant advances in its management, the prognosis of BC remains poor due to the lack of advanced diagnosis, frequent distal metastasis and resistance to chemotherapy.²⁵

Studies have implicated linc-ROR in the initiation and development of BC (Figure 1) and its role in subverting drug-resistance has been confirmed. Many lincRNAs, including linc-ROR, regulate EMT by modifying the functions of associated proteins. Recent studies have indicated that linc-ROR promotes the invasion and migration of BC cells by inducing EMT.^{13,14} Tamoxifen, a selective modulator of estrogen receptors (ER), blocks estrogen-induced DNA transcription to inhibit the cell cycle and promote

apoptosis.^{26,27} However, drug-resistance decreases the efficacy of such treatments in up to 40% of BC patients. Evidence shows that linc-ROR participates in the oncogenesis, progression, metastasis and multidrug resistance of BC cells. Decreasing the expression of linc-ROR attenuates the resistance towards tamoxifen^{9,28,29} and suppresses the EMT process in BC cells.⁹ Overexpression of linc-ROR in BC tissues promotes EMT and tamoxifen resistance by acting as a molecular sponge of miR-205 and increasing the expression of (zinc-finger E-box binding homeobox 1) ZEB1 and ZEB2.⁹ linc-ROR can also promote estrogen independency and tamoxifen resistance by activating MAPK/ERK signaling pathways.²⁸ Given that Dual Specificity Phosphatase 7 (DUSP7) is an important inhibitor of the MAPK/ERK signaling pathway, linc-ROR activates ERK by facilitating DUSP7 degradation leading to tamoxifen resistance and tumor growth.²⁸ These findings not only associate MAPK/ERK signaling pathway to linc-ROR, but also reveal that estrogen deprivation induces ERK activation. Elsewhere, it was found that decreasing the expression of linc-ROR reversed resistance to tamoxifen and gemcitabine in BC.^{29,30} Hou et al indicated that overexpression of linc-ROR promotes the proliferation and invasion of BC cells.³¹ Moreover, linc-ROR activates TGF- β signaling pathway by increasing TGF- β levels, which further upregulates the expression of factors involved in the development of BC such as α -SMA and Smad2.^{31,32}

Triple-negative BC (TNBC) accounts for approximately 20% of all BC cases and is characterized by lack of HER2 gene, progesterone receptors or the amplification of estrogen.³³ TNBC is also associated with high rates of relapse and metastasis necessitating the search for more effective molecular targets for this cancer.³⁴ linc-ROR is upregulated in the tissues and cells of triple-negative BC (TNBC).³⁵ Given that the 3'UTR of Mucin1 (MUC1) is tightly linked to miR-145, the binding of linc-ROR and miR-145 increases expression of MUC1. High MUC1 expression in turn increases membrane localization of E-cadherin, contributing to invasion and migration of MDA-MB-231 BC cells. ADP-ribosylation factor 6 (ARF6) is also a target of miR-145. Interaction between linc-ROR and miR-145 increases ARF6 expression. Increased ARF6 expression promotes EMT in TNBC thereby accelerating cancer development.³⁶ Therefore, linc-ROR promotes BC by stimulating TGF- β signaling pathway or inducing EMT in an miR-145-dependent manner. It can, therefore, be targeted to treat BC.

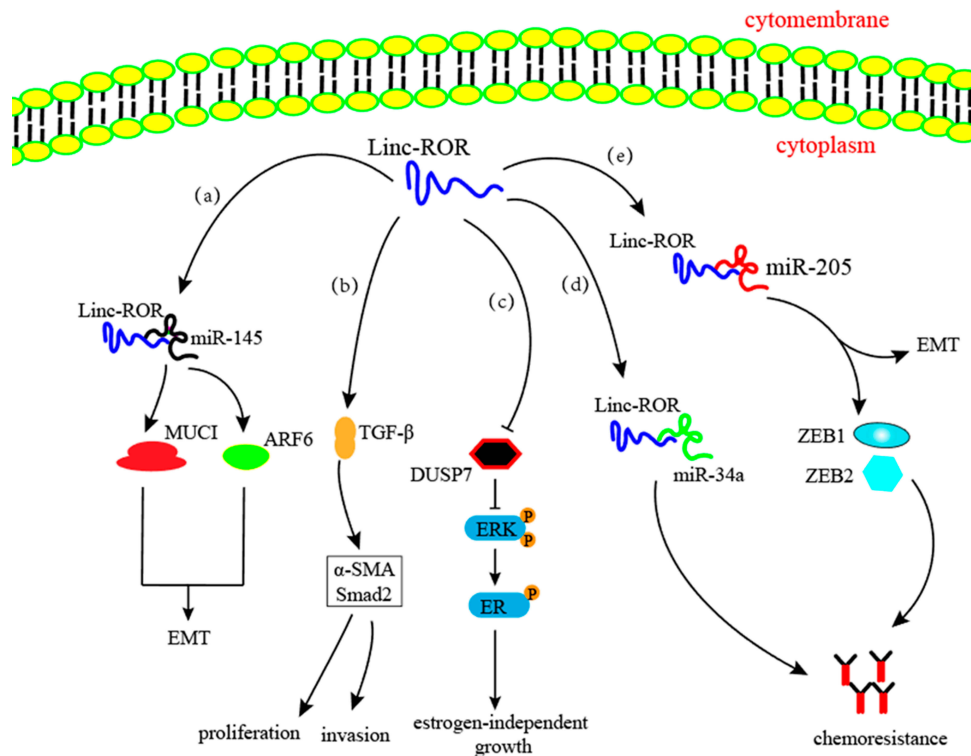


Figure 1 The regulatory mechanisms of linc-ROR in breast cancer. (a) Given that the binding sites between miR-145 and mRNAs of MUC1 and ARF6, linc-ROR promotes the EMT process in breast cancer by targeting miR-145; (b) linc-ROR overexpression activates TGF- β signaling pathway via upregulating the expression level of TGF- β , which triggers α -SMA and Smad2 expression, and thereby promotes the proliferation and invasion of breast cancer cells. (c) by facilitating the degradation of DUSP7, an inhibitor of MAPK/ERK signaling pathway, linc-ROR activates ERK and promotes the tamoxifen resistance and tumor growth of breast cancer. (d) linc-ROR down-regulates the sensitivity of tamoxifen in breast cancer cells by binding to the 3'UTR of miR-34a. (e) linc-ROR promotes EMT and tamoxifen resistance in breast cancer via acting as a molecular sponge of miR-205 and increasing the expression of ZEB1 and ZEB2.

Abbreviations: MUC1, Mucin1; ARF6, ADP-ribosylation factor 6; EMT, epithelial-mesenchymal transition; DUSP7, dual specificity phosphatase 7; MAPK/ERK, mitogen-activated protein kinase/extracellular regulated protein kinases; ZEB1, zinc finger E-box binding homeobox 1.

Previous studies showed that linc-ROR is a potential clinical biomarker in BC. Results from receiver operating characteristic (ROC) curve analyses show that linc-ROR is a superior diagnostic biomarker that outperform CEA and CA153 in BC.^{37,38} The expression level of linc-ROR in plasma may also be utilized to dynamically monitor the progress of BC.³⁵

Hepatocellular Carcinoma

Liver cancer is the fourth leading cause of cancer-related deaths worldwide and the fifth leading cause of deaths in China in 2017.^{1,20} Radiotherapy and chemotherapy are frequently used to treat cancers but their effects are limited by high drug-resistance as seen in hepatocellular carcinoma (HCC).³⁹ Thus, surgical resection remains the primary strategy for liver cancer treatment. Therefore, defining the mechanisms underlying HCC radio-resistance and chemoresistance, as well as identifying novel and effective targets to improve early diagnosis is imperative.

Zhi et al reported that Forkhead Box M1b (FOXM1) promotes the expression of linc-ROR in HCC. They identified a positive feedback loop among linc-ROR, miR-876-5p and FOXM1. In addition, they noted that elevated linc-ROR upregulated FOXM1 by acting as a molecular sponge of miR-875-5p,⁴⁰ thus enhancing resistance to chemotherapy and promoting HCC progression. Whereas overexpression of linc-ROR and FOXM1 enhances resistance to sorafenib in HCC, miR-876-5p inhibits proliferation, promotes apoptosis and increases sensitivity of HCC cells to drugs. The positive feedback loop promotes HCC progression and reduces sensitivity to sorafenib. Thus, silencing linc-ROR may be an effective approach for suppressing radio-resistance. Although the therapeutic value of sorafenib in HCC is limited by frequent chemoresistance, sorafenib improves HCC prognosis to some extent.⁴¹⁻⁴³ Therefore, characterizing the molecular mechanisms of sorafenib resistance is critical. linc-ROR/miR-876-5p/FOXM1 axis can be targeted to reverse sorafenib resistance in HCC. Besides,

the high radio-resistance rate substantially limits the effect of radiotherapy as a treatment of HCC. linc-ROR promotes radio-resistance in HepG2 and SMMC-7721 HCC cells.⁴⁴ Chen et al reported that linc-ROR upregulates the expression of RAD18 by acting as a competing endogenous RNA (ceRNA) of miR-145 thereby promoting radiation resistance in HCC cells.⁴⁴ Increasing the expression of linc-ROR also significantly reduces the γ -H2AX protein levels, a vital marker of DNA damage. This indicates that linc-ROR upregulates DNA repair activity and promotes radiation resistance.⁴⁵ It was reported that linc-ROR increases the expression of ZEB2 is also a target gene of miR-145 and by its expression, can also promote the proliferation, invasion and EMT in HCC cells. This effect has been also been confirmed by *in vivo* experiments.⁴⁵ A study by Takahashi et al reported that linc-ROR was up-regulated in hypoxic regions within tumor cell xenografts *in vivo*.⁴⁶ Knockdown of linc-ROR by small interfering RNA (siRNA) suppressed the proliferation of HCC cells whereas introduction of extracellular linc-ROR into HCC cells promoted cell survival under hypoxia conditions.⁴⁶ Extracellular linc-ROR was found to be associated with TGF- β -mediated chemoresistance in HCC.⁴⁷ In their study, knockdown of linc-ROR enhanced HCC chemosensitivity.⁴⁷ This indicates that linc-ROR reverses resistance to radiotherapy and chemotherapy in HCC patients.

Colorectal Cancer

CRC is the second most common cause of cancer-related mortality worldwide.¹ Numerous studies show that linc-ROR is overexpressed in CRC tissues compared with normal tissues. Some of the findings from recent studies that highlight the relationship between linc-ROR and CRC are discussed here.

It has been shown that miR-145 inhibits and even delays the development of CRC.^{16,48} linc-ROR was upregulated in both CRC tissue samples and cell lines. Evidence suggests that it is involved in the initiation and progression of CRC mainly by acting as a molecular sponge of miR-145.^{8,49,50} linc-ROR shares the miRNA binding sites of miR-145 with transcription factors, such as Sox2, Oct4 and Nanog.⁴⁹ Therefore, increasing the expression of miR-145 will significantly suppress the expression of linc-ROR, Sox2, Oct4, c-Myc and Nanog, thereby inhibit colon cancer stem cell proliferation and chemoresistance.⁴⁹ In colon cancer tissues,

overexpression of linc-ROR increased the expression of Sox2, Oct4, c-Myc and Nanog by tightly binding to the miRNA response elements (MREs) of miR-145, thus promoting CRC progression and decreasing sensitivity to radiotherapy and chemotherapy.^{8,49} In addition, linc-ROR knockdown inhibited tumor growth in nude mice after radiotherapy treatment suggesting that linc-ROR modulates sensitivity to CRC therapy. Yang et al reported that overexpression of linc-ROR increased p53-mediated (a tumor inhibitor) DNA damage, thereby promoting CRC oncogenesis.⁸ Thus, linc-ROR can be used as a prognostic biomarker and a target for the treatment of CRC.⁵⁰

Pancreatic Cancer

With less than 6% of overall 5-year survival, pancreatic cancer (PC) is one of the most aggressive malignancies.^{1,51} Gemcitabine is the mainstay drug for patients with PC ineligible for surgical resection.⁵² However, resistance to gemcitabine significantly limits its therapeutic effects. Li et al reported that linc-ROR promotes gemcitabine-resistance.⁵³

Studies have demonstrated that miR-124 can modulate the expression of PKM by switching PKM isoform expression from PKM2 to PKM1.^{54,55} Given the close link between PTBP1 (a key regulator of PKM splicing) and miR-124, linc-ROR enhances the expression of PKM1 and inhibits PKM2 expression in a miR-124-dependent manner. This cascade of events confer gemcitabine resistance to PANC-1, MIAPaCa-2 and SW1990 PC cells.⁵³ In addition, linc-ROR induces EMT by upregulating ZEB1, which then promotes the aggressive behaviors of PC cells.⁵⁶ Although studies have reported that linc-ROR exerts important effects by modulating ZEB1 in many types of cancer,^{9,57} the specific miRNA mediating ZEB1 and linc-ROR remains unclear.

Papillary Thyroid Carcinoma

Thyroid carcinoma is the ninth most commonly diagnosed malignancy worldwide with approximately 567,000 cases.¹ It is also the most abundant cancer in Korea, with more prevalence in females.^{1,58} Despite the advances in the management of thyroid cancer which have significantly improved its prognosis, its etiology remains unclear. So far, ionizing radiation is the only risk factor associated with the development of cancer. Therefore, exploring its etiology is central to improving the quality of life for patients with thyroid cancer. linc-ROR was unregulated

in papillary thyroid carcinoma (PTC) tissues. A negative association between linc-ROR and miR-145 has been identified in TPC-1 cells.⁵⁹ By measuring the expression level of linc-ROR after treatment with or without TGF- β , it was found that linc-ROR promoted the progression of PTC by modulating the expression of TGF- β and EMT process. Since TGF- β is an important regulator of the EMT process in PTC,⁵⁹ silencing linc-ROR may be an effective measure to control PTC.

Esophageal Squamous Cell Carcinoma

Esophageal carcinoma is a serious threat to public health, ranking seventh among all malignancies worldwide.¹ Esophageal squamous cell carcinoma (ESCC) is most prevalent among Eastern Asian men. ESCC is a subtype of esophageal cancer derived from esophageal epithelial cells.⁶⁰ Currently, the molecular mechanisms underlying development of esophageal cancer are not well understood.

linc-ROR has been shown to modulate proliferation, invasion, migration and chemoresistance in EC9706 ESCC cells. It increases the expression of Sox9 by regulating miRNAs such as miR-145, miR-206, miR-15b, miR-33a and miR-129.⁶¹ It was noted that knockdown of Sox9 reversed the effects of linc-ROR overexpression in EC9706 ESCC cells. Utilizing a target prediction tool,⁶² 12 miRNAs were identified to share binding sites with linc-ROR and Sox9. Finally, it was reported that miR-206, miR-145, miR-129, miR-33a and miR-15b modulated Sox9 via linc-ROR. Moreover, knockdown of linc-ROR in vivo not only suppressed tumor growth, but also reduced expression levels of Sox9 and CSC-marker CD44.

Gastric Cancer

GC is one of the most common life-threatening cancers globally. According to global cancer statistics, GC was the fifth most commonly diagnosed cancer and third leading cause of cancer-related deaths in 2018.¹ Exploring the molecular mechanisms of GC progression may lead to the development of strategies and treatments to improve the prognosis of GC patients. Numerous lncRNAs, including linc-ROR, have shown promising potential in GC management.

Wang et al found that knockdown of linc-ROR suppressed the proliferation and invasion of gastric cancer stem cells,¹⁵ suggesting that it may be a potential treatment target for GC. Moreover, they found that the expression of linc-ROR was positively associated with multiple-drug resistance and poor prognosis of GC patients.⁶³ Silencing

linc-ROR increased apoptosis of Adriamycin- and vincristine-resistance GC cells, and thereby subverting tumor development.⁶³

Urologic Cancer

Urologic cancers (UC) are some of the most common malignant tumors. It comprises renal cancer, bladder cancer and prostate cancer. Previous studies showed that prevalence of renal cancer (RC) has been on the rise and its prognosis is poor.^{1,64} Prostate cancer is the second commonest malignancy in men and the fifth leading cause of mortality.¹ Bladder cancer is the seventh most prevalent malignancy in men. Further research is required to elucidate its pathogenesis, identify novel biomarkers and effective therapeutic targets to improve early diagnostic rates and prognosis of UC.¹

Shi et al showed that RC cells and tissues overexpressing linc-ROR exhibited aggressive cancer phenotypes, as well as poor prognosis.⁶⁵ In vitro studies showed that silencing linc-ROR increased cell apoptosis and decreased the expression of c-Myc while increasing the expression of p53.⁶⁵ Other studies found that linc-ROR promotes RC by modulating miR-206/VEGF axis.⁶⁶ On the other hand, overexpression of miR-206 can reverse linc-ROR-induced proliferation and invasion of RC cells. linc-ROR upregulates the expression VEGF by sponging miR-206 thereby promoting RC cell proliferation, invasion and migration. These findings suggest that silencing linc-ROR may be a potential target for RC treatment.

By modulating the miR-145/Oct4 axis, linc-ROR also promotes human prostate cancer stem cell proliferation and invasion. Previous studies have demonstrated that Oct4, a marker of stem cells, is closely associated with the prognosis of cancer.^{67,68} Given the close link between the 3'UTR of miR-145 and Oct4 mRNA, the binding of linc-ROR and miR-145 increases expression level of Oct4.⁶⁹ Liu et al reported that curcumin suppresses the proliferation of human prostate cancer stem cells in nude mice by inhibiting the expression of linc-ROR and enhancing miR-145 expression.⁶⁹ Thus, linc-ROR/miR-145 may be a good therapeutic target for the treatment of prostate cancer. Besides, linc-ROR was upregulate in bladder cancer tissues compared to adjacent normal tissues and bladder cancer cell lines.⁵⁷ ZEB1 is not only increased in bladder cancer but also positively correlated with the expression of linc-ROR. Therefore, overexpression of linc-ROR promotes the proliferation, invasion and migration of bladder cancer cells in a ZEB1-dependent manner.⁵⁷

Other Cancers

Other studies have indicated that linc-ROR can also regulate the initiation and progression of ovarian cancer (OC),^{70,71} gallbladder cancer,⁷² nasopharyngeal carcinoma (NPC)⁷³ or glioma.⁷⁴

It was reported that linc-ROR activates of Wnt/ β -catenin signaling pathway and induces EMT, thereby promoting OC development.⁷⁰ Increasing the expression of linc-ROR up-regulates TGF- β 1 expression and further induces EMT process in gallbladder cancer.⁷² In this way, linc-ROR promotes the proliferation, invasion and migration of gallbladder cancer cells suggesting its importance as a prognostic biomarker. Resistance to chemotherapy and radiotherapy leads to poor NPC prognosis.⁷⁵ Therefore, exploring new targets that help reverse resistance to treatment may be a novel therapeutic strategy. Li et al reported that expressing linc-ROR not only enhanced the proliferation, invasion and migration of NPC cells but promoted chemoresistance.⁷³ Evidence indicates that linc-ROR suppresses the sensitivity to chemotherapy through the p53 pathway in cancer treatment. It was found that linc-ROR enhanced resistance to chemotherapy in NPC cells by modulating p53 levels.⁷³ Although linc-ROR exerts oncogenic effects in almost all types of cancer, it was also found to be a tumor suppressor in glioma by inhibiting the proliferation of glioma cells and the self-renewal of glioma stem cells.⁷⁴

Perspective

Numerous studies have investigated the roles of lincRNAs in cancer development. However, the full landscape of lincRNAs with diagnostic and therapeutic value is yet to be unraveled. linc-ROR is a novel functional lincRNA that is associated with the onset and progression of many types of cancers except glioma. linc-ROR promotes the proliferation, invasion, migration and drug-resistance of cancer cells through a variety of mechanisms at pre-transcription and post-transcription levels. Mechanistically, linc-ROR modulates miRNAs, gene transcription and protein translation. linc-ROR indirectly regulates the activities of oncogenic proteins by interacting with microRNAs which influences tumor progression. Of note, linc-ROR may directly exert its oncogenic effects by either directly reducing miR-145 levels,⁵⁰ or regulating the expression of miR-145's downstream targets, such as ZEB2,⁴⁵ p53⁸ or TGF- β ,⁵⁹ and c-Myc.⁶⁵ Investigation of the mechanisms underlying cancer initiation and progression is important in order to improve the therapeutic strategies. These crucial linc-ROR functions define potential for clinical application in cancer treatment.

Growing evidence has demonstrated that linc-ROR may not only be used as a diagnostic or prognostic biomarker,^{23,37} but also as a therapeutic target for cancer treatment. Among all the clinical applications, the most promising is its role in reversing resistance to chemotherapy. In LC, up-regulation of linc-ROR attenuates the sensitivity to docetaxel and cisplatin.^{12,24} Although tamoxifen is an important supplement for surgery in the treatment of BC, studies have found that linc-ROR overexpression can reduce its therapeutic effects.^{9,28,29} Moreover, reducing the expression of linc-ROR in BC cells can partly reverse resistance to tamoxifen, further suggesting that linc-ROR may be a chemotherapeutic target in the treatment of BC. Besides, linc-ROR also enhances chemotherapeutic sensitivity in HCC,⁴⁴ CRC,⁴⁹ PC,⁵³ ESCC⁶¹ and NPC.⁷³ However, these findings are derived from cell experiments. Further clinical studies employing different methodologies are needed to investigate additional mechanisms and roles of linc-ROR in cancer. This will lead to the identification of lincRNA-based therapies for cancer.

Conclusion

We have reviewed recent studies interrogating the relationship between linc-ROR and malignant tumors, as well as analyzed the underlying characteristics defining the effects of linc-ROR on cancer. Significant oncogenic effects of linc-ROR were observed in various cancers with the most important being that linc-ROR promotes cancer progression by acting as a sponge of miR-145. linc-ROR pathway may be exploited to develop diagnostic or prognostic biomarkers and therapeutic targets. Besides, linc-ROR can be used to reverse chemotherapeutic resistance.

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Disclosure

The authors report no conflicts of interest regarding this work.

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